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DNA-based identification of forensically important Diptera, 2005-DA-BX-K102, Ronald W. DeBry

Abstract

Flies (Diptera) from three families (Calliphoridae, Sarcophagidae, Muscidae) are normally the first insect visitors to an exposed body. Many species of dipterans either lay eggs or deposit live larvae into a body within hours of death. By determining both the species identity and the age of larvae present at a human death scene at the time of discovery, it is possible to estimate the post-mortem interval. With regard to species identity, common practice requires collection of live larvae that are maintained until they achieve adulthood, at which time they can be identified to species using morphological characteristics.

Purpose: The purpose of this project is to develop a reference database of DNA sequences that is suitable for species-level identification of forensically relevant Diptera from the 48 contiguous United States. We focus here on flies in the families Calliphoridae and Sarcophagidae. This reference database will be suitable for querying DNA sequences obtained from larvae collected at a death scene.

Goals and Objectives: Goals of this project were to: 1) Collect reference adult dipteran specimens from across the 48 contiguous states. Specific attention was to be given to collecting specimens from widely spread geographic localities, with an emphasis on the margins of the relevant area. 2) Identify a genetic locus that is suitable for discrimination of relevant species in a phylogenetic analysis. 3) Develop a reference DNA database. The database will have associated with it all original physical specimens preserved as vouchers, with sufficient morphological features retained so as to allow future re-examination of the specimens.

Research subjects: The research subjects for this project are flies (Diptera) of the families Calliphoridae, Sarcophagidae, and Muscidae.

Research Design: The project is divided into two phases. In Phase I, we examine a range of possible genetic loci from a limited number of species and individuals, in order to assess the relative performance of the different loci (and regions therein) at discriminating among species. In Phase II, data are collected from a large number of species and individuals for the locus determined in Phase I to be best for discrimination among forensically important species.

Methods: Adult flies were collected both using traps baited with meat and by sweep netting. Specimens were identified using morphological characteristics. Those flies from the forensically relevant families were identified to the species level. A small portion of each specimen (usually 1-3 legs) was removed and used for DNA extraction. Loci of interest were amplified by the polymerase chain reaction, and the DNA sequence was determined for the resulting product. The utility of the final reference database was assessed by phylogenetic analysis with bootstrap resampling using 1000 pseudoreplicates.

Results: Over 2,500 flies were collected and identified from 84 different counties in 20 different states. In Phase I, it was determined that any region of the mitochondrial genome was likely to be sufficient for discrimination among most forensically relevant species, but that a region within the mitochondrially encoded gene Cytochrome oxidase I was best for discrimination between a particularly closely related pair of species. In Phase II, DNA

sequence for the chosen locus was obtained for a total of 504 individual flies representing 106 distinct species.

Conclusions: The reference DNA database is suitable (based on bootstrap support values >95%) for correct species identification of 16 of the 18 species that are most important for forensic investigations. One pair (*Lucilia coeruleiviridis* and *Lucilia mexicana*) are indistinguishable, apparently for any mitochondrial locus. A further 16 of 20 species represented in our database that are deemed to be of lesser forensic utility can be unambiguously identified using phylogenetic analysis of the reference database, with two species pairs (both in the genus *Ravinia*) being indistinguishable.

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Executive Summary

Synopsis of the problem. A fundamental task in death investigation is determination of the post-mortem interval (PMI). One tool that can sometimes be used for PMI estimation is the developmental stage of carrion-feeding insects (forensic entomology). During the relatively early stages of decomposition, the primary entomological evidence is obtained by determining the developmental stage and species identification of larval flies (Diptera), primarily of the families Calliphoridae, Sarcophagidae, and Muscidae. Species identification is crucial, because different species develop at different rates under the same environmental conditions. Species identification by gross morphology requires adult flies in a large proportion of cases, because larvae of many species at many age ranges often do not carry sufficient distinguishing characteristics.

Existing practice requires collection of live larvae, with (generally) transfer of those larvae to an established forensic entomologist, who will rear the larvae to adulthood (ca. 1-2 weeks in most cases) and then make the species determination. Possibly in large part because of this step, entomological evidence is generally under-utilized in death investigation in the United States. For a number of years, an important goal in forensic entomology has been to

develop a method capable of providing accurate species identification directly from the dipteran larvae.

Purpose. The goal of this project is to provide the tools needed for quick, reliable species identification of forensically important flies directly from larvae. To accomplish this goal, we have developed a database of DNA sequences from a known-identity reference collection. The original reference collection consists of adult flies that have been identified to species by an expert in the taxonomy of the group, and is maintained as vouchers for future morphological study, should the need arise. The functional part of the identification database is the set of DNA sequences generated from those vouchers. It is presented as an appendix to this report. We have further developed PCR primers that are useful for amplification of a DNA segment that is appropriate for discrimination among the large majority of the species of Diptera that are likely to be encountered in an early-stage forensic setting in the contiguous United States.

Research design.

Development of a list of the primary forensically important species of Diptera: We examined a large variety of relevant literature for lists of dipteran species that had been identified either as being encountered in a forensic investigation or as having a life cycle that is associated with large animal carrion, with the focus being on flies that are likely to be among the earliest colonizers of a corpse. That process led us to develop the list of forensically most important species, which are shown in red boldface in Table 1 of the main body of this report. In total, we identified 11 species of Calliphoridae (blow flies), 6 species of Sarcophagidae (flesh flies), and 2 species of Muscidae (house flies) as being the dipterans most likely to be relevant for a forensic death investigation in the contiguous United States.

Sampling strategy: Our sampling strategy had the following goals:

1. Maximize the probability that we will obtain samples for all of the species of primary forensic importance in the contiguous United States.
2. Ensure that we also sample species that, while not of primary importance, have been reported as being associated with large-animal carrion.
3. Maximize the geographic coverage for those species of primary forensic importance that have large geographic ranges.
4. Maximize the total species coverage of the reference database. A reference database for species identification should include as many species as possible that both are found in the relevant geographic area and are phylogenetically related to the species of interest.

Goals 1 & 2 were addressed in two ways. First, we undertook a number of collecting trips to a wide variety of regions of the country, and to a diversity of habitats. Second, we focused our collection procedure on trapping adult flies that were attracted to a meat bait (normally chicken meat obtained from a local grocery store). This bait was not only highly attractive to those species that are of greatest forensic interest, it would also attract adult flies that were simply searching for a protein meal (as compared to the flies of primary forensic interest, which would be attracted to the bait to deposit eggs or larvae). Such “incidental” visitors to the carrion bait would likely include many or most of the species represented by Goal 2. Goal 3 was addressed by our choice of regions in which to collect reference samples. We focused our effort on the margins of the continent. So, for example, we undertook collecting trips to New York, Maine, New Mexico, Florida, Wisconsin and Oregon. If a widespread species of primary forensic importance did have geographic variation in mtDNA

haplotype, those differences would be expected to be largest between widely separated populations. Goal 4 is important for the reduction of instances in which a query to the database is made that is, in fact, a species not represented by any reference specimens. Thus, in addition to the primary collection effort using bait and traps, adult sarcophagid and calliphorid flies were also collected at most locations using sweep nets.

Collection of fly reference specimens We undertook 6 major collecting trips. Flies were collected at a total of 84 different counties in 20 different states over a three year period. As most species can only be reliably identified using morphology as adults, we collected only adult flies. The geographic coverage of these trips was specifically designed to maximize the geographic diversity of our samples, particularly for those forensically important species that can be found throughout North America. In this way, we would maximize our chance to observe geographic variation in DNA type within such widespread species. The primary means of capture used for this project were traps set above a bait (usually consisting of raw chicken). In addition, adult flies were caught by sweep netting in the vicinity of the chicken bait.

Flies were identified by our consultant and collaborator, Dr. Gregory Dahlem of Northern Kentucky University. Some particularly problematic specimens (primarily in the genus *Lucilia*) were sent to Dr. Terry Whitworth for confirmation. More than 2,500 individual flies were collected and identified for this project (Table 3), and we also included a small number of additional flies that we had collected prior to the initiation of this project. Once identified, each fly either: had 3 legs removed (with the legs placed in EtOH for later DNA extraction), while the remainder of the specimen was pinned as a standard insect specimen or, in some cases, the entire fly was preserved in EtOH. DNA was extracted from individual legs, using either a commercial kit (DNEasy, QIAGEN) or by sonication for 30-300 s in 50ul water in a centrifuge tube placed into a sonicating water bath. All flies identified for use in this study have been retained as voucher specimens (either pinned or in EtOH). All vouchers for individuals for which we obtained DNA sequence data and, thus, are part of the reference database will be deposited in the Florida State Collection of Arthropods by July 2012. A complete list of specimens and their identifications, collecting localities, and associated information is provided as an Excel spreadsheet attached to this report.

Molecular methods. Loci of interest were amplified by the polymerase chain reaction (PCR), using primer sequences either obtained from the literature or newly designed for this project. PCR products were sent to the High-Throughput Genomics Lab at the University of Washington for DNA sequencing.

Phase I: Phase I of the project was intended to identify the specific locus on the mitochondrial genome (mtDNA) that would provide the best level of species discrimination. In Phase I, data collection consisted of a small collection of 28 individuals representing 8 commonly encountered, forensically relevant species. We amplified and determined the DNA sequence for these specimens for the entire coding region of the mtDNA genes Cytochrome Oxidase I (COI), Cytochrome Oxidase II (COII) and NADH dehydrogenase subunit 4 (ND4). The Phase I data were analyzed in sliding 600-bp windows to determine if there were substantial differences across regions in the ability of a 600-bp region to allow successful discrimination among species. For most species pairs in the Phase I analysis, any 600-bp segment resulted in complete segregation of the individuals belonging to each species. The exception to that pattern was the pair of species *Lucilia cuprina* and *L. sericata*. These two species were known from previously published work (Wells, Wall, Stevens 2007) to be extremely closely related and particularly difficult to distinguish using mtDNA data. For *L. cuprina* and *L. sericata*, we

found that only the 3' end of the COI gene provided nucleotide characters that could reliably distinguish between our samples of those two species. We therefore chose a pair of primers that produce an amplicon that includes the best 600-bp window for use in Phase II.

Phase II. Phase II consisted of amplification and sequencing of the target locus for a total of 504 individual flies representing 106 distinct species. Our original PCR primers covering the 600-bp target locus actually amplify a 1,100-bp amplicon, so the full 1,100-bp region was amplified and sequenced for Phase II. Raw sequence reads were collated and aligned with the aid of the computer program BioEdit (Hall 1999). Because the target locus encodes part of the Cytochrome oxidase protein complex, there was essentially zero variation in the length of the target region, so generation of the multiple alignment was trivial.

The aligned data were transferred to two different standard data formats: Nexus (used by the computer package Mesquite, among others) and MEGA (used only by the computer program MEGA 4). Mesquite was used primarily for data storage and retrieval. MEGA 4 was used for most of the phylogenetic analyses reported below.

Bootstrap resampling followed by phylogenetic analysis of each pseudoreplicate data matrix was the primary analysis performed on our data and the primary means used herein for assessing whether or not the DNA sequence database achieves the stated goal of being sufficient for the identification of the forensically important Calliphoridae and Sarcophagidae of the contiguous United States. Bootstrap resampling is a standard non-parametric statistical technique for assessment of levels of variability when the underlying sampling distribution is not known, and is a generally accepted approach for species identification in forensic entomology (Wells, Williams 2007; Wells, Stevens 2008; Meiklejohn, Wallman, Dowton 2011b). For the purpose of this report, we define successful species discrimination in the following way: The reference samples of the species in question must form a *monophyletic group* on the phylogenetic tree. That means that all sampled individuals of that species must be traceable back to a single node on the tree, and that there must be reference samples from no other species subtending that same node. In addition, the ancestral node so defined must be supported in the bootstrap resampling analyses in at least 95% of the pseudoreplicates generated and analyzed. Potential shortcomings and alternatives to this approach are discussed later in this report.

Findings

Main findings

Classification of species according to forensic importance: Based on literature reports, we designated 18 species as being of primary forensic importance (Table 1). We designated an additional 20 species as being of possible forensic importance.

Traceability of this database: A major feature of the database we have compiled is that all of the reference specimens were collected specifically for this project. No data were obtained from GenBank, or other publications. All species determinations were performed based on adult morphology by Dr. Greg Dahlem, a widely recognized expert in the groups of flies involved (see below), and all reference specimens are maintained in a single collection. This last point is extremely important should any of the species identifications come into question, or should future taxonomic changes affect the use of entomological evidence, and is the primary reason why we used only specimens that we have physically in our possession.

Collection success: We successfully collected samples for 18 of the 19 species that we designate as being of primary forensic importance. Of the 22 species of lesser forensic importance, we were able to collect or obtain specimens for all but 2 species. The forensically important species that are reported to occur in the 48 contiguous United States that we do not have represented in our reference database are: *Comptosyiops callipes*, which is found in the southwestern U.S., *Lucilia eximia*, which is commonly found in the forensic setting in Central and South America, but is only rarely found in the United States (Whitworth 2006); and *Sarcophaga ruficornis*, which is commonly found in Hawaii, but is rare in the western contiguous United States (Whitworth 2006). A summary of the samples included in the reference database for species of primary and secondary forensic importance can be found in Table 4, in the main body of this report. For most of these “most forensically important” species, we were able to collect specimens from widely distributed geographic locations. This latter point is critical. The presence of substantial variation in the DNA sequence within a species across its geographic range would be of considerable importance. On the positive side, it would, perhaps, be possible to infer the general geographic origin of an individual fly, based on its DNA sequence (although this presumably would be of forensic relevance only in the case in which a victim was transported across a substantial fraction of the continent). On the negative side, finding substantial geographic variation would necessitate development of a much larger, more comprehensive reference database, in order to ensure adequate geographic coverage. From that perspective, it is fortunate that we identified only a very small number of instances in which there is substantial geographic variation within a single species.

In addition to those species denoted as being of primary or secondary forensic importance, we also collected specimens from an additional 68 species. These additional species vary in their direct relevance to the forensic entomology database. Some have been rarely reported as having been found either in an actual forensic investigation or as fauna associated with large-animal carrion. Those reports may be the result of incidental appearance at a crime scene, or the result of misidentification, or the result of taxonomic confusion on the part of the authors of the report in question. Although they are unlikely to be of direct value in forensic investigations, it is critically important that these species be represented in a DNA database for forensic entomology. Others of the “additional” species probably are of no forensic importance. Presumably, their capture (by us) was incidental to the presence of the carrion bait. These samples, however, are also important to have represented in the database, because they could also be collected as incidental visitors at a death scene. We captured them either directly using traps baited with meat or by sweep net in the vicinity of such a trap (a similar procedure might be followed as part of a forensic investigation). Consequently, their inclusion in the reference database will greatly reduce the chance that an unknown sample being queried to the database will, in fact, be a species that is not represented.

Phase I

Results of Phase I: Phase I of the project had the goal of choosing an appropriate genetic locus on which to base our reference database. For our in-depth analysis of a larger region of sequence data, we examined a series of 600-bp sliding windows across the complete COI+COII region. We found, for most comparisons in the limited taxonomic sample, that any 600-bp region is adequate for discrimination. Only for the very closely related species pair *Lucilia cuprina* – *L. sericata* does the ability to discriminate vary across 600-bp windows (Fig. 1). For that pair, a region extending from position 1800 – 3000 is superior (with the numbering system being based on the GenBank sequence for the complete mtDNA genome of *Drosophila yakuba*). We therefore undertook to collect data from that region for the larger part of the project (Phase II).

Phase II

Overall scope and success of the DNA sequence reference database: For Phase II, we collected, identified, and obtained the DNA sequence for the 1800-3000 region of the COI gene for 504 individual flies, representing 106 different species (33 species from the family Calliphoridae; 71 species from the family Sarcophagidae; and 2 species from the family Muscidae). The full phylogenetic analysis will be presented as Fig. 10 in the main body of this report. For the 18 species of primary forensic importance, 16 of them appear to be reliably distinguishable using the database. Of the 20 species of secondary forensic importance, 16 appear to be reliably distinguishable using the database.

Geographic variation: If there is substantial geographic variation within any of the forensically important species, then any reference database would need to be much larger (sufficiently large so as to include a representative sample from a large number of populations across the full species range. We find only two species to have substantial levels of intra-specific variation with any hint of a geographic pattern. Larger samples from these two species (*Lucilia mexicana* and *Sarcophaga bullata*) would be warranted.

Species-specific findings: Nearly all of the forensically important species in our database exhibit reciprocal monophyly with respect to all other species in the database. Here, we discuss the exceptions to that generalization:

Lucilia cuprina – L. sericata: These two species were known in the existing literature to present a particularly difficult problem for DNA-based identification (Stevens, Wall, Wells 2002; Wells, Wall, Stevens 2007; DeBry et al. 2010). We collected and sequenced both species intensively, and find that our preferred locus is successful at discriminating between these two species (see Fig. 10 in the main body of this report). In our main phylogenetic analysis, each is supported as being monophyletic with bootstrap values > 95%. We have reported on this specific issue in a publication – DeBry et al. (2010).

Lucilia coeruleiviridis – L. mexicana: Of all the species of primary forensic importance, these are the two that cannot be distinguished using DNA sequence from the mtDNA locus that we chose for construction of our database. As seen in Fig. 10, a large preponderance of the mtDNA haplotypes for these two species are intermingled. In many regions of the country, only one of these two species is found (with *L. coeruleiviridis* having the much larger range). They do co-occur in the Southwest, so many of our sequenced specimens are from New Mexico. The impact of the inability to distinguish between these two species will depend primarily on two factors: 1) how common they are in the forensic setting (if they are only of minor importance, it is likely that other species (that can be identified using mtDNA) will be present as well, so any PMI estimates could be derived from the species with the more certain identification; 2) how different are the developmental profiles of the two species in the area in which they co-occur.

Ravinia: There are two pairs of species within this genus that present considerable problems for DNA-based identification using our database. The impact is likely to be minimal, however, as the species in this genus have larvae that primarily feed on animal dung and they are of limited forensic utility. In addition, according to our consultant (Dr. Greg Dahlem, a

recognized expert on the biology of *Ravinia*), even the morphologically defined limits of species are uncertain in these particular cases.

Sarcophaga bullata: one specimen, our only example of this species from California, sometimes groups with a different species, *S. polistensis*. This causes the bootstrap support for monophyly of *S. bullata* to fall to ~83%. This result is, however, very unlikely to be of any import in the forensic context. This is because *S. polistensis* is a highly specific parasitoid whose larvae only develop within the bodies of large wasps of the genus *Polistes*.

Therefore, of the 18 species of greatest forensic importance, only the pair *Lucilia coeruleiviridis* – *L. mexicana* cannot be identified to a reasonable degree of confidence using our database.

Additional findings

Sonication of legs is a fast and inexpensive method for extraction of genomic DNA for use in PCR.

Main Body of the Final Technical Report

I. Introduction

STATEMENT OF THE PROBLEM

A primary task in death-scene investigation is estimation of the post-mortem interval (PMI). Beyond approximately 24-48 hours post mortem (Fatteh 1973), a major forensic tool is the presence and developmental staging of larvae deposited on the corpse by various species of carrion-feeding flies (Hall 1990). By applying information regarding species-specific colonization times and maturation rates, it is possible to estimate of a window within which the corpse was initially exposed to colonization (Greenberg 2002).

Accurate species identification of dipteran specimens is the first step in using those specimens for PMI estimation (Greenberg 1985; Smith 1986). Without this information, there is the significant risk that any further insight that could be gained from the specimen will be flawed. For example, very closely related carrion fly species can have substantially different growth rates (Kamal 1958) or diapause response (Ash, Greenberg 1975). Misidentification of one species as another might then lead to substantially different estimates of time since colonization, which in turn is often used to provide a minimum estimate of PMI. Diagnostic anatomical differences to identify carrion fly larvae to the correct species are simply not known.

Traditional practice in forensic entomology has been to collect live larvae at the death scene, and transfer that evidence to a qualified forensic entomologist, who will rear the larvae until they achieve adulthood, and then identify the individuals to the species level using traditional morphological characteristics (Amendt et al. 2007). This practice is both time-consuming and time-critical (the live larvae will not wait for a crime lab to find the time to process the transfer of the evidence to an entomologist).

Beginning with the pioneering work of Sperling, Anderson, Hickey (1994), it has been recognized that DNA could be used to identify forensic specimens to species. Because a larva

has the same DNA that it will have as an eventual adult, DNA-based species identification can be applied directly to the larvae. Further, DNA sequencing does not require live larvae. So long as the larvae are correctly preserved at the time of collection, they can be stored indefinitely.

DNA-based species identification requires that there be a suitable reference database against which to compare the unknown specimen. In order for the species identification protocol to have the maximum utility, the reference database must meet the following criteria:

1. All species likely to be encountered in a forensic setting for the geographic region of interest should be represented.
2. Each forensically important species should be represented by a large number of reference specimens, from throughout the geographic range of the species, or throughout the geographic range covered by the reference database.
3. The physical reference specimens should be maintained in a publicly available collection

Point 1 is important because most approaches to DNA-based species identification can sometimes produce ambiguous or unpredictable results if presented with a specimen from a species that is not included in the reference database. If a reference database is intended to be used over a specified geographic area, then reference samples should be included from all target (forensically important) species *and* from all closely related species that might be attracted to carrion, even if not to breed. Therefore, any species that might be encountered as an accidental visitor to carrion should be included in the reference collection. Most existing DNA databases for forensic entomology include only the forensically most-relevant species.

Point 2 is critical (Wells, Stevens 2008) because of the variability inherent in natural populations. A larger number of reference specimens will have a correspondingly greater probability of including low-frequency genotypes. Further, the variability within a species might represent geographic structure (genotypes that are rare in one part of a species' range might be common in another). Early DNA databases for forensic entomology typically included only a single exemplar for each forensically important species, but more recent studies have substantially increased the number of reference samples per target species.

Point 3 is also critical. The original specimens should be available for examination, should anyone wish to challenge the correct species identification of a specimen used to construct the reference database, or should future taxonomic work change the number of recognized species, or change the morphological definition of a species, or even simply change the name applied to a species. The existing record for forensic entomology DNA databases is mixed. Most papers describing databases make no mention of the current disposition of the reference specimens.

Our goal in the present project is to address all three of these important/critical points. We include in our reference database all but 2 species found in the contiguous United States that are generally considered to be of primary forensic importance (and neither of those species are widespread or common). We also include all species within the families Calliphoridae and Sarcophagidae that were attracted to our baited traps, whether or not they are recognized as species of forensic interest. For those species that are of primary forensic importance, we include a relatively large number of reference samples. In addition, we specifically target specimen collection for a wide diversity of geographic locations across the contiguous U.S., thus maximizing our chances to discover any geographic pattern to intra-species variation. Finally, we maximize the future utility of our reference specimens by using only 3 legs for DNA extraction (thus preserving the body parts that are most important for

morphological species identification), and by retaining all voucher specimens. We will deposit all reference specimens into a recognized, publicly available natural history museum (the Florida State Collection of Arthropods).

REVIEW OF RELEVANT LITERATURE

A primary task in death-scene investigation is estimation of the post-mortem interval (PMI). Beyond approximately 24-48 hours post mortem (Fatteh 1973), a major forensic tool is the presence and developmental staging of larvae deposited on the corpse by various species of carrion-feeding flies (Hall 1990). Existing information regarding species-specific colonization times and maturation rates allow estimation of a window within which the corpse was initially exposed to colonization (Greenberg 2002).

Three groups (taxonomic families) of carrion flies are of primary importance in PMI investigations, particularly in the early stages of decomposition: blow flies (Diptera: Calliphoridae), flesh flies (Diptera: Sarcophagidae) and house flies (Diptera: Muscidae) (Byrd, Castner 2001). Among the fly fauna of the contiguous U.S., 14 species in the family Calliphoridae, 25 in the family Sarcophagidae, and 3 in the family Muscidae have been reported in the literature as being of forensic importance (Smith 1986; Byrd, Castner 2001; Greenberg 2002).

Accurate identification of dipteran specimens is the first step in using those specimens in a forensic setting (Smith 1986; Greenberg 2002). If the identification is incorrect, there is the significant risk that any further insight that could be gained from the specimen will be flawed. For example, even very closely related carrion fly species can have substantially different growth rates (Kamal 1958) or diapause response (Ash, Greenberg 1975). Misidentification of one species as another might then lead to substantially different estimates of time since colonization, which is often the primary variable used in the calculation of a minimum estimate of time since death (PMI, or the postmortem interval). Species-diagnosis using anatomical differences for the larval and even adult forms of many carrion flies simply are not known.

Because of the critical nature of the questions being asked from these specimens, there have been increasing attempts to utilize molecular data to increase the accuracy of species determination in the field of forensic entomology. Such work began in the mid-1990s (Sperling, Anderson, Hickey 1994; Benecke 1998) and has grown substantially in recent years (Benecke 1998; Malgorn, Coquoz 1999; Stevens, Wall 2001; Wallman, Adams 2001; Wallman, Donnellan 2001; Wells et al. 2001; Wells, Pape, Sperling 2001; Stevens, Wall, Wells 2002; Harvey et al. 2003a; Harvey, Dadour, Gaudieri 2003; Harvey et al. 2003b; Chen, Hung, Shiao 2004; Zehner et al. 2004; Campobasso et al. 2005; Harvey 2005; Wallman, Leys, Hogendoorn 2005; Stevens et al. 2006; Nelson, Wallman, Dowton 2007; Wells, Wall, Stevens 2007; Wells, Williams 2007; Harvey et al. 2008; Nelson, Wallman, Dowton 2008; Wells, Stevens 2008; McDonagh et al. 2009; Park et al. 2009; Picard, Wells 2009; Mazzanti et al. 2010; Picard, Wells 2010; Meiklejohn, Wallman, Dowton 2011b; Singh, Kurahashi, Wells 2011).

While other techniques have been tried, a majority of these attempts to identify species utilize direct sequencing of mitochondrial DNA, usually cytochrome c oxidase subunit one (COI), to produce molecular-based phylogenies. COI is well studied throughout Arthropoda, and in particular Insecta (Hogg, Hebert 2004; Cywinska, Hunter, Hebert 2006; Meier et al. 2006; Smith et al. 2006; Costa et al. 2007; Nelson, Wallman, Dowton 2007; Smith et al. 2007; Burns et al. 2008; Foottit et al. 2008; Janzen et al. 2009; Zhou et al. 2009; Boehme et al. 2010; Ekrem, Stur, Hebert 2010; Park et al. 2010; Dewaard, Hebert, Humble 2011; Meiklejohn, Wallman, Dowton 2011b; Meiklejohn, Wallman, Dowton 2011a; Zaidi et al.

2011). The locus is used so much that the 5' end has been adopted by the DNA Barcoding Consortium as the standard for barcoding all of life (Hebert et al. 2003; Hebert, Ratnasingham, Dewaard 2003; Hebert, Gregory 2005; Meier et al. 2006). Unfortunately, the 5' end of COI does not seem serviceable to forensic entomology (Meier et al. 2006), as also indicated by our own Phase I results (below).

Previous studies in Diptera There is no established precedent as to which markers are most appropriate for discriminating among Diptera of forensic importance, even though a suitable genetic marker is essential for the success of any molecular identification or taxonomy study. Most studies of identification of this group have concentrated on the use of different fragments of the COI gene, including the full barcoding fragment (Wallman, Donnellan 2001; Wells, Pape, Sperling 2001; Wells, Sperling 2001; Stevens, Wall, Wells 2002; Harvey, Dadour, Gaudieri 2003; Harvey et al. 2003b; Stevens 2003; Chen, Hung, Shiao 2004; Nelson, Wallman, Dowton 2007; Caine et al. 2009), or, sometimes, even shorter fragments (<600bp) of the standard barcode region (Wells, Lunt, Villet 2004; Zehner et al. 2004) (Ames, Turner, Daniel 2006; Wells, Williams 2007) (Tourle, Downie, Villet 2009). Other genes that have been used for discriminating forensically important Diptera include the mitochondrially encoded genes COII (Wallman, Donnellan 2001; Wallman, Leys, Hogendoorn 2005) (Stevens, Wall, Wells 2002; Stevens 2003), NADH dehydrogenase 4 (ND4) (Wallman, Leys, Hogendoorn 2005), ND5 (Zehner et al. 2004), cytochrome oxidase b (cytb) (Vincent, Vian, Carlotti 2000) and the mitochondrial control region (Azeredo-Espin, Lessinger, Lessinger 2006), as well as the nuclear-encoded ribosomal genes 28S rRNA (Stevens 2001; Stevens, Wall, Wells 2002; Stevens 2003; Tourle, Downie, Villet 2009).

Methods for analysis of DNA sequence data for species identification. In the forensic context, the most commonly applied approach to DNA-based species identification is to perform a phylogenetic analysis using a known-identity reference collection (Wiens, Penkrot 2002; Morando, Avila, Sites 2003; Meier et al. 2006; Ross, Murugan, Li 2008; Ross et al. 2010; Berger, Krompass, Stamatakis 2011). An unknown sample is deemed to belong to whichever species it clusters with in that analysis. Because all reference individuals are included in the phylogenetic analysis, this approach naturally accommodates the variable nature of populations (provided, of course, that the reference collection includes multiple individuals per species). The phylogenetic method relies on the species in question exhibiting a pattern known as reciprocal monophyly (Wells, Wall, Stevens 2007; Wells, Williams 2007). If monophyly holds, then all the alleles found in individuals of one species will share an inferred common ancestor on the phylogenetic tree that is not shared by alleles possessed by a member of any different species (Felsenstein 2002).

Phylogenetic methods. A phylogenetic tree can be inferred either by a simple algorithm or by a method that is based on some optimality criterion, with those methods using optimality being widely regarded as superior (Felsenstein 2004). Those methods, in turn, can utilize either the original ("discrete") characters (e.g., the nucleotide states that comprise a set of DNA sequences) or a complete set of pair-wise distances. In this project, we use the distance approach (using the Minimum-Evolution method), because such methods are more efficient (i.e., able to support a strong inference despite having only a short segment of DNA sequence information; Kumar, Gadagkar 2000).

It is common practice in phylogenetic analysis to assess the reliability of a phylogenetic inference through a non-parametric resampling technique known as bootstrap resampling (Efron 1979), which was first applied to phylogenetics by (Felsenstein 1985). In

this method, the original data are repeatedly (1,000x, in our case) resampled with replacement to create a series of “pseudo-replicate” data sets. Each pseudo-replicate is then analyzed by the same method used in the original analysis. For any specific node on the final tree, the “Bootstrap percentage”, or BP, represents the fraction of the pseudo-replicates in which that same relationship was inferred. Interpretation of specific BP values is not straightforward (Hillis, Bull 1993). There is general agreement, however, that values >95% can be safely taken as indicating very strong support (Felsenstein, Kishino 1993).

Reciprocal monophyly in previously published databases The success at finding reciprocal monophyly among forensically important Diptera in previously published DNA databases is mixed. Among the recent, large-scale studies, Meiklejohn, Wallman, Dowton (2011b) found > 95% of Australian species of Sarcophagidae exhibit reciprocal monophyly, with, consequently, only a few cases of non-monophyly. Similarly, Harvey et al. (2008) found primarily reciprocal monophyly, with a few cases of paraphyly and a single pair of species for which the mitochondrial haplotypes were completely intermingled. In contrast, Wells, Wall, Stevens (2007) recently highlighted particular difficulties in DNA-based species identification for a number of species in the blow fly genus *Lucilia*. Members of this genus are frequently known by the common name “greenbottle flies,” and they include a number of species of substantial forensic importance. However, successful DNA-based identification may be problematic for the genus *Lucilia*. Wells et al. (2007) found reciprocal monophyly to be the exception, rather than the norm, within the subfamily that contains *Lucilia*, the Luciliinae. They analyzed Cytochrome oxidase I (COI) sequences from multiple individuals of five *Lucilia* species, and all five exhibited some form of non-monophyly. Wells et al. (2007) caution that the task of obtaining a reference data set that meets the criterion of having primarily or exclusively reciprocal monophyly may appear to be a simple task when the number of specimens examined per species is small, and that cases of non-monophyly might only become apparent when a large number of individuals are included for each species.

Geographic scope of the reference database. A reference database could either attempt to be global in scope, or be focused on a specific geographic region. Some recent papers describing DNA-based species identification of forensically important Diptera have taken each approach. Among those with a worldwide scope are Harvey et al. (2008), who generated some new data and combined that with existing data from GenBank to develop a worldwide reference database for the family Calliphoridae, and one paper that was a product of the present project (DeBry et al. 2010). The latter paper examined in detail the status of two very closely related and difficult to distinguish species, *Lucilia cuprina* and *L. sericata*. The samples generated from this project are all from the contiguous U.S., but additional data obtained from published sources were added to our own, in order to provide a global perspective on that specific issue. Other recent DNA-based studies adopt a more limited scope, often at the level of a country or, sometimes, a continent. These include studies focused on: Taiwan (Chen, Hung, Shiao 2004); Australia (Meiklejohn, Wallman, Dowton 2011b); (Tan et al. 2010), and the United States (Wells, Williams 2007)

Recent large-scale studies on DNA-based species identification in forensic entomology. A number of species-identification papers have been published on forensically important Diptera in the past several years. Of these, the largest in scope are discussed in the following paragraphs.

Harvey et al. (2008) used essentially the same segment of the COI gene as is used in the present study. They focused on the Calliphoridae, and obtained data from 42 individuals representing 22 species. They performed a phylogenetic analysis using Bayesian Inference.

Along with the 42 new sequences, they also included an additional 72 sequences that they obtained from GenBank. Altogether, they included 26 different species. Their study was worldwide in scope, with data coming from Australia, Africa, Europe, Asia, North and South America. They reported finding paraphyly for 4 of the 22 species in their study, while the remaining 18 species appear to be identifiable based on their COI sequence data. Only 7 of their 26 species occur in the contiguous U.S. and, thus, are included in the database presented here.

Meiklejohn, Wallman, Dowton (2011b) obtained approximately 850bp worth of sequence data from the 5' end of COI (the "barcode" fragment) for 16 species of Sarcophagidae from Australia. Because of their focus on the Australian fauna, only a single species, *Sarcophaga crassipalpis*, is in common between the Meiklejohn et al. data set and ours. They found that two of the 16 species formed widely disjunct, paraphyletic clusters, while the rest exhibited reciprocal monophyly.

One of the foundational papers in DNA-based species identification was the study on the blow fly subfamily Chrysomyinae by Wells, Sperling (2001), in which a region of COI was proposed for use in species identification. Recently, Wells, Williams (2007) published a follow-up study, in which they obtained a small amount of COI sequence data (304 bp) for nearly the complete North American chrysomyine fauna for large number of individuals (a total of 245 test sequences). Wells and Williams then tested each short sequence read, one at a time, in a phylogenetic analysis using the original Wells and Sperling (2001) data as the reference database. Even though the Wells and Sperling reference database consists of only a single reference specimen per species, Wells and Williams found that every short query sequence from all species but one paired with the correct reference specimen in a parsimony analysis with bootstrap support $\geq 95\%$, and sequences from the remaining species paired with its correct reference sequence with $\geq 92\%$ bootstrap support. The Wells and Williams study focused on the U.S. and Canada, and included one species that is not included in the current project. It is now recognized, however, that a proper reference database should include considerably more reference specimens per target species – and that is the approach we have adopted in the present work.

A series of papers has explored the utility of DNA-based species identification in the blow fly subfamily Luciliinae (which is represented in North America only by members of the genus *Lucilia*). Early reports (Stevens 2001; Stevens, Wall, Wells 2002) indicated an unusual pattern with respect to the very widespread species *L. cuprina* in terms of its relationship to the closely related and similarly widespread species *L. sericata*. Data from nuclear DNA showed reciprocal monophyly between the two species, but data from the mitochondrial genome indicated that *L. cuprina* is paraphyletic, with some *L. cuprina* being more closely related to *L. sericata* than they are to other *L. cuprina*. Stevens et al. (2002) concluded that the paraphyly probably did not pose a problem for DNA-based identification of the two species, however, for two reasons. First, those *L. cuprina* that were more closely related to *L. sericata* were, originally, found only in Hawaii. Second, the *sericata*-like *L. cuprina* did form a distinct cluster, such that an unknown specimen could, potentially, be identified as either of two types of *L. cuprina*, either of which is distinct from *L. sericata*. In 2007, however, Wells et al. (2007) published a "cautionary tale for forensic species identification." In that report, they found many (indeed, most) species of *Lucilia* they examined to be paraphyletic, with a correspondingly significant potential for misidentification. Further, the *sericata*-like *L. cuprina* were discovered to not be limited to Hawaii, as they were also to be found in Taiwan. Thus, the situation in *Lucilia* may be in stark contrast to that reported in the Chrysomyinae validation paper. Our results (DeBry et al. 2010) speak directly to this issue, and will be discussed in detail later in this report.

Existing publicly available DNA sequence data: The above-mentioned studies and others have resulted in a fairly large collection of DNA sequence data being deposited in GenBank for the Sarcophagidae and, especially, for the Calliphoridae. These data certainly have the potential to be used as part of a DNA-based identification system. However, we do not include those data in our reference collection or our database, for two related reasons. First, many of the sequences in GenBank are for flies that were collected outside of the United States. We do not yet have a good understanding of the levels of geographic variability in mtDNA sequence in these flies. Therefore, it would be very difficult to justify use of (for example) *Lucilia cuprina* data from Africa to help identify an unknown larval calliphorid from California. Second, we believe that a crucial and indispensable component of a DNA sequence database for use in species identification is the association between the DNA sequence and the original specimen (the “voucher specimen”). Morphological identification of the adult flies used in the reference collection should be performed by qualified experts, but few GenBank entries provide any information regarding the person who performed the species determination, and most entries provide no information regarding the current whereabouts or disposition of the specimen used. For this reason, we use as reference samples only those flies that we collected pursuant to this project. The particulars regarding morphological identification and disposition of the voucher specimens is discussed later in this report.

Choosing the best tool for the job of species identification.

The problem posed in species identification is fundamentally different from that of personal identification, and requires the correct molecular markers. The question we are addressing in this study (“to which species does this unknown larval specimen belong?”) is fundamentally different from the forensic question most frequently addressed with molecular markers (e.g., “did the specimen collected at a crime scene come from suspect X?”). In the latter case, with all but the most hyper-variable marker systems, there must be an exact match between the genetic profile of the specimen and that of suspect X (else, suspect X would already be excluded). The question really being asked in that context is “what is the probability that another individual, randomly chosen from the relevant population, would also be an exact match to the genetic profile of the specimen?” There are two key differences between that forensic application of genetic data and the application of genetic data to the problem of species identification. First, in the case of species identification there is no exclusion step. A mismatch between the genetic profile of the unknown specimen and the database of known individuals of a species cannot eliminate that known species from consideration, because the unknown sample may simply represent a genetic variant of that species that has not yet been incorporated into the database. Second, if the question was constructed to parallel the question posed in associating a suspect with a DNA specimen, it would be phrased as “what is the probability that an individual, randomly chosen from *a different species*, would also match the target species. With human personal identification, random matches have an extremely low probability because allele frequencies in the reference population are known with great precision. To achieve an equivalent discriminatory power for genotypes across so many species of flies would require an enormous sampling effort. For these reasons, I argue below that a phylogenetic analysis provides the most appropriate framework for DNA-based species identification.

“DNA Barcoding” is similar in spirit and execution to species identification in forensic entomology. The use of short DNA sequencing reads for species identification has become generally accepted within the broader scientific community. The so-called “DNA Barcode”

initiative has, as a main focus, the use of short (ca. 600bp) pieces of DNA sequence that allow placement of an unknown sample into a species (reviewed above). This idea, however, has its origin within the science of forensic entomology. The pioneering work of Wells, Pape, Sperling (2001) was based on the notion that larvae contain the same complement of DNA that the fly would have if it attained adulthood. This led to the insight that the DNA sequence from an unknown larva recovered from a crime scene could be identified to species by using a reference database of DNA sequences from a series of identified adult specimens.

Following the demonstration by Hebert et al. (2003) that even relatively short stretches of mtDNA are sufficient to place a query sample in the correct species, DNA-based species identification has spread quickly. That study marked the beginning of the Barcoding of Life initiative, which proposes the use of a ~600bp fragment near the 5' end of the mitochondrial cytochrome c oxidase subunit one (COI) gene for identifying animal life (Hebert et al. 2003a, b). The barcoding approach has been used successfully to identify species across a range of taxa (Hebert et al. 2004a; Hebert et al. 2004b; Ward et al. 2005; Elsasser et al. 2009; Robinson et al. 2009; Rougerie et al. 2009).

Standardization on a single barcode region is most appropriate for broad survey or inventory work, especially if the investigator is generating only the query sequences while relying on the existing public database for the reference samples. Forensic entomology differs from survey work in that, in any given geographic region, there exists a restricted and relatively well-studied fauna that the investigators need to identify to species. Morphological species identification for adults is generally well characterized in the forensically important species (see, for example, the morphological key to species in the family Calliphoridae of Whitworth 2010), even though morphological identification of larvae is typically not possible. For forensic entomology, the specific need (to correctly identify any larvae that might be encountered during the course of a death investigation) must take precedence over the general desire to contribute sequence data for the standardized barcode region. For that reason, Phase I of this project involved a survey of levels of inter- and intra-species variation for several regions of the mitochondrial genome

It is possible that the standard barcode region might be sufficient for identification of forensically important Diptera in the contiguous U.S., but that must first be demonstrated. Successful species identification requires a locus at which inter-specific divergence substantially exceeds, and does not overlap with, intra-species variation. Roe, Sperling (2007) examined sliding 600bp windows across the entire COI plus COII region in several species of Lepidoptera and Diptera. They found that the region of maximal utility for identification varied, and would likely need to be established separately for any particular study. Consequently, in this study we considered several pairs of closely related species of forensic importance in the contiguous United States, including one pair, *Lucilia cuprina* and *L. sericata*, that is known to be very difficult to distinguish using mtDNA (Wells, Wall, Stevens 2007).

The mitochondrial genome is expected to provide the best data for robust species identification. It is certain that the best approach for using DNA sequence data for accurate species identification would be to obtain data from multiple markers spread across both the nuclear and mitochondrial genomes. Practical limitations instead dictate that crime lab practitioners are unlikely to amplify and sequence anything beyond a single PCR amplicon of modest size. Therefore, it becomes important to identify a marker that will be expected to give the most reliable results from a single amplicon of not much more than 1 kb in size.

Regardless of the source of genetic markers, the biggest barrier to correct species identification occurs when two (or more) species share alleles. Such a pattern could occur in at least two ways. First, closely related species might retain ancestral alleles. In other words, a

potential indicator locus was, of course, homogeneous in the past, when the two species were a single, ancestral species. In such a case, the genetic marker will not discriminate those two species. Species identification can only be accomplished once sufficient time has passed such that new alleles become fixed in one or both descendant species. The rate at which alleles are fixed by genetic drift is a function of the effective population size, N_e , with smaller N_e resulting in more rapid fixation and, therefore, greater power to discriminate closely related species. mtDNA is a haploid genetic system, so its N_e is approximately $\frac{1}{4}$ that of the diploid nuclear genome in the same species (Moore, 1995). It has long been recognized that, on this basis, mtDNA data are superior to nuclear markers for phylogenetic analysis at the species level (Moore, 1995). A second circumstance that can cause distinct species to share one or more alleles is hybridization. Unfortunately, mtDNA is more prone to crossing the species barrier in cases of limited hybridization than is the nuclear genome. ***unfortunate, but does not outweigh the benefits of using mtDNA for species identification.

1. **Statement of rationale for the research:**

The rationale for this research follows from a logical progression through the following steps:

- 1) Flies are usually the first visitors to arrive at exposed human remains;
- 2) Flies lay eggs (Calliphoridae, Muscidae) or deposit 1st-instar larvae (Sarcophagidae), the larvae feed on the remains;
- 3) A minimum Post-Mortem Interval (PMI) can be estimated from the developmental stage of a larva, with the oldest specimen found being most useful because its age will be closest to the true PMI;
- 4) Different species develop at different rates, even under identical environmental conditions;
- 5) Therefore, a necessary first step in using entomological evidence in a death investigation is to establish the species identity of the sampled larvae;
- 6) Immature (larval) specimens of many forensically important dipteran species cannot reliably be assigned to the correct species on the basis of morphology;
- 7) Therefore, there is a need for DNA-based identification of forensically important Diptera;
- 8) DNA-based species identification can only be as good as is the reference collection of morphologically identified samples upon which it is based;
- 9) Therefore, we undertook to collect, identify, and obtain DNA sequence data from as many different species of forensically important flies that occur in the contiguous United States as possible. This allowed us to construct what is, by far, the most comprehensive reference database for use in DNA-based species identification of forensically important Diptera.

II. **Methods**

Collection of specimens

The following major collecting trips were undertaken through the course of this project: 2006: Florida; Georgia/South Carolina/Tennessee; Oregon;

2007: Wisconsin/Minnesota; New York; Oregon/Washington/some in California; Texas/New Mexico.

Each collecting trip included at least one week of active collection of flies. Additional specimens were caught using similar methods on shorter, local trips. A small number of specimens were donated by outside individuals.

Flies were collected either by net or trapped with the aid of meat bait (aged chicken thigh and liver). Collected specimens were frozen and taken to the lab for morphological identification, and for molecular analysis. To maintain all morphological structures important for species identification in the voucher specimens, only legs were utilized as the source of specimen DNA. Three legs were removed from each specimen and transferred to a vial of 95% EtOH and stored at -80°C until DNA extractions. The remainder of the specimen was retained as a voucher. Voucher specimens were either pinned or placed in a separate vial of ethanol and stored at -80°C.

All specimens identified and potentially available for DNA sequence analysis in this project are given identification codes (Table 3) maintained as physical voucher specimens. These are available for further study or for confirmation of identifications, if necessary. In the case of specimens for which DNA sequence data are reported (Table 4), 1-3 legs have been removed for DNA analysis. In a very few cases, some abdominal material has been removed for DNA extraction. In those cases, the terminalia have been preserved, as nearly all species in this study have definitive morphological characters in the male genitalia. Pinned specimens are stored in standard museum cases at room temperature. When necessary, additional legs can be removed from a pinned specimen for subsequent DNA extraction. Specimens in alcohol are stored in 1.5 ml snap-cap tubes at -20° C. All alcohol-stored specimens include within the storage tube an embossed label giving the project identification code. Most of the alcohol-stored specimens for which DNA sequence data are reported have also had 1-3 legs removed. A small number may have had portions of the abdomen removed for DNA extraction. All post-preparation manipulations of specimens (removal of legs or abdomen for DNA analysis) was conducted using sterile, UV-irradiated forceps and scalpels.

Identification of specimens. All specimens used to develop the reference database were adults, and were identified to species using standard morphological characteristics. All identifications were performed by Dr. Greg Dahlem of Northern Kentucky University. For the Calliphoridae, the reference work used to determine specific characteristics was the key from Whitworth (2006).

Choice of study species – A relatively small number of dipteran species are both common (sometimes regionally) and well known for being of forensic importance (see Table 1). However, there is a much larger suite of species that *might* be encountered in the forensic setting, either because the adults utilize carrion as a protein source or simply as accidental captures. Trying to query a sequence database with a species that is not represented in the reference collection will produce unpredictable and, certainly, unreliable results. Thus, it is important that the reference database include not only the species of direct forensic importance, but also a good representation of related species. We surveyed the existing literature on which species of Diptera have the potential to be associated with large-animal carrion, as well as species that had been specifically mentioned in the forensic literature. This review was complicated by numerous changes over the years in taxonomic nomenclature, by some cases of likely misidentification, and by the anecdotal nature of some of the literature. We took a generally inclusive approach (as there is little harm in including an irrelevant species in the reference database), and developed the species list given in Table 1 as our target for the database. In total, we identified 40 species as being of at least potential forensic importance.

Rationale for using only specimens collected during his project and DNA sequences generated directly by this project – There exist a substantial number of DNA sequences in GenBank for some of the species in Table 1. However, we feel that it is vitally important for the reference database produced through this project to have the physical voucher specimen available for future study, should there ever arise the need to confirm the morphological species identification, or should future taxonomic changes require re-examination of the original voucher specimens. We therefore decided to include in the reference database only specimens that we collected ourselves (or, in a handful of cases, for which the specimen and associated information was provided by a collaborator, and for which we retain the physical specimen). As all of our collecting effort occurred within the 48 contiguous United States, our database is only directly relevant to that geographic area.

Molecular methods

DNA extractions were performed sometimes on abdomens, but primarily on individual legs. One of two methods was used. The first used the QIAGEN DNeasy kit, and followed the manufacturer's protocol. QIAGEN extraction was used for all abdomen DNA extractions. QIAGEN-extracted DNA was resuspended in ~50ul molecular grade (DNase- and RNase-free) water (hereafter referred to simply as "water") and stored at either 4° C or -20° C or -80° C. The second method used was sonication. Sonication is non-destructive, so the leg can be retained for further DNA extraction, which, if necessary, could be by another method. Individual legs were washed 3x in water, then placed in a 1.5 ml snap-cap tube in 30ul water. This tube was placed in a Branson model 1510 sonicating water bath, and exposed to ultrasonic vibrations for 3 bouts of 30 seconds each. The leg was then removed and placed back in alcohol for storage. The water was subsequently used, as-is, as the genomic DNA template in subsequent PCR amplification reactions. Extracted DNA is stored in water at -80° C.

PCR and collection of DNA sequence data: Phase I - The goal of Phase I was to identify a region of the mtDNA protein-coding genome that provided the best discrimination among the forensically important species. We began with preliminary analyses that examined data from COI, COII and ND4. These data showed that all regions were roughly equally suitable for most species pairs, and that a region of COI provided adequate discrimination for an extremely difficult species pair (see Results), so other regions of the mitochondrial genome were not used in this project.

DNA extractions were performed using a single fly leg, cut in half, and then subjected to DNA extraction using QIAGEN's DNeasy kit. A fragment of approximately 2.3kb, comprising tRNA tyrosine, the cytochrome oxidase (CO) I gene, tRNA leucine, the COII gene and tRNA lysine, was amplified. This fragment spans nucleotides 1460 to 3775 in the *Drosophila yakuba* mitochondrial genome (Clary & Wolstenholme 1985), which was used for numbering nucleotide positions. Amplification was performed using four primer pairs: TY-J-1460/C1-N-2191, C1-J-2183/C2-N-3014, C1-J-2792/C2-N-3389 and TL-J-3043/3775 (Simon et al. 1994). In addition, a 690bp fragment of the ND4 gene, was amplified using primers N4-J-8502/H3 (Simon et al. 1994; Yu et al. 1999). Amplification reactions were performed in a total volume ranging between 10-50µl and contained 200µM of each dNTP, 1.5mM MgCl₂, 1µM of each primer, 0.25 units Platinum *Pfx* DNA polymerase (Invitrogen), 10x buffer (Idaho Technology) and 1/10th volume of template DNA. Thermocycling profiles were as follows: 95°C for 3 min, followed by 35 cycles of 94°C for 15s, 53°C for 15s and 72°C for 30s. A final extension period of 72°C for 5min completed the reaction. Products were visualized with UV trans-illumination using 1.25% agarose gels. PCR products were purified

using the QIAquick PCR purification kit (QIAGEN). Both forward and reverse strands were sequenced directly with the amplification primers using BigDye chemistry (PE Applied Biosystems) on an automated ABI 3730 XL instrument. Cycle sequencing was performed by the High Throughput Genomics Unit at the Department of Genome Sciences, University of Washington. For each specimen, an initial 10ul PCR reaction was run with the appropriate primer pair. This reaction was subjected to agarose gel electrophoresis in the presence of ethidium bromide, and subsequently visualized by exposure to UV light. A size standard was included, and the correct size of the PCR product was confirmed in each case. If the initial small-scale PCR reaction was successful, then the reaction was scaled up to 50ul total. 1ul of this reaction was visualized as above

Sequences were edited manually by confirming the identity of all polymorphic bases using the original chromatograms in BioEdit version 7.0.9.0 (Hall 1999). There were no indel differences within the protein-coding regions (tRNA regions were discarded prior to phylogenetic analysis), so sequences were aligned by eye and using the ClustalW accessory application in BioEdit.

A number of different data sets, based on the COI (*Dy#* 1460-3010), COII (*Dy#* 3100-3760) and/or ND4 (*Dy#* 8502-9194) fragments, were analyzed. The total data set consisted of the three concatenated genes, and each gene was considered individually. In addition, the three most commonly analyzed regions of the COI gene (the barcoding fragment *Dy#* 1490-2198, as well as fragments *Dy#* 2183-3014 and *Dy#* 2792-3389), were considered. The COI-COII region was also examined as a single data set and within this region, sliding windows, consisting of 600bp, were analyzed. Window sizes of 600bp were chosen to allow comparison with the ~600bp barcoding fragment. Each sliding window was referred to by its nucleotide midpoint.

Nucleotide substitution patterns were analyzed using DNAsp 4.90.1 (Rozas et al. 2003), using default settings. To calculate variability within species, nucleotide diversity (p) (Nei, 1987), was estimated. Variability between sister-species pairs was quantified as the nucleotide divergence K (Tajima 1983). A Jukes-Cantor correction was used for both estimates. For sliding window analysis, a step size of 5bp was used to analyze nucleotide substitution patterns. Nucleotide divergence was also expressed relative to total COI-COII divergence to facilitate comparisons among all sister pairs. To determine the effect of sample sizes on nucleotide diversity, regression analysis was performed using SPSS for Windows version 11.

Phase II - We generated our database using the primer pair *Dy#* 1751 / *Dy#* 3014, which amplifies a segment comprising approximately 1,100 bp in the 5' half of COI. PCR protocols were, otherwise, identical to those used for Phase I. After trimming the ends of the sequencing reads, the region used to construct the final Phase II reference database is referred to as "1800-3000".

Alignment - No insertion/deletion polymorphisms were detected for the "Phase II region" during this project, so multiple sequence alignment did not present a technical challenge. Alignment was accomplished using the manual sequence editor program BioEdit (Hall 1999). Forward and reverse sequencing reactions for an individual PCR product were first aligned to each other and combined to form a consensus sequence for that amplicon. Any apparent discrepancies between the forward and reverse reads were reconciled by reference to the original chromatogram trace files. Any remaining disagreements between the forward and reverse reads were resolved by re-sequencing.

The final consensus sequence for each PCR product was then aligned to the complete mtDNA genome sequence for *Lucilia sericata* (GenBank [NC_009733](#)) which, in turn, we had

aligned to the complete mtDNA sequence of *Drosophila yakuba* (GenBank NC 001322). All sequence position numbering is referenced back to the *D. yakuba* sequence. The final, complete alignment is maintained in BioEdit, and exported in FASTA format for entry the phylogenetic analysis program MEGA 4.0.

Phylogenetic analyses – Due to the large size of the database, and the fact that the database contains many species pairs that are expected to have rather low levels of sequence divergence, we chose to perform our primary phylogenetic analyses using a method based on pair-wise genetic distances. We used the Tamura-Nei 1993 (TN93) (Tamura, Nei 1993) distance correction, to account for unobserved substitution events. TN93 is a 3-parameter model. One parameter is used to account for all transversion substitutions, and two transition parameters separately account for A-G and C-T transitions. Pair-wise distances were calculated in the computer program MEGA 4.0 (Tamura et al. 2007), using pair-wise deletion to handle missing data. The phylogenetic tree was inferred using the Minimum Evolution (ME) approach. This method begins by applying the Neighbor-Joining star-decomposition algorithm, followed by branch swapping. The tree found that imputes the least overall evolutionary changes across the entire tree is chosen as the point estimate of the phylogeny. The ME method was chosen because it is expected to be among the most efficient methods available for closely related sequences (Nei 1991). The robustness of each node on the phylogeny was assessed by bootstrap resampling (Efron 1979; Felsenstein 1985), using 1000 pseudo-replicates.

III. Results

1. Statement of Results

Phase I

We examined 600bp sliding windows across the full COI-COII region, plus the approximately 600bp ND4 gene, and assessed each window's suitability for robust species identification for these species. For each window, we performed a number of analyses to characterize levels of within-species variability and between-species divergence.

The main result of Phase I is that any region is suitable for distinguishing most of the species pairs in the Phase I data set, but only a ca. 1100 bp region near the 3' end of the COI gene is able to provide robust discrimination between *Lucilia cuprina* and *L. sericata* (Fig. 1). We denoted the region of high discrimination as 1800-3000, in accordance with the approximate locations of the forward and reverse PCR primers, numbered according to the *Drosophila yakuba* mitochondrial genome numbering system. The 1800-3000 fragment is able to distinguish between *L. cuprina* and *L. sericata* with bootstrap values > 95%. At this point, we could have continued to examine additional mitochondrial loci. However, it was apparent that other loci would, at best, be equivalent to the 1800-3000 fragment. We felt, at that point, that it would be a more efficient to proceed to Phase II. Consequently, the 1800-3000 region, was chosen for use in Phase II of this project.

Detailed Results from Phase I

Intraspecific diversity

Since levels of interspecific divergence may be dependent on sample sizes, regression analysis was performed to determine their relationship. Intraspecific divergence was not significantly related to species sample sizes for any of the three genes analyzed or the total data set (COI: $R^2 = 0.055$, $p = 0.575$, COII: $R^2 = 0.042$, $p = 0.628$, ND4: $R^2 = 0.054$, $p = 0.580$, total data set: $R^2 = 0.052$, $p = 0.588$).

Nucleotide diversity for the total data set consisting of concatenated COI, COII and ND4 sequences was calculated as 7.877% (Table 5), with intraspecific diversity estimates ranging between 0.090 (*L. sericata*) and 1.804 (*S. bullata*). Intraspecific nucleotide diversity was consistently highest within *S. bullata* for all regions and genes analyzed. However, minimum intraspecific diversity varied with the region and genes analyzed. Of the three genes analyzed, COI yielded the highest level of total intraspecific nucleotide diversity (8.407%), followed by COII (8.028%). The ND4 gene displayed the lowest level of diversity (7.752%). This pattern of diversity based on individual genes was evident within most species, with COI gene outperforming both the COII and ND4 genes for intraspecific diversity calculations. Nucleotide diversity within the COI gene relative to that of the concatenated COI, COII and ND4 data set was greater than 100% for all species analyzed. For the ND4 gene, only three species displayed mean nucleotide diversity levels greater than that of the total data set. Relative nucleotide diversity within the COII gene was less than 100% than that of the total data set for all species.

Of the three regions of the COI gene most commonly sequenced, the “2183” fragment provided the highest level of nucleotide divergence (8.806%, Table 6), followed by the barcoding and “2792” fragments (7.982% and 7.076% respectively). This pattern of diversity for the three fragments was reflected by intraspecific nucleotide diversity values for individual species, where the “2183” fragment provided the highest and the “2792” fragment the lowest values for most species (Table 6). Thus, it is evident that the barcoding fragment is not the most suitable fragment of the COI gene for determining maximum levels of intraspecific nucleotide diversity.

Sliding window profiles (600bp) based on the COI-COII region are illustrated in Fig. 2. Profiles differed substantially among the seven species analyzed. Maximum nucleotide diversity among 600bp windows varied between 0.222% (*L. sericata* and *S. africa*) and 2.556% (*S. bullata*) (Fig. 2). The location of maximum diversity was variable among species, with *S. bullata* and *C. vomitoria* displaying multiple regions of maximum diversity (Fig. 2). Minimum nucleotide diversity was 0 in four of the seven species analyzed (*C. vicina*, *L. cuprina*, *L. sericata* and *S. africa*). Locations of minimum diversity for most species occurred most frequently at the 3' end of the COI gene, although ranges of maximum diversity for most species were more widely spread. Thus, different regions of the COI and COII genes are more efficient at determining intraspecific nucleotide diversity of each of the species analyzed.

Interspecific diversity

The ultimate discrimination among closely related species in this project will come from demonstration of reciprocal monophyly. However, it is also informative to examine interspecific divergence compared to intraspecific divergence across sliding windows, as a guide to which genomic regions may be most suitable for discrimination. Maximum intra- and

minimum interspecific diversity was examined across the COI-COII region for all species investigated. Since these divergence values did not overlap for any sister pair of species included in the Phase I analysis (Tables 5 and 7).

Interspecific nucleotide diversity for five sister species pairs was extremely variable, ranging between 0.900% (*L. sericata* and *L. cuprina*) and 9.274% (*S. africa* and *S. bullata*) (Table 7). Nucleotide diversity was consistently highest between *S. africa* and *S. bullata* and lowest between *L. sericata* and *L. cuprina*, regardless of the region analyzed or primer pair used. The especially close relationship between *L. sericata* and *L. cuprina* is in agreement with that of previous studies, where a lineage of *L. cuprina* has been found to be paraphyletic with regard to *L. sericata* (cit).

For all sister species pairs, nucleotide diversity within the COI gene relative to that of the total COI-COII-ND4 data set was greater than 100%. Although the COI gene yielded the highest level of nucleotide diversity (5.349%), it was closely followed by that of the ND4 gene (5.314%). The COII gene yielded a lower level of diversity (4.236%). These results confirm those of intraspecific diversity, where the COI gene appears to be more efficient than the COII and ND4 genes at detecting maximum levels of nucleotide divergence. For the commonly sequenced COI regions, the 2183 fragment once again provided the highest level of diversity, followed by the barcoding and 2792 fragments (Table 8). These results indicate once again that the barcoding fragment is not ideal for identifying forensically important Diptera.

Sliding window profiles of species pairs were less variable than those of individual species profiles and many were remarkably similar (Fig. 3), such as those between *L. sericata* and *L. illustris* and *L. illustris* and *L. cuprina*. The location of minimum and maximum diversity for each of the five species pairs occurred within a fairly narrow region (midpoints Dy# 2971-3275 and Dy# 2371-2555 respectively). It is thus possible to pinpoint midpoints Dy# 2371-2555 of the COI gene as being most suitable for discrimination of the eight species included in analyses. This pattern of diversity was also evident in the sliding window profile of mean relative divergence for all five species pairs (Fig. 4). From variance analyses, it appears that the optimal fragment of a 600bp window of the COI-COII region appears to be at midpoint Dy# 2500 (positions Dy# 2200-2800). This region encompasses much of the maximum levels of interspecific divergence as well as diversity within the 2183 fragment (midpoint Dy# 2598).

Phylogenetic analysis for species discrimination

Relationships among the seven species analyzed, based on concatenated sequences of the COI, COII and ND4 genes, are shown in Fig. 5. The tree topology reflects that of accepted phylogenies, with bootstrap analysis supporting the majority of internal nodes, and species from the same genus forming monophyletic clusters with 100% bootstrap support. All species of Calliphoridae and Sarcophagidae clustered together within their respective families. Within the genus *Lucilia*, *L. cuprina* and *L. sericata* appeared to be more closely related to each other than to *L. illustris*, which confirms previous results (Wells, Williams 2007). Analysis of individual COI, COII and ND4 genes each provided the same basic tree topology. Genetic distances were similar for all three genes as well as the COI-COII and COI-COII-ND4 regions (Table 9). However, for the COI, COII and ND4 genes bootstrap support for *L. cuprina* and *L. sericata* decreased to 96/100, 96/98 and 69/95 respectively. The number of informative characters varied with the region analyzed (Fig. 6), with the largest

numbers of informative characters appearing towards the 3' end of the COI gene (midpoints *Dy*# 2460-2860). These results are in agreement with those of nucleotide divergence, where this region appeared most suitable for species discrimination.

Each 600bp window of the COI-COII region was able to discriminate between species, providing 100% bootstrap support for *C. vicina*, *C. vomitoria*, *L. illustris*, *S. africa*, *S. bullata* and *P. regina* (not shown). However, bootstrap support for *L. cuprina* and *L. sericata* varied widely among different windows (Fig. 1), with support for *L. cuprina* as low as 62%. It is thus clear that no 600bp fragment of the COI-COII region is sufficient to provide robust identification of *L. cuprina* and *L. sericata*. Therefore, it may be likely that fragment length must be increased when examining species that are very closely related.

Fragment length

The effect of fragment length on nucleotide divergence and phylogeny was investigated by examining fragments with 200bp increments, starting at the 5' end of the COI gene, the 3' end of the total data set and the midpoint of the total data set. Patterns of mean relative divergence were rather dissimilar for the three starting locations (Fig. 7). Starting at the 5' end of COI, the initial mean relative divergence at 200bp was 90%, decreased slightly to 88% at 400bp and increased to steadily to 100% at 1200bp. Relative divergence peaked at 105% at 1400bp and remained at about 100% for increased lengths. Starting at the 3' end of ND4, divergence was highest at 200bp (120%), indicating that a region of high divergence was being sampled. Divergence decreased to 100% at 600bp and decreased further to its lowest point at 95% at 800bp before increasing to 100% at 1000 and 1200bp. Diversity decreased to 96% at 1400bp and 1600bp, remained steady at 100%-102% thereafter. Starting at the midpoint of the COI-COII-ND4, initial divergence was 80% and decreased further to 75% at 400bp. Divergence increased to 100% at 1000bp and peaked at 105% at 1400bp and 1800bp. These results show that fragment lengths of at least 1000bp are necessary to ensure that close to 100% total diversity is sampled.

Bootstrap support for phylogenetic analysis based on the Minimum-Evolution algorithm was used to evaluate the effect of fragment length on monophyly support for each species. A 200bp fragment was sufficient to resolve many of the terminal nodes in the tree, with bootstrap support above 90% for most species, although phylogenetic signal was decreased (Fig. 8). However, these short fragments could not distinguish between *L. cuprina* and *L. sericata* at any of the three starting points or between *S. africa* and *S. bullata* at the 5' starting end. For most species, a fragment length of 400bp was sufficient to yield a bootstrap support of 100%. These results show why studies using even short fragments of DNA have been so successful. The exceptions were *L. cuprina* and *L. sericata*, where bootstrap values were more dependent on fragment length. For *L. sericata* bootstrap support for all three starting points stabilized at 100% between 800bp and 1000bp. For *L. cuprina* fragment starting point had a remarkable effect on initial bootstrap support (Fig. 9). At the 5' end of the COI gene, bootstrap support was only 36% at 200bp, increased sharply to 66% at 400 and thereafter rose steadily to 100% at 2000bp, after which bootstrap values flattened off at 100%. At the 3' end of the ND4 gene, bootstrap support for *L. cuprina* monophyly increased sharply from 66% to 95% between 200bp and 400bp and increased to remain steady at 98%-100% at 800bp until the entire COI-COII-ND4 fragment was included. Starting at the midpoint of the COI-COII-ND4 data set, the bootstrap value at 200bp was 57%, which increased to 60% at 400bp and rose sharply to 97% at 600bp, where it remained steady until 1000bp, after which values decreased dramatically to 86% at 1200bp before climbing to 100% at 1800bp, after which values remained steady. Thus, it is clear that a minimum fragment length of 1000bp is needed to ensure an unambiguous phylogenetic signal for both *L. sericata* and *L. cuprina*.

The use of longer sequences is often impractical in large studies (and for degraded samples, such as might be encountered in casework). However, a sequence length of 1000bp can usually be obtained in a single reaction. Therefore, we constructed the reference DNA database for this project by extending the optimal 600bp fragment (*Dy#* 2200-2800 of the COI gene) by 400bp towards the 5' end of the COI gene and 200bp in the 3' direction (to positions *Dy#* 1800-3000), to ensure greater diversity coverage. Further tests might be conducted in the future, assessing the ability to place shorter unknown sequences into the phylogenetic framework of the full reference database.

Phase II

The primary result from this project is the production of a reference database of COI DNA sequences for the 1800-3000 region comprising 504 individuals from a total of 106 species. This database can be used in a phylogenetic analysis to infer species-level identification of unknown dipteran larvae of forensic significance.

Table 4 provides a listing of every specimen for which we obtained DNA sequence data for the Phase II locus. In total, we include in the reference database 504 individuals representing 106 species. The database includes data for 38 of the 40 species that which we determined from the literature to be of likely forensic significance.

Forensically important species for which we have no data: We were unable to obtain samples for the following 5 species: *Comptosyriops callipes* – this species can be common in

parts of the southwestern U.S., but we did not encounter any during our collecting efforts, and collections from southern California sent to us by a colleague did not include any specimens of this species. Addition of *Comptosyiops callipes* to the reference database should be a high priority going forward. *Lucilia eximia* – this species is common in Central and South America; it is reported as “rare” in Texas, Oklahoma and Florida (Whitworth 2006), but we did not collect any despite extensive collecting in Texas and Florida; *Sarcophaga melanura* – this species is simply very uncommon, and we have not been successful at collecting any (and, presumably, its rarity reflects a very low probability that this species will turn up in a forensic investigation); *Sarcophaga ruficornis* – this species is common in Hawaii, but rare in the western contiguous U.S., and we did not collect in Hawaii; *Cynomya mortuorum* – this species is found only north of the Arctic Circle (Whitworth 2006), another region where we did no collecting.

Species not of forensic interest that are included in the database: In addition to the 36 species of highest forensic importance, we include another 68 species, albeit with fewer individuals per species. We feel that this is an extremely important aspect of the construction of a reference database for DNA-based species identification, but that it is an issue that has not been discussed previously in the forensic entomology literature. It is important that closely related, but non-target species be present in a reference database for species identification, because the phylogenetic analysis methods we use to query the database will place any sample on the tree, even if the unknown is an individual from a species that is not represented in the reference database. It is difficult to predict the behavior of a query sample for which there is not a corresponding reference sample in the database. One would hope that the query sequence would not join with a forensically important species to form a monophyletic group with strong bootstrap support, but such a result is certainly not out of the question. By including a large number of non-target species, all of which were collected during the same collection trips and using the same methodologies, we expect to have included most of the species that might be encountered as “accidentals” already present in the reference database. A final reason for inclusion of the 68 “not forensically important” species is that many of these specimens were collected at baited traps or by sweep netting near a baited trap. Consequently, at least some of those species could, potentially, be of unrecognized forensic utility.

Phylogenetic analysis

The reference database developed in Phase II is the fundamental deliverable of this project; those data are presented in Table 10. The full, bootstrapped phylogenetic tree for the complete database is shown in Fig. 10. The most important feature of the tree overall is that the large majority of species are reciprocally monophyletic vs. all other species with high bootstrap values. Specifically, we include multiple individuals for 57 species. 50 of those are monophyletic with high bootstrap values.

Of the 18 species that are of greatest forensic interest, only two cannot be distinguished from other forensically important species, based on the DNA segment used in our reference database.

The main result from this project is the production of a DNA database that can be used to successfully identify 89% (16 of 18) of the species of carrion flies most likely to be encountered in the forensic setting in the contiguous United States.

Calliphoridae Within the blow flies (Calliphoridae), we find that 23 of the 25 species for which we obtained samples are distinctly different from all other species, as indicated by ME bootstrap. Only a single species pair within the Calliphoridae (*Lucilia coeruleiviridis* – *L. mexicana*) cannot be identified using our database, based on the results obtained in this project.

The Calliphorid genus *Lucilia* deserves particularly close attention, because Wells et al. (2007) suggested that it might be difficult, or even impossible, to identify many species of *Lucilia* using only DNA data from a relatively small region of the mitochondrial genome.

Lucilia coeruleiviridis – *L. mexicana*: *L. coeruleiviridis* is a very commonly encountered and forensically important fly throughout much of the United States. *L. mexicana* is somewhat more restricted in its distribution, being found primarily in the Southwestern United States. These two species have a range of overlap running through much of Texas and New Mexico (and, possibly, Arizona). Our samples from the region of overlap are limited to several collecting localities in New Mexico and Texas. We sequenced the Phase II region for a large number of each of these two species, and found nearly complete overlap in their haplotypes. Some of our data are consistent with the possibility that the overlap in haplotypes may be a result of hybridization between the two species, followed by introgression of the *L. coeruleiviridis* mitochondrial genome into *L. mexicana*. Although closely related, these two species are not too difficult to distinguish on the basis of adult morphology, and our collections from New Mexico and Texas do not show signs of being morphologically intermediate. Even more interesting is the observation that the mtDNA sequences from *L. mexicana* collected in southern California are distinctly different from those of both *L. mexicana* and *L. coeruleiviridis* from New Mexico and Texas, with an average pairwise difference of 5%. In contrast, all of our samples of *L. coeruleiviridis* from throughout its range are very similar genetically to those found in the region of overlap, with a maximum pairwise divergence of only 0.5%. This would be consistent with a scenario in which the haplotypes we found for *L. mexicana* in southern California represent the original *L. mexicana* mtDNA lineage, while the *L. mexicana* in New Mexico possess introgressed *L. coeruleiviridis* mtDNA. The inability to distinguish between these two species in parts of their common range should be further investigated. In particular, growth studies should be undertaken in order to assess whether or not there are differences in developmental timing in relation to temperature between the two species in regions where the mtDNA sequences cannot be used to correctly identify the species. If there turn out to be no significant developmental differences, then it will be sufficient to use mtDNA evidence to identify larvae as “either *L. mexicana* or *L. coeruleiviridis*.” However, if there are developmental differences, then, when at least one of these two species is involved in their range of overlap, either forensic evidence must include traditional rearing of larvae to adulthood for morphological identification, or, if possible, other species found at the scene that could be positively identified on the basis of DNA sequence should be used for estimation of PMI.

L. cuprina and *L. sericata* – As reported in a publication based on this project (DeBry et al. 2010), data from the 5' end of COI are sufficient to discriminate between *L. cuprina* and *L. sericata* in a phylogenetic analysis. In the fully inclusive analysis, we find *L. cuprina* and *L. sericata* to be reciprocally monophyletic, based on bootstrap values >95%. There is no doubt, however, that these two species have mtDNA haplotypes that are very similar to each other. We see no evidence for geographic variation within either *L. cuprina* or *L. sericata* across the

contiguous United States. Average intra-species pairwise divergences are low for both species (Table 11), but the average inter-species divergence is also low. For this reason, further validation is warranted to determine whether or not hitherto undiscovered, divergent haplotypes exist that would compromise the discriminatory power of this gene region for these two species.

L. elongata, *L. silvarum*, and *L. thatuna* – These three species appear to be the most closely related to *L. cuprina* and *L. sericata*, although deeper phylogenetic relationships cannot reliably be inferred from a limited sample of a single genetic locus. We were only able to collect *L. elongata* from two localities in Oregon, while our only two specimens of *L. thatuna* come from a single locality in northern California. In keeping with our collecting success, Whitworth (2010) characterized each as being either rare or uncommon. Consequently, neither species is likely to be encountered commonly in forensic investigations. Nevertheless, it is important that all reference specimens available be included in any phylogenetically based species identification procedure. *L. silvarum*, in contrast, is characterized by Whitworth (2010) as being a common, widespread species, and we collected it from a number of widely separated localities. All three of these species are well differentiated from each other and from all other species, and each is strongly monophyletic in our bootstrap resampling analysis.

L. illustris – In some parts of the world, *L. illustris* cannot be distinguished from *L. caesar* using mtDNA data (Wells, Wall, Stevens 2007). In the United States, however, *L. caesar* does not occur, and *L. illustris* is easily distinguished from all other *Lucilia* based on DNA sequence data from the 5' end of COI. We see no evidence that there is any geographic variation within *L. illustris*, and the average pairwise intraspecies divergence is very low (0.5%; Table 11).

L. coeruleiviridis and *L. mexicana* – A large number of *L. mexicana* individuals carry a COI haplotype that is either identical to or extremely similar to haplotypes found in *L. coeruleiviridis* (Fig.10). Consequently, these two species cannot be distinguished using this locus. We do see substantial geographic variation within *L. mexicana*. All of the sequences with the *L. coeruleiviridis*-like COI haplotypes were collected in either Texas or New Mexico. In contrast, all of the individuals we sampled from California form a distinctly separate clade in the phylogenetic analysis (Fig. 10). The average pairwise divergence between the California and the Texas/New Mexico populations (5%) also substantially exceeds the level of intraspecies divergence observed for any other *Lucilia* species. We did examine a different mtDNA fragment (from the ND4 gene) for a limited number of individuals of both *L. coeruleiviridis* and *L. mexicana* from New Mexico. Those data also failed to differentiate between the two species (data not shown). While there are a number of possible explanations for the pattern we observed, perhaps the most likely is that there has been some hybridization between *L. coeruleiviridis* and *L. mexicana*, either ongoing or in the relatively recent past, somewhere in their region of geographic overlap. Subsequently, *L. coeruleiviridis*-like mtDNA has been introgressing into *L. mexicana*. Because the two species maintain their distinctive morphological characteristics across this region, it would appear that nuclear alleles are not crossing the species barrier, although we do not have any nuclear data with which to test that hypothesis. It is not clear what will be the consequence of this pattern for forensic purposes. If the two species were to be found to have similar developmental characteristics, then it would be sufficient to characterize an unknown larva as “either *L. coeruleiviridis* or *L. mexicana*”. Lacking data to that effect, however, a DNA-based identification alone will not be adequate for PMI estimation for these two species.

L. cluvia – This species is restricted to the southeastern U.S., and even there it is uncommon (Whitworth 2010). We have only 4 exemplars, all from Florida. These specimens are reciprocally monophyletic compared to all other species, at > 95% bootstrap support.

All of the remaining 15 calliphorid species are clearly and distinctly monophyletic with respect to all other taxa included in the reference database (Fig. 10). The only specific case requiring mention is that we find there to be two distinct subtypes of *Calliphora livida*. Although the two subtypes do not form a monophyletic grouping, each is strongly monophyletic compared to all other species in the reference database.

Sarcophagidae Within the flesh flies (Sarcophagidae), there are three pairs of species for which we observe a failure to achieve reciprocal monophyly. *However, we believe that none of these three pairs will represent a significant problem for DNA-based species identification for the forensically important flies.* In two cases, both species involved are of only limited forensic importance. In the third case, one species is of high forensic importance, but the confounding species is very unlikely to be encountered in a forensic investigation.

Ravinia lherminieri / *R. floridensis* According to our consultant, Dr. Greg Dahlem, an expert on the biology and identification of sarcophagid flies, these may not actually be separate species. They might, instead, represent two subspecies within a single, more inclusive species. In addition, both are primarily dung feeders as larvae, so their actual forensic utility is limited. There are two reasons to include *Ravinia* species as being of potential forensic importance. First, *Ravinia* do frequently appear at large animal carrion, increasing that chance that they might be found as visitors during a forensic investigation. Second, several species of *Ravinia* are known to feed on human feces. If a death scene includes any human excrement, then larvae of *Ravinia* might, indeed, be present.

Ravinia anxia / *R. querula* Another problematic pair within the genus *Ravinia*. As above, the larvae of these two species feed primarily on animal dung, and, thus, they are of relatively minor forensic importance.

Sarcophaga bullata / *S. polistensis* Again, the lack of strict reciprocal monophyly for this species pair is unlikely to have an adverse impact on forensic investigations. Of the 23 individuals of *S. bullata* that we sampled, only one (AF37) falls outside of the main cluster. In some bootstrap pseudoreplicates, this individual grouped with our single sample of *S. polistensis*. However, the reason that we sampled only a single individual of *S. polistensis* is that this species is highly specialized, with its larvae being obligate endoparasites on paper wasps of the genus *Polistes*. Thus, *S. polistensis* is quite unlikely to be encountered in a forensic investigation. When the full reference database is analyzed with *S. polistensis* deleted, all samples of *S. bullata* are monophyletic with bootstrap support of 99% (result not shown). There is some biological interest in the highly divergent mtDNA haplotype found in specimen AF37. As described in the subsection of geographic variation, AF37 represents our only sample of *S. bullata* from California. It may well be that other individuals of *S. bullata* in California will share that divergent haplotype.

Intra-specific variation

Substantial intra-specific variation would have at least two potential consequences. First, it would require the reference database to include a larger number of individuals of each species. This is because observing a large range of variation within a relatively small sample would indicate that a large amount of variation remains un-sampled in natural populations. Second, if there were both intra-specific variation and a geographic pattern to that variation, then we would need to increase not only the total number of individuals sampled, but also the number of collection localities and the coverage of the geographic range for that species that those localities represent. On the other hand, geographic variation might allow inference regarding the geographic origin of an individual fly, which might be useful in some cases. In general, however, we observed only low levels of intra-specific variation, and we could identify only a single case in which there appears to be geographic structure to the intra-specific variation.

Remaining Issues

1. Increase sampling and geographic coverage

We earlier justified the use of a phylogenetic approach (as opposed to a simple definition-based method or a “signature” method) on the basis that natural populations are inherently variable. Consequently, a single DNA sequence definition cannot be expected to capture that variation. Indeed, even if a “signature sequence” sufficed to identify an individual as being a member of a particular species now, the nature of population variability and the processes of mutation and evolution would render that approach incorrect in the future. Although the phylogenetic method does allow for species identification even in the presence of natural variation, that same natural variability also always ensures that a sample, no matter how large, does not capture the full range of variation present in nature. The database developed in this project is the largest fully vouchered database yet to be developed for species identification in forensic entomology. Nevertheless, it will always be desirable to add more specimens, both from the current range of populations included and from additional geographic localities. In much the same way as for reference databases for human DNA profiling, increase the number of reference samples will lead to an increase in statistical power for determination of species identity of an unknown sample.

2. Validation

To date, the reference database developed through this project has been validated only against itself (the bootstrap resampling procedure can be viewed as a form of internal validation). See “Implications for further research” in the last section of the report for additional details.

1. **Tables:**

Table 1. Species of forensic interest in the 48 contiguous United States.

Species of greatest forensic interest		
Family Sarcophagidae		Family Calliphoridae
<i>Blaesoxipha plinthopyga</i>		<i>Calliphora vicina</i>
<i>Sarcophaga africa</i>		<i>Calliphora vomitoria</i>
<i>Sarcophaga argyrostoma</i>		<i>Cochliomyia macellaria</i>
<i>Sarcophaga bullata</i>		<i>Comptosyiops callipes</i>
<i>Sarcophaga cooleyi</i>		<i>Cynomya cadaverina</i>
<i>Sarcophaga crassipalpis</i>		<i>Lucilia coeruleiviridis</i>
		<i>Lucilia cuprina</i>
Family Muscidae		<i>Lucilia illustris</i>
<i>Musca domestica</i>		<i>Lucilia mexicana</i>
<i>Musca autumnalis</i>		<i>Lucilia sericata</i>
		<i>Phormia regina</i>
Species of occasional or possible forensic utility		
Family Sarcophagidae	Family Sarcophagidae	Family Calliphoridae
<i>Boettcheria latisterna</i>	<i>Ravinia querula</i>	<i>Calliphora terraenovae</i>
<i>Helicobia rapax</i>	<i>Ravinia stimulans</i>	<i>Cynomya mortuorum</i>
<i>Mecynocorpus salyum</i>	<i>Sarcophaga sarracenioides</i>	<i>Lucilia chuvia</i>
<i>Oxysarcodexia ventricosa</i>	<i>Sarcophaga subvicina</i>	<i>Lucilia eximia</i>
<i>Ravinia derelicta</i>	<i>Sarcophaga triplasia</i>	<i>Lucilia silvarum</i>
<i>Ravinia floridensis</i>	<i>Sarcophaga utilis</i>	
<i>Ravinia lherminieri</i>	<i>Wohlfahrtia vigil</i>	
<i>Ravinia pusiola</i>		

Table 2. List of specimens for which Phase II DNA sequence data are included in the reference database produced through this project.	
Species (forensically relevant in red , Forensically most important in red and boldface)	COI 1800-3000 Completed sequences
Family Sarcophagidae	
<i>Agria housei</i>	1
<i>Amobia eyrthrura</i>	1
<i>Blaesoxipha alcedo</i>	1
<i>Blaesoxipha arizona</i>	1
<i>Blaesoxipha caridei</i>	1
<i>Blaesoxipha cessator</i>	1
<i>Blaesoxipha eleodis</i>	1
<i>Blaesoxipha masculina</i>	1
<i>Blaesoxipha plinthopyga</i>	8
<i>Blaesoxipha sp.</i>	3
<i>Boettcheria bisetosa</i>	1
<i>Boettcheria cimbicis</i>	1
<i>Boettcheria latisterna</i>	11
<i>Boettcheria literosa</i>	1
<i>Brachicoma sarcophagina</i>	1
<i>Emblemasoma brachicoma</i>	1
<i>Fletcherimyia fletcheri</i>	1
<i>Helicobia rapax</i>	15
<i>Helicobia resinata</i>	1
<i>Macronychia nigrifunus</i>	1
<i>Macronychia pulcra</i>	1
<i>Mecynocorpus salvum</i>	2
<i>Metopia inermis</i>	1
<i>Oxysarcodexia cingarus</i>	1
<i>Oxysarcodexia ventricosa</i>	14
<i>Peckia chrysostoma</i>	1
<i>Peckia intermutans</i>	1
<i>Peckia uncinata</i>	1
<i>Ravinia anxia</i>	3
<i>Ravinia derelicta</i>	6
<i>Ravinia floridensis</i>	8
<i>Ravinia lherminieri</i>	5
<i>Ravinia plantifrons</i>	4
<i>Ravinia pusiola</i>	10
<i>Ravinia querula</i>	9
<i>Ravinia stimulans</i>	10
<i>Ravinia vagabunda</i>	1
<i>Sarcodexia lambens</i>	4
<i>Sarcophaga acrophila</i>	1

<i>Sarcophaga aldrichi</i>	3
<i>Sarcophaga africa</i>	7
<i>Sarcophaga argyrostoma</i>	10
<i>Sarcophaga bancroftorum</i>	1
<i>Sarcophaga bullata</i>	22
<i>Sarcophaga carnaria</i>	1
<i>Sarcophaga cooleyi</i>	3
<i>Sarcophaga crassipalpis</i>	6
<i>Sarcophaga froggatti</i>	1
<i>Sarcophaga georgiana</i>	1
<i>Sarcophaga hinei</i>	1
<i>Sarcophaga houghi</i>	1
<i>Sarcophaga idonea</i>	1
<i>Sarcophaga iota</i>	1
<i>Sarcophaga johnsoni</i>	1
<i>Sarcophaga lehmanni</i>	2
<i>Sarcophaga mimoris</i>	1
<i>Sarcophaga nearctica</i>	12
<i>Sarcophaga omikron</i>	1
<i>Sarcophaga polistensis</i>	1
<i>Sarcophaga sarracenioides</i>	4
<i>Sarcophaga seagoi</i>	1
<i>Sarcophaga shermani</i>	4
<i>Sarcophaga subvicina</i>	6
<i>Sarcophaga taenionota</i>	1
<i>Sarcophaga triplasia</i>	10
<i>Sarcophaga utilis</i>	5
<i>Sarcophaga zeta</i>	1
<i>Senotainia trilineata</i>	1
<i>Sphenometopa nebulosa</i>	1
<i>Tricharaea femoralis</i>	1
<i>Tripanurga importuna</i>	1
<i>Villegasia postuncinnata</i>	1
<i>Wohlfahrtia vigil</i>	1
Family Calliphoridae	
<i>Calliphora coloradensis</i>	6
<i>Calliphora grahami</i>	2
<i>Calliphora latifrons</i>	12
<i>Calliphora livida</i>	6
<i>Calliphora terraenovae</i>	3
<i>Calliphora vicina</i>	16
<i>Calliphora vomitora</i>	17
<i>Chrysomya megacephala</i>	7
<i>Chrysomya rufifacies</i>	14
<i>Cochliomyia macellaria</i>	30

<i>Compsomyiops callipes</i>	0
<i>Cynomya cadaverina</i>	9
<i>Cynomya mortuorum</i>	0
<i>Lucilia cluvia</i>	4
<i>Lucilia coeruleiviridis</i>	22
<i>Lucilia cuprina</i>	23
<i>Lucilia eximia</i>	0
<i>Lucilia elongata</i>	3
<i>Lucilia illustris</i>	21
<i>Lucilia mexicana</i>	23
<i>Lucilia sericata</i>	16
<i>Lucilia silvarum</i>	8
<i>Phormia regina</i>	20
<i>Pollenia angustigena</i>	1
<i>Pollenia labialis</i>	1
<i>Pollenia pediculata</i>	8
<i>Protocalliphora bennetti</i>	2
<i>Protocalliphora occidentalis</i>	1
<i>Protocalliphora rugosa</i>	2
<i>Protocalliphora sialia</i>	2
<i>Pollenia rudis</i>	5
<i>Protophormia terranovae</i>	3
Family Muscidae	
<i>Musca domestica</i>	2
<i>Musca autumnalis</i>	1

Table 3. Specimen code, identification, collection locality, and associated information for all fly specimens collected and identified as a result of work on the project.

ID	FAMILY	GENUS	SUBGENUS	SPECIES	SEX	STATE	COUNTY	LOCALITY1	LOCALITY 2	DATE	TRAP	NOTES
AA01	Sarcophagidae	Ravinia		floridensis (Aldrich)		Florida	Duval	Roadside rest	west of Jacksonville	V.7.2006		
AA02	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Duval	Roadside rest	west of Jacksonville	V.7.2006		
AA03	Sarcophagidae	Ravinia		floridensis (Aldrich)		Florida	Duval	Roadside rest	west of Jacksonville	V.7.2006		
AA04	Sarcophagidae	Ravinia		floridensis (Aldrich)		Florida	Duval	Roadside rest	west of Jacksonville	V.7.2006		
AA05	Sarcophagidae	Lucilia		cuprina Wiedemann	M	Florida	Duval	Rest Stop	west of Jacksonville	V.7.2006		
AA06	Sarcophagidae	Lucilia		cuprina Wiedemann	M	Florida	Duval	Rest Stop	west of Jacksonville	V.7.2006		
AA07	Muscidae				F	Florida	Columbia	O'Leno State Park	Limestone trail	V.7.2006		
AA08	Sarcophagidae	Boettcheria		sp.	F	Florida	Columbia	O'Leno State Park	Limestone trail	V.7.2006		
AA09	Sarcophagidae	Boettcheria		latisterna Parker	F	Florida	Columbia	O'Leno State Park	Limestone trail	V.7.2006		
AA10	Sarcophagidae	Ravinia		derelicta (Walker) M	F	Florida	Columbia	O'Leno State Park	Limestone trail	V.7.2006		
AA11	Sarcophagidae	Boettcheria		latisterna Parker	M	Florida	Columbia	O'Leno State Park	Limestone trail	V.7.2006		
AA12	Sarcophagidae	Boettcheria		latisterna Parker	M	Florida	Columbia	O'Leno State Park	Limestone trail	V.7.2006		
AA13	Sarcophagidae	Ravinia		derelicta (Walker) F	F	Florida	Columbia	O'Leno State Park	Riverwalk	V.7.2006		
AA14	Sarcophagidae	Ravinia		sp.	F	Florida	Duval	Roadside rest	west of Jacksonville	V.7.2006		
AA15	Sarcophagidae	Fletcheriomyia		rileyi (Aldrich)	F	Florida	Liberty	SR-65 roadside	North of Sumatra on S. flava	V.08.2006		
AA16	Sarcophagidae	Fletcheriomyia		rileyi (Aldrich)	M	Florida	Liberty	SR-65 roadside	North of Sumatra on S. flava	V.08.2006		
AA17	Sarcophagidae	Lucilia		cuprina Wiedemann	F	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.8.2006		
AA18	Sarcophagidae	Lucilia		cuprina Wiedemann	F	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.8.2006		
AA19	Sarcophagidae	Lucilia		cuprina Wiedemann	F	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.8.2006		
AA20	Calliphoridae	Lucilia		cuprina Wiedemann	F	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.8.2006		
AA21	Calliphoridae	Lucilia		cuprina Wiedemann	M	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.8.2006		
AA22	Calliphoridae	Stomoxys		calitrans (Linnaeus)	F	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.8.2006		
AA23	Calliphoridae	Oxysarcodexia		ventricosa (Wulp)	F	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.08.2006		
AA24	Calliphoridae	Oxysarcodexia		ventricosa (Wulp)	F	Florida	Columbia	I-10 Rest Stop	near I-75 intersection	V.8.2006		
AA25	Sarcophagidae	Ravinia		derelicta (Walker)	F	Florida	Liberty	Torrey State Park		V.8.2006		
AA26	Sarcophagidae	Ravinia		derelicta (Walker)	F	Florida	Liberty	Torrey State Park		V.8.2006		
AA27	Sarcophagidae	Sarcophaga	Neobellieria	triplesia Wulp	M	Florida	Liberty	Torrey State Park		V.8.2006		
AA28	Calliphoridae	Cochliomyia		macellaria (Fabricius) M	M	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA29	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA30	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA31	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA32	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA33	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA34	Calliphoridae	Lucilia		cuprina Wiedemann	M	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA35	Sarcophagidae	Helicobia		monilella (Aldrich)	M	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA36	Sarcophagidae	Ravinia		Iherminieri (Robineau-Desvoidy)	M	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA37	Calliphoridae	Oxysarcodexia		ventricosa (Wulp)	F	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.08.2006		
AA38	Sarcophagidae	Ravinia		derelicta (Walker)	F	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA39	Muscidae	Musca		domestica Linnaeus	F	Florida	Madison	I-10 Rest Stop	East of Tallahassee	V.8.2006		
AA40	Sarcophagidae	Lucilia		cuprina Wiedemann	F	Florida	Bay	Panama City	Sunset Inn motel	V.9.2006		
AA41	Sarcophagidae	Lucilia		cuprina Wiedemann	F	Florida	Bay	Panama City	Sunset Inn motel	V.9.2006		
AA42	Calliphoridae	Lucilia		cuprina Wiedemann	M	Florida	Bay	Panama City	Sunset Inn motel	V.9.2006		
AA43	Calliphoridae	Lucilia		cuprina Wiedemann	F	Florida	Bay	Panama City	Sunset Inn motel	V.9.2006		
AA44	Calliphoridae	Lucilia		cuprina Wiedemann	F	Florida	Bay	Panama City	Sunset Inn motel	V.9.2006		
AA45	Calliphoridae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA46	Calliphoridae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA47	Sarcophagidae	Spirobolomyia		singularis (Aldrich)	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA48	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA49	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Florida	Walton	Greyton Beach State Park		V.9.2006		ID checked
AA50	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA51	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA52	Sarcophagidae	Senotania		sp.	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA53	Sarcophagidae	Senotania		sp.	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA54	Sarcophagidae	Helicobia		sp.	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA55	Sarcophagidae	Eumacronychia		sp.	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA56	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Sarasota	Myakka River State Park	Dam area	V.11.2006		
AA57	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Sarasota	Myakka River State Park	Dam area	V.11.2006		
AA58	Sarcophagidae	Amobia		floridensis (Townsend)	M	Florida	Sarasota	Myakka River State Park	Dam area	V.11.2006		
AA59	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA60	Sarcophagidae	Ravinia		floridensis (Aldrich)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA61	Sarcophagidae	Oxysarcodexia		galeata (Aldrich)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA62	Sarcophagidae	Amobia		floridensis (Townsend)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA63	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	F	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA64	Calliphoridae	Chrysomya		rufficeps (Macquart)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA65	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA66	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA67	Sarcophagidae	Amobia		floridensis (Townsend)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA68	Sarcophagidae	Amobia		floridensis (Townsend)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA69	Sarcophagidae	Sarcodexia		lambens Wiedemann	M	Florida	Sarasota	Myakka River State Park	Dam area	V.11.2006		
AA70	Sarcophagidae	Helicobia		sp.	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA71	Sarcophagidae	Amobia		floridensis (Townsend)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA72	Sarcophagidae	Amobia		floridensis (Townsend)	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA73	Sarcophagidae	Helicobia		sp.	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA74	Sarcophagidae	Helicobia		sp.	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA75	Sarcophagidae	Helicobia		sp.	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA76	Sarcophagidae	Helicobia		sp.	M	Florida	Sarasota	Myakka River State Park	Birdwalk	V.11.2006		
AA77	Sarcophagidae	Sarcophaga		sp.	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA78	Sarcophagidae	Sarcophaga		sp.	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA79	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AA80	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AB01	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AB02	Calliphoridae	Lucilia		coeruleiviridis Macquart	M	Florida	Walton	Greyton Beach State Park		V.9.2006		
AB03	Sarcophagidae	Sarcophaga		sp.	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AB04	Calliphoridae	Lucilia		coeruleiviridis Macquart	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AB05	Sarcophagidae	Ravinia		derelicta (Walker)	F	Florida	Walton	Greyton Beach State Park		V.9.2006		
AB06	Sarcophagidae	Spirobolomyia		singularis (Aldrich)	M	Florida	Washington	Falling Waters State Park		V.10.2006		
AB07	Calliphoridae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Washington	Falling Waters State Park		V.10.2006		
AB08	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB09	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB10	Sarcophagidae	Sarcophaga	Wohlfahrtiopsis	georgiana Dodge	M	Florida	Washington	Falling Waters State Park		V.10.2006		
AB11	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB12	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB13	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB14	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB15	Sarcophagidae	Ravinia		floridensis (Aldrich)	M	Florida	Washington	Falling Waters State Park		V.10.2006		
AB16	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB17	Sarcophagidae	Spirobolomyia		singularis (Aldrich)	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB18	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB19	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB20	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB21	Sarcophagidae	Spirobolomyia		singularis (Aldrich)	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB22	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB23	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB24	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB25	Sarcophagidae	Sarcophaga	Bercaeopsis	sp.	M	Florida	Washington	Falling Waters State Park		V.10.2006		
AB26	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB27	Sarcophagidae	Spirobolomyia		sp.	M	Florida	Washington	Falling Waters State Park		V.9.2006		
AB												

AE22	Sarcophagidae	Sarcophaga	Wohlfahrtiopsis	utilis Aldrich	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE23	Sarcophagidae	Sarcophaga	Wohlfahrtiopsis	utilis Aldrich	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE24	Sarcophagidae	Sarcophaga	Wohlfahrtiopsis	utilis Aldrich	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE25	Sarcophagidae	Amobia		oculata (Zetterstedt)	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE26	Calliphoridae	Phormia		regina (Meigen)	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE27	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE28	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE29	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE30	Sarcophagidae	Blaesoxipha		sp.	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE31	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE32	Calliphoridae	Phormia		regina (Meigen)	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE33	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE34	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE35	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE36	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE37	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE38	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Picnic Area & garbage cans	V.23.2006
AE39	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	M	South Carolina	Newberry	Sumter NF: Enoree Dist.	Firetower Area	V.22.2006
AE40	Calliphoridae	Phormia		regina (Meigen)	F	South Carolina	Newberry	Sumter NF: Enoree Dist.	Firetower Area	V.23.2006
AE41	Calliphoridae	Calliphora		vomitiora (Linnaeus)	M	Georgia	Rabun	Rabun Bald	base	V.24.2006
AE42	Calliphoridae	Calliphora		vomitiora (Linnaeus)	F	Georgia	Rabun	Rabun Bald	base	V.24.2006
AE43	Sarcophagidae	Tripanurga		sp.	M	Oregon	Klamath	US-97 Rest stop	near Beaver Marsh	VI.7.2006
AE44	Sarcophagidae	Tripanurga		sp.	M	Oregon	Klamath	US-97 Rest stop	near Beaver Marsh	VI.7.2006
AE45	Calliphoridae	Ravinia		planifrons (Aldrich)	M	Oregon	Klamath	US-97 Rest stop	near Beaver Marsh	VI.7.2006
AE46	Sarcophagidae	Ravinia		anxia (Walker)	F	Oregon	Klamath	US-97 Rest stop	near Beaver Marsh	VI.7.2006
AE47	Sarcophagidae	Tripanurga		sp.	F	Oregon	Klamath	US-97 Rest stop	near Beaver Marsh	VI.7.2006
AE48	Sarcophagidae	Sarcophaga		sp.	M	California	Glenn	I-5 Rest Stop	Near Artois	VI.7.2006
AE49	Sarcophagidae	Sarcophaga		sp.	M	California	Glenn	I-5 Rest Stop	Near Artois	VI.7.2006
AE50	Sarcophagidae	Blaesoxipha	Gigantotheca	plinthopyga (Wiedemann)	M	California	Glenn	I-5 Rest Stop	Near Artois	VI.7.2006
AE51	Sarcophagidae	Ravinia		errabunda (Wulp)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE52	Sarcophagidae	Ravinia		stimulans (Walker)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE53	Sarcophagidae	Blaesoxipha		sp.	F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE54	Sarcophagidae	Ravinia		anxia (Walker)	F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE55	Sarcophagidae	Sarcophaga		sp.	F	California	Glenn	I-5 Rest Stop	Near Artois	VI.7.2006
AE56	Sarcophagidae	Wohlfahrtia		vigili (Walker)	F	California	Glenn	I-5 Rest Stop	Near Artois	VI.7.2006
AE57	Sarcophagidae	Ravinia		derelecta (Walker)?	F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE58	Muscidae				F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE59	Calliphoridae	Lucilia		sericata (Meigen)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE60	Calliphoridae	Lucilia		mexicana Macquart	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE61	Calliphoridae	Lucilia		cuprina Wiedemann	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE62	Calliphoridae	Lucilia		sericata (Meigen)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE63	Calliphoridae	Lucilia		mexicana Macquart	F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE64	Calliphoridae	Lucilia		sericata (Meigen)	F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE65	Calliphoridae	Lucilia		sericata (Meigen)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE66	Calliphoridae	Lucilia		sericata (Meigen)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE67	Calliphoridae	Phormia		regina (Meigen)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE68	Calliphoridae	Lucilia		sericata (Meigen)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE69	Calliphoridae	Lucilia		sericata (Meigen)	M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE70	Muscidae				F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE71	Muscidae				F	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE72	Muscidae				M	California	Glenn	I-5 Rest Stop	near Artois	VI.7.2006
AE73	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	California	Shasta	I-5 Rest Stop	Shasta Lake overview	VI.7.2006
AE74	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	California	Shasta	I-5 Rest Stop	Shasta Lake overview	VI.7.2006
AE75	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	California	Shasta	I-5 Rest Stop	Shasta Lake overview	VI.7.2006
AE76	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	F	California	Shasta	I-5 Rest Stop	Shasta Lake overview	VI.7.2006
AE77	Sarcophagidae	Helicobia		sp.	F	California	Shasta	I-5 Rest Stop	Shasta Lake overview	VI.7.2006
AE78	Calliphoridae	Calliphora		latifrons Hough	M	California	Siskiyou	US-97 Rest stop	Grass Lake	VI.7.2006
AE79	Sarcophagidae	Ravinia		anxia (Walker)	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AE80	Sarcophagidae	Ravinia		anxia (Walker)	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AE81	Sarcophagidae	Ravinia		anxia (Walker)	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF01	Sarcophagidae	Brachicoma		setosa Coquillett	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF02	Sarcophagidae	Brachicoma		setosa Coquillett	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF03	Sarcophagidae	Ravinia		querula (Walker)	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF04	Sarcophagidae	Tripanurga		sp.	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF05	Sarcophagidae	Ravinia		planifrons (Aldrich)	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF06	Sarcophagidae	Ravinia		planifrons (Aldrich)	M	California	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF07	Conopidae				M	California	Siskiyou	US-97 Rest Stop	Grass Lake	VI.7.2006
AF08	Tachinidae				M	Oregon	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF09	Tachinidae				M	Oregon	Siskiyou	US-97 Rest stop	Grass lake	VI.7.2006
AF10	Calliphoridae	Protophormia		terraenovae (Robineau-Desvoidy)	F	California	Siskiyou	US-97 Rest stop	Grass Lake	VI.7.2006
AF11	Calliphoridae	Calliphora		vomitiora (Linnaeus)	M	California	Siskiyou	US-97 Rest stop	Grass Lake	VI.7.2006
AF12	Calliphoridae	Phormia		regina (Meigen)	M	California	Siskiyou	US-97 Rest stop	Grass Lake	VI.7.2006
AF13	Calliphoridae	Phormia		regina (Meigen)	M	California	Siskiyou	US-97 Rest stop	Grass Lake	VI.7.2006
AF14	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Clackamas	Mini Crater Lake	VI.9.2006	
AF15	Calliphoridae	Calliphora		latifrons Hough	M	Oregon	Clackamas	Mini Crater Lake	VI.9.2006	
AF16	Sarcophagidae	Ravinia		querula (Walker)	M	Oregon	Clackamas	Mini Crater Lake	VI.9.2006	
AF17	Calliphoridae	Ravinia		querula (Walker)	M	Oregon	Clackamas	Mini Crater Lake	VI.9.2006	
AF18	Sarcophagidae	Sarcophaga	Robineauella	nearctica Parker	M	Oregon	Crook	Ococho Reservoir	Fish cleaning station	V.8.2006
AF19	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Washington	Clark	I-5 Rest Stop	Welcome Center	VI.9.2006
AF20	Calliphoridae	Lucilia		illustris (Meigen)	F	Washington	Clark	I-5 Rest Stop	Welcome center	VI.9.2006
AF21	Calliphoridae	Calliphora		latifrons Hough	F	Washington	Clark	I-5 Rest Stop	Welcome Center	VI.9.2006
AF22	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Washington	Clark	I-5 Rest Stop	Welcome Center	VI.9.2006
AF23	Calliphoridae	Phormia		nudis (Fabricius)	M	Washington	Clark	I-5 Rest Stop	Welcome Center	VI.9.2006
AF24	Calliphoridae	Phormia		regina (Meigen)	M	Washington	Clark	I-5 Rest Stop	Welcome Center	VI.9.2006
AF25	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF26	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF27	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF28	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF29	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF30	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF31	Sarcophagidae	Sarcophaga		sp.	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF32	Sarcophagidae	Sarcophaga		sp.	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF33	Sarcophagidae	Sarcophaga		sp.	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF34	Sarcophagidae	Sarcophaga		sp.	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF35	Sarcophagidae	Sarcophaga		sp.	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF36	Sarcophagidae	Sarcophaga		sp.	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF37	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF38	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF39	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF40	Sarcophagidae	Sarcophaga		sp.	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF41	Calliphoridae	Phormia		regina (Meigen)	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF42	Muscidae				F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF43	Muscidae				M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF44	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF45	Calliphoridae	Phormia		regina (Meigen)	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF46	Calliphoridae	Phormia		regina (Meigen)	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF47	Calliphoridae	Lucilia		sericata (Meigen)	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF48	Calliphoridae	Phormia		regina (Meigen)	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF49	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF50	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF51	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF52	Calliphoridae	Phormia		regina (Meigen)	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF53	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF54	Calliphoridae	Phormia		regina (Meigen)	F	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF55	Calliphoridae	Phormia		regina (Meigen)	M	California	Yolo	Rest Stop	South of Caluso	VI.9.2006
AF56	Sarcophagidae	Sarcophaga		sp.	M	Oregon	Linn	I-5 Rest Stop	south of Harrisburg	VI.10.2006
AF57	Calliphoridae	Lucilia		silvarum (Meigen)	M	Oregon	Linn	I-5 Rest Stop	south of Harrisburg	VI.10.2006

AF58	Calliphoridae	Lucilia		silvarum (Meigen)	F	Oregon	Linn	I-5 Rest Stop	south of Harrisburg	VI.10.2006
AF59	Calliphoridae	Lucilia		silvarum (Meigen)	M	Oregon	Linn	I-5 Rest Stop	south of Harrisburg	VI.10.2006
AF60	Calliphoridae	Lucilia		silvarum (Meigen)	M	Oregon	Linn	I-5 Rest Stop	south of Harrisburg	VI.10.2006
AF61	Sarcophagidae	Sarcophaga	Neobellieria	cooley Parker	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AF62	Sarcophagidae	Sarcophaga		sp.	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AF63	Calliphoridae	Calliphora		vomitiora (Linnaeus)	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF64	Muscidae				F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF65	Calliphoridae	Calliphora		terraenovae Macquart	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF66	Calliphoridae	Lucilia		sericata (Meigen)	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF67	Muscidae				M	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF68	Calliphoridae	Pollenia		rudis (Fabricius)	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF69	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF70	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF71	Muscidae				F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF72	Muscidae				F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF73	Calliphoridae	Lucilia		sp.	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF74	Calliphoridae	Lucilia		sericata (Meigen)	F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF75	Muscidae				F	Oregon	Crook	Ococho Reservoir	Fish cleaning station	VI.8.2006
AF76	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AF77	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AF78	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AF79	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AF80	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AF81	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG01	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG02	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG03	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG04	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG05	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG06	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG07	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG08	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG09	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG10	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG11	Sarcophagidae	Sarcophaga		sp.	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG12	Muscidae				F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG13	Muscidae				F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG14	Muscidae				F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG15	Calliphoridae	Cynomya		cadaverina Robineau-Desvoidy	M	Oregon	Benton	Mary's Peak		VI.10.2006
AG16	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Benton	Mary's Peak		VI.10.2006
AG17	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Benton	Mary's Peak		VI.10.2006
AG18	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Benton	Mary's Peak		VI.10.2006
AG19	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Oregon	Clatsop	Astoria	Best Western parking lot fences	VI.10.2006
AG20	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Oregon	Clatsop	Astoria	Best Western parking lot fences	VI.10.2006
AG21	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Oregon	Clatsop	Astoria	Best Western parking lot fences	VI.10.2006
AG22	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Clatsop	Astoria	Best Western parking lot fences	VI.10.2006
AG23	Calliphoridae	Calliphora		latifrons Hough	M	Oregon	Clatsop	Astoria	Best Western parking lot fences	VI.10.2006
AG24	Calliphoridae	Calliphora		latifrons Hough	M	Oregon	Clatsop	Astoria	Best Western parking lot fences	VI.10.2006
AG25	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	M	Oregon	Clatsop	Astoria	Best Western parking lot fences	VI.10.2006
AG26	Sarcophagidae	Sarcophaga	Neobellieria	itorosa (Raintard)	M	Oregon	Crook	Haystack Reservoir	Best Western parking lot fences	VI.10.2006
AG27	Sarcophagidae	Sarcophaga	Neobellieria	cooley Parker	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG28	Sarcophagidae	Ravinia		querula (Walker)	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG29	Sarcophagidae	Blaesoxipha	Gigantotheca	plinthopyga (Wiedemann)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AG30	Sarcophagidae	Ravinia		querula (Walker)	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG31	Sarcophagidae	Ravinia		querula (Walker)	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG32	Sarcophagidae	Ravinia		querula (Walker)	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG33	Sarcophagidae	Ravinia		querula (Walker)	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG34	Sarcophagidae	Ravinia		planifrons (Aldrich)	M	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG35	Sarcophagidae	Metopia		sp.	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG36	Sarcophagidae	Sarcophaga	Neobellieria	cooley Parker	M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG37	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Benton	Mary's Peak		VI.10.2006
AG38	Calliphoridae	Calliphora		latifrons Hough	F	Oregon	Benton	Mary's Peak		VI.10.2006
AG39	Muscidae				M	Oregon	Lane	Coburg		VI.10.2006
AG40	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG41	Sarcophagidae	Blaesoxipha		sp.	F	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG42	Calliphoridae	Calliphora		latifrons Hough	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG43	Calliphoridae	Calliphora		latifrons Hough	M	Oregon	Crook	Powell Butte	6645 SW Steffey Ln.	VI.8.2006
AG44	Calliphoridae	Lucilia		sericata (Meigen)	M	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG45	Tachinidae				M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG46	Tachinidae				M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG47	Calliphoridae	Lucilia		sericata (Meigen)	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG48	Sarcophagidae	Ravinia		anxia (Walker)	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG49	Sarcophagidae	Ravinia		sp.	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG50	Tachinidae				M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG51	Sarcophagidae	Ravinia		anxia (Walker)	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG52	Tachinidae				M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG53	Calliphoridae	Calliphora		coloradensis Hough	M	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG54	Tachinidae				M	Oregon	Jefferson	Haystack Reservoir		VI.8.2006
AG55	Sarcophagidae	Blaesoxipha		sp.	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG56	Sarcophagidae	Blaesoxipha		sp.	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AG57	Calliphoridae	Pollenia		rudis (Fabricius)	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG58	Sarcophagidae	Sarcophaga		sp.	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG59	Calliphoridae	Lucilia		illustris (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG60	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG61	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG62	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG63	Calliphoridae	Lucilia		illustris (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG64	Sarcophagidae	Ravinia		sp.	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG65	Calliphoridae	Lucilia		illustris (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG66	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG67	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG68	Calliphoridae	Lucilia		illustris (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG69	Calliphoridae	Calliphora		vomitiora (Linnaeus)	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG70	Muscidae				F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG71	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG72	Calliphoridae	Lucilia		elongata Shannon	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG73	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG74	Sarcophagidae	Sarcophaga		sp.	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG75	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG76	Calliphoridae	Lucilia		elongata Shannon	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG77	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG78	Calliphoridae	Phormia		regina (Meigen)	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG79	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG80	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AG81	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH01	Calliphoridae	Lucilia		sericata (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH02	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH03	Calliphoridae	Calliphora		latifrons Hough	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH04	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH05	Sarcophagidae	Sarcophaga		sp.	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH06	Calliphoridae	Pollenia		rudis (Fabricius)	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH07	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH08	Calliphoridae	Pollenia		rudis (Fabricius)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH09	Muscidae			domestica Linnaeus	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH10	Calliphoridae	Pollenia		rudis (Fabricius)	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH11	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH12	Calliphoridae	Phormia		regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006

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AH13	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH14	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH15	Muscidae	Phormia		F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH16	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH17	Sarcophagidae	Senotainia	sp.	F	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH18	Sarcophagidae	Boettcheria	illirosa (Reinhard)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH19	Sarcophagidae	Sarcophaga	sp.	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH20	Sarcophagidae	Sarcophaga	sp.	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH21	Sarcophagidae	Sarcophaga	nearctica Parker	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH22	Sarcophagidae	Sarcophaga	shermani Parker	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH23	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH24	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Douglas	I-5 Rest Stop	near Oakland	VI.11.2006
AH25	Sarcophagidae	Sarcophaga	sp.	F	Oregon	Douglas	Umpqua National Forest	Crashing rivers viewpoint	VI.11.2006
AH26	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH27	Tachnidae	Phormia	regina (Meigen)	M	Oregon	Douglas	Umpqua National Forest	Crashing rivers viewpoint	VI.11.2006
AH28	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH29	Calliphoridae	Lucilia	illustris (Meigen)	M	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH30	Calliphoridae	Lucilia	illustris (Meigen)	M	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH31	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH32	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH33	Calliphoridae	Calliphora	latifrons Hough	M	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH34	Calliphoridae	Calliphora	latifrons Hough	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH35	Calliphoridae	Pollenia	angustigena Wainwright	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH36	Calliphoridae	Pollenia	angustigena Wainwright	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH37	Calliphoridae	Lucilia	elongata Shannon	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH38	Calliphoridae	Lucilia	illustris (Meigen)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH39	Calliphoridae	Lucilia	illustris (Meigen)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH40	Calliphoridae	Lucilia	illustris (Meigen)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH41	Calliphoridae	Calliphora	vomitiora (Linnaeus)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH42	Sarcophagidae	Ravinia	querula (Walker)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH43	Sarcophagidae	Boettcheria	illirosa (Reinhard)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH44	Sarcophagidae	Sarcophaga	shermani Parker	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH45	Sarcophagidae	Sarcophaga	sp.	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH46	Sarcophagidae	Sarcophaga	sp.	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH47	Sarcophagidae	Sarcophaga	shermani Parker	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH48	Sarcophagidae	Sarcophaga	shermani Parker	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH49	Sarcophagidae	Sarcophaga	sp.	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH50	Sarcophagidae	Metopia	sp.	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH51	Sarcophagidae	Ravinia	querula (Walker)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH52	Sarcophagidae	Ravinia	anxia (Walker)	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AH53	Muscidae	Phormia		F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH54	Muscidae	Muscidae		F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH55	Muscidae	Muscidae		F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH56	Calliphoridae	Pollenia	rudis (Fabricius)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH57	Calliphoridae	Calliphora	livida Hall	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH58	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH59	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH60	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH61	Calliphoridae	Calliphora	vicina Robineau-Desvoidy	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH62	Calliphoridae	Calliphora	vicina Robineau-Desvoidy	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH63	Calliphoridae	Calliphora	vicina Robineau-Desvoidy	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH64	Calliphoridae	Calliphora	vicina Robineau-Desvoidy	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH65	Calliphoridae	Calliphora	vicina Robineau-Desvoidy	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH66	Calliphoridae	Calliphora	livida Hall	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH67	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH68	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH69	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH70	Calliphoridae	Calliphora	latifrons Hough	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH71	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH72	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH73	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH74	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH75	Sarcophagidae	Sarcophaga	cooley Parker	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH76	Sarcophagidae	Sarcophaga	cooley Parker	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH77	Sarcophagidae	Blaesoxipha	plinthopyga (Wiedemann)	M	Oregon	Crook	Ococho Reservoir		VI.8.2006
AH78	Calliphoridae	Calliphora	vomitiora (Linnaeus)	F	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH79	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH80	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AH81	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Umpqua Natl. Forest	Crashing rivers viewpoint	VI.11.2006
AJ01	Calliphoridae	Calliphora	vomitiora (Linnaeus)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ02	Calliphoridae	Calliphora	vomitiora (Linnaeus)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ03	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ04	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ05	Calliphoridae	Protosphormia	terraenovae (Robineau-Desvoidy)	M	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ06	Calliphoridae	Protosphormia	terraenovae (Robineau-Desvoidy)	M	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ07	Calliphoridae	Calliphora	latifrons Hough	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ08	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ09	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ10	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ11	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ12	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ13	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ14	Muscidae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ15	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ16	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ17	Calliphoridae	Phormia	regina (Meigen)	F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ18	Calliphoridae	Phormia	regina (Meigen)	M	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ19	Muscidae	Muscidae		F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ20	Muscidae	Muscidae		F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ21	Muscidae	Muscidae		F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ22	Muscidae	Muscidae		F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ23	Muscidae	Muscidae		F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ24	Muscidae	Muscidae		F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ25+	Muscidae	Muscidae		F	Oregon	Douglas	Diamond Lake		VI.11.2006
AJ01	Calliphoridae	Calliphora	latifrons Hough	F	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ02	Calliphoridae	Calliphora	grahami Aldrich	F	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ03	Tachnidae	Phormia		F	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ04	Tachnidae	Phormia		F	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ05	Sarcophagidae	Brachicoma	setosa Coquillett	F	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ06	Calliphoridae	Calliphora	latifrons Hough	F	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ07	Sarcophagidae	Brachicoma	setosa Coquillett	F	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ08	Calliphoridae	Calliphora	grahami Aldrich	M	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ09	Sarcophagidae	Brachicoma	setosa Coquillett	M	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ10	Calliphoridae	Calliphora	grahami Aldrich	M	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ11	Sarcophagidae	Blaesoxipha	sp.	M	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ12	Sarcophagidae	Sarcophaga	africa (Wiedemann)	M	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ13	Sarcophagidae	Sarcophaga	africa (Wiedemann)	M	Oregon	Coos	Suislaw Natl. Forest	Coastal dunes S of Lakeside	VI.13.2006
AJ14	Muscidae	Muscidae		M	Oregon	Curry	I-101 Rest Stop	Welcome Center North of Brookings	VI.13.2006
AJ15	Calliphoridae	Calliphora	terraenovae Macquart	F	Oregon	Curry	I-101 N Rest stop	Welcome Center North of Brookings	VI.13.2006
AJ16	Sarcophagidae	Ravinia	anxia (Walker)	F	Oregon	Curry	I-101 Rest Stop	Welcome Center North of Brookings	VI.13.2006
AJ17	Sarcophagidae	Sarcophaga	cooley Parker	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AJ18	Sarcophagidae	Ravinia	anxia (Walker)	M	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AJ19	Sarcophagidae	Ravinia	querula (Walker)	M	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AJ20	Calliphoridae	Calliphora	latifrons Hough	M	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AJ21	Sarcophagidae	Sarcophaga	sp.	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AJ22	Sarcophagidae	Sarcophaga	sp.	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006
AJ23	Calliphoridae	Calliphora	querula (Walker)	F	Oregon	Jefferson	Haystack Reservoir		VI.14.2006

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AK60	Tachnidae			M	Georgia	Rabun	Rabun Bald		Summit	1:30-2:30	V.24.2006
AK61	Tachnidae			M	Georgia	Rabun	Rabun Bald		Summit	1:30-2:30	V.24.2006
AK62	Tachnidae			M	Georgia	Rabun	Rabun Bald		Summit	1:30-2:30	V.24.2006
AK63	Sarcophagidae	Sarcophaga	Bercaeopsis	sp.	M	Georgia	Rabun	Rabun Bald	Summit	1:30-2:30	V.24.2006
AK64	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	1:30-2:30	V.24.2006
AK65	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	1:30-2:30	V.24.2006
AK66	Sarcophagidae	Brachionoma		seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	1:30-2:30	V.24.2006
AK67	Sarcophagidae	Boettcheria		bisetosa Parker	F	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK68		Calliphora		vomitoria (Linnaeus)	F	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK69		Calliphora		vicina Robineau-Desvoidy	F	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK70		Calliphora		vicina Robineau-Desvoidy	F	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK71	Calliphoridae	Calliphora		vomitoria (Linnaeus)	F	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK72	Calliphoridae	Calliphora		vomitoria (Linnaeus)	F	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK73	Sarcophagidae	Boettcheria		bisetosa Parker	M	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK74	Sarcophagidae	Sarcophaga	Neobellieria	triplesia Wulp	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AK75	Sarcophagidae	Sarcophaga	Bercaeopsis	pulla Aldrich	M	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AK76	Tachnidae				F	Georgia	Rabun	Rabun Bald	Trail to summit 2:30-3:30		V.24.2006
AK77	Tachnidae				M	Georgia	Rabun	Rabun Bald	Trail to summit 2:30-3:30		V.24.2006
AK78	Tachnidae				M	Georgia	Rabun	Rabun Bald	Trail to summit 2:30-3:30		V.24.2006
AK79	Sarcophagidae	Lepidodexia	Neophyto	sheldoni (Coquillett)?	M	Georgia	Rabun	Rabun Bald	Trail from summit 2:30-3:30		V.24.2006
AK80	Sarcophagidae	Macronychia		aurata (Coquillett)	F	Georgia	Rabun	Rabun Bald	Trail from summit 2:30-3:30		V.24.2006
AK81	Sarcophagidae	Brachionoma		sarcophagina (Townsend)	M	Georgia	Rabun	Rabun Bald	Trail from summit 2:30-3:30		V.24.2006
AL01	Calliphoridae	Calliphora		livida Hall	M	Georgia	Rabun	Rabun Bald summit			V.24.2006
AL02	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	M	Georgia	Rabun	Rabun Bald summit			V.24.2006
AL03	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL04	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL05	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL06	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL07	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL08	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL09	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL10	Sarcophagidae	Sarcophaga	Neobellieria	triplesia Wulp	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL11	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL12	Sarcophagidae	Sarcophaga	Liopygia	crassipalpis Macquart	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL13	Sarcophagidae	Blaesoxipha		sp.	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL14	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL15	Sarcophagidae	Sarcophaga	Neobellieria	triplesia Wulp	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL16	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL17	Sarcophagidae	Sarcophaga	Bercaeopsis	sp.	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL18	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL19	Sarcophagidae	Sarcophaga	Bercaeopsis	hesterna Reinhard	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL20	Sarcophagidae	Blaesoxipha		sp.	M	Georgia	Rabun	Rabun Bald	Summit		V.24.2006
AL21	Sarcophagidae	Blaesoxipha		sp.	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Picnic Area		V.23.2006
AL22	Sarcophagidae	Boettcheria		bisetosa Parker	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Picnic Area & garbage cans		V.23.2006
AL23	Sarcophagidae	Boettcheria		cimbicis (Townsend)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Picnic Area		V.23.2006
AL24	Sarcophagidae	Blaesoxipha		sp.	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Picnic Area		V.23.2006
AL25	Tachnidae				M	South Carolina	Newberry	Sumter NF; Enoree Dist.	Picnic Area		V.23.2006
AL26	Sarcophagidae	Boettcheria		latisterna Parker	M	South Carolina	Newberry	Sumter NF; Enoree Dist.	Picnic Area		V.23.2006
AL27	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL28	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL29	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL30	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL31	Sarcophagidae	Sarcophaga	Bercaeopsis	africa (Wiedemann)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL32	Sarcophagidae	Ravinia		querula (Walker)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL33	Sarcophagidae	Sarcophaga	Liopygia	crassipalpis Macquart	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL34	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL35	Sarcophagidae	Sarcophaga	Liopygia	crassipalpis Macquart	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL36	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL37	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL38	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL39	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL40	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL41	Sarcophagidae	Sarcophaga	Mehria	houghi Aldrich	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL42	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL43	Sarcophagidae	Sarcophaga	Liopygia	crassipalpis Macquart	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL44	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL45	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL46	Sarcophagidae	Sarcophaga	Bercaeopsis	acrophila (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL47	Sarcophagidae	Blaesoxipha		sp.	M	Georgia	Rabun	Rabun Bald	Summit	11:30-12:30	V.24.2006
AL48	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL49	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL50	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL51	Tachnidae				F	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL52	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL53	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL54	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL55	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL56	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL57	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL58	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL59	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL60	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL61	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL62	Tachnidae				F	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL63	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL64	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL65	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL66	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL67	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL68	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL69	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL70	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL71	Calliphoridae	Calliphora		vomitoria (Linnaeus)	M	Georgia	Rabun	Rabun Bald	Trail to summit		V.24.2006
AL72	Sarcophagidae	Sarcophaga	Bercaeopsis	atarsata Aldrich	M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL73	Sarcophagidae	Brachionoma		sarcophagina (Townsend)	M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL74	Sarcophagidae	Ravinia		stimulans (Walker)	M	Georgia	Rabun	Rabun Bald	Summit	12:30-1:30	V.24.2006
AL75	Tachnidae				M	Georgia	Rabun	Rabun Bald	Summit	10:50-11:30	V.24.2006
AL76	Tachnidae				F	Georgia	Rabun	Rabun Bald	Summit	10:50-11:30	V.24.2006
AL77	Sarcophagidae	Spirobolomyia		sp.	M	Georgia	Rabun	Rabun Bald	Summit	10:50-11:30	V.24.2006
AL78	Sarcophagidae	Blaesoxipha		sp.	M	Georgia	Rabun	Rabun Bald	Summit	10:50-11:30	V.24.2006
AL79	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	10:50-11:30	V.24.2006
AL80	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit	10:50-11:30	V.24.2006
AL81	Sarcophagidae	Sarcophaga	Liopygia	crassipalpis Macquart	M	Georgia	Rabun	Rabun Bald	Summit	10:50-11:30	V.24.2006
AM01	Muscidae	Phormia		regina (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoolic Head summit		VII.15.2006
AM02	Calliphoridae	Phormia		regina (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoolic Center		VII.15.2006
AM03	Calliphoridae	Phormia		regina (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoolic Center		VII.15.2006
AM04	Calliphoridae	Phormia		regina (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoolic Center		VII.15.2006
AM05	Tachnidae				M	Maine	Hancock	Acadia N.P.	Schoolic Head summit		VII.15.2006
AM06	Tachnidae				M	Maine	Hancock	Acadia N.P.	Schoolic Head summit		VII.15.2006
AM07	Sarcophagidae	Agria		housei (Shewell)	F	Maine	Hancock	Acadia N.P.	Schoolic Head summit		VII.15.2006
AM08	Sarcophagidae	Sarcophaga	Mehria	hinei Aldrich	F	Maine	Hancock	Acadia N.P.	Schoolic Head summit		VII.15.2006
AM09	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	Maine	Hancock	Acadia N.P.	Schoolic Head summit		VII.15.2006
AM10	Sarcophagidae	Agria		housei (Shewell)	M	Maine	Hancock	Acadia N.P.	Schoolic Head summit		VII.15.2006
AM11	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Maine	Hancock	Acadia N.P.	Schoolic Center		VII.15.2006
AM12	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Maine	Hancock	Acadia N.P.	Schoolic Center		VII.15.2006
AM13	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Maine	Hancock	Acadia N.P.	Schoolic Center		VII.15.2006
AM14	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	F	Maine	Hancock	Acadia N.P.	Schoolic Center		VII.15.2006

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AM15	Calliphoridae	Phormia		regina (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM16	Calliphoridae	Phormia		regina (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM17	Calliphoridae	Phormia		regina (Meigen)	M	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM18	Calliphoridae	Phormia		regina (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM19	Calliphoridae	Lucilia		sericata (Meigen)	F	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM20	Muscidae					Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM21	Muscidae					Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM22	Muscidae					Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM23	Muscidae					Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM24	Muscidae					Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM25	Calliphoridae	Cynomya		cadaverina Robineau-Desvoidy	M	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM26	Calliphoridae	Cynomya		cadaverina Robineau-Desvoidy	M	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM27	Calliphoridae	Calliphora		vomitiora (Linnaeus)	M	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM28	Calliphoridae	Lucilia		illustrius (Meigen)	M	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM29	Calliphoridae	Phormia		regina (Meigen)	M	Maine	Hancock	Acadia N.P.	Schoodic Center	VII.15.2006
AM30	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM31	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM32	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM33	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM34	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM35	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM36	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM37	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM38	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM39	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM40	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM41	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM42	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM43	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM44	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM45	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM46	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM47	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM48	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM49	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM50	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM51	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM52	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM53	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM54	Calliphoridae	Pollenia		vagabunda (Meigen)	M	Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM55	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM56	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM57	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM58	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM59	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM60	Tachinidae					Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM61	Sarcophagidae	Agria		housei (Shewell)	F	Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM62	Sarcophagidae	Mehria		hini Aldrich	M	Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM63	Oestridae	Cephenemyia		sp.	M	Maine	Hancock	Acadia N.P.	Schoodic Head summit	VII.15.2006
AM64	Tachinidae					Maine	Hancock	Acadia N.P.	Jordan's Pond	VII.15.2006
AM65	Tachinidae					Maine	Hancock	Acadia N.P.	Jordan's Pond	VII.15.2006
AM66	Tachinidae					Maine	Hancock	Acadia N.P.	Dorr Mt. summit	VII.15.2006
AM67	Tachinidae					Maine	Hancock	Acadia N.P.	Dorr Mt. summit	VII.15.2006
AM68	Tachinidae					Maine	Hancock	Acadia N.P.	Dorr Mt. summit	VII.15.2006
AM69	Tachinidae					Maine	Hancock	Acadia N.P.	Dorr Mt. summit	VII.15.2006
AM70	Sarcophagidae	Boettcheria		latisterna Parker	M	Maine	Hancock	Acadia N.P.	Dorr Mt. summit	VII.15.2006
AM71	Tachinidae					Maine	Hancock	Acadia N.P.	Dorr Mt. summit	VII.15.2006
AM72	Tachinidae					Maine	Hancock	Acadia N.P.	Dorr Mt. summit	VII.15.2006
AM73	Tachinidae					Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM74	Calliphoridae	Calliphora		vomitiora (Linnaeus)	M	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM75	Calliphoridae	Calliphora		vomitiora (Linnaeus)	F	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM76	Calliphoridae	Calliphora		vomitiora (Linnaeus)	F	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM77	Calliphoridae	Calliphora		vomitiora (Linnaeus)	F	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM78	Calliphoridae	Calliphora		vomitiora (Linnaeus)	M	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM79	Calliphoridae	Calliphora		vomitiora (Linnaeus)	F	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM80	Calliphoridae	Calliphora		vomitiora (Linnaeus)	M	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AM81	Calliphoridae	Calliphora		vomitiora (Linnaeus)	M	Georgia	Rabun	Rabun Bald	Trail to summit	V.24.2006
AN01	Sarcophagidae	Sarcophaga	Liopygia	crassipalpis Macquart	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN02	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN03	Sarcophagidae	Sarcophaga	Mehria	pleomenda Reinhard	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN04	Sarcophagidae	Sarcophaga	Mehria	houghi Aldrich	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN05	Sarcophagidae	Blaesoxipha		sp.	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN06	Sarcophagidae	Macronychia		n.sp. (spinosa of Downes)	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN07	Sarcophagidae	Sarcophaga	Bercaeopsis	seagoi (Dodge)	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN08	Sarcophagidae	Sarcophaga		sp.	M	Georgia	Rabun	Rabun Bald	Summit 10:50-11:30	V.24.2006
AN09	Tachinidae					Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN10	Tachinidae					Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN11	Sarcophagidae	Ravinia		stimulans (Walker)	F	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN12	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN13	Sarcophagidae	Ravinia		derelicta (Walker)	F	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN14	Sarcophagidae	Ravinia		Iherminieri (Robineau-Desvoidy)	F	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN15	Sarcophagidae	Ravinia		Iherminieri (Robineau-Desvoidy)	M	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN16	Sarcophagidae	Ravinia		Iherminieri (Robineau-Desvoidy)	M	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN17	Sarcophagidae	Ravinia		Iherminieri (Robineau-Desvoidy)	M	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN18	Sarcophagidae	Ravinia		derelicta (Walker)	M	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN19	Sarcophagidae	Helicobia		rapax (Walker)	M	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN20	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	M	Tennessee	Jefferson	I-40 Rest Stop	near I-81 intersection	V.22.2006
AN21	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN22	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN23	Sarcophagidae	Oxysarcodexia		sp.	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN24	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN25	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN26	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN27	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN28	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN29	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN30	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN31	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN32	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN33	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN34	Sarcophagidae	Oxysarcodexia		sp.	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN35	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN36	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN37	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN38	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN39	Sarcophagidae	Oxysarcodexia		sp.	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN40	Sarcophagidae	Oxysarcodexia		sp.	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN41	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN42	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN43	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN44	Sarcophagidae	Ravinia		derelicta (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN45	Sarcophagidae	Ravinia		derelicta (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN46	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN47	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN48	Sarcophagidae	Oxysarcodexia		ventricosa (Wulp)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN49	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006
AN50	Sarcophagidae	Ravinia		stimulans (Walker)	F	South Carolina	Newberry	Sumter NF; Enoree Dist.	Firetower Area	V.23.2006

ID checked TW

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AP39	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP40	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP41	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP42	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP43	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP44	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP45	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP46	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP47	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP48	Calliphoridae	Phormia	regina (Meigen)		New York	Chautauqua	Long Point State Park		nr Bemus Pt.	VI.08.2007	
AP49	Tachinidae				New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP50	Tachinidae				New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP51	Tachinidae				New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP52	Tachinidae				New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP53	Tachinidae				New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP54	Tachinidae				New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP55	Muscidae	Musca	sp.		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP56	Muscidae	Musca	sp.		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP57	Calliphoridae	Calliphora	vomitiora (Linnaeus)		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP58	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP59	Calliphoridae	Lucilia	coeruleiviridis Macquart		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP60	Calliphoridae	Pollenia	pediculata Macquart		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP61	Calliphoridae	Lucilia	illustrius (Meigen)		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP62	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP63	Sarcophagidae	Macronychia	sp. 1 nigrifurcis of Downes		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP64	Calliphoridae	Calliphora	livida Hall		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP65	Sarcophagidae	Sarcophaga	aldrichi Parker		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP66	Sarcophagidae	Sarcophaga	nearctica Parker		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP67	Sarcophagidae	Oxysarcodexia	cingarus (Aldrich)		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP68	Sarcophagidae	Oxysarcodexia	cingarus (Aldrich)		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP69	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP70	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP71	Sarcophagidae	Helicobia	rapax (Walker)		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP72	Sarcophagidae	Sarcophaga	subvicina Rohdendorf		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP73	Calliphoridae	Lucilia	coeruleiviridis Macquart		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP74	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP75	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP76	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP77	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AP78	Sarcophagidae	Sarcophaga	sp.		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP79	Sarcophagidae	Boettcheria	latisterna Parker		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP80	Calliphoridae	Oxysarcodexia	sp.		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AP81	Sarcophagidae	Oxysarcodexia	sp.		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AQ01	Tachinidae	Musca	sp.		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AQ02	Muscidae	Musca	sp.		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AQ03	Muscidae	Musca	sp.		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AQ04	Muscidae	Musca	sp.		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AQ05	Sarcophagidae	Sarcophaga	Neobellieria		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AQ06	Sarcophagidae	Sarcophaga	Neobellieria		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AQ07	Calliphoridae	Calliphora	livida Hall		New York	Schuyler	Watkins Glen State Park			VI.09.2007	
AQ08	Sarcophagidae	Sarcophaga	nearctica Parker		M	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AQ09	Sarcophagidae	Sarcophaga	sp.		F	New York	Schuyler	Watkins Glen State Park		VI.09.2007	
AQ10	Tachinidae				New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ11	Tachinidae	Calliphora	vicina Robineau-Desvoidy		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ12	Tachinidae	Calliphora	vicina Robineau-Desvoidy		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ13	Tachinidae	Calliphora	vomitiora (Linnaeus)		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ14	Calliphoridae	Calliphora	vomitiora (Linnaeus)		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ15	Calliphoridae	Calliphora	vomitiora (Linnaeus)		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ16	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ17	Calliphoridae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ18	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ19	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ20	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ21	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ22	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ23	Sarcophagidae	Sarcophaga	aldrichi Parker		F	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ24	Sarcophagidae	Calliphora	vomitiora (Linnaeus)		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ25	Sarcophagidae	Sarcophaga	crassipalpis Macquart		M	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ26	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ27	Sarcophagidae	Sarcophaga	berceaia Wiedemann		M	New York	Rockland	Bear Mountain State Park		VI.10.2007	
AQ28	Tachinidae				New York	Rockland	Bear Mountain State Park			VI.10.2007	
AQ29	Calliphoridae	Calliphora	vomitiora (Linnaeus)		New York	Rockland	Bear Mountain State Park			VI.10.2007	
AQ30	Calliphoridae	Calliphora	vomitiora (Linnaeus)		New York	Rockland	Bear Mountain State Park			VI.10.2007	
AQ31	Calliphoridae	Cynomyia	cadaverina Robineau-Desvoidy		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ32	Calliphoridae	Cynomyia	cadaverina Robineau-Desvoidy		New York	Rockland	Bear Mountain State Park		summit of Bear Mt.	VI.10.2007	
AQ33	Calliphoridae	Pollenia	angustigena Wainwright		F	New York	Richmond	Clay Pit Ponds State Park		VI.11.2007	
AQ34	Sarcophagidae	Metopia	sp.		F	New York	Richmond	Clay Pit Ponds State Park		VI.11.2007	
AQ35	Sarcophagidae	Metopia	sp.		F	New York	Richmond	Clay Pit Ponds State Park		VI.11.2007	
AQ36	Sarcophagidae	Metopia	sp.		F	New York	Richmond	Clay Pit Ponds State Park		VI.11.2007	
AQ37	Sarcophagidae	Senotania	trilineata (Wulp)		F	New York	Richmond	Clay Pit Ponds State Park		VI.11.2007	
AQ38	Sarcophagidae	Senotania	trilineata (Wulp)		M	New York	Richmond	Clay Pit Ponds State Park		VI.11.2007	
AQ39	Sarcophagidae	Lucilia	sericata (Meigen)		New York	Suffolk	Heckscher State Park		Long Island; along beach	VI.11.2007	
AQ40	Sarcophagidae	Sarcophaga	johnsoni Aldrich		M	New York	Richmond	Heckscher State Park	Long Island; along beach	VI.11.2007	
AQ41	Tachinidae				New York	Richmond	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ42	Calliphoridae	Calliphora	vicina Robineau-Desvoidy		New York	Richmond	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ43	Tachinidae				New York	Richmond	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ44	Tachinidae				New York	Richmond	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ45	Tachinidae				New York	Richmond	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ46	Muscidae	Musca	sp.		New York	Richmond	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ47	Muscidae	Musca	sp.		New York	Richmond	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ48	Calliphoridae	Lucilia	sericata (Meigen)		New York	Dutchess	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ49	Sarcophagidae	Sarcophaga	sarracenioides Aldrich (?)		F	New York	Richmond	James Baird State Park		VI.12.2007	
AQ50	Sarcophagidae	Helicobia	rapax (Walker)		F	New York	Richmond	James Baird State Park		VI.12.2007	
AQ51	Calliphoridae	Phormia	regina (Meigen)		New York	Dutchess	James Baird State Park		E. of Poughkeepsie	VI.12.2007	
AQ52	Sarcophagidae	Sarcophaga	nearctica Parker		M	New York	Richmond	James Baird State Park		VI.12.2007	
AQ53	Sarcophagidae	Sarcophaga	nearctica Parker		M	New York	Richmond	James Baird State Park		VI.12.2007	
AQ54	Sarcophagidae	Lucilia	coeruleiviridis Macquart		New York	Dutchess	James Baird State Park		E. of Poughkeepsie	VI.13.2007	
AQ55	Tachinidae				New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ56	Tachinidae				New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ57	Sarcophagidae	Tripanurga	importuna (Walker)		F	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ58	Sarcophagidae	Tripanurga	importuna (Walker)		F	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ59	Sarcophagidae	Tripanurga	importuna (Walker)		F	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ60	Sarcophagidae	Metopia	sp.		F	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ61	Calliphoridae	Pollenia	rudis (Fabricius)		New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ62	Calliphoridae	Lucilia	sericata (Meigen)		New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ63	Calliphoridae	Phormia	regina (Meigen)		New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ64	Sarcophagidae	Sarcophaga	aldrichi Parker		M	New York	Onondaga	Green Lakes State Park		nr Syracuse	VI.14.2007
AQ65	Sarcophagidae	Sarcophaga	aldrichi Parker		M	New York	Onondaga	Green Lakes State Park		nr Syracuse	VI.14.2007
AQ66	Sarcophagidae	Helicobia	rapax (Walker)		M	New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007
AQ67	Sarcophagidae	Tripanurga	importuna (Walker)		M	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ68	Sarcophagidae	Tripanurga	importuna (Walker)		M	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ69	Sarcophagidae	Tripanurga	importuna (Walker)		M	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ70	Sarcophagidae	Tripanurga	importuna (Walker)		M	New York	Saratoga	Moreau Lake State Park		VI.13.2007	
AQ71	Tachinidae				New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ72	Tachinidae				New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ73	Tachinidae				New York	Saratoga	Moreau Lake State Park		SE of Glens Falls	VI.13.2007	
AQ74	Sarcophagidae	Metopia	sp.		F	New York	Saratoga	Moreau Lake State Park		VI.13.2007	

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AQ75	Sarcophagidae	Metopia	sp.	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AQ76	Sarcophagidae	Tripanurga	importuna (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AQ77	Sarcophagidae	Tripanurga	importuna (Walker)	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AQ78	Sarcophagidae	Oxysarcodexia	sp.	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AQ79	Calliphoridae	Pollenia	rudis (Fabricius)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	ID checked TW
AQ80	Sarcophagidae	Sarcophaga	aldrichi Parker	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AQ81	Sarcophagidae	Sarcophaga	aldrichi Parker	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR01	Sarcophagidae	Sarcophaga	Liosarcophaga saracenooides Aldrich	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR02	Sarcophagidae	Metopia	argyrocephala (Meigen)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR03	Sarcophagidae	Tripanurga	importuna (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR04	Sarcophagidae	Tripanurga	importuna (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR05	Calliphoridae	Calliphora	vomitoria (Linnaeus)	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR06	Calliphoridae	Calliphora	vomitiora (Linnaeus)	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR07	Calliphoridae	Phormia	regina (Meigen)	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR08	Calliphoridae	Phormia	regina (Meigen)	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR09	Calliphoridae	Phormia	regina (Meigen)	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR10	Calliphoridae	Phormia	regina (Meigen)	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR11	Calliphoridae	Pollenia	pediculata Macquart	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR12	Muscidae	New York		F	New York	Richmond	James Baird State Park	E. of Poughkeepsie	VI.12.2007	
AR13	Muscidae	New York		F	New York	Richmond	James Baird State Park	E. of Poughkeepsie	VI.12.2007	
AR14	Muscidae	New York		F	New York	Richmond	James Baird State Park	E. of Poughkeepsie	VI.12.2007	
AR15	Sarcophagidae	Sarcophaga	aldrichi Parker	F	New York	Richmond	James Baird State Park	E. of Poughkeepsie	VI.12.2007	
AR16	Sarcophagidae	Sarcophaga	aldrichi Parker	F	New York	Richmond	James Baird State Park	E. of Poughkeepsie	VI.12.2007	
AR17	Calliphoridae	Lucilia	coeruleovittata Macquart	M	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR18	Calliphoridae	Lucilia	sericata (Meigen)	F	New York	Dutchess	James Baird State Park	E. of Poughkeepsie	VI.12.2007	carriion
AR19	Tachinidae	New York		F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR20	Sarcophagidae	Sarcophaga	Neobellieria tripiliasia Wulp	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR21	Sarcophagidae	Sarcophaga	Robineauella nearctica Parker	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR22	Calliphoridae	Phormia	regina (Meigen)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR23	Calliphoridae	Lucilia	silvarum (Meigen)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR24	Sarcophagidae	Sarcophaga	aldrichi Parker	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR25	Sarcophagidae	Tripanurga	importuna (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR26	Sarcophagidae	Sarcophaga	bulbata Parker	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR27	Sarcophagidae	Sarcophaga	bulbata Parker	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR28	Sarcophagidae	Tripanurga	importuna (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR29	Sarcophagidae	Tripanurga	importuna (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR30	Sarcophagidae	Tripanurga	importuna (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR31	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR32	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR33	Sarcophagidae	Ravinia	stimulans (Walker)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR34	Sarcophagidae	Metopia	argyrocephala (Meigen)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR35	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR36	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR37	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR38	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR39	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR40	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR41	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR42	Sarcophagidae	Senotania	trilineata (Wulp)	M	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR43	Sarcophagidae	Sarcophaga	sp.	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	same specimen listed twice?
AR44	Sarcophagidae	Tripanurga	importuna (Walker)	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR45	Sarcophagidae	Sarcophaga	aldrichi Parker	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR46	Sarcophagidae	Tripanurga	importuna (Walker)	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR47	Sarcophagidae	Tripanurga	importuna (Walker)	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR48	Calliphoridae	Phormia	regina (Meigen)	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR49	Sarcophagidae	Ravinia	querula (Walker)	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR50	Muscidae	Musca	sp.	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR51	Sarcophagidae	Senotania	trilineata (Wulp)	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR52	Sarcophagidae	Metopia	sp.	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR53	Sarcophagidae	Sarcophaga	sp.	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR54	Sarcophagidae	Sarcophaga	aldrichi Parker	F	New York	Saratoga	Moreau Lake State Park	SE of Glens Falls	VI.13.2007	
AR55	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR56	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR57	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR58	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR59	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR60	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR61	Sarcophagidae	Sarcophaga	Liopygia crassipalpis Macquart	M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR62	Sarcophagidae	Sarcophaga	Liopygia crassipalpis Macquart	M	Wisconsin	Waukesha	Lapham Peak State Park	lookout tower	VI.30.2007	
AR63	Calliphoridae	Phormia	regina (Meigen)	M	Wisconsin	Waukesha	Lapham Peak State Park	VI.01.2007	carriion	
AR64	Sarcophagidae	Sarcophaga	Robineauella nearctica Parker	F	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AR65	Sarcophagidae	Lepidodexia	sp.	F	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AR66	Sarcophagidae	Brachicomma	sarcophagina (Townsend)	F	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AR67	Sarcophagidae	Sarcophaga	Liopygia crassipalpis Macquart	M	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AR68	Sarcophagidae	Brachicomma	sarcophagina (Townsend)	M	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AR69	Sarcophagidae	Brachicomma	sarcophagina (Townsend)	M	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AR70	Sarcophagidae	Ravinia	derelicta (Walker)	M	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AR71	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	ID checked
AR72	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AR73	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AR74	Calliphoridae	Pollenia	pediculata Macquart	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	ID checked TW
AR75	Calliphoridae	Pollenia	rudis (Fabricius)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AR76	Calliphoridae	Pollenia	rudis (Fabricius)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	ID checked TW
AR77	Calliphoridae	Pollenia	rudis (Fabricius)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	ID checked TW
AR78	Sarcophagidae	Sarcophaga	Robineauella nearctica Parker	F	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AR79	Sarcophagidae	Boettcheria	bisetososa Parker	F	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AR80	Sarcophagidae	Ravinia	stimulans (Walker)	F	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VI.03.2007	
AS01	Sarcophagidae	Boettcheria	latisterna Parker	F	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS02	Calliphoridae	Phormia	regina (Meigen)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS03	Calliphoridae	Phormia	regina (Meigen)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS04	Sarcophagidae	Metopia	inermis Allen (?)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS05	Sarcophagidae	Metopia	inermis Allen (?)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS06	Sarcophagidae	Metopia	inermis Allen (?)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS07	Sarcophagidae	Metopia	inermis Allen (?)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS08	Sarcophagidae	Metopia	inermis Allen (?)	M	Wisconsin	Waukesha	Lapham Peak State Park	along trails	VI.01.2007	
AS09	Oestridae	Cephenemyia	sp.	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS10	Oestridae	Cephenemyia	sp.	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS11	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower: tree top level	VI.01.2007	
AS12	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower: tree top level	VI.01.2007	
AS13	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower: tree top level	VI.01.2007	
AS14	Tachinidae	Wisconsin		M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower: tree top level	VI.01.2007	
AS15	Sarcophagidae	Macronychia	sp. 1 nigrifunus of Downes	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS16	Sarcophagidae	Macronychia	sp. 1 nigrifunus of Downes	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS17	Sarcophagidae	Macronychia	sp. 1 nigrifunus of Downes (?)	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS18	Sarcophagidae	Macronychia	sp. 1 nigrifunus of Downes	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS19	Sarcophagidae	Macronychia	sp. 1 nigrifunus of Downes	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS20	Sarcophagidae	Macronychia	sp. 1 nigrifunus of Downes	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS21	Sarcophagidae	Macronychia	sp. 1 nigrifunus of Downes	M	Wisconsin	Waukesha	Lapham Peak State Park	observation tower at tree top level	VI.01.2007	
AS22	Sarcophagidae	Sarcophaga	Liopygia crassipalpis Macquart	M	Wisconsin	Waukesha	Lapham Peak State Park	base of observation tower	VI.01.2007	
AS23	Tachinidae	Wisconsin		M	Wisconsin	Marathon	Rib Mountain State Park	along trails	VI.02.2007	
AS24	Tachinidae	Wisconsin		M	Wisconsin	Marathon	Rib Mountain State Park	along trails	VI.02.2007	
AS25	Sarcophagidae	Ravinia	querula (Walker)	F	Wisconsin	Marathon	Rib Mountain State Park	along trails	VI.02.2007	
AS26	Sarcophagidae	Ravinia	stimulans (Walker)	F	Wisconsin	Marathon	Rib Mountain State Park	along trails	VI.02.2007	
AS27	Sarcophagidae	Boettcheria	latisterna Parker	F	Wisconsin	Marathon	Rib Mountain State Park	along trails	VI.02.2007	
AS28	Tachinidae	Wisconsin		F	Wisconsin	Marathon	Rib Mountain State Park	along trails	VI.02.2007	
AS29	Sarcophagidae	Boettcheria	latisterna Parker	F	Wisconsin	Marathon	Rib Mountain State Park	along trails	VI.02.2007	

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AS30		Calliphora		vicina Robineau-Desvoidy	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS31		Calliphora		vomitoria (Linnaeus)	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS32		Calliphora		vomitoria (Linnaeus)	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS33	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS34	Calliphoridae	Musca		sp.	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS35	Calliphoridae	Musca		sp.	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS36	Sarcophagidae	Sarcophaga	Liopygia	sargostoma (Robineau-Desvoidy)	M	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007	
AS37	Sarcophagidae	Sarcophaga	Robineauella	nearctica Parker	M	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007	
AS38	Sarcophagidae	Sarcophaga	Robineauella	nearctica Parker	M	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007	
AS39	Sarcophagidae	Blaesoxipha		uncata (Wulp)	M	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007	
AS40	Calliphoridae	Calliphora	Servaisia	latifrons Hough	M	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007	
AS41	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS42	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS43	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS44	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS45	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS46	Tachinidae				Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS47	Tachinidae				Wisconsin	Marathon	Rib Mountain State Park	along trails	VII.02.2007		
AS48	Sarcophagidae	Sarcophaga	Bercaeopsis	idonea Aldrich	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS49	Sarcophagidae	Sarcophaga	Bercaeopsis	idonea Aldrich	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS50	Sarcophagidae	Emblemasoma		albicomis Reinhard	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS51	Sarcophagidae	Emblemasoma		albicomis Reinhard	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS52	Sarcophagidae	Macronychia		sp. 2 pulcra of Downes	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS53	Sarcophagidae	Macronychia		sp. 2 pulcra of Downes	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS54	Sarcophagidae	Brachicoma		sarcophagina (Townsend)	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS55	Sarcophagidae	Brachicoma		sarcophagina (Townsend)	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS56	Sarcophagidae	Macronychia		sp. 1 nigrifrons of Downes	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS57	Sarcophagidae	Macronychia		sp. 1 nigrifrons of Downes	M	Wisconsin	Marathon	Rib Mountain State Park	observation tower	VII.02.2007	
AS58	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007		
AS59	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007		
AS60	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007		
AS61	Calliphoridae	Phormia	Robineauella	regina (Meigen)	F	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007	
AS62	Calliphoridae	Phormia		nearctica Parker	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007		
AS63	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007		
AS64	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007		
AS65	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007		
AS66	Sarcophagidae	Helicobia		rapax (Walker)	M	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007	
AS67	Sarcophagidae	Ravinia		stimulus (Walker)	M	Wisconsin	Barron	Rest stop on US-53	dog dung & dead ground squirrel	VII.02.2007	
AS68	Sarcophagidae	Sarcophaga	Sarcophaga	subvicina Rohdendorf	M	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007	carrión
AS69	Sarcophagidae	Sarcophaga	Robineauella	nearctica Parker	F	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007	carrión
AS70	Calliphoridae	Pollenia		pediculata Macquart	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		ID checked TW
AS71	Calliphoridae	Pollenia		pediculata Macquart	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		ID checked TW
AS72	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		
AS73	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		
AS74	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		
AS75	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		
AS76	Calliphoridae	Lucilia		coeruleiviridis Macquart	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		carrión
AS77	Calliphoridae	Lucilia		illustris (Meigen)	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		ID checked TW
AS78	Calliphoridae	Lucilia		coeruleiviridis Macquart	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		ID checked TW
AS79	Calliphoridae	Lucilia		coeruleiviridis Macquart	Wisconsin	Waukesha	Lapham Peak State Park	nature trail	VII.01.2007		ID checked TW
AS80	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	waterfall area	VII.03.2007		
AS81	Calliphoridae	Lucilia		illustris (Meigen)	Wisconsin	Douglas	Amnicon Falls State Park	waterfall area	VII.03.2007		
AT01	Calliphoridae	Pollenia		pediculata Macquart	Wisconsin	Douglas	Amnicon Falls State Park	waterfall area	VII.03.2007		ID checked TW
AT02	Calliphoridae	Pollenia		labialis Robineau-Desvoidy	Wisconsin	Douglas	Amnicon Falls State Park	waterfall area	VII.03.2007		
AT03	Calliphoridae	Cynomyia		cadaverina Robineau-Desvoidy	Wisconsin	Douglas	Amnicon Falls State Park	waterfall area	VII.03.2007		
AT04	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	waterfall area	VII.03.2007		
AT05	Sarcophagidae	Tripanurga		sp.	F	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007	
AT06	Sarcophagidae	Sarcophaga		sp.	F	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007	
AT07	Sarcophagidae	Sarcophaga		sp.	F	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007	
AT08	Sarcophagidae	Boettcheria		latisterna Parker	F	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007	
AT09	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT10	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT11	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT12	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT13	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT14	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT15	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT16	Calliphoridae	Calliphora		vomitoria (Linnaeus)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT17	Calliphoridae	Cynomyia		cadaverina Robineau-Desvoidy	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT18	Calliphoridae	Phormia		regina (Meigen)	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007		
AT19	Calliphoridae	Phormia		regina (Meigen)	M	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007	
AT20	Calliphoridae	Sarcophaga	Wohlfahrtiopsis	utilis Aldrich	M	Wisconsin	Douglas	Amnicon Falls State Park	nature trail	VII.03.2007	
AT21	Tachinidae		Minnesota		Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007		
AT22	Tachinidae		Minnesota		Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007		
AT23	Tachinidae		Minnesota		Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007		
AT24	Tachinidae		Minnesota		Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007		
AT25	Tachinidae		Minnesota		Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007		
AT26	Tachinidae		Minnesota		Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007		
AT27	Sarcophagidae	Sarcophaga	Neobellieria	triplesia Wulp	M	Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007	
AT28	Sarcophagidae	Sarcophaga	Neobellieria	triplesia Wulp	M	Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007	
AT29	Sarcophagidae	Sarcophaga	Bercaeopsis	idonea Aldrich	M	Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007	
AT30	Sarcophagidae	Sarcophaga	Bercaeopsis	idonea Aldrich	M	Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007	
AT31	Sarcophagidae	Sarcophaga	Bercaeopsis	hesterna Reinhard	M	Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007	
AT32	Sarcophagidae	Sarcophaga	Bercaeopsis	hesterna Reinhard	M	Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007	
AT33	Sarcophagidae	Brachicoma		sarcophagina (Townsend)	M	Minnesota	Winona	Great River Bluffs State Park	along trails	VII.07.2007	
AT34	Tachinidae		Minnesota		Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007		
AT35	Tachinidae		Minnesota		Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007		
AT36	Sarcophagidae	Helicobia		rapax (Walker)	F	Minnesota	Kandiyohi	Sibley State Park	along trails	VII.06.2007	
AT37	Sarcophagidae	Metopia		sp.	F	Minnesota	Kandiyohi	Sibley State Park	along trails	VII.06.2007	
AT38	Calliphoridae	Phormia		regina (Meigen)	F	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT39	Calliphoridae	Phormia		regina (Meigen)	F	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT40	Tachinidae		Minnesota		Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007		
AT41	Tachinidae		Minnesota		Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007		
AT42	Tachinidae		Minnesota		Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007		
AT43	Tachinidae		Minnesota		Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007		
AT44	Sarcophagidae	Ravinia		anxia (Walker)	F	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT45	Sarcophagidae	Ravinia		anxia (Walker)	F	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT46	Sarcophagidae	Ravinia		anxia (Walker)	F	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT47	Sarcophagidae	Ravinia		anxia (Walker) (?)	F	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT48	Sarcophagidae	Ravinia		derelicta (Walker)	F	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT49	Sarcophagidae	Ravinia		anxia (Walker)	M	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT50	Sarcophagidae	Ravinia		anxia (Walker)	M	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT51	Sarcophagidae	Brachicoma		sarcophagina (Townsend)	M	Minnesota	Kandiyohi	Sibley State Park	horse trails and Mt. Tom	VII.06.2007	
AT52	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	North Dakota	Trail	Rest stop on I-29	south of Hillsboro	VII.05.2007	
AT53	Calliphoridae	Calliphora		vicina Robineau-Desvoidy	M	North Dakota	Trail	Rest stop on I-29	south of Hillsboro	VII.05.2007	
AT54	Calliphoridae	Lucilia		illustris (Meigen)	M	North Dakota	Trail	Rest stop on I-29	south of Hillsboro	VII.05.2007	
AT55	Calliphoridae	Lucilia		illustris (Meigen)	M	North Dakota	Trail	Rest stop on I-29	south of Hillsboro	VII.05.2007	
AT56	Calliphoridae	Lucilia		illustris (Meigen)	M	North Dakota	Trail	Rest stop on I-29	south of Hillsboro	VII.05.2007	
AT57	Calliphoridae	Lucilia		illustris (Meigen)	M	North Dakota	Trail	Rest stop on I-29	south of Hillsboro	VII.05.2007	

AZ47	Sarcophagidae	Oxysarcodexia	ventricosa (Wulp)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ48	Sarcophagidae	Oxysarcodexia	ventricosa (Wulp)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ49	Calliphoridae	Lucilia	sericata (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ50	Calliphoridae	Lucilia	sericata (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ51	Calliphoridae	Phormia	regina (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ52	Calliphoridae	Phormia	regina (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ53	Calliphoridae	Phormia	regina (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ54	Calliphoridae	Phormia	regina (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ55	Calliphoridae	Phormia	regina (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ56	Calliphoridae	Phormia	regina (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ57	Calliphoridae	Phormia	regina (Meigen)		West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	
AZ58	Sarcophagidae	Ravinia	therminieri (Robineau-Desvoidy)?	M	West Virginia	Kanawha	Rest area on I-64I-77	south of Standard	VI.19.2008	ID checked
AZ59	Tachinidae				Virginia	Bland	Welcome rest stop on I-77		VI.19.2008	
AZ60	Sarcophagidae	Ravinia	stimulans (Walker)	F	Virginia	Bland	Welcome rest stop on I-77		VI.19.2008	
AZ61	Sarcophagidae	Ravinia	querula (Walker)	M	Virginia	Bland	Welcome rest stop on I-77		VI.19.2008	
AZ62	Sarcophagidae	Ravinia	dereicta (Walker)		Virginia	Bland	Welcome rest stop on I-77		VI.19.2008	
AZ63	Calliphoridae	Pollenia	rudis (Fabricius)		North Carolina	Durham	Duke University campus		VI.20.2008	
AZ64	Sarcophagidae	Mecynocorpus	salvum (Aldrich)	M	North Carolina	Durham	Duke University campus		VI.20.2008	
AZ65	Sarcophagidae	Helicobia	rapax (Walker)		North Carolina	Orange	Duke University campus		VI.21.2008	
AZ66	Tachinidae				North Carolina	Orange	Motel parking lot border		VI.21.2008	
AZ67	Sarcophagidae	Helicobia	rapax (Walker)		North Carolina	Orange	Motel parking lot border		VI.21.2008	
AZ68	Sarcophagidae	Helicobia	rapax (Walker)		North Carolina	Orange	Motel parking lot border		VI.21.2008	
AZ69	Tachinidae				North Carolina	Davie	Rest stop on I-40		VI.21.2008	
AZ70	Calliphoridae	Lucilia	sericata (Meigen)		North Carolina	Davie	Rest stop on I-40		VI.21.2008	
AZ71	Calliphoridae	Lucilia	sericata (Meigen)		North Carolina	Davie	Rest stop on I-40		VI.21.2008	
AZ72	Calliphoridae	Lucilia	coeruleiviridis Macquart		North Carolina	Davie	Rest stop on I-40		VI.21.2008	
AZ73	Calliphoridae	Calliphora	terraenovae Macquart		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ74	Calliphoridae	Calliphora	vornitoria (Linnaeus) (?)		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ75	Calliphoridae	Calliphora	terraenovae Macquart		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ76	Calliphoridae	Calliphora	terraenovae Macquart		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ77	Calliphoridae	Calliphora	latifrons Hough		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ78	Calliphoridae	Calliphora	livida Hall		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ79	Calliphoridae	Calliphora	latifrons Hough		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ80	Calliphoridae	Calliphora	livida Hall		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
AZ81	Calliphoridae	Calliphora	livida Hall		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA01	Calliphoridae	Calliphora	livida Hall		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA02	Calliphoridae	Calliphora	latifrons Hough		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA03	Calliphoridae	Calliphora	livida Hall or coloradensis Hough	F	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA04	Calliphoridae	Calliphora	coloradensis Hough		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA05	Calliphoridae	Calliphora	latifrons Hough		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA06	Calliphoridae	Calliphora	latifrons Hough (?)		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA07	Calliphoridae	Calliphora	latifrons Hough		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA08	Calliphoridae	Calliphora	latifrons Hough (?)		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA09	Calliphoridae	Calliphora	latifrons Hough		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA10	Calliphoridae	Pollenia	pediculata Macquart		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA11	Calliphoridae	Pollenia	pediculata Macquart		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA12	Calliphoridae	Pollenia	pediculata Macquart		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA13	Calliphoridae	Pollenia	pediculata Macquart		New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA14	Sarcophagidae	Boettcheria	cimbicis (Townsend)	M	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA15	Sarcophagidae	Ravinia	vagabunda	M	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA16	Sarcophagidae	Ravinia	querula (Walker)	M	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA17	Sarcophagidae	Ravinia	vagabunda	M	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA18	Sarcophagidae	Ravinia	pusiola (Wulp)	M	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA19	Sarcophagidae	Ravinia	pusiola (Wulp)	M	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA20	Sarcophagidae	Sarcophaga	bercea	M	New Mexico	Grant	Gila National Forest	Cherry Creek Forest Camp	VIII.13.2007	carrion
BA21	Sarcophagidae	Oxysarcodexia	ventricosa (Wulp)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA22	Sarcophagidae	Oxysarcodexia	ventricosa (Wulp)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA23	Sarcophagidae	Helicobia	rapax (Walker)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA24	Sarcophagidae	Helicobia	rapax (Walker)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA25	Sarcophagidae	Boettcheria	cimbicis (Townsend)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA26	Sarcophagidae	Boettcheria	cimbicis (Townsend)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA27	Sarcophagidae	Ravinia	stimulans (Walker)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA28	Sarcophagidae	Ravinia	stimulans (Walker)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA29	Sarcophagidae	Ravinia	dereicta (Walker)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA30	Sarcophagidae	Ravinia	dereicta (Walker)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA31	Sarcophagidae	Mecynocorpus	salvum (Aldrich)	M	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA32	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA33	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA34	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA35	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA36	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA37	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA38	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA39	Tachinidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA40	Muscidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA41	Muscidae				Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA42	Calliphoridae	Pollenia	rudis (Fabricius)	F	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA43	Calliphoridae	Pollenia	rudis (Fabricius)	F	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA44	Calliphoridae	Lucilia	coeruleiviridis Macquart	F	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA45	Calliphoridae	Lucilia	coeruleiviridis Macquart	F	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA46	Calliphoridae	Lucilia	illustis (Meigen)	F	Ohio	Hamilton	Anderson Twp.		VII.6.2008	Malaise
BA47	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA48	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA49	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA50	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA51	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA52	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA53	Calliphoridae	Phormia	regina (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA54	Calliphoridae	Phormia	regina (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA55	Calliphoridae	Phormia	regina (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA56	Calliphoridae	Phormia	regina (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA57	Calliphoridae	Phormia	regina (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA58	Calliphoridae	Phormia	regina (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA59	Calliphoridae	Lucilia	illustis (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA60	Calliphoridae	Lucilia	illustis (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA61	Calliphoridae	Lucilia	illustis (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA62	Calliphoridae	Lucilia	illustis (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA63	Calliphoridae	Lucilia	illustis (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA64	Calliphoridae	Lucilia	illustis (Meigen)	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA65	Calliphoridae	Lucilia	illustis (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA66	Calliphoridae	Lucilia	illustis (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA67	Calliphoridae	Lucilia	illustis (Meigen)	M	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA68	Calliphoridae	Lucilia	coeruleiviridis Macquart	F	Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA69	Sarcophagidae	Sarcophaga	Neobellieria		Minnesota	Winona	Great River Bluffs State Park		VII.7.2007	carrion
BA70	Calliphoridae	Cochliomyia	macellaria (Fabricius)	M	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA71	Calliphoridae	Cochliomyia	macellaria (Fabricius)	F	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA72	Calliphoridae	Pollenia	pediculata Macquart	F	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA73	Calliphoridae	Pollenia	pediculata Macquart	F	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA74	Muscidae				Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA75	Muscidae				Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA76	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA77	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA78	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA79	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA80	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BA81	Calliphoridae	Phormia	regina (Meigen)	M	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion
BB01	Calliphoridae	Phormia	regina (Meigen)	F	Minnesota	Kandiyohi	Sibley State Park		VII.6.2007	carrion

BD74	Calliphoridae	Phormia		regina (Meigen)	M	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BD75	Calliphoridae	Phormia		regina (Meigen)	F	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BD76	Calliphoridae	Phormia		regina (Meigen)	F	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BD77	Calliphoridae	Phormia		regina (Meigen)	F	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BD78	Calliphoridae	Phormia		regina (Meigen)	F	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BD79	Calliphoridae	Phormia		regina (Meigen)	F	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BD80	Calliphoridae	Phormia		regina (Meigen)	F	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BD81	Muscidae				F	New York	Richmond	Clay Pit Ponds State Park	Staten Island	VI.11.2007	carrión
BE01	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	Kentucky	Campbell	Highland Heights		IX.25.2008	reared from chicken
BE02	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	Kentucky	Kenton	Erlanger		X.7.2008	reared from chicken
BE03	Sarcophagidae	Sarcophaga		sericata (Meigen)	F	New York	Hamilton	Anderson Twp.		VI.1.2008	Malaise
BE04	Sarcophagidae	Sarcophaga		aldrichi Parker	M	Ohio	Hamilton	Anderson Twp.		V.1.2008	Malaise
BE05	Calliphoridae	Lucilia		coeruleiviridis Macquart	F	Ohio	Hamilton	Anderson Twp.		V.1.2008	Malaise
BE06	Calliphoridae	Lucilia		coeruleiviridis Macquart	F	Ohio	Hamilton	Anderson Twp.		V.1.2008	Malaise
BE07	Calliphoridae	Phormia		regina (Meigen)	F	New York	Schuyler	Watkins Glen State Park		VI.9.2007	carrión
BE08	Calliphoridae	Phormia		regina (Meigen)	F	New York	Schuyler	Watkins Glen State Park		VI.9.2007	carrión
BE09	Calliphoridae	Lucilia		illustis (Meigen)	F	New York	Schuyler	Watkins Glen State Park		VI.9.2007	carrión
BE10	Calliphoridae	Lucilia		illustis (Meigen)	F	New York	Schuyler	Watkins Glen State Park		VI.9.2007	carrión
BE11	Calliphoridae	Lucilia		coeruleiviridis Macquart	F	New York	Schuyler	Watkins Glen State Park		VI.9.2007	carrión
BE12	Calliphoridae	Lucilia		sericata (Meigen)	F	New York	Schuyler	Watkins Glen State Park		VI.9.2007	carrión
BE13	Calliphoridae	Calliphora		vomitoria (Linnaeus)	M	New York	Schuyler	Watkins Glen State Park		VI.9.2007	carrión
BE14	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	Kentucky	Campbell	Highland Heights		IX.25.2008	reared from chicken
BE15	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	Kentucky	Campbell	Highland Heights		IX.25.2008	reared from chicken
BE16	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	F	Kentucky	Campbell	Highland Heights		IX.25.2008	reared from chicken
BE17	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	M	Kentucky	Kenton	Erlanger		X.7.2008	reared from chicken
BE18	Sarcophagidae	Sarcophaga	Liopygia	argyrostoma (Robineau-Desvoidy)	F	Kentucky	Kenton	Erlanger		X.7.2008	reared from chicken
BE19	Calliphoridae	Chrysomya		megacephala (Fabricius)	M	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE20	Calliphoridae	Chrysomya		megacephala (Fabricius)	M	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE21	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE22	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE23	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE24	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE25	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE26	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE27	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE28	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE29	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE30	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE31	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE32	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE33	Calliphoridae	Lucilia		cuprina Wiedemann	M	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE34	Calliphoridae	Lucilia		cuprina Wiedemann	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE35	Calliphoridae	Lucilia		cuprina Wiedemann	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE36	Muscidae	Musca		sp.	F	Texas	Cameron	Howard Johnson Hotel		VIII.07.2007	carrión
BE37	Sarcophagidae	Sarcophaga	Neobellieria	libera Aldrich	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE38	Sarcophagidae	Sarcophaga	Neobellieria	libera Aldrich	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE39	Sarcophagidae	Microcerella		scrofa (Aldrich)	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE40	Sarcophagidae		Bercaeopsis	pulla Aldrich	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE41	Sarcophagidae	Mecynocorpus		salvum (Aldrich)	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE42	Sarcophagidae			sp.	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE43	Sarcophagidae	Helicobia		rapax (Walker)	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE44	Sarcophagidae	Titanogrypa		metampyga (Aldrich)	M	Ohio	Adams	Edge of Appalachia Preserve		V.22.2009	
BE45	Sarcophagidae				M	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE46	Sarcophagidae				F	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE47	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE48	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE49	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE50	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE51	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE52	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Fort Bend	Brazos Bend State Park		VIII.08.2007	carrión
BE53	Sarcophagidae				M	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE54	Sarcophagidae				F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE55	Sarcophagidae				F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE56	Sarcophagidae				F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE57	Sarcophagidae				F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE58	Sarcophagidae				F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE59	Sarcophagidae				M	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE60	Sarcophagidae				M	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE61	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE62	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE63	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE64	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE65	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE66	Calliphoridae	Lucilia		mexicana Macquart	F	Texas	Kimble	South Llano River State Park	Junction	VIII.06.2007	carrión
BE67	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Jackson	Lake Texana State Park		VIII.07.2007	carrión ID checked TW
BE68	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Jackson	Lake Texana State Park		VIII.07.2007	carrión
BE69	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Jackson	Lake Texana State Park		VIII.07.2007	carrión
BE70	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Jackson	Lake Texana State Park		VIII.07.2007	carrión
BE71	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Jackson	Lake Texana State Park		VIII.07.2007	carrión
BE72	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Jackson	Lake Texana State Park		VIII.07.2007	carrión
BE73	Muscidae	Musca		sp.	F	Texas	Jackson	Lake Texana State Park		VIII.07.2007	carrión
BE74	Muscidae	Musca		domestica (Linnaeus)	M	Texas	McMullen	Choque Canyon State Park		VIII.06.2007	carrión
BE75	Muscidae	Musca		domestica (Linnaeus)	F	Texas	McMullen	Choque Canyon State Park		VIII.06.2007	carrión
BE76	Muscidae	Musca		domestica (Linnaeus)	M	Texas	McMullen	Choque Canyon State Park		VIII.06.2007	carrión
BE77	Muscidae	Musca		domestica (Linnaeus)	F	Texas	McMullen	Choque Canyon State Park		VIII.06.2007	carrión
BE78	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	McMullen	Choque Canyon State Park		VIII.06.2007	carrión
BE79	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	McMullen	Choque Canyon State Park		VIII.06.2007	carrión
BE80	Sarcophagidae	Musca			M	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BE81	Sarcophagidae	Musca			M	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF01	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF02	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF03	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF04	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF05	Calliphoridae	Phormia		regina (Meigen)	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF06	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF07	Calliphoridae	Chrysomya		ruffiacis (Macquart)	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF08	Calliphoridae	Chrysomya		mexicana Macquart or eximia ?	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF09	Calliphoridae	Lucilia		cuprina Wiedemann	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF10	Calliphoridae	Lucilia		cuprina Wiedemann	M	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF11	Calliphoridae	Lucilia		cuprina Wiedemann	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF12	Calliphoridae	Lucilia		cuprina Wiedemann	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF13	Calliphoridae	Lucilia		cuprina Wiedemann	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF14	Calliphoridae	Lucilia		cuprina Wiedemann	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF15	Calliphoridae	Lucilia		sericata (Meigen)	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF16	Calliphoridae	Lucilia		sericata or cuprina	F	Texas	Dallas	Dallas	Canine Country Club	VIII.09.2007	carrión
BF17	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF18	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF19	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF20	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF21	Calliphoridae	Chrysomya		megacephala (Fabricius)	M	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF22	Calliphoridae	Chrysomya		megacephala (Fabricius)	M	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF23	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF24	Calliphoridae	Chrysomya		megacephala (Fabricius)	F	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF25	Muscidae	Musca		sp.	F	Texas	Kenedy	Roadside Rest on Highway 77	S. of Corpus Christi	VIII.07.2007	carrión
BF26	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Ellis	Ennis	Jacob Way's house	VIII.10.2007	carrión
BF27	Calliphoridae	Cochliomyia		macellaria (Fabricius)	M	Texas	Ellis	Ennis	Jacob Way's house	VIII.10.2007	carrión
BF28	Calliphoridae	Cochliomyia		macellaria (Fabricius)	F	Texas	Ellis	Ennis	Jacob Way's house	VIII.10.2007	carrión

This resource was prepared by the author(s) using Federal funds provided by the U.S. Department of Justice. Opinions or points of view expressed are those of the author(s) and do not necessarily reflect the official position or policies of the U.S. Department of Justice.

BG65	Sarcophagidae	Sarcophaga	Neobellieria	bullata Parker	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG66	Sarcophagidae	Blaesoxipha	Gigantotheca	plinthopyga (Wiedemann)	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG67	Sarcophagidae	Oxy sarcodexia		ventricosa (Wied.)	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG68	Sarcophagidae	Sarcodexia		lambens (Wiedemann)	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG69	Calliphoridae	Lucilia		cuprina Wiedemann	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG70	Sarcophagidae	Tricharea		sp.	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG71	Sarcophagidae	Tricharea		sp.	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG72	Calliphoridae	Lucilia		cuprina Wiedemann	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG73	Calliphoridae	Lucilia		cuprina Wiedemann	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG74	Calliphoridae	Lucilia		cuprina Wiedemann	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG75	Muscidae	Musca		domestica (Linnaeus)	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG76	Muscidae	Musca		domestica (Linnaeus)	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG77	Muscidae	Musca		domestica (Linnaeus)	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG78	Muscidae	Musca		domestica (Linnaeus)	M	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG79	Sarcophagidae	Sarcodexia		lambens (Wiedemann)?	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG80	Sarcophagidae	Sarcodexia		lambens (Wiedemann)?	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BG81	Sarcophagidae	Helicobia		sp.	F	Florida	Broward	John U. Lloyd State Park	Fl. Lauderdale	VII.13.2009	
BH01	Calliphoridae	Lucilia		sericata (Meigen)	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH02	Calliphoridae	Lucilia		sericata (Meigen)	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH03	Calliphoridae	Lucilia		sericata (Meigen)	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH04	Calliphoridae	Lucilia		sericata (Meigen)	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH05	Calliphoridae	Lucilia		mexicana Macquart	M	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH06	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH07	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH08	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH09	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH10	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH11	Calliphoridae	Lucilia		mexicana Macquart	M	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH12	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH13	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH14	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH15	Calliphoridae	Lucilia		mexicana Macquart	M	California	Los Angeles	Angeles National Forest	Gold Creek Rd. 1 mi E of Little Tujunga Cyn. Rd.	VIII.23.2009	
BH16	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH17	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH18	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH19	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH20	Calliphoridae	Lucilia		mexicana Macquart	M	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH21	Calliphoridae	Lucilia		mexicana Macquart	M	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH22	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH23	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH24	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH25	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH26	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Leo Carrillo State Beach	Arroyo Sequit Cr.	IX.7.2009	rotting tilapia fish
BH27	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH28	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH29	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH30	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH31	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH32	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH33	Calliphoridae	Lucilia		mexicana Macquart	M	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH34	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH35	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BH36	Calliphoridae	Lucilia		mexicana Macquart	F	California	Los Angeles	Tarzana Caballero Cyn.	Santa Monica Mts	VIII.30.2009	rotting tilapia fish
BI01	Calliphoridae	Lucilia		thatusa	F	California	Del norte	Botanical Trail	trail	VI.8.2009	Whitworth donation
BI02	Calliphoridae	Lucilia		thatusa	F	California	Del norte	Botanical Trail	trail	VI.8.2009	Whitworth donation
BI03	Calliphoridae	Calliphora		loewi	F	Alaska	Denali Park	Denali Park	Nenana River	VIII.10.2008	Whitworth donation
BI04	Calliphoridae	Calliphora		loewi	F	Alaska	Denali Park	Denali Park	Nenana River	VIII.10.2008	Whitworth donation
BI05	Calliphoridae	Protocalliphora		bennetti	F						Whitworth donation
BI06	Calliphoridae	Protocalliphora		bennetti	F						Whitworth donation
BI07	Calliphoridae	Protocalliphora		sialia	F						Whitworth donation
BI08	Calliphoridae	Protocalliphora		sialia	F						Whitworth donation
BI09	Calliphoridae	Protocalliphora		sialia	F						Whitworth donation
BI10	Calliphoridae	Protocalliphora		rugosa	F						Whitworth donation
BI11	Calliphoridae	Trypocalliphora		braueri	F						Whitworth donation
BI12	Calliphoridae	Protocalliphora		occidentalis	F						Whitworth donation
BI13	Calliphoridae	Protocalliphora		occidentalis	F						Whitworth donation
BI14	Calliphoridae	Protocalliphora		occidentalis	F						Whitworth donation
OH01	Sarcophagidae	Eumacronychia		sp.	M	Florida	Martin	Jonathon Dickinson State Park	trail	V.13.2006	Whitworth donation

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Table 4. List of specimen codes for samples included in the final Phase II reference database.

Family Calliphoridae		Family Sarcophagidae	
Code	Species	Code	Species
AG53	<i>C. coloradensis</i>	AM10	<i>Ag. housei</i>
AJ79	<i>C. coloradensis</i>	AY09	<i>B. alcedo</i>
AZ01	<i>C. coloradensis</i>	AW32	<i>B. arizona</i>
BA04	<i>C. coloradensis</i>	AW36	<i>B. arizona</i>
BF63	<i>C. coloradensis</i>	AW29	<i>B. caridei</i>
AJ02	<i>C. grahami</i>	AW27	<i>B. cessator</i>
AJ10	<i>C. grahami</i>	AY05	<i>B. eleodis</i>
AF14	<i>C. latifrons</i>	AC79	<i>B. plinthopyga</i>
AF15	<i>C. latifrons</i>	AE50	<i>B. plinthopyga</i>
AG16	<i>C. latifrons</i>	AG29	<i>B. plinthopyga</i>
AG22	<i>C. latifrons</i>	AH77	<i>B. plinthopyga</i>
AG37	<i>C. latifrons</i>	AK02	<i>B. plinthopyga</i>
AK48	<i>C. latifrons</i>	BG46	<i>B. plinthopyga</i>
AS40	<i>C. latifrons</i>	BG47	<i>B. plinthopyga</i>
AZ31	<i>C. latifrons</i>	AJ80	<i>B. sp</i>
AZ33	<i>C. latifrons</i>	AJ81	<i>B. sp</i>
BA02	<i>C. latifrons</i>	AK01	<i>B. sp</i>
BA05	<i>C. latifrons</i>	A37	<i>Bo. bisetosa</i>
BA09	<i>C. latifrons</i>	E35	<i>Bo. cimbicis</i>
BG58	<i>C. latifrons</i>	AA11	<i>Bo. latisterna</i>
AH57	<i>C. livida</i>	AA12	<i>Bo. latisterna</i>
AH66	<i>C. livida</i>	AL26	<i>Bo. latisterna</i>
AP64	<i>C. livida</i>	AM70	<i>Bo. latisterna</i>
AQ07	<i>C. livida</i>	AR81	<i>Bo. latisterna</i>
AZ32	<i>C. livida</i>	AW17	<i>Bo. latisterna</i>
BA01	<i>C. livida</i>	E22	<i>Bo. latisterna</i>
BI03	<i>C. loewi</i>	E49	<i>Bo. latisterna</i>
BI04	<i>C. loewi</i>	E53	<i>Bo. latisterna</i>
AF65	<i>C. terranovae</i>	AU69	<i>Bo. latsiterna</i>
AJ15	<i>C. terranovae</i>	AG26	<i>Bo. litorosa</i>
AK47	<i>C. terranovae</i>	AR69	<i>Br. sarcophagina</i>
AF19	<i>C. vicina</i>	AS51	<i>Em. albicoma</i>
AG19	<i>C. vicina</i>	E76	<i>Fl. fletcheri</i>
AH61	<i>C. vicina</i>	AP71	<i>H. rapax</i>
AK45	<i>C. vicina</i>	AQ50	<i>H. rapax</i>
AK70	<i>C. vicina</i>	AS66	<i>H. rapax</i>
AM11	<i>C. vicina</i>	AT36	<i>H. rapax</i>
AM12	<i>C. vicina</i>	AU12	<i>H. rapax</i>
AP58	<i>C. vicina</i>	AU34	<i>H. rapax</i>
AP69	<i>C. vicina</i>	AU72	<i>H. rapax</i>
AS41	<i>C. vicina</i>	AW05	<i>H. rapax</i>
AT53	<i>C. vicina</i>	AZ65	<i>H. rapax</i>
AV57	<i>C. vicina</i>	AZ68	<i>H. rapax</i>
AV70	<i>C. vicina</i>	BA23	<i>H. rapax</i>
AV74	<i>C. vicina</i>	BA24	<i>H. rapax</i>
BB52	<i>C. vicina</i>	E4	<i>H. rapax</i>
AF11	<i>C. vomitoria</i>	E55	<i>H. rapax</i>

AF63	C. vomitoria	E77	H. rapax
AK68	C. vomitoria	AZ64	Me. salvum
AM27	C. vomitoria	BA31	Me. salvum
AM75	C. vomitoria	AV26	Mi. adelphe
AM77	C. vomitoria	AP68	Ox. cingarus
AO01	C. vomitoria	AA23	Ox. ventricosa
AO02	C. vomitoria	AA37	Ox. ventricosa
AP08	C. vomitoria	AA59	Ox. ventricosa
AP57	C. vomitoria	AB44	Ox. ventricosa
AQ29	C. vomitoria	AB48	Ox. ventricosa
AS31	C. vomitoria	AC72	Ox. ventricosa
AS32	C. vomitoria	AC73	Ox. ventricosa
AT10	C. vomitoria	AD56	Ox. ventricosa
BB53	C. vomitoria	AZ47	Ox. ventricosa
BC50	C. vomitoria	BA21	Ox. ventricosa
BE19	Ch. megacephala	BD59	Ox. ventricosa
BE20	Ch. megacephala	E62	Ox. ventricosa
BF21	Ch. megacephala	E8	Ox. ventricosa
BG23	Ch. megacephala	AE81	R. anxia
BG24	Ch. megacephala	AT45	R. anxia
BG59	Ch. megacephala	AT50	R. anxia
BG60	Ch. megacephala	AT48	R. derelicta
AA64	Ch. ruffacies	AZ09	R. derelicta
AC70	Ch. ruffacies	AZ62	R. derelicta
AC77	Ch. ruffacies	BA29	R. derelicta
AC78	Ch. ruffacies	BA30	R. derelicta
AD19	Ch. ruffacies	BB21	R. derelicta
AD43	Ch. ruffacies	AA01	R. floridensis
AY76	Ch. ruffacies	AA03	R. floridensis
AY78	Ch. ruffacies	AA60	R. floridensis
BE50	Ch. ruffacies	A22	R. lherminieri
BE65	Ch. ruffacies	AA36	R. lherminieri
BE72	Ch. ruffacies	AN14	R. lherminieri
BF07	Ch. ruffacies	AN15	R. lherminieri
BF40	Ch. ruffacies	AN16	R. lherminieri
BF49	Ch. ruffacies	AN17	R. lherminieri
AA28	Co. macellaria	AQ58	R. lherminieri
AA29	Co. macellaria	AZ44	R. lherminieri
AA30	Co. macellaria	AZ58	R. lherminieri
AA49	Co. macellaria	AE45	R. planifrons
AA50	Co. macellaria	AF05	R. planifrons
AA51	Co. macellaria	AF06	R. planifrons
AB39	Co. macellaria	AK08	R. planifrons
AB61	Co. macellaria	AW64	R. pusiola
AB67	Co. macellaria	AW73	R. pusiola
AD22	Co. macellaria	AX61	R. pusiola
AE09	Co. macellaria	AX62	R. pusiola
AE10	Co. macellaria	AX63	R. pusiola
AE27	Co. macellaria	AY36	R. pusiola
AE29	Co. macellaria	AZ05	R. pusiola

AT52	Co. macellaria	BA18	R. pusiola
AT70	Co. macellaria	BA19	R. pusiola
AY46	Co. macellaria	AW30	R. querula
AY48	Co. macellaria	AZ61	R. querula
AY50	Co. macellaria	BA16	R. querula
AY54	Co. macellaria	AR33	R. stimulans
BA70	Co. macellaria	AZ10	R. stimulans
BE27	Co. macellaria	AZ45	R. stimulans
BE28	Co. macellaria	AZ46	R. stimulans
BE47	Co. macellaria	AZ60	R. stimulans
BE67	Co. macellaria	BA27	R. stimulans
BE78	Co. macellaria	E7	R. stimulans
BF02	Co. macellaria	BA15	R. vagabunda
BF44	Co. macellaria	BA17	R. vagabunda
BF52	Co. macellaria	AL34	S. acrophila
AG15	Cy. cadaverina	AJ12	S. africa
AM25	Cy. cadaverina	AJ13	S. africa
AM26	Cy. cadaverina	AL31	S. africa
AQ31	Cy. cadaverina	AL31	S. africa
AT03	Cy. cadaverina	AW42	S. africa
AT17	Cy. cadaverina	AW43	S. africa
AV55	Cy. cadaverina	BA20	S. africa
AV69	Cy. cadaverina	AV03	S. aldrichi
E40	Cy. Cadaverina	E23	S. aldrichi
AD18	L. cluvia	AE73	S. argyrostoma
AD20	L. cluvia	AE74	S. argyrostoma
AD49	L. cluvia	AE76	S. argyrostoma
BF79	L. cluvia	AS36	S. argyrostoma
AB02	L. coeruleiviridis	BE01	S. argyrostoma
AB04	L. coeruleiviridis	BE02	S. argyrostoma
AB37	L. coeruleiviridis	BE14	S. argyrostoma
AB38	L. coeruleiviridis	BE16	S. argyrostoma
AB66	L. coeruleiviridis	BE18	S. argyrostoma
AB70	L. coeruleiviridis	AA02	S. bullata
AP59	L. coeruleiviridis	AA45	S. bullata
AP73	L. coeruleiviridis	AA46	S. bullata
AP74	L. coeruleiviridis	AA48	S. bullata
AR17	L. coeruleiviridis	AA57	S. bullata
AS76	L. coeruleiviridis	AB07	S. bullata
AS78	L. coeruleiviridis	AC71	S. bullata
AS79	L. coeruleiviridis	AD09	S. bullata
AU64	L. coeruleiviridis	AD70	S. bullata
AV59	L. coeruleiviridis	AE39	S. bullata
AW01	L. coeruleiviridis	AF37	S. bullata
AW14	L. coeruleiviridis	AQ05	S. bullata
AX13	L. coeruleiviridis	AQ06	S. bullata
AX16	L. coeruleiviridis	AR26	S. bullata
AX19	L. coeruleiviridis	AR27	S. bullata
BA68	L. coeruleiviridis	BA69	S. bullata
AA05	L. cuprina	BB19	S. bullata

AA18	L. cuprina	BB20	S. bullata
AA20	L. cuprina	BC10	S. bullata
AA21	L. cuprina	BC12	S. bullata
AA34	L. cuprina	BC14	S. bullata
AA40	L. cuprina	BG64	S. bullata
AA42	L. cuprina	BG65	S. bullata
AE61	L. cuprina	AF61	S. cooleyi
BE33	L. cuprina	AH76	S. cooleyi
BE34	L. cuprina	AJ17	S. cooleyi
BE35	L. cuprina	AL12	S. crassipalpis
BF09	L. cuprina	AL33	S. crassipalpis
BF10	L. cuprina	AL35	S. crassipalpis
BF11	L. cuprina	AQ25	S. crassipalpis
BF12	L. cuprina	AR61	S. crassipalpis
BF13	L. cuprina	E74	S. crassipalpis
BG02	L. cuprina	AB10	S. georgiana
BG72	L. cuprina	AV61	S. hinei
AG72	L. elongata	AO50	S. houghi
AG76	L. elongata	AT30	S. idonea
AH37	L. elongata	AQ40	S. johnsoni
AF20	L. illustris	E39	S. mimoris
AG59	L. illustris	AH21	S. nearctica
AG63	L. illustris	AQ52	S. nearctica
AG65	L. illustris	AQ53	S. nearctica
AH38	L. illustris	AR21	S. nearctica
AM28	L. illustris	AR64	S. nearctica
AO24	L. illustris	AR78	S. nearctica
AP13	L. illustris	AS37	S. nearctica
AP15	L. illustris	AS38	S. nearctica
AS81	L. illustris	AS62	S. nearctica
AT54	L. illustris	AU24	S. nearctica
AT55	L. illustris	AU70	S. nearctica
AT56	L. illustris	BB67	S. nearctica
AT57	L. illustris	E43	S. polistensis
AT58	L. illustris	BC36	S. saracenioides
AV54	L. illustris	AQ49	S. sarracenioides
AX11	L. illustris	AR01	S. sarracenioides
BA61	L. illustris	AX80	S. sarracenioides
BB44	L. illustris	AL80	S. seagoi
BB68	L. illustris	AH22	S. shermani
BE09	L. illustris	AH44	S. shermani
BE10	L. illustris	AH47	S. shermani
AE60	L. mexicana	AH48	S. shermani
AE63	L. mexicana	AV79	S. sp
AX39	L. mexicana	AP06	S. subvicina
AX40	L. mexicana	AP07	S. subvicina
AX41	L. mexicana	AP72	S. subvicina
AX42	L. mexicana	AS68	S. subvicina
AX43	L. mexicana	AV23	S. subvicina
AX44	L. mexicana	AW02	S. subvicina

AX45	L. mexicana	A23	S. triplasia
AX46	L. mexicana	AA27	S. triplasia
AX47	L. mexicana	AL10	S. triplasia
AX48	L. mexicana	AL15	S. triplasia
AX49	L. mexicana	AR20	S. triplasia
AX50	L. mexicana	AT27	S. triplasia
AX51	L. mexicana	AT28	S. triplasia
AX53	L. mexicana	E24	S. triplasia
AX56	L. mexicana	E25	S. triplasia
BE66	L. mexicana	E27	S. triplasia
BF35	L. mexicana	AE22	S. utilis
BF55	L. mexicana	AE23	S. utilis
BF56	L. mexicana	AE24	S. utilis
BF73	L. mexicana	AT19	S. utilis
BF74	L. mexicana	E42	S. utilis
BH05	L. mexicana	AV21	Sb. flavipalpis
BH06	L. mexicana	BG79	Sx. lambens
BH16	L. mexicana	BG80	Sx. lambens
BH17	L. mexicana	AC25	Ti. alata
BH27	L. mexicana	F63	Ti. luculenta
BH28	L. mexicana	AQ68	Tp. importuna
AE59	L. sericata	AE56	W. vigil
AF47	L. sericata		
AJ76	L. sericata		
AM19	L. sericata		
AO51	L. sericata		
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AX36	L. sericata		
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AT77	L. silvarum		
AU62	L. silvarum		
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BI02	L. thatuna		
AB62	P. regina		
AB64	P. regina		
AD71	P. regina		

Family Muscidae

E46	Mu. atumnalis
BG75	Mu. domestica
BG76	Mu. domestica

AD72	P. regina
AE67	P. regina
AF39	P. regina
AG01	P. regina
AI17	P. regina
AI18	P. regina
AJ29	P. regina
AM03	P. regina
AS01	P. regina
AS03	P. regina
AT18	P. regina
AT38	P. regina
AT39	P. regina
AT63	P. regina
AT64	P. regina
AU35	P. regina
BF05	P. regina
BF36	P. regina
E51	P. regina
AT02	Po. labialis
AP60	Po. pediculata
AR74	Po. pediculata
AS70	Po. pediculata
AS71	Po. pediculata
AU78	Po. pediculata
BA12	Po. pediculata
BB37	Po. pediculata
BG57	Po. pediculata
AQ33	Po. rudis
AR75	Po. rudis
AV81	Po. rudis
AW13	Po. rudis
AZ63	Po. rudis
AF10	Pr. terranovae
AI04	Pr. terranovae
AI05	Pr. terranovae

Table 5. Intraspecific nucleotide diversity (%) for seven species with multiple specimens (N), calculated as the average diversity between all specimens within a species (Pi). Sliding window analysis was calculated for 600bp windows of the COI-COII region.

Species	N	Sliding window analysis				COI (%)	COII (%)	COI-COII (%)	ND4 (%)	COI-COII-ND4 (%)
		Max (%)	Midpoint (s)	Min (%)	Midpoint (s)					
<i>C. vicina</i>	5	0.300	2309-2509	0	2909-3146	0.162	0.063	0.127	0.062	0.122
<i>C. vomitoria</i>	2	0.500	2701-2871, 3248-3268	0.167	1759-1863, 2506-2661, 3308-	0.334	0.154	0.267	0.156	0.237
<i>L. illustris</i>	4	0.833	1990-2030	0.167	2755-2900	0.463	0.230	0.380	0.312	0.356
<i>L. sericata</i>	3	0.222	1767-1776	0	1781-1841, 2451-3462	0.130	0	0.088	0.104	0.090
<i>L. cuprina</i>	6	0.244	2033-2490	0	2640-3176	0.174	0.162	0.162	0.116	0.183
<i>S. Africa</i>	3	0.222	1891-2026	0	2509-2724, 3339-3404	0.088	0.101	0.116	0.104	0.113
<i>S. bullata</i>	3	2.556	2151-2176, 2351-2386	1.0	2761-2766	2.018	1.118	1.740	1.996	1.804
<i>P. regina</i>	4	1.667	2511-2571	0.083	3118-3488	0.753	0.076	0.514	0.572	0.385
<i>Total</i>	-	-	-	-	-	8.407	8.028	8.124	7.752	7.877

Table 6. Intraspecific nucleotide diversity (%) for seven species with multiple specimens (N), calculated as the average diversity between all specimens within a species (Pi) for 3 commonly sequenced regions of the COI gene.

Species	N	Barcoding	"2183"	"2792"
<i>C. vicina</i>	5	0.088	0.0223	0
<i>C. vomitoria</i>	2	0.296	0.374	0.339
<i>L. illustris</i>	4	0.495	0.502	0.184
<i>L. sericata</i>	3	0.094	0	0
<i>L. cuprina</i>	6	0.207	0.179	0
<i>S. africa</i>	3	0.095	0.082	0.113
<i>S. bullata</i>	3	2.194	1.958	1.478
<i>P. regina</i>	4	0.143	1.288	0.085
Total absolute divergence	-	7.982	8.806	7.076
Total relative divergence	-	98.25%	108.39%	87.10%

Table 7. Interspecific nucleotide diversity (Pi%) for 5 species pairs for the COI-COII region. Sliding window analysis was calculated for 600bp windows of the COI-COII region.

Species	N	Sliding window analysis				COI (%)	COII (%)	COI-COII (%)	ND4 (%)	COI-COII-ND4 (%)
		Max (%)	Midpoint (s)	Min (%)	Midpoint (s)					
<i>C. vicina</i> x <i>C. vomitoria</i>	7	5.820	2487-2497	1.651	3169-3184	4.712	2.757	3.952	4.272	4.024
<i>L. sericata</i> x <i>L. illustris</i>	7	7.856	2551	3.803	3098, 3118-3123	5.804	4.856	5.362	5.330	5.226
<i>L. sericata</i> x <i>L. cuprina</i>	9	1.345	2371-2436	0.502	2971-3137	0.940	1.015	0.925	0.873	0.900
<i>L. illustris</i> x <i>L. cuprina</i>	10	7.825	2555	3.584	3147-3157	5.742	4.856	5.315	4.995	5.099
<i>S. Africa</i> x <i>S. bullata</i>	6	12.353	2529	6.921	2989-3008, 3275	9.548	7.694	8.737	11.098	9.274
Mean	-	-	-	-	-	5.349	4.236	4.858	5.314	4.905

Table 8. Interspecific nucleotide diversity (%) for 5 species pairs for the COI-COII region.

Species	Barcoding	2183	2792
<i>C. vicina</i> x <i>C. vomitoria</i>	4.630	4.793	1.711
<i>L. sericata</i> x <i>L. illustris</i>	4.943	6.775	3.597
<i>L. sericata</i> x <i>L. cuprina</i>	0.804	1.026	0.508
<i>L. illustris</i> x <i>L. cuprina</i>	5.056	6.598	3.403
<i>S. africa</i> x <i>S. bullata</i>	7.623	11.386	7.708

Table 9. Estimates of overall genetic distance and mean intra- and interspecific diversity for COI, COII and ND4 genes.

	Overall genetic distance
COI	0.084
COII	0.084
ND4	0.079
COI-COII	
COI-COII-ND4	0.080

Table 10. DNA sequence data for the reference database for species identification of forensically important Diptera

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AAT TTT ATT ACT ACA GTT AAT ATA CGA TCT ACT TTT GAC CGA ATA CCT TTA TTT GGA TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT CTA CCT GTA CTT GCC GGA GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGA GAC CCA ATT TTA TAT CAA CAC TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGT TCA CTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGT TTA TTA GGT TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTG GGA ATG GAT GTA GAT ACT CGA GCT TAT TTT ACA TCT GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA CTA GCA ACT CTA TAT GCA CAA CTT AAT TAT TCT CCA GCA ATT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTT GTT CTA GCT AAT TCT TCA ATT GAT ATC ATT TTA CAT GAT ACT
TAT TAT GTA GTA GCA CAT TTC CAT TAT GTA TTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACT GGT TTA ACA TTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGT CTT GCT GGA ATA CCT CGA CGT TAT TCT GAC TAC CCA GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCA ACT ATT GGT TCA ACA ATT TCA TTA TTA GGA ATT CTA TTC TTC TTC TTT ATT ATT TGA GAA
AGT TTA ACT TCT CAA CGT CAA GTT ATA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA --- --- --- ---
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#B._eleodis_AY05_NM_(Grant)_USA
TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGT TGA ACT GTA TAT CCA CCT CTT TCA TCT AAC ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCT GGA ATT TCT TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGT ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAT CCA GCA GGA GGA GGT GAC CCA ATT TTA TAT CAA CAC TTA TTT TGA TTT TTT GGT CAC
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCT GGA AAA AAG GAA
ACA TTT GGA TCA TTA GGA ATA ATT TAT GCT ATG CTA GCA ATT GGT TTA TTA GGT TTT ATT GTC TGA GCT CAT CAC ATA
TTT ACT GTA GGA ATA GAC GTA GAT ACT CGA GCC TAT TTT ACT TCT GCA ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACA CAA CTT AAT TAT TCC CCA GCA ATT TTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGT GGA TTA ACA GGA TTA CTA TTG GCT AAT TCA TCA ATT GAT ATT TTT TTA CAT GAT ACA
TAC TAT GTA GTA GCA CAT TTT CAT TAC GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAC CCA TTA TTT ACT GGT TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTT CCG CAA CAT TTT TTA GGT CTT GCT GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCT TAC ACA
TCT TGA AAT GTT GTT TCA ACT ATT GGT TCA ACA ATT TCC TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTG ACT TCT CAA CGT CAA GTA TTA TTC CCA ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCT CCA
GC

#B._arizona_AW36_NM_(Grant)_USA
ATA CTA TTA ATA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT CTT TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCT GGT ATT TCT TCT ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTC CTT TCT TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GAC CCA ATT TTA TAT CAA CAC TTA TTT TGA TTC TTT GGT CAT
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGT ATA ATT TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAA GAA
ACA TTT GGA TCT TTA GGA ATA ATT TAT GCC ATA CTA GCA ATT GGT TTA TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAT GTT GAT ACT CGA GCT TAT TTT ACT TCT GCC ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTT AGA TGA CTA GCA ACT CTT TAT GGT ACA CAA CTT AAT TTC TCT CCA GCA ATT TTA TGA GCC CTT GGA TTT
GTA TTT TTA TTT ACC GTA TTA ACA GGT GGA TTA GCA GGA TTA GCT AAT TCT TCA ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTT TTA GCA CAT TTT CAC TAT GTC CTT TCT ATA GGA GCA GTG TTT GCA ATT ATA GCA GGA TTT GAT CAT TGA
TAT TCT TTA TTC ACT GGT TTA ACC CTA AAT GTT AAT ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTT TTA GCA CTT GCT GGA ATA CCT CGA CGT TAC TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATT TCA ACT ATT GGT TCA ACA ATT TCA TTC CTA GGA ATT CTA TAT TTC TTT TAC ATT ATT TGA GAA
AGT CTA ACT TCT CAA CGT CAA GTA ATA TTT CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA TTT CAA AAT ACT CCG CCA
GC

#B._plinthopyga_AC79_FL_(Highland)_USA
TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT CTT TCA TCT TCT AAT ATT
GCC CAT GGA GGA GCA TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCT GGA ATT TCA TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTG CCT GTA CTT GCT GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GCA ATT CCA ATT CTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACA TTT GGT TCA TTA GGG ATA ATT TAT GCA ATG TTA GCA ATT GGT TTA TTA GGT TTC ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAT GAT ACT CGT TAT TTT ACA TCT GCA ACA ATA ATT ATT GCT GTT CCT ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCA ATT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTT GCA CAT TTT CAC TAT GCA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTC GAT CAC TGA
TAC CCT CTA TTT ACT GGA CTA ACA TTA AAT AGA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCT CAA CAC TTC TTA GGA CTT GCT GGA ATA CCT CGA CGT TAT TCA GAC TAT CCA GAT GCT TAT ACT
ACT TGA AAT GTT ATT TCA ACT ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCC CAA CGT CAA GTG ATA TTC CCT ATC CAA CTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCT
GC

#B._plinthopyga_AE50_CA_(Glen)_USA

GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCT GTA CTT GCT GGT GCA ATT ACT ATA TTA TTT ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAC TTA TTC TGA TTC TTT GCA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACA TTT GGT TCA TTA GGG ATA ATT TAT GCA ATG TTA TTA GCA ATT GGT TTA TTA GGT TTC ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAC GTA GAT ACT CGT GCT TAT TTT ACA TCT GCA ACA ATA ATT ATT GCT GTT CCT ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCA ATT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA GCT GTT TTA GCT AAT TCT TCT ATT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTT GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTC GTA CAC TGA
TAC CCT CTA TTT ACT GGA CTA ACA TTA AAT AGA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCT CAA CAC TTC TTA GGG CTT GC- --- --- --- --- --- --- --- --- --- --- --- --- --- ---

#B._plinthopyga_AG29_OR_(Crook)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCA TCT GTT GAT TTA GCA ATT TTC TCT CTT CAC TTA GCT GGA ATT TCA TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA CTT CTT TCT TTA CCT GTA CTT GCT GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAC TTA TTC TGA TTC TTT GCA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACA TTT GGT TCA TTA GGG ATA ATT TAT GCA ATG TTA GCA ATT GGT TTA TTA GGT TTC ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA RTG GAC GTA GAT ACT CGT GCT TAT TTT ACA TCT GCA ACA ATA ATT ATT GCT GTT CCT ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCA ATT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTT GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTC GTA CAC TGA
TAC CCT CTA TTT ACT GGA CTA ACA TTA AAT AGA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCT CAA CAC TTC TTA GGA CTT GCT GGA ATA CCT CGA CGT TAT TCA GAC TAT CCA GAT GCT TAT ACT
ACT TGA AAT GTT ATT TCA ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCC CAA CGT CAA GTA ATA TTC CCT ATT CAA CTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCT
GC

#B._plinthopyga_AH77_OR_(Crook)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCA TCT GTT GAT TTA GCA ATT TTC TCT CTT CAC TTA GCT GGA ATT TCA TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT CTA TTA TTA CTT CTT TCT TTA CCT GTA CTT GCT GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAC TTA TTC TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GCA GGA GGA GGA GAT CCA ATT TCC CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACA TTT GGT TCA TTA GGG ATA ATT TAT GCA ATG TTA GCA ATT GGT TTA TTA GGT TTC ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAC GTA GAT ACT CGT GCT TAT TTT ACA TCT GCA ACA ATA ATT ATT GCT GTT CCT ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCA ATT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTT GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTC GTA CAC TGA
TAC CCT CTA TTT ACT GGA CTA ACA TTA AAT AGA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCT CAA CAC TTC TTA GGA CTT GCT GGA ATA CCT CGA CGT TAT TCA GAC TAT CCA GAT GCT TAT ACT
ACT TGA AAT GTT ATT TCA ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCC CAA CGT CAA GTA ATA TTC CCT ATT CAA CTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCT
GC

#B._sp_AJ80_OR_(Jefferson)_USA

TTA TTA TTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCC CTT TCA TCT AAT ATT
GCA CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCT GGA ATT TCT TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCC TTA TTT GTT TGA TCT GTA
GTA ATT ACC GCC TTA TTA TTA CTT TCT TTA CCA GTC CTT GCT GGA GCA ATT ACT ATA TTA TTA ACT GAC CGA AAC
ATT AAT ACT TCT TTC TTT GAT CCT GCA GGA GGA GGG GAT CCA ATT TTA TAT CAA CAC TTA TTT TGA TTT TTT GGT CAC
CCA GAA GTT TAC ATT TTA ATT TTA CCG GGA TTT GGT ATA ATT TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGT TTA TTA GGT TTT ATT GTT TGA GCT CAT CAC ATA
TTT ACT GTA GGA ATA GAT GTA GAT ACT CGA GCA TAT TTT ACT TCT GCT ACT ATA ATT ATT GCT GTC CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGT ACA CAA CTT AAT TAC TCT CCA GCA ATT TTA TGG GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGT GGA TTA ACA GGA GTA GTT TTA GCT AAC TCT TCA ATT GAT ATT TTA CAT GAT ACC
TAT TAT GTA GTA GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT TTA TTT ACT GGA TTA ACA TTA AAT AAA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTC ATA GGA GTT AAT
TTA ACT TTC TTT CCA CAA CAT TTT TTA GGT CTT GGT GGA ATA CCT CGA CGT TAC TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATT TCA ACT ATC GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCT CAA CGT CAA GTA ATA TTT CCT GTA CAA TTA AAT TCT TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA
GC

#B._sp_AK01_OR_(Jefferson)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCC CTT TCA TCT AAT ATT
GCA CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCT GGA ATT TCT TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCC TTA TTT GTT TGA TCT GTA
GTA ATT ACC GCC TTA TTA TTA CTT TCT TTA CCA GTC CTT GCT GGA GCA ATT ACT ATA TTA TTA ACT GAC CGA AAC
ATT AAT ACT TCT TTC TTT GAT CCT GCA GGA GGA GGG GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTT TTT GGT CA-

ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA CTA GCA ATT GGT TTA TTA GGT TTT ATT GTT TGA GCT CAT CAC ATA
TTT ACT GTA GGA ATA GAT GTA GAT ACT CGA GCA TAT TTT ACT TCT GCT ACT ATA ATT ATT GCT GTC CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGT ACA CAA CTT AAT TAC TCC CCA GCA ATT TTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGT GGA TTA ACA GGA GTA GTT TTA TTA GCT AAC TCT TCA ATT GAT ATT TTA CAT GAT ACC
TAT TAT GTA GTA GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTC ATA GGA GTT AAT
TTA ACT TTC TTT CCA CAA CAT TTT TTA GGT CTT GGT GGA ATA CCT CGA CGT TAC TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATT TCA ACT ATC GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCT CAA CGT CAA GTA ATA TTT CCT GTA CAA TTA AAT TCT TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA
AC

#B._plinthopyga_AK02_OR_(Jefferson)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCA TCT GTT GAT TTA GCA ATT TTC TCT CTT CAC TTA GCT GGA ATT TCA TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTG CCT GTA CTT GCT GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT

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ATT AAT ACT TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAC TTA TTC TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GCA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACA TTT GGT TCA TTA GGG ATA ATT TAT CCA GCA ATG TTA GCA ATT GGT TTA TTA GGT TTC ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAC GTA GAT ACT CGT GCT TAT TTT ACA TCT GCA ACA ATA ATT ATT GCT GTT CCT ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GCA ACT CAA TTA AAT TAT TCT CCA GCA ATT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA TTA TTA ACA GGA GTT GTT TTT TTA GCT AAT TCT TCT ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTT GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTC GTA CAC TGA
TAC CCT CTA TTT ACT GGA CTA ACA TTA AAT AGA AGT GAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA CTA AAT
TTA ACT TTT TTC CCT CAA CAC TTC TTA GGA CTT GGT GGA ATA CCT CGA CGT TAT TCA GAT TAT CCA GAT GCT TAT ACT
ACT TGA AAT GTT ATT TCA ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCC CAA CGT CAA GTA ATA TTC CCT ATC CAA CTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCT
GC

#B._sp_AJ81_OR_(Jefferson)_USA
CTC CTA CTA GTA AGC AGA ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAT CCT CCT TTA TCC TCT AAC ATC
GCC CAT GGA GGA GCA TCT GTA GAT CTA GCA ATT TTC TCA TTA CAC TTA GCT GGA ATT TCT TCA ATT CTT GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT TCA GGA ATT CCT ACT TTC GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA CTT CTT CTA CTT TCA TCC GTT CTT GCA GGA GCA ATT ACT ATA CTA TTA ACT GAC GCA AAT
ATC AAT ACC TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGG CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCG GGA TTT GTT ATA ATT TCT CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACC TTT GGT TCA CTA GGA ATA ATT TAC GCA ATA ATT GCA ATT GGA TTA TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATG GAT GTG GAT ACT CGA GCT TAC TTT ACT TCA GCT ACT ATA ATT ATT GCT GTT CCA ACC GGA ATT
AAA ATT TTT AGA TGA TTA GCT ACT CTT TAC GGA ACA CAA CTA AAT TAT TCC CCA GCA ATT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA AAT GGT TTA TTA GCT AAC TCT TCT ATT GAT ATT ATT TTA CAT GAT GCA ACT
TAT TAT GTA GTT GCA CAT TTC CAT TAT GTA CTT TCT ATA GGA GCC GTT TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTC ACA GGT TTA GCT TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA TTA AAT
CTA ACT TTC TTC CCT CAA CAT TTT TTA GGA CTA CCT CGG GAT CCA GAT TAC TCT GGC TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATC TCA ACT ATT GGA TCT ACA ATT TCA CTA CTA GGA ATT TTA TTC TTC TTT TTC ATT ATT TGA GAA
AGT TTA ACT TCC CAA CGT CAA ATT TTA TAC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCT CCA
TC

#B._plinthopyga_BG47_NM_(Socorro)_USA
TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCA TCT GTT GAT TTA GCA ATT TTC TCT CTT CAC TTA GCT GGA ATT TCA TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA COT GTA CTT GCT GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GAT CCA ATT CTA TAC CAA CAC TTA TTC TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACA TTT GGT TCA TTA GGG ATA ATT TAT GCA ATG TTA GCA ATT GGT TTA TTA GGT TTC ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GGA GAT GAT ACT CGT GCT TAT TTT ACA TCT GCA ACA ATA ATT ATT GCT GTT CCT ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCA ATT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTT GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTC GTA CAC TGA
TAC CCT CTA TTT ACT GGA CTA ACA TTA AAT AGA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCT CAA CAC TTC TTA GGA CTT GCT GGA ATA CCT CGA CGT TAT TCA GAC TAT CCA GAT GCT TAT ACT
ACT TGA AAT GTT ATT TCA ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCC CAA CGT CAA GTA ATA TTC CCT ATC CAA CTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCT
GC

#B._plinthopyga_BG46_NM_(Socorro)_USA
TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCA TCT GTT GAT TTA GCA ATT TTC TCT CTT CAC TTA GCT GGA ATT TCA TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTG CCT GTA CTT GCT GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAC TTA TTC TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACA TTT GGT TCA TTA GGG ATA ATT TRT GCA ATG TTA GCA ATT GGT TTA TTA GGT TTC ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAC GTA GAT ACT CGT GCT TAT TTT ACA TCT GCA ACA ATA ATT ATT GCT GTT CCT ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCA ATT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTT GCA CAT TTT CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATT ATA GCT GGA TTC GTA CAC TGA
TAC CCT CTA TTT ACT GGA CTA ACA TTA AAT AGA AAG ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCT CAA CAC TTC TTA GGA CTT GCT GGA ATA CCT CGA CGT TAT TCA GAC TAT CCA GAT GCT TAT ACT
ACT TGA AAT GTT ATT TCA ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCC CAA CGT CAA GTA ATA TTC CCT ATC CAA CTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA C--
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#Bo._bisetosa_OH_A37_(Hocking)_USA
TTA CTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAT CCC CCC CTA TCC TCT AAC ATT
GCT CAT GGA GGA GCC TCA GTA GAT TTA GCA ATT TTC TCA TTA CAT TTA GCA GGA ATT TCA TTA ATT CTA GGA GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA TTT GAT CGG ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT CTT TTA TTA CTT CTT TCC CTA CCA GTA CTT GCT GGA GCT ATT ACA ATA TTA TTA ACT GAT CGA AAC
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CTT ATT CTA TAT CAA CAC CTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TT- --- --- --- --- --- --- --- --- TCT CAC ATT ATT AGA CAA GAA TCA GGA AAA AGG GAA
ACC TTC GGT TCT TTA GGA ATA ATT TAT GCT ATA CTA GCA ATT GGA TTA TTA GGA TTT ATT GTA TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACC CGA GCT TAT TTT ACC TCA GCT ACT ATA ATT ATT GCT TGA CCA ACA GGT ATT
AAA ATT TTT AGT TGA TTA GCT ACC CTA TAC GCA ACT CAA TTA AAC TAT TCT CCA GCT ACT TTA TGA GCC TTA GGG TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTG ACG GGA GTA GTA TTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAC TAT GTT TTA TCA ATA GGA GCA GTA TTC GCA ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCC CTA TTT ACA GGA TTA ACC TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ACA ATT ATA TTT ATA GGA GTA AAT
CTA ACT TTT TTC CCC CAA CAT TTC TTA GGG CTT GCT GGA ATA CCT CGT CGA TAT TCT GAC TAT CCA GAT GCC TAT ACT
GCT TGA AAT GTA ATT TCA ACT ATT GGT TCA ACA ATT TCA CTT CTA GGA ATT TTA TTT TTT TTC TAT ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTA ATA TAC CCA GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCT CCA
GC

#Bo._cimbicis_E35_OH_(Claremont)_USA
CTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT CTA TCT TCT AAT ATT
GCA CAT GGA GGG GCT TCA GTA GAT TTA GCA ATT TTT TCA TTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGA GCC GTA
AAC TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCT TTA TTC GTA TGA TCT GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTT TCT TTA CCA GTA CTT GCT GGA GCT ATT ACA ATA TTA TTA ACT GAT GCA AAC
ATT AAT ACG TCA TTT TTT GAY CCT GCA GGA GGA GGA GAY CCA ATT TTA TAC CAA CAT TTA TTT TGG TTC TTT GGA CAC
CCC GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGG ATA ATC TCC CAT ATT ATT AGA CAA GAA TCA GGG AAA AAG GAA
ACT TTT GGT TCT TTA GGA ATA ATT TAT GCT ATT TTA GCT ATT GGA CTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACT GTA GGA ATA GAT GTA GAT ACA CGA GCT TAT TTT ACT TCA GCT ACT ATA ATC ATT GCT GTA CCA ACA GGT ATT
AAA ATT TTT AGT TGA CTA GCT ACA TTA TAC GGA ACC CAA TTA AAT TAC TCT CCA GCC ACT TTA TGA GCA TTA GGG TTT

GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA GTA GTT CTA GCT AAT TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAT TAT GTA GTA GCC CAT TTT CAT TAT TTA TTA TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCT CAA CAT TTT TTA GGR CTT GCT GGT ATA CCT CGT CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
GCC TGA AAC GTA ATT TCA ACC ATT GGT TCA ACA AAT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAT ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAC ACA CCC CCA
GC

#Bo._latisterna_E22_OH_(Hocking)_USA

TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CC? ??? ??? ??? ??? ??? ??? ??? ??? GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGC TCA ACA AAT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
GC

#Bo._latsiterna_AU69_NY_(Onondaga)_USA

TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTG CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGA TCA ACA AAT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
GC

#Bo._latisterna_AW17_NY_(Monroe)_USA

TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTG CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGA TCA ACA AAT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
G-

#Bo._latisterna_AA11_FL_(Columbia)_USA

TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTG CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGA TCA ACA AAT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
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#Bo._latisterna_AA12_FL_(Columbia)_USA

--- --- -TA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTG CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTG GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGC TCA ACA AAT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA

AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA GC

#Bo._latisterna_AL26_SC (Newberry)_USA
TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGC TCA ACA ATT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
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#Bo._latisterna_AM70_ME (Hancock)_USA
TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGC TCA ACA ATT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
GC

#Bo._latisterna_E49_TN (Washington)_USA
TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGC TCA ACA ATT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
GC

#Bo._latisterna_E53_TN (Washington)_USA
TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTG
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGC TCA ACA ATT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
GC

#Bo._latisterna_AR81_WI (Waukesha)_USA
--- --- --- GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA CTA CAT CTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTT CTA TTA CTT CTC TCT TTA CCA GTT CTT GCT GGA GCC ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGA CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATG
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCC ACT ATA ATC ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA TTA TAT GGA ACC CAA TTG AAT TAC TCT CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTT TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTC CTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCC CAT TTT CAT TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAG TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTC CCT CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGA CGA TAT TCA GAT TAT CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACC ATT GGC TCA ACA ATT TCA CTT CTA GGA ATT TTA TTT TTC TTC TAC ATT ATC TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTA ATA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAG TGA CTT CAA AAT ACA CCT CCA
GC

#Bo._litorosa_AG26_OR (Clatsop)_USA
TTA TTA TTA GTA AGT AGT ATA GTG GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT CTA TCC TCT AAT ATC
GCC CAT GGA GGA GCT TCA GTA GAT TTA GCA ATT TTT TCA TTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGA GCT GTA

AAT TTT ATC ACT ACA GTT ATT AAT ATA CGA TCG ACA GGA ATT ACA TTC GAT CGA ATA CCT TTA TTT GTT TGA TCT GTT
GTA ATT ACA GCT CTT CTC TTA CTC TCT TTA CCA GTA CTT GCT GGA GCT ATT ACA ATA TTA TTA ACT GAT CGA AAT
ATT AAC ACT TCA TTT TTT GAC CCT GCG GGA GGA GCA GAC CCA ACT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGG CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCT CAT ATT ATC AGA CAA GAA TCA GGG AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT CAT GCT ATG TTA TTA GCT ATT GGA TTA TTA GGA TTT ATT TTA TGA GCT CAC CAT ATA
TTT ACT GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCT ACT ATA ATT ATT GCT GTA CCA ACA GGT ATT
AAA ATT TTT AGT TGA CTA GCT ACA CTA TAT GGA ACT CAA TTA AAT TAT TCT CCA GCC ACT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACA GGA TTA GCT AAC TCT TCT GTA GAT ATT ATT TTA CAT GAT ACT
TAC TAT GTA GTA GCT CAT TTT CAC TAT GTC TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTC CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ACA ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTC CCC CAA CAT TTT TTA GGA CTT GCT GGT ATA CCT CGC CGA TAC TCA GAT TAC CCA GAC GCT TAT ACA
ACT TGA AAC GTA ATT TCA ACT ATT GGT TCA ACA ACT TCA CTC CTA GGA ATT TTA TTT TTC TAT ATT ATT TGA GAA
AGT TTA GTA TCC CAA CGA CAA GTA ATA TAC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACA CCC CCA
GC

#Br._sarcophagina_AR69_WI_(Waukesha)_USA

--- --- GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCT GGA ATT TCT TCT TTT TTA GGA GCT GTT
AAT TTT ATT ACA ACA GTA ATT AAT ATA CGT GCT ACA GGA ATT TCA TTC GAC CGT ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATC ACA GCT TTA TTA CTA CTT TTA TCT CTT CCA GTT TTA GCA GGA GCA ATT ACA ATA TTA TTA ACA GAC CGA AAT
CTT AAT ACT TCA TTT TTT GAT CCA GGA GGG GGA GGT TAT CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GCT CAC
CCA GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA TTA GTT TCT CAC GTT ATT AGC CAA GAA TCT GGA AAA AAG GAA
GCT TTT GGT TCT TTA GGA ATA ATT TAT GCT ATA TTA ACT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCC CAC CAT ATA
TTT ACT GTA GGA ATA GAT GAT ACA CGA GCT TAT TTT ACA TCT GCT ACT ACT ATA GTT ATT GCT ATC CCA ACT GGA ATT
AAA ATT TTT AGT TGA ATA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCG GCT ATA ATA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTA ATT CTT GCT AAT TCA TCA ATT GAT ATT GTA CTT CAT GAT ACA
TAC TAT GTA GTA GCT CAT TTT CAC TAC TCA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA ACT GGA TTT GTA CAT TGA
TAC CCT TTA TTT ACT GGA CTA ATA TTA AAT ACA AAA ATA TTA AAA AAT CAA TTC ATC ATT ATA TTT ATT GGA GTA AAT
ATA ACA TTT TTC CCT CAA CAT TTT TTA GGT CTT GCT GGA ATA CCT CGA CGA TAC TCA GAT TAT CCT GAT GCT TAT ACA
ACA TGA AAT ATT TCT ACT AAT GGT TCA ACA ATT TCT TCT TTA ATA GGA ATT TTA TTT TTT TTA ATT ATT TGA GAA
AGT TTA ATT TCT CAA CGT AAA ATT TTA TTT CCA ATT CAA TTA AAT TCA TCA ATT GAA TGA TTC CAA AAT ACC CCT CCT
GC

#C._coloradensis_AG53_OR_(Jefferson)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATT
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAT CCT GCA GGA GGA GGA GAT CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTG GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ATT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATG CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATC TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAT CCT GTT CAA TTA AAT TCT TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA
GC

#C._coloradensis_AJ79_OR_(Jefferson)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATT
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAT CCT GCA GGA GGA GGA GAT CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTG GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATG CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAT CCT GTT CAA TTA AAT TCT TCA ATT GAA TGA CTA CAA AAT ACT CCG CC--
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#C._coloradensis_AZ01_NM_(Grant)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GWT TAT CCT CCT TTA TCT TCT AAT ATT
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAT CCT GCA GGA GGA GGA GAT CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT TTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTG GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATG CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GT- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#C._coloradensis_BA04_NM_(Grant)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATT
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAT CCT GCA GGA GGA GGA GAT CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA

TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
 AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
 GTA TTT TTA TTC ACA GTG GCA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
 TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
 TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT CAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
 ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATA CCT CGA CAA TAT TCA GAT TAC CCA GAT GCT TAT ACA
 ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATT TGA GAA
 AGT TTA GTT TCA CAA CGT CAA GTT TTA TAT CCT GTT CAA TTA AAT TCT TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA
 GC
 #C._coloradensis_BF63_NM_(Sante_Fe)_USA
 TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATT
 GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGA GCT GTA
 AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
 GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CCA AAT
 CTT AAT ACT TCA TTT TTT GAT CCT GCA GGA GGA GGA GAT CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
 CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
 ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCC CAC CAT ATA
 TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
 AAA ATT TTT AGT TGA TTA GCA ACT CTT TAC GGA ACG CAA TTA AAT TCT TCC CCA GCT ACT CTA TGA GCT TTA GGG TTT
 GTA TTT TTA TTT ACA GTA GGG GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
 TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
 TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
 ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATT CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
 ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATT TGA GAA
 AGT TTA GTT TCA CAA CGT CAA GTT TTA TAT CCT GTT CAA TTA AAT TCT TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA
 GC
 #C._grahami_AJ02_OR_(Coos)_USA

 --- -TT ATT ACA ACA GTA ATT AAT ATA CGA TCA ACT GGA ATT ACC TTT GAT CGA ATA CCA TTA TTT GTA TGA TCA GTA
 GTA ATC ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAC CGA AAT
 CTT AAT ACT TCA TTT TTT GAT CCC GCC GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGT CAC
 CCT GAA GTT TAT ATT CTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
 ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCC CAC CAT ATA
 TTC ACA GTA GGA ATA GAC GTT GAT ACC CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCC GTT CCA ACA GGA ATT
 AAA ATT TTT AGT TGA TTA GCA ACT CTT TAC GGA ACG CAA TTA AAT TCT TCC CCA GCT ACT CTA TGA GCT TTA GGG TTT
 GTC TTT TTA TTT ACA GTA GGG GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
 TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
 TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AAT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
 ATC ACA TTT TTC CCT CAA CAT TTC TTA GGT TTA GCA GGA ATA CCT CGA CGA TAC TCA GAT TAT CCT GAT GTT TAT ACA
 ACT TGA AAT GTA TTT TCT ACT ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT TTA TGA GAA
 AGT TTA ATT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAT TTA AAT TCA TCA ATT GAA TGA TTA CCA AAT TAC TCC AC-
 --
 #C._grahami_AJ10_OR_(Coos)_USA
 TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
 GCA CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGA GCT GTA
 AAT TTT ATT ACA ACA GTA ATT AAT ATA CGA TCA ACT GGA ATT ACC TTT GAT CGA ATA CCA TTA TTT GTA TGA TCA GTA
 GTA ATC ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAC CGA AAT
 CTT AAT ACT TCA TTT TTT GAT CCC GCC GGA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGT CAC
 CCT GAA GTT TAT ATT CTA ATT TTA CCK GGA TTT GGA ATA ATT TCT CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
 ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCC CAC CAT ATA
 TTC ACA GTA GGA ATA GAC GTT GAT ACC CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCC GTT CCA ACA GGA ATT
 AAA ATT TTT AGT TGA TTA GCA ACT CTT TAC GGA ACG CAA TTA AAT TCT TCC CCA GCT ACT CTA TGA GCT TTA GGG TTT
 GTC TTT TTA TTT ACA GTA GGG GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
 TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
 TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AAT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
 ATC ACA TTT TTC CCT CAA CAT TTC TTA GGT TTA GCA GGA ATA CCT CGA CGA TAC TCA GAT TAT CCT GAT GTT TAT ACA
 ACT TGA AAT GTA TTT TCT ACT ATT GGA TCA ACA ATC TCT TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT TTT TGA GAA
 AGT TTA ATT TCA CAA CGT CAA --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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 #C._latifrons_AF14_OR_(Clackamas)_USA
 --- -AT GGA TGA CCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
 AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
 GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCT GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
 CTT AAT ACC TCA TTT TTT GAC CCT GCA GGA GGA GGA GAC CCT ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGT CAT
 CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCC CAT ATT AGT CAA GAA TCA GGA AAG AAG GAA
 ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCC CAC CAT ATA
 TTT ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
 AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA TTT
 GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
 TAT TAT GTA GTT GCT CAT TTC CAT TAT GCT CTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
 TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
 ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCA CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
 ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
 AGC TTA GTT TCA CAA CGT CAA --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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 #C._latifrons_AF15_OR_(Clackamas)_USA
 TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
 GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
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 GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCT GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
 CTT AAT ACC TCA TTT TTT GAC CCT GCA GGA GGA GGA GAC CCT ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGT CAT
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 AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA TTT
 GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
 TAT TAT GTA GTT GCT CAT TTC CAT TAT GCT CTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
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 ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCA CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
 ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
 AGC TTA GTT TCA CAA CGT CAA --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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ATT ACA TTT GTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCA CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT TCA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
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#C._latifrons_BG58_AF21_WA(Clark)_USA
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GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
CTT AAT ACC TCA TTT TTT GAC CCT GCA GGA GGA GGA GAC CCT ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGT CAT
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ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGG CTA TTA GGA TTC ATT GTA TGA GCC CAC CAT ATA
TTC ACA GTA GGG ATA GAT GCA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
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ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACC CCA CCA
GC
#C._latifrons_AG16_OR_(Benton)_USA
TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
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CCG GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGG ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
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TTC ACA GTA GGG ATA GAT GCA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
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ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCG GGA ATA CCG CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA A-- --- ---
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#C._latifrons_AG22_OR_(Clatsop)_USA
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GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
CTT AAT ACC TCA TTT TTT GAC CCT GCA GGA GGA GGA GAC CCT ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGT CAT
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TTC ACA GTA GGG ATA GAT GCA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
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AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACC CCA CCA
GC
#C._latifrons_AG37_OR_(Benton)_USA
TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
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GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
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CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGG ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
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ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCG GGA ATA CCG CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACC CCA CCA
GC
#C._latifrons_AK48_OR_(Curry)_USA
TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
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GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
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AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
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AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACC CCA CCA
GC
#C._latifrons_AS40_WI_(Madison)_USA

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--- -AT GGA GGA CCT TCT GTT GAT TTA GCA CTT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT AAA ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
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CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGG ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGG CTA TTA GGA TTC ATT GTA TGA GCC CAC CAT ATA
TTC ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
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ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCA CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
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#C._latifrons_AZ31_NM_(Grant)_USA

--- -AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
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CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGG ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGG CTA TTA GGA TTC ATT GTA TGA GCC CAC CAT ATA
TTC ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
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#C._latifrons_AZ33_NM_(Grant)_USA

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ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGG CTA TTA GGA TTC ATT GTA TGA GCC CAC CAT ATA
TTC ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
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ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA A-- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---

#C._latifrons_BA02_NM_(Grant)_USA

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AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
CTT AAT ACC TCA TTT TTT GAC CCT GCA GGA GGA GAC CCT ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGG ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGG CTA TTA GGA TTC ATT GTA TGA GCC CAC CAT ATA
TTC ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
TAT TAT GTA GTT GCT CAT TTT CAC TAC GTT CTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCA CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACC CCA CCA
GC

#C._latifrons_BA05_NM_(Grant)_USA

--- -TTC CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTT TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCC GGT GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
CTT AAT ACC TCA TTT TTT GAC CCT GCA GGA GGA GAC CCT ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGG ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGG CTA TTA GGA TTC ATT GTA TGA GCC CAC CAT ATA
TTC ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACC TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
TAT TAT GTA GTT GCT CAT TTT CAC TAC GTT CTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA GCG GGA ATA CCA CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACC CCA CCA
GC

#C._latifrons_BA09_NM_(Grant)_USA

--- -A TAC CAA CAT TTA TTT TGA TTC TTT GGT CAT

CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGG ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTC GGT TCA TTA GGA ATA ATT TAT GCT ATT TCC CAT TTA GGA TTT ATT GTA TGA GCC CAC CAT ATA
TTC ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACC TGA GCT ACT ATA TTT ATT GCT GTF CCA ACA GAA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCT CCA GCT ACA TTA TGA GCA TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GCT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
TAT TAT GTA GTT GCT CAT TTT CAC TAC GTT CTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCC CAA CAT TTC TTA GGA TTA CGC GGA ATA CCA CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGG TCA ACA ATC TCA TTA CTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACC CCA CCA
GC

#C._livida_AH57_OR_(Crook)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATC
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTC ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGG TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA TAT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGG ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCC CCA GCT ACT TTA TGA GCT TTA GGG TTT
GTA TTT CTA TTC ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATG CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACT ATT TCA CTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TT- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#C._livida_AH66_OR_(Crook)_USA

TTA TTG TTA GTA AGT AGC ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATC
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTC ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGG TCA GTA
GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAG TCA GGA AAA AAG GAA
ACT TTC GGG TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTG TTA GGA TTT ATT GTA TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCC CCA GCT ACT TTA TGA GCT TTA GGG TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAG AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATA CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCT TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#C._livida_AP64_NY_(Schuyler)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATC
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTC ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGG TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA TAT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGG ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCC CCA GCT ACT TTA TGA GCT TTA GGG TTT
GTA TTT CTA TTC ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATG CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACT ATT TCA CTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TT- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#C._livida_AQ07_NY_(Schuyler)_USA

TTA TTG TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATC
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTC ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAY CCT GCA GGA GGA GGA GAT CCA ATT TTA TAY CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGG TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGG ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCC CCA GCT ACT TTA TGA GCT TTA GGG TTT
GTA TTT CTA TTC ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAC GGT AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATG CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACT ATT TCA CTA TTA GGA ATT TTA TTT TTC TTC TTC ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#C._livida_AZ32_NM_(Grant)_USA

TTA TTG TTA GTA AGT AGC ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATC
GCA CAT GGA GGA GCT TCA GTA GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTC ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGG TCA GTA
GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT TTA TAT GAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGG TCA TTA GGA ATA TAT TAT GCT ATA TTA GCT ATT GGT TTG TTA GGA TTT ATT GTA TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCG CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA

TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAG AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATA CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAT CCT GTT CAA TTA AAT TCT TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA
GC
#C._livida_BA01_NM_(Grant)_USA
TTA TTG TTA GTA AGT AGC ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCT CCT TTA TCT TCT AAT ATC
GCA CAT GGA GGA GCT TCA GTA GAT TTA GST ATT TTT TCT CTA CAT TTA GCA GGA ATT TCC TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAC CGA ATA CCA TTA TTT GTT TGG TCA GTA
GTA ATT ACA GCT TTA TTA CTT CTA TTA TCT TTA CCA GTA CTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAG TCA GGA AAA AAG GAA
ACT TTC GGG TCA TTA GGA ATA ATT TAT GCT ATT GGT TTG TTA GGA TTT ATT GTA TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAC TCT TCA CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC TCA ACA TTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAG AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTT TTC CCT CAA CAC TTT TTA GGA TTA GCA GGA ATA CCT CGA CGA TAT TCA GAC TAC CCA GAT GCT TA- ---

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#C._loewi_BI03_AK_USA

ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ACT ATA ATT ATT GCT GTC CCA ACA GAA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA TCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT ACA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTC CAT TGA
TAC CCT TTA TTT ACT GGA TTG ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATC GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA
GC
#C._loewi_BI04_AK_USA

ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ACT ATA ATT ATT GCT GTC CCA ACA GAA ATT
AAA ATT TT- AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA TCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT ACA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTC CAT TGA
TAC CCT TTA TTT ACT GGA TTG ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATC GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA
GC
#C._terranovae_AF65_OR_(Crook)_USA
TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCC GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCT GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCT GGG TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGT TCA TTA GGG ATA ATC TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCA CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACT CTT TAT GGT ACT CAA TTA AAT TCT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA TTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACC TTA AAT GGT AAA ATG CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTA AAT
ATT ACA TTT TTC CCT CAA CAT TTC TTA GGA TTA GCA GGA ATA CCT CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA CTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC
#C._terranovae_AK47_OR_(Curry)_USA
TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCC GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCA TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTC TTA TTA TCT TTA CCA GTA TTA GCT GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCA ATC TTA TAT CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCT GGG TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGT TCA TTA GGG ATA ATC TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCA CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACT CTT TAT GGT ACT CAA TTA AAT TCT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA TTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACC TTA AAT GGT AAA ATG CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTA AAT
ATT ACA TTT TTC CCT CAA CAT TTC TTA GGA TTA GCA GGA ATA CCT CGA CGA TAT TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA CTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GCA GAC CCA ATC TTG TAC CAA CAT TTA TTT TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT GCT ACT ATA ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA ACC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA TTA ACA GGA GGT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TCC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#C._vicina_AM11_ME_(Hancock)_USA

--- --CA CCT TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GCA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TCC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA TGT CAA GTT TTA TAC CCT GTT TAT TTA ACT TCA TCA A--- --- --- --- --- --- --- ---

#C._vicina_AM12_ME_(Hancock)_County)_USA

TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GGA GAC CCA ATC TTG TAC CAA CAT ??? ??? ??? ??? ??? ???
?CT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTG TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TCC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#C._vicina_AP58_NY_(Schuyler)_USA

TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GCA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TCC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA --- --- --- ---

#C._vicina_AP69_NY_(Schuyler)_USA

TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GCA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TCC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#C._vicina_AS41_WI_(Marathon)_USA

TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GCA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT

AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACC GTA GGA GGA TTA ACA GGA GCT ATT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT TTA TTT ATT GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GAA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC
#C._vicina_AT53_ND_(Trail)_USA

GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA --- --- --- --- ---
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#C._vicina_AV57_NY_(Niagara)_USA

GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA TAC CCT GTT CAT TGA AAT TCA TCA ATT GAA TGA TTA --- --- --- --- ---
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#C._vicina_AV70_NY_(Niagara)_USA

GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA ACC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGT CAA GTT TTA T-- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#C._vicina_AV74_NY_(Niagara)_USA

--T CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA ACC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGT CAA GTT TT- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#C._vicina_BB52_MN_(Koochiching)_USA

TTA CAT TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAC CGA ATA CCA TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGT GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
CTT AAT ACA TCA TTC TTT GAC CCA GCA GGA GGA GAC CCA ATC TTG TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCA CAT ATT ATT AGC CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGT TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAC GTT GAT ACT CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAG ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT ATT TTA GCT AAT TCT TCA GTA GAC ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTC GTT CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGG GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGA TTG GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGT CAA GTT TT- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---

ACT TGA AAC GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
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TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAT CCA CCT TTA TCT TCT AAT ATT
GCA CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCA GGA ATT TCT TCA ATT CTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGT ATT ACC TTC GAC CGA ATA CCG TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCC TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
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TTT ACA GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCT GCA ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACT CAA TTA AAT TCT TCC CCA GCC ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACT GGA GTT ATT TTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAC ACA
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AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGT ATT ACC TTC GAC CGA ATA CCG TTA TTT GTT TGA TCA GTA
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AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGT ATT ACC TTC GAC CGA ATA CCG TTA TTT GTT TGA TCA GTA
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GTA ATT ACA GCC TTA TTA CTT TTA TTA TCT TTA CCA GTA TTA GCA GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
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TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAT CCA CCT TTA TCT TCT AAT ATT
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TTA CTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAT CCA CCT TTA TCT TCT AAT ATT

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TTC CCT CTA TTT ACT GGA TTA ACT TTA AAT AGC AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATC GGA GTA AAT
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GCT TGA AAT GTA ATT TCT ACA ATT GGT TCA ACA ATT TCA TTA TTA GGA ATT TTA TTC TTC TTT TTC ATT ATT TGA GAA
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TTA CTA TTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT TTA TCA TCT AAT ATT
GCA CAT GGT GGA GCA TCA GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA TCT GGA ATT TCA TCA ATT TTA GGG GCC GTA
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ACC TTT GGA TCT TTA GGA ATA ATT TAT CGA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
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ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
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#Ch._rufifacies_AC70_FL_(Highland)_USA

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GCA CAT GGT GGA GCA TCA GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCT GGA ATT TCA TCA ATT TTA GGG GCC GTA
AAT TTT ATT ACA ACT GTT ATT AAT ATA CGA TCT ACA GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GTA TTA GCA GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
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CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATC ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACC TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GGA GGA TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC CCA TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT CTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
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TTA CTA TTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT TTA TCA TCT AAT ATT
GCA CAT GGT GGA GCA TCA GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCT GGA ATT TCA TCA ATT TTA GGG GCC GTA
AAT TTT ATT ACA ACT GTT ATT AAT ATA CGA TCT ACA GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GTA TTA GCA GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGG GGA GAC CCT ATT TTA TAT CAA CAC TTA TTT TGA TTC TTT GGT CAT
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATC ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACC TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA TTA GTA TTT GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC CCA TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT CTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
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TTA CTA TTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT TTA TCA TCT AAT ATT
GCA CAT GGT GGA GCA TCA GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCT GGA ATT TCA TCA ATT TTA GGG GCC GTA
AAT TTT ATT ACA ACT GTT ATT AAT ATA CGA TCT ACA GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GTA TTA GCA GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGG GGA GAC CCT ATT TTA TAT CAA CAC TTA TTT TGA TTC TTT GGT CAT
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATC ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACC TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA TTA GTA TTT GCT AAT TCA TCT ATW GAT ATT ATT TTA CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC CCA TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT CTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
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TTA CTA TTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT TTA TCA TCT AAT ATT
GCA CAT GGT GGA GCA TCA GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCT GGA ATT TCA TCA ATT TTA GGG GCC GTA
AAT TTT ATT ACA ACT GTT ATT AAT ATA CGA TCT ACA GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GTA TTA GCA GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGG GGA GAC CCT ATT TTA TAT CAA CAC TTA TTT TGA TTC TTT GGT CAT
CCA GRA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATC ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACC TTK GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC CCA TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT CTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
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GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
TTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGG GGA GAC CCT ATT TTA TAT CAA CAC TTA TTT TGA TTC TTT GGT CAT
CCA GAA GTT TAT ATT TTA ATT TTA CYT GGA TTC GGA ATA ATT TCT CAT ATC ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACC TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC CCA TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT CTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
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GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GTA TTA GCA GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT


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GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA GTA GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAC TAT GTA GTA GCT CAC TTC CAT TAT GAT CTT TCA ATA GCA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC COA TTA TTT ACT ACC GGA TTA ACC TAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA ACA GTT CTA TTA CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
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GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GTA TTA GCA GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGG GGA GAC CCT ATT TTA TAT CAA CAC TTA TTT TGA TTC TTT GGT CAT
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATC ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACC TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA GTA GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAC TAT GTA GTA GCT CAC TPC CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC COA TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA --- --- --- --- --- --- --- --- --- ---
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AAT TTT ATT ACA ACT GTT ATT AAT ATA CGA TCT ACA CGA GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTT ATT ACT GCT CTT CTT TTA TTA TTA TCA TTA CCA GTA TTA GCA GGT GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGG GGA GAC CCT ATT TTA TAT CAA CAC TTA TTT TGA TTC TTT GGT CAT
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATC ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACC TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTC ACT GTA GGA ATG GAT GTA GAT ACT CGA GCA TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA GTA GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAT TGA
TTC COA TTA TTT ACT GGA TTA ACC TTA AAT AAT AAA ATA CTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGA CTA GCT GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA --- --- --- --- --- --- --- --- --- ---
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TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GAT ATT TPC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TPC CAC TAT TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA ATA CAA AAT ACT CCA CCA
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C?T GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TPC CAC TAT TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA ATA CAA AAT ACT CCA CCA
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#Co._macellaria_AA30_FL_(Madison)_USA
TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TPC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TPC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT ATT ATT TGA GAA

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TTA ACT TTC GTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTC ACT TCT CAC ATA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT AAC CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
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GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT TTT TFC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGG ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA ATA CTC CAC CAG
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--- --- --- --- --- AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA CCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTT GGA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTT GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA --- --- --- --- --- ---
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#Co._macellaria_BE27_TX_(Cameron)_USA
TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACT TTA A-- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---

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#Co._macellaria_BE28_TX_(Cameron)_USA
TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA CTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC
#Co._macellaria_BE47_TX_(Fort_Bend)_USA
TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTT GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
TAC CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA CTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC
#Co._macellaria_BE67_TX_(Jackson)_USA

TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACC TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC

#Co._macellaria_BE78_TX_(McMullen)_USA

TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAC CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC

#Co._macellaria_BF02_TX_(Dallas)_USA

TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAT CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAT CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC

#Co._macellaria_BF44_NM_(Dona Ana)_USA

TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAC CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC

#Co._macellaria_BF52_TX_(El_Paso)_USA

TTA TTA TTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCA GTT GAT CTA GCT ATT TTC TCT CTT CAT TTA GCC GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTA ATT ACT GCT CTT TTA CTT TTA TTA TCT TTA CCA GTT TTA GCC GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAT CCA GCC GGA GGA GGA GAC CCA ATT TTA TAC CAA CAC TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTA TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCA CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAC ATG
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACA CTA TGA GCT TTA GGG TTC
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAC TCA TCA ATT GAT ATT ATC TTA CAT GAT ACT
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACT TTA AAT AGA AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACT TTC TTC CCT CAA CAC TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAT ACT
ACT TGA AAT GTA ATC TCT ACA ATT GGT TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTT ACT CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC

#Ch._megacephala_BE19_TX_(Cameron)_USA

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GCT CAT GGA GGA GCA TCA GTT GAT TTA GCT ATT TTC TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACA ACT GTA ATT AAT ATA CGA TCA ACA GGA ATT ACA TTC GAT CGA ATG CCT TTA TTC GTA TGA TCA GTA
GTT ATT ACT GCT CTA TTA TTA TTA TCT TTA CCA GTA TTA GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
CTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GAT CCT ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAT

CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TCA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAC CAC ATG
TTT ACT GTT GGA ATA GAC GTA GAC ACA GCA GCT TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCA ACT CTT TAC GGA ACA CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGA GTT GTT TCA TCA ATT GAC ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAC TTC CAT TAT GCT CTA TCA ATG GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TTC CCT CTA TTT ACT GGA TTA ACT TTA AAT AGC AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATC GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAC GCT TAC ACA
GCT TGA AAT GTA ATT TCT ACA ATT GGT TCA ACA ATT TCA TTA TTA GGA ATT TTA TTC TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CGA GTT TTA TTC CCT GTT CAA CTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT C-- ---

#Ch._megacephala_BE20_TX_(Cameron)_USA

TTA TTA TTA GTA AGT AGT ATA GTA GAA AAT GGG GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCA TCA GTT GAT TTA GCT ATT TTC TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACA ACT GTA ATT AAT ATA CGA TCT ACA GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTT ATT ACT GCT CTA TTA TTA TTA TTA TCT TTA CCA GTA TTA GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
CTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAC CAC ATG
TTT ACT GTT GGA ATA GAC GTA GAC ACA CGA GCT TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCA ACT CTT TAC GGA ACA CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAC ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAC TTC CAT TAT GCT CTA TCA ATG GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TTC CCT CTA TTT ACT GGA TTA ACT TTA AAT AGC AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATC GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAC GCT TAC ACA
GCT TGA AAT GTA ATT TCT ACA ATT GGT TCA ACA ATT TCA TTA TTA GGA ATT TTA TTC TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CGA GTT TTA TTC CCT GTT CAA CTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#Ch._megacephala_BF21_TX_(Cameron)_USA

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GCT CAT GGA GGA GCA TCA GTT GAT TTA GCT ATT TTC TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACA ACT GTA ATT AAT ATA CGA TCT ACA GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTT ATT ACT GCT CTA TTA TTA TTA TTA TCT TTA CCA GTA TTA GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
CTA AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTC GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCT TTA GGA ATG ATT TAT GCT ATA CTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAC CAC ATG
TTT ACT GTT GGA ATA GAC GTA GAC ACA CGA GCT TAT TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAG ATT TTC AGT TGA TTA GCA ACT CTT TAC GGA ACA CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAC ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAC TTC CAT TAT GCT CTA TCA ATG GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TTC CCT CTA TTT ACT GGA TTA ACT TTA ACT AGC AAG TTA TTA AAG AGT CAA TTT GCT ATT ATA TTT ATC GGA TTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCA GGT ATA CCT CGA CGA TAC TCA GAC TAT CCA GAC GCT TAC ACA
GCT TGA AAT GTA ATT TCT ACA ATT GGT TCA ACA ATT TCA TTA TTA GGA ATT TTA TTC TTC TTT TTC ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CGA GTT TTA TTC CCT GTT CAA CTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#Cy._cadaverina_AG15_OR_(Benton)_USA

TTA CTA TTA GTA AGT AGT ATA GTG GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACA ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCA CTA TTC GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT TTA CTA TCT TTA CCT GTT CTA GCT GGT GCT ATT ACA ATA TTA TTA ACA GAC CGA AAC
CTT AAT ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAC CCA ATT CTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGT TCA CTA GGG ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACC GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACA TTA GGA GGG TTA ACT GGC ATT TTT CTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTT GCC CAT TTC CAT TAT GTC CTA TCT ATA GGA GCT G-- --- --- --- --- --- --- --- --- ---

GC

#Cy._cadaverina_AM25_ME_(Hancock)_USA

TTA CTA TTA GTA AGT AGT ATA GTG GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACA ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCA CTA TTC GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT TTA CTA TCT TTA CCT GTT CTA GCT GGT ATT ACA ATA TTA TTA ACA GAC CGA AAC
CTT AAT ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAC CCA ATT CTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGT TCA CTA GGG ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACC GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTT ACA TTA GGA GGG TTA ACT GGC ATT TTT CTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTT GCC CAT TTC CAT TAT GTC CTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT AAA ATG TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGG TTA GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACT
ACT TGA AAT GTA ATT TCT ACT ATT GGG TCA ACA ATC TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATC TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT CTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA
GC

#Cy._cadaverina_AM26_ME_(Hancock)_USA

TTA CTA TTA GTA AGT AGT ATA GTG GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACA ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCA CTA TTC GTT TGA TCA GTA
GTA ATT ACA GCT TTA TTA CTT TTA CTA TCT TTA CCT GTT CTA GCT GGT GCT ATT ACA ATA TTA TTA ACA GAC CGA AAC
CTT AAT ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAC CCA ATT CTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGT TCA CTA GGG ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACC GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGG TTA ACT GGA GTT ATT CTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA

GC

#Cy_cadaverina_E40_OH_(Adams)_USA

TTA CTA TTA GTA AGT AGT ATA GTG GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACA ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCA CTA TTC GTT TGA TCT GTA
GTA ATT ACA GCT TTA TTA CTT TTA CTA TCT TTA COT GTT CTA GCT GGT GCT ATT ACA ATA TTA TTA ACA GAC CGA AAC
CTT AAT ACT TCA TTC TTC GAC CCA GCA GGA GGA GAC CCA ATT CTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAT
CC- --- --- --- ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGT TCA CTA GGG ATA ATT TAT GCT ATA CTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACC GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACC CAA TTA AAT TCT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACA TTA GGA GGG TTA ACT GGA GTT ATT CTA GCT AAT TCT TCA GTA GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTT GCC CAT TTC CAT TAT GTC CTA TCT ATA GGA GCT GTA TTC GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACT TTA AAT GGT TTA ACA ATG TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
ATT ACA TTC TTC CCT CAA CAT TTC TTA GGG TTA GCA GGA ATA CCT CGA CGA TAC TCA GAT TAC CCT GAT GCT TAC ACT
ACT TGA AAT GTA ATT TCT ACT ATT GGG TCA ACA ATC TCA TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATC TGA GAA
AGT TTA GTT TCA CAA CGT CAA GTT CTA TAC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCG CCA

GC

#Em_albicoma_A551_WI_(Marathon)_USA

CTG CTA CTT ATT AGT AGC ATA GTA GAA AAT GGA GCT GGA ACT GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATC
GCT CAT GGT GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC CTC CAT TTG GCA GGA ATC TCT TCT ATT TTA GGA GCA GTA
AAT TTT ATT ACA ACA GTT ATT AAT ATG CGA TCT TCA GGA ATT ACC TTT GAT CGA ATA CCT CTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA CTA TTA CTT CTT TCT TCA GGA TTA GCA GGA GCA ATT ACA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCA GGA GGG GGG GAT CCT ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGC CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCA TTA GGG ATA ATC TAT GCC ATA TTA ACA ATT GGA TTA TTA GGA TTT ATT GTT TGA GCC CAT CAT ATA
TTT ACT GTA GGT ATA GAT GTA GAT ACT CGA GCC TAC TTT ACT TCA ACT ACA ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AGA TGA CTA GCT ACT TTA TAT GGA ACA CAA CTA AAT TAT TCC CCA GCT --- --- --- --- --- --- ---
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AGT TTA GTT TCA CAA CGT CAA RKW ATA TAT ACT GTA CAA TTA AAC TCA TCA ATT GAA TGA TTT CAG AAC ACT CCA CCA

TC

#Fl_fletcheri_Ontario_Canada_E76

TTA TTA TTG GTA AGT AGA ATA GTG GAA AAC GGA GCC GGT ACA GGA TGA ACT GTT TAT CCA CCT TTA TCT TCT AAT ATT
GCT CAC GGA GGA GCT TCT GTT GAT TTA ACA ATT TTT TCT CTT CAC TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAC TTT ATT ACA ACA GTT ATT AAT ATA CGA TCT GCA GGA ATT ACT TTT GAT CGT ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATT ACA GCT TTA TTG CTT CTT CTT TCT TCA CCA GTA CTT GCA GGA GCA ATT ACT ATA TTA TTA ACC GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCT GCA GGA SGT GAC CCA ATT ??? ??? ??? ??? ??? ??? ??? ??? ??? ??? ???
??? ??? ??? ??? ??? ??? ??? ??? ??? ??? GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCC GGA AAA AAA GAA
ACA TTC GGA GCT TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGA CTA TT? ??? ??? ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACA GCA GCT TAT TTT ACT TCT GCT ACT ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGC TGA TTA GCC ACT CTT TAC GGA ACA CAA CTA AAT TAT TCT CCC GCT ATT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATC GAT ATT ATT TTA CAT GAT ACT
TAT TAT GTA GTA GCC CAT TTC CAT TAT TCA CTT TCA ATA GGA GCT GTA TTT GCA ATT ATA GCT GGA TTT GTC CAT TGA
TAT CCT CTA TTT ACT GGA TTA ACT TTA AAT AAA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTA AAT
TTA ACT TTT TTT CCC CAA CAT TTT TTA GGG CTT GCA GGT ATA CCT CGT CGA TAC TCT GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT ATT ATT TCA ACA ATC GGG TCT ACA ATT TCT CTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACT TCA CAA CGT CAA GTT ATA TTT CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA

GC

#H_rapax_E4_OH_(Hamilton)_USA

CTA TTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTT TTT GAC CCT GCA GGA GGA GAC CCA ATT --- --- --- --- --- --- --- --- --- --- ---
--A GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA GCA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACT CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAT TAC GTT GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATC ATG CCA GGA TTT GTC CAT TGA
TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATA CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTG CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA

GC

#H_rapax_BA23_OH_(Hamilton)_USA

CTA TTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GAC CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA GCA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACT CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAT TAC GTT GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATC ATG CCA GGA TTT GTC CAT TGA
TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATA CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTG CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA

GC

#H_rapax_BA24_OH_(Hamilton)_USA

-TA TTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA

GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTT TTT GAC CCT GCA GGA GGA GCA GAC CCA ATT CTT TAC CAA CAT TTA TTA TTA TTT TTT TTT GCA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA GCT ACT TTT ACA TCA GCT GCT ACT ATA ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAT TAC GTT GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATC ATG GCA GGA TTT GTC CAT TGA
TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATA CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTC TTT TTT TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTG CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
GC

#H._rapax_E55_TN_(Washington)_USA

--- --- -TG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GGA GAC CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAT TAC GTT GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATC ATG GCA GGA TTT GTC CAT TGA
TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATA CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTC TTT TTT TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
GC

#H._rapax_AZ65_NC_(Durham)_USA

CTA TTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GGA GAC CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAT TAC GTT GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATC ATG GCA GGA TTT GTC CAT TGA
TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATA CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTC TTT TTT TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
GC

#H._rapax_AZ68_NC_(Orange)_USA

-TA TTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GGA GAC CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAT TAC GTT GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATC ATG GCA GGA TTT GTC CAT TGA
TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATA CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTC TTT TTT TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTG CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
GC

#H._rapax_AP71_NY_(Schuyler)_USA

--- --- -TG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GGA GAC CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAT TAC GTT GTT GCT CAT -T- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---

#H._rapax_AS66_WI_(Barron)_USA

CTA TTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC CTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTA GAT TTA GCT ATC TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GGA GAC CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT TTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT

GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTC TTT TTC TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
GC

#H. rapax_AU12_MN (Kittson)_USA
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AAT TTT ATT ACG ACA GTA ATT AAT ATG CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CCA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCA GGA GGA GCA GAC CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGG ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
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AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA ACT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
TAT TAC GTT GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCA GTA TTT GCA ATC ATG GCA GGA TTT GTC CAT TGA
TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC TTA GGA TTA GCT GGA ATA CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTC TTT TTC TTT ATT ATT TGA GAA
AGT CTT GCA TCT CAA CGA CAA GTA ATA TTC CCA GTG CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCT CCA
GC

#H. rapax_AU34_MN (Koochiching)_USA
CTA TTA TTG GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCC TTA TCT TCT AAT ATC
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AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT TCA GGA ATT ACT TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTG TTA TTA TCG TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CCA AAT
ATT AAT ACT TCT TTC TTT GAC CCT GCG GGA GGA GGA GAT CCA ATT CTT TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCA GAA GTT TAC ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACA TTT GGA TCA CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTT GAT ACA CGA GCT TAT GTT ACA TCA GCT ACT ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTC AGT TGA TTA GCA ACC CTT TAT GGA ACT CAA CTA AAT TAT TCC CCT GCT ACA CTT TGA GCA TTA GGA TTT
GTT TTC TTA TTC ACT GTA GGA GGA TTA ACA GGA GTA GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACA
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TAC CCT CTT TTC ACA GGA TTA ACT CTA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAT TCT GAC TAT CCA GAC GCT TAT ACT
GCA TGA AAT GTA ATT TCT ACT ATT GGA TCA ACA ATT TCA TTA CTA GGA ATT TTA TTC TTT TTC TTT ATT ATT TGA GAA
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GC

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TCAACACTTC TTAGGATTTG CCGGAATACC ACGACGATAT TCAGACTACC CAGATGCCTA TACAACTTGA AATGTAATTT TACTATTGTT GTCTCAATTT TCTTTATTAG
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#L. cluvia_BF79_FL_Highland
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TCAACATTTT TTAGGATTAG CAGGAATACC ACGACGATAT TCAGACTACC CAGATGCTTA CACAACCTGA AATGTAATTT CTACAATTGG GTCAACAATT TCTTTATTAG
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TCAACATTTT TTAGGATTAG CAGGAATACC ACGACGATAT TCAGACTACC CAGATGCTTA CACAACCTGA AATGTAATTT CTACAATTGG GTCAACAATT TCTTTATTAG
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#L_cuprina_BG02_FL_Monroe -----ATTAG TTAGTAGTAT AGTAGAAAAC GGAGCTGGAA CAGGATGAAC AGTTTACCCT CCCCTATCTT
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Genetic marker data for various species including Clark, Douglas, Crook, Hamilton, and Chautauqua. Each section lists individuals and their genotypes across multiple loci.

TCAACATTTT TTAGGACTAG CAGGAATACC GCGACGATAC TCAGATTACC CAGATGCTTA CACAACCTGA AATGTAATTT CTACAATTGG GTCAACAATT TCCTATTAG
 GAATTTTATT TTCTTTTTTC ATTTATTTAG AAAGCTCTGT AACTCAACGT CAAGTTTTAT TCCTGTTCCTA ATTAATTTCA TCAATTGAAT GACTACAAAA TACTCCA--- --
 #L_illustris_AX11_OH Hamilton

 -----TAGCTG GAGCTATAAC TACTTTTTTA
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 #L_illustris_BA61_MN Winona

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 #L_illustris_BB44_MN Koochichin

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 #L_illustris_BB68_MN Kitson

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 #L_illustris_BE09_NY Schuyler

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 #L_illustris_BE10_NY Schuyler

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 #L_mexicana_AE60_CA Glenn

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 #L_mexicana_AE63_CA Glenn

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#L_mexicana_AX39_NM_Grant

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#L_mexicana_AX40_NM_Grant

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#L_mexicana_AX41_NM_Grant

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#L_mexicana_AX42_NM_Grant

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#Me_salvum_BA31_OH_(Hamilton)_USA

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#Me_salvum_AZ64_NC_(Durham)_USA

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#Mi_adelphie_AV26_NY_(Onandaga)_USA

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AAG ATT TTT AGT TGA TTA GCT ACA TTA TAC GGA ACT CAA CTA ACT TAT TCT CCA GCT ATT TTA TGA GCT TTA GGA TTC
GTA TTC TTA TTT ACT GTA GGA GGT TTA ACA GGA GTA GTA CTA GCT AAC TCA TCT GTT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GAT CAT TGA
TAC CCT CTA TTT ACT GGA TTA ACT CTA AAT AAT AAA CTT TTA AAA AGT CAA TTT GTT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTC TTA GGA TTA GCC GGA ATA CCT CGA CGA TAT TCT GAT TAT CCT GAT GCT TAT ACA
GCA TGA AAT GTA ATT TCA ACA ATC GGT TCA ACA ATT TCA TTA TTA GGA ATT TTA TAT TTA TTC TAT ATT ATC TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTA ATT TTC CCA ATT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA C--
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#Mu._domestica_BG76_FL_(Broward)_USA
CTA TTA TTA GTA AGA AGT ATA GTA GAA AAG GGA GCT GGA ACA GGT TGA ACT GTT TAT CCA CCT TTA TCA TCA ATT ATT
GCT CAT GGT GGA GCT TCA GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACA ACT GTT ATT AAC ATA CGA TCA ACA GGT ATT ACA TTC GAT CGA AT- CCT TTA TTT GTT TGA TCA GTT
GTA ATT ACT GCA TTA TTA TTA TTA TTA TCT CTT CCF GTT CTT GCT GGA GCT ATT ACT ATA CTA TTA ACT GAT CGA AAT
TTA AAT ACT TCA TTC TTT GAC C-A GCT GGA GGA GGT GAT CCA ATT CTT TAT CAA CAC TTA TTC TGA TTC TTT GGA CAT
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACA TTC GGT TCT TTA GGA ATA ATT TAT GCT ATG TTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACT CGA GCT TAC TTC ACT TCA GCT ACA ATA ATT ATT GCT GTA CCT ACT GGA ATT
AAG ATT TTT AGT TGA TTA GCT ACA TTA TAC GGA ACT CAA CTA ACT TAT TCT CCA GCT ATT TTA TGA GCT TTA GGA TTC
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TAT TAT GTA GTT GCT CAT TTC CAC TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GAT CAT TGA
TAC -CT CTA TTT ACT GGA TTA ACT CTA AAT AAT AAA CTT TTA AAA AGT CAA TTT GTT ATT ATA TTT ATT GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCC GGA ATA CCT CGA CGA TAT TCT GAT TAT CCT GAT GCT TAT ACA
GCA TGA AAT GTA ATT TCA ACA ATC GGT TCA ACA ATT TCA TTA TTA GGA ATT TTA TAT TTA TTC TAT ATT ATC TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTA ATT TTC CCA ATT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA C--
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#Ox._cingarus_AP68_NY_(Schuyler)_USA
CTT TTA CTA GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATC TCA TCT ATT TTA GGA GCT GTT
AAT TTT ATT ACA ACA GTA ATT AAT AAT GCA TCA ACA GGT ATT ACT TTT GAT CGA ATA CCA TTA TTT GTC TGA TCA GTA
GTA ATT ACT GCA TTA TTG TTA CTT CTT TCT TTA COT GTT CTA GCA GGA GCT ATT ACT ATA CTA TTA ACG GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCC GCA GGA GGA GGA GAC CCA ATT CTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACA GCA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCA TGT CCT ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT TAC CAA TTA AAT TAT TCT CCA GCT ACC TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACT GTT GGA GGA TTG ACT GGA GTA GTT TTA GCT AAT TCT TCT GTT GAT ATT ATT CTT CAT GAT ACT
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ACT TGA AAT GTT ATT TCA ACA ATT GGT TCA ACA ATT TCT CTA TTA GGA ATT TTA TTT TTC TTC TAC ATC ATT TGA GAA
AGT TTA GCC TCT CAA CGA CAA GTA ATA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC
#Ox._ventricosa_E8_OH_(Hamilton)_USA
CCT CTA CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATC TCA TCT ATT TTA GGA GCT GTT
AAT TTT ATT ACA ACA GTA ATT AAC ATA CGA TCA ACA GGT ATT ACT TTC GAT CGA ATA CCA TTA TTT GTC TGA TCA GTA
GTA ATT ACT GCA TTA TTA TTA CTT CTT TCT TTA COT GTT CTA GCA GGA GCT ATT ACT ATA CTA TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAC CCT GCA GGA GGA GGA GAC CCA ATT CTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCT GA- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ---- ----
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACG CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCG GTT CCT ACT GGA ATT
AAA ATT TTT AGT TGA TTT GCA ACT CTT TAT GGT ACA CAA TTA AAT TAT TCT CCA GCT ACC TTA TGA GCT CAT CAC ATA
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TAC CCA TTA TTT ACA GGA TTA ACA TTA AAC TCA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
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ACT TGA AAT GTT ATT TCA ACA ATT GGT TCA ACA ATT TCT CTA TTA GGA ATT TTA TTT TTC TTC TAC ATC ATT TGA GAA
AGT TTA GCC TCT CAA CGA CAA GTA ATA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
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CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT TGT TTA TTA GGA TTT ATT GGT TGA GCT CAT CAC ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACG GCA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCG GTF CCT ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGT ACA CAA TTA AAT TAT TCT CCA GCT ACC TTA TGA GCT TTA GAA TTT
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TAC CCA TTA TTT ACA GGA TTA ACA TTA AAC TCA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTT CCA CAA CAC TTC TTA GGA CTA GCT GGG ATA CCT CGA CGA TAC TCT GAC TAT CCT GAT GCC TAT ACA
ACT TGA AAT GGT ATT TCA ACA ATT GGT TCA ACA ATT TCT CTA TTA GGA ATT TTA TTT TTT TTC TAC ATC ATT TGA GAA
AGT TTA GCC TCT CAA CGA CAA GTG ATA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AA- --- --- ---

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#Ox._ventricosa_E62_TN_(Washington)_USA

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-TT CTA CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATC TCA TCT ATT TTA GGA GCT GTT
AAT TTT ATT ACA ACA GTA ATT AAC ATA CGA TCA ACA GGT ATT ACT TTC GAT CGA ATA CCA TTA TTT GTC TGA TCA GTA
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ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT TTA TTA GGA TTT ATT GGT TGA GCT CAT CAC ATA
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ACT TGA AAT GGT ATT TCA ACA ATT GGT TCA ACA ATT TCT CTA TTA GGA ATT TTA TTT TTT TTC TAC ATC ATT TGA GAA
AGT TTA GCC TCT CAA CGA CAA GTG ATA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC

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#P._regina_AB62_FL_(Washington)_USA

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-TT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
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AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TT- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---

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#P._regina_AB64_FL_(Washington)_USA

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--T ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
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AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT T-- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---

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#P._regina_AD71_SC_(Newberry)_USA

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GCT CAT GGA GGA GCA TCT GTT GAT CTA GCT ATT TTC TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT ATT ACT ATA TTA TTA ACT GAT CCA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CT ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?
??? ??? GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT TGA TGA GCT CAT CAT ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
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GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GCA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT ACA ATT GGT TCA ACA ATC TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGT CAA GTT TTA TTC CCT GTA CAA TTA AAT TCA TCT ATT GAA TGA TTA CAA AAT ACT CCA CCA
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#P._regina_AD72_SC_(Newberry)_USA

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AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CCA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
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AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA

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TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
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TTA ACA TTC TTC CCT CAA CAT TTC TTT GGA ATT GGT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT ACA ATT GGT TCA ACA ATC TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
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#P._regina_AE67_CA_(Glenn)_USA
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GCT CAT GGA GGA GCA TCT GTT GAT CTA GCT ATT TTC TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA TTA CCT GTA TTA GCC GGT ATT ACT ATA TTA TTA ACT GAT CCA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT C?? ??? ??? ??? ??? ??? ??? ??? ???
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TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT ACA ATT GGT TCA ACA ATC TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGT CAA GTT TTA TTC CCT GTA CAA TTA AAT TCA TCT ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC
#P._regina_AF39_CA_(Yolo)_USA
GCT CAT GGA GGA GCA TCT GTT GAT CTA GCT ATT TTC TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CCA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT ACA ATT GGT TCA ACA ATC TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGT CAA --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#P._regina_AG01_OR_(Crook)_USA
GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CCA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT AC- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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#P._regina_AI17_OR_(Douglas)_USA
GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CCA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG

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#P._regina_AI18_OR_(Douglas)_USA
GTT GAT CTA GCT ATT TTC TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CCA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATC GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT ACA ATT GGT TCA ACA ATC TCA --- --- --- --- --- --- --- --- --- --- --- --- --- ---
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AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GAT GAA TTT
GTA TTT TTA TTT ACT GTA GGA GAA TTA ACT GGT TTT GTT TCA TCA ACT GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTC AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GCA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT ACA ATT GGT TGA ACA ATC TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TGA GTA TCT CAA CGT CAA GTT TTA TTC CCT GTA CAA TTA AAT TCA TCT ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#P._regina_BF36_NM_(Dona_Ana)_USA

CTA TTG TTA GTT AGT AGT ATA GTA GAA AAT GGG GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCA TCT GTT GAT CTA GCT ATT TTC TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA TCA CCT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTW WTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT GAA GTT TAT ATT TTA ATT CTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAG AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA TCA TCT TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AGT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT TTT GTT TTA GCT AAT TCA TCA ACT GAT ATT ATT CTT CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TTA AAT AAT AAG TTA TTA AAA AGT CAA TTT GCT ATT ATA TTT ATT GGG GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACG
GCT TGA AAC GTA ATC TCT ACA ATT GGT TCA ACA ATC TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGT CAA GTT TTA TTC CCT GTA CAA TTA AAT TCA TCT ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#P._regina_E51_TN_(Washington)_USA

CTA TTG TTA GTT AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCA TCT GTT GAT CTA GCT ATT TTC TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTC ATT ACA ACT GTA ATT AAT ATA CGA TCA ACT GGA ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTT ATT ACT GCT CTA TTA CTT TTA TTA TCT TTA CCT GTA TTA GCC GGT GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
TTA AAC ACT TCA TTC TTT GAC CCA GCA GGA GGA GAT CCT ATT TTA TAT CAA CAC TTA TTC TGA TTC TTT GGT CAC
CCT G-- ---
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTT GGA ATA GAC GTT GAT ACA CGA GCT TAC TTT ACT TCA GCA ACT ATA ATT ATT GCT GTA CCA ACT GGA ATT
AAA ATT TTC AAT TGA CTA GCA ACT CTT TAT GGA ACT CAA TTA AAT TAC TCT CCA GCA ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA TTA ACT GGT GTT GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAC GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA TTA TCA ATA GGA GCT GTA TTC GCT ATT ATA GCT GGA TGT GTC CAC TGA
TTC CCT TTA TTT ACT GGA TTA ACA TT-- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- --- ---
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA TTG GCT GGA ATA CCT CGA CGA TAC TCA GAT TAC CCA GAT GCT TAC ACA
GCT TGA AAC GTA ATC TCT ACA ATT GGT TCA ACA ATC TCA TTA TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGT CAA GTT TTA TTT CCT GTA CAA TTA AAT TCA TCT ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#Po._labialis_AT02_WI_(Douglas)_USA

CTA CTG TTA GTA AGT AGT ATA GTG GAA AAC GGA GCT GGA ACA GGA TGA ACT GTC TAC CCC CCA CTA TCA TCT AAC ATT
GCT CAT GGA GGA GCC TCT GTT GAT TTA GCT ATT TTT TCC CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACT GTA ATT AAT ATA CGA TCT ACA GGT ATT ACA CTT GAC CGA ATA CCC TTG TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTG TTA CTT TTT TTG TCT TTA CCA GTA TTA GCA GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
TTA AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCA GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGG ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAA GAA
ACT TTT GGG GCT TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT TTA CTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATA GAT GTT GAT ACT CGA GCT TAT TTT ACA TCT GCT ACT ATA GTA ATT GCA GTT CCT ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACC TTA TAT GGA ACT CAA ATA AAT CAT TCC CCA GCT ACT TTA TGG TCT TTG GGA TTT
ATT TTT TTG TTC ACA GTA GGA GGG TTG ACA GGA TTT GTA CTA GCT AAC TCT TCA CTA GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTT CAC TAT GTA CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCC GGA TTT GTA CAT TGA
TAC CCT CTT TTT ACA GGT TTA ACA ATA AAT TCA ATA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTT TTC CCC CAA CAT TTC TTA GGA TTA GCA GGA ATA CCT CGT CGA TAT TCA GAT TAT CCT GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT TCA ATT GGA TCA ACA ATT TCA TT-- --- --- --- --- --- --- --- --- --- --- --- ---

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#Po._pediculata_AP60_NY_(Schuyler)_USA

TTA TTA TTG GTG AGC AGT ATA GWG GAA AAC GGA GCW GGG ACA GGA TGA ACT GTT TAC CCA CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGG GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACA ACT GTA ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCA TTA TTA CTT CTT TTA TCC TTA CCT GTA TTA GCA GGA GAT ATC ACT ATA TTA TTA ACA GAT CGA AAT
TTA AAT ACT TCC TTT TTT GAT CCT GCA GGA GGA GAC CCA ATT TTA TAT GAA CAT TTA TTT TGA TTT TTT GGA CAC
CCA GAA GTG TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGA GCT TTA GGA ATA ATT TAT GCA ATA CTA GCT ATT GGC TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGG ATA GAT GTA GAT ACT CGA GCT TAT TTT ACA TCT GCT ACA TCA GAT TTT ATT GCT CCT ACT GGA ATT
AAA ATT TTT AGC TGA TTA GCA ACC CTT TAC GGA ACT CAA ATA AAT CAC TCC CCA GCC ACA TTA TGA TCT TTA GGA TTT
ATT TTT TTA TTT ACA TTA GGA GGA CTA ACT GGT GTT GTT CTT GCT AAT TCT TCC CTA GAC ATT ATT CTT CAT GAT ACT
TAC TAT GTA GTT GCT CAT TTT CAC TAT GTA CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAC CCT CTT TTT ACA GGA TTA ACA ATA AAT ACA ATA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTT TTT CCT CAA CAT TTC TTA GGA TTA GCA GGA ATA CCT CGT CGA TAT TCA GAT TAT CCT GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCG ACA ATT TCA TTT TTA GGA ATT CTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA ATA TCA CAA CGT CAA GTT TTA TTC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACA CCA CCA
GC

#Po._pediculata_AR74_WI_(Waukesha)_USA

TTA TTA TTG GTG AGC AGT ATA GTG GAA AAC GGA GCT GGG ACA GGA TGA ACT GTT TAC CCA CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGG GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACA ACT GTA ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCA TTA TTA CTT CTT TTA TCC TTA CCT GTA TTA GCA GGA GAT ATC ACT ATA TTA TTA ACA GAT CGA AAT
TTA AAT ACT TCC TTT TTT GAT CCT GCA GGA GGA GAC CCA ATT TTA TAT GAA CAT TTA TTT TGA TTT TTT GGA CAC
CCA GAA GTG TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGA GCT TTA GGA ATA ATT TAT GCA ATA CTA GCT ATT GGC TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGG ATA GAT GTA GAT ACT CGA GCT TAT TTT ACA TCT GCT ACA ATA GAT TTT ATT GCT CCT ACT GGA ATT
AAA ATT TTT AGC TGA TTA GCA ACC CTT TAC GGA ACT CAA ATA AAT CAC TCC CCA GCC ACA TTA TGA TCT TTA GGA TTT
ATT TTT TTA TTT ACA GGA GGA CTA ACT GGT GTT GTT CTT GCT AAT TCT TCC CTA GAC ATT ATT CTT CAT GAT ACT
TAC TAT GTA GTT GCT CAT TTT CAC TAT GTA CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAC CCT CTT TTT ACA GGA TTA ACA ATA AAT ACA ATA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTT TTT CCT CAA CAT TTC TTA GGA TTA GCA GGA ATA CCT CGT CGA TAT TCA GAT TAT CCT GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCG ACA ATT TCA TTT TTA GGA ATT CTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA ATA TCA CAA CGT CAA GTT TTA TTC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACA CCA CCA
GC

GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACA ACT GTA ATT AAT AAT ATA CGA TCT AAT GGT ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCA CTA TTA CTT CTT TTA TCC TTA COT GTA TTA GCA GGA GCT ATC ACT ATA TTA TTA ACA GAT CGA AAT
TTA AAT ACT TCC TTT TTT GAT CCT GCA GGA GGA GAC CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCA GAA GTG TAT ATT TTA ATT TTA CCT GCA GGA TTT GCA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTT GGA GCT TTA GGA ATA ATT TAT GCA ATA CTG GCT ATT GGC TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATG
TTT ACT GTA GGA ATA GAT GFA GAT ACT CGA GCT TAT TTT ACA TCT GCT ACA ATA GTT ATT GCA GTT CCT ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCA ACC CTT TAC GCA ACT CAA ATA AAT CAC TCC CCA GCC ACA TTA TGA TCT TTA GGA TTT
ATT TTT TTA TTT ACA GTA GGA GGA CTA ACT GGT GTT GTT CTT GCT AAT TCT TCC CTA GAC ATT ATT CTT CAT GAT ACT
TAC TAT GTA GTT GCT CAT TTT CAC TAT GTA CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAC CCT CTT TTT ACA GGA TTA ACA ATA AAT ACA ATA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTT TTT CCT CAA CAT TTC TTA GGA CTA GCA GGT ATA CCA CGA CGA TAC TCA GAT TAT CCT GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCT ACT ATT GGA TCG ACA ATT TCA TTT TTA GGA ATT CTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA ATA TCA CAA CGT CAA GTT TTA TTT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACA CCA CCA
GC

#Po._rudis_AQ33_NY_(Richmond)_USA

CTA CTG TTA GTG AGC AGT ATA GTG GAA AAT GGA GCT GGG ACA GGA TGA ACT GTT TAC CCT CCA CTA TCT TCT AAC ATT
GCT CAT GGA GGA GCT TCT GTT GAC TTA GCT ATT TTT TCC CTT CAT TTA RCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACA ACT GTA ATT AAT ATA CGA TCC ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
KTA ATT ACA GCA CTA TTA CTT CTT TTA TCT TTA COT GTA TTA GCC GGG GCT ATT ACT ATA TTA TTA ACA GAT CGA AAC
TTA AAT ACT TCC TTC TTT GAT CCT GCA GGA GGA GGA GAC CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GST CAC
CCA SAA GTA TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAA GAA
ACT TTT GGA GCT TTA GGT ATA ATT TAT GCA ATA TTA GCA ATT TGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATA GAT GTA GAT ACT CGA GCT TAC TTT ACA TCC GCT ACA ATA GTT ATT GCA GTT CCT ACT GGT ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACT CAA ATA AAT TAC TCC CCA GCT ACA TTA TGA TCT TTA GGT TTT
ATT TTT TTA TTT ACT GTA GGA GGA TTA ACA GGT GTC GTC CTT GCT AAT TCT TCC CTA GAT ATT ATC CTT CAT GAT ACT
TAT TAT GTA GTT GCC CAT TTT CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCC GGA TTT GTT CAT TGA
TAC CCT CTT TTT ACA GGA TTA ACA ATA AAT ACC ATG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTT TTC CCT CAA CAT TTT TTA GGA TTA GCA GGT ATA CCA CGA CGA TAC TCA GAT TAT CCT GAT GCT TAT ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGA TCA ACA ATT TCA TTC TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA ATG TCC CAA CGT CAA GTT TTA TTT CCC ATT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
GC

#Po._rudis_AR75_WI_(Waukesha)_USA

CTA CTA TTG GTG AGC AGT ATA GTG GAA AAT GGA GCT GGG ACA GGA TGA ACT GTT TAC CCT CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATY TTT TCC CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACA ACT GTT ATT AAT ATG CGA TCT ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCA TTA TTA CTC CTT TTR TCC TTA COT GTA TTA GCA GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
TTA AAT ACT TCA TTT TTT GAY CCT GCA GGA GGT GAT TTA TTA TAY GAY CAT TTA TTT TGA TTT TTT GWG CAC
CCA GAA GTA TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAA GAA
ACT TTT GGG GCT TTA GGA ATA ATT TAC GCA ATA TTA GCA ATT GGC TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATA GAC GFA GAT ACT CGA GCT TAT TTT ACA TCC GCT ACA ATA GTT ATT GCA GTT CCT ACT GGA ATT
AAA ATT TTT AGT TGA CTA GCA ACC CTT TAT GGA ACT CAA ATA AAT CAC TCC CCA GCC ACA TTA TGA TCT TTA GGA TTT
ATT TTT TTA TTT ACA GTA GGA GGA CTA ACA GGT GTT GTT CTT GCT AAT TCT TCC CTA GAT ATT ATT CTC CAT GAT ACT
TAT TAT GTA GTC GCT CAT TTT CAC TAT GCA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAC CCT CTT TTT ACA GGA TTA ACA ATA AAT ACA ATA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTC TTT CCT CAA CAT TTC TTA GGA TTA GCA GGA ATT CCT CGA CGA TAC TCA GAT TAT CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGG TCA ACA ATT TCA TTC TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA ATA TCA CAA CGT CAA GTT TTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACA CCA CCA
GC

#Po._rudis_AV81_NY_(Niagara)_USA

CTA CTA TTA GTG AGC AGT ATA GTG GAA AAT GGA GCT GGG ACA GGA TGA ACT GTT TAC CCT CCA CTA TCT TCT AAC ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATC TTT TCC CTT CAT TTA RCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACA ACT GTA ATT AAT ATA CGA TCC ACA GGT ATT ACA TTT GAT CGA ATG CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCA CTA TTA CTT CTT TTA TCT TTA COT GTA TTA GCC GGG GCT ATT ACT ATA TTA TTA ACA GAT CGA AAC
TTA AAT ACT TCC TTC TTT GAT CCT GCA GGA GGA GGA GAC CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCA GAA GTA TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAA GAA
ACT TTT GGA GCT TTA GGT ATA ATT TAT GCA ATA TTA GCA ATT GGT TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATA GAT GTA GAT ACT CGA GCT TAC TTT ACA TCC GCT ACA ATA GTT ATT GCA GTT CCT ACT GGT ATT
AAA ATT TTT AGT TGA TTA GCA ACT CTT TAT GGA ACC CAA ATA AAT TAC TCC CCA GCT ACA TTA TGA TCT TTA GGG TTT
ATT TTT TTA TTT ACT GTA GGA GGA TTA ACA GGT GTT GTC CTT GCT AAT TCT TCC CTA GAT ATT ATC CTT CAT GAT ACT
TAT TAT GTA GTT GCC CAT TTT CAT TAT GTT CTT TCA ATA GGA GCT GTA TTT GCT ATT ATA GCC GGA TTT GTT CAT TGA
TAC CCT CTT TTT ACA GGA TTA ACA ATA AAT ACC ATG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTT TTC CCT CAA CAT TTT TTA GGA TTA GCA GGT ATA CCA CGA CGA TAC TCA GAT TAT CCT GAT GCT TAT ACA
ACT TGA AAT GTA GTT TCT ACT ATT GGA TCA ACA ATT TCA TTC TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA ATT TCC CAA CGT CAA GTT TTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
GC

#Po._rudis_AW13_NY_(Monroe)_USA

CTA CTA TTG GTG AGC AGT ATA GTG GAA AAC GGA GCT GGG ACA GGA TGA ACT GTT TAC CCT CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATC TTT TCC CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACA ACT GTT ATT AAT ATG CGA TCT ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCA TTA TTA CTC CTT TTG TCC TTA CCT GTA TTA GCA GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
TTA AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCA GAA GTA TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAA GAA
ACT TTT GGG GCT TTA GGA ATA ATT TAC GCA ATA TTA GCA ATT GGC TTA TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATA GAC GFA GAT ACT CGA GCT TAT TTT ACA TCC GCT ACA ATA GTT ATT GCA GTT CCT ACT GGA ATT
AAA ATT TTT AGT TGA CTA GCA ACC CTT TAT GGA ACT CAA ATA AAT CAC TCC CCA GCC ACA TTA TGA TCT TTA GGA TTT
ATT TTT TTA TTT ACA GTA GGA GGA CTA ACA GGT GTT GTT CTT GCT AAT TCT TCC CTA GAT ATT ATT CTC CAT GAT ACT
TAT TAT GTA GTC GCT CAT TTT CAC TAT GTA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAC CCT CTT TTT ACA GGA TTA ACA ATA AAT ACA ATA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATT GGA GTT AAT
TTA ACT TTC TTT CCT CAA CAT TTC TTA GGA TTA GCT GGA ATG CCT CGA CGA TAC TCA GAT TAT CCT GAT GCT TAC ACA
ACT TGA AAC GTA ATT TCT ACT ATT GGG TCA ACA ATT TCA TTC TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA ATA TCA CAA CGT CAA GTT TTA TTT CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA TTA CAA AAT ACT CCA CCA
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#Po._rudis_AZ63_NC_(Durham)_USA

CTA CTA TTG GTG AGC AGT ATA GTG GAA AAC GGA GCT GGG ACA GGA TGA ACT GTT TAC CCT CCT CTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATC TTT TCC CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACA ACT GTT ATT AAT ATG CGA TCT ACA GGA ATT ACA TTT GAC CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCA TTA TTA CTC CTT TTG TCC TTA CCT GTA TTA GCA GGA GCT ATT ACT ATA TTA TTA ACA GAT CGA AAT
TTA AAT ACT TCA TTT TTT GAC CCT GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGT CAC
CCA GAA GTA TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGA AAA AAA GAA

TAT CCT TTA TTT ACA GGA CTA ACA TTA AAT AGT AAA ATA TTA AAA AGT CAA TTT ACT ATC ATA TTC ATA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC TTA GCT CTA GCA GGA ATA CTT CGA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATT TCT ACA ATT GGA TCA ACA ATT TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT TAT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTA TTA TTT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC

#R._anxia_AT50_MN_(Kandiyohi)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTC CAT TTA GCA GGG ATT TCT TCA TCA ATT TTA GGG GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAC CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTA TTA TTA CTT TTA TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GAC CCT ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GCA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTC GGG TCA CTA GGT ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT GCA GCT TAC TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGA TGA TTA GCC ACA CTT TAT GGA ACT CAA TTA AAC TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAC TAT GTA CTA TCT TTA GGA GCA GTA TTT GCT ATT ATA GCA GGA TFC GAT CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACA TTA AAT AGC AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC TTA GGT CTT GCG GGA ATA CCT CGA CGA TAC TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATC TCT ACA ATT GGA TCA ACA ATT TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT TAT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTA TTA TTC CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCC CCA
GC

#R._derelicta_BA30_OH_(Hamilton)_USA
TTA TTA CTA GTT AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTC CAT TTA GCA GGG ATT TCT TCT ATT TTA GGA GCA GTT
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACT TTC GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATG CTT TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCT GCC GGA GGG GGT GAT CCA ATT TTA TAC CAA CAT TTA TTC TGA TTT TTC GGA CAT
CCT GAA GTA TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTT GGA TCA TTA GGT ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAC CAC ATG
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA CTA AAT TAT TCA CCT GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTT TTA TTC ACA GTA GGA GGT TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAC ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT TTA TCC ATA GGA GCT GTA TTT GCA ATT ATA GCC GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC CTT GGT CTA GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCC TAT ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA CTA GGT ATT TTA TTC TTT TTT TAC ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCT CCA
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#R._derelicta_AT48_MN_(Kandiyohi)_USA
--- --A CTA GTT AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTT CAT TTA GCA GGG ATT TCT TCT ATT TTA GGA GCA GTT
AAT TTT ATC ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACT TTC GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATG CTT TTA ACA GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAT CCT GCC GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTC TGA TTT TTC GGA CAT
CCT GAA GTA TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGT ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAC CAC ATG
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA CTA AAT TAT TCA CCT GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGT TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAC ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCC GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC CTT GGT CTA GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCC TAT ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA CTA GGT ATT TTA TTC TTT TTT TAC ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCT CCA
GC

#R._derelicta_AZ09_OH_(Hamilton)_USA
-TA TTA CTA GTT AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTT CAT TTA GCA GGG ATT TCT TCT ATT TTA GGA GCA GTT
AAT TTT ATC ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACT TTC GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATG CTT TTA ACA GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAT CCT GCC GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTC TGA TTT TTC GGA CAT
CCT GAA GTA TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGT ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAC CAC ATG
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA CTA AAT TAT TCA CCT GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGT TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAC ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCC GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC CTT GGT CTA GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCC TAT ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA CTA GGT ATT TTA TTC TTT TTT TAC ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA A-- --- --- ---
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#R._derelicta_AZ62_VA_(Bland)_USA
--A TTA CTA GTT AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTC CAT TTA GCA GGG ATT TCT TCT ATT TTA GGA GCA GTT
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACT TTC GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATG CTT TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCT GCC GGA GGG GAT CCA ATT TTA TAC CAA CAT TTA TTC TGA TTT TTC GGA CAT
CCT GAA GTA TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTT GGA TCA TTA GGT ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAC CAC ATG
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA CTA AAT TAT TCA CCT GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTT TTA TTC ACA GTA GGA GGT TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAC ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCC GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TCT CCT CAA CAT TTC CTT GGT CTA GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCC TAT ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA CTA GGT ATT TTA TTC TTT TTT TAC ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT A-- --- --- ---
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#R. derelicta_BB21_MN_(Kandiyohi)_USA

-TA TTA CTA GTT AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT CTA GCT ATC TTC TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGT GCT GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACT GCT TTA TTA CTT CTT TTA TCA TCA CCA GTT CTT GCT GGA GCT ATT ACT ATG CTT TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCT GCC GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTC TGA TTT TTC GGA CAT
CCT GAA GTA TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAA GAA
ACC TTT GGA TCA TTA GGT ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAC CAC ATG
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA CTA AAT TAT TCA CCT GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTT TTA TTC ACA GTA GGG GGT TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAC ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCC GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTC CTT GGT CTA GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCC TAT ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA CTA GGT ATT TTA TTC TTT TTT TAC ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT A-- --- ---

#R. derelicta_BA29_OH_(Hamilton)_USA

TTA TTA CTA GTT AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT TTA GAA AAT TTC TCT CTT CAT TTA GCA GGA ATT TCT TCT ATT TTA GGA GCA GTT
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACT TTC GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA CTT CTT TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATG CTT TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCT GCC GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTC TGA TTT TTC GGA CAT
CCT GAA GTA TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTT GGA TCA TTA GGT ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAC CAC ATG
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA CTA AAT TAT TCA CCT GCT ACT TTA TGA GCC TTA GGG TTT
GTA TTT TTA TTC ACA GTA GGG GGT TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAC ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCC GGA TTT GTA CAC TGA
TAC CCT CTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTC CTT GGT CTA GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCC TAT ACA
ACA TGA AAT GTT ATT TCA ACA ATT GGA TCA ACA ATT TCA TTA CTA GGT ATT TTA TTC TTT TTT TAC ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACT CCT CCA

#R. floridensis_AA01_FL_(Duval)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT CTA GCT ATC TTC TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGT GCT GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATC ACT GCT TTA TTA CTT CTT TTA TCA TTA CCA GTT CTT GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCT TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTA CCC ACA GGA ATT
AAA ATC TTC AGA TGA TTA GCC ACA CTT TAC GCA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTC ACA GTT GGA GGA TTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT TCT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGA CGT TAC TCA GAT TAT CCA GAT GCC TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTC TTT TTT TTC ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGT CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA

#R. floridensis_AA03_FL_(Duval)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT CTA GCT ATC TTC TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGT GCT GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATC ACT GCT TTA TTA CTT CTT TTA TCA TTA CCA GTT CTT GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCT TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTA CCC ACA GGA ATT
AAA ATC TTC AGA TGA TTA GCC ACA CTT TAC GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTC ACA GTT GGA GGA TTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT TCT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGA CGT TAC TCA GAT TAT CCA GAT GCC TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTC TTT TTT TTC ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGT CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA

#R. floridensis_AA60_FL_(Sarasota)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT CTA GCT ATC TTC TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGT GCT GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATC ACT GCT TTA TTA CTT CTT TTA TCA TTA CCA GTT CTT GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCT TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTA CCC ACA GGA ATT
AAA ATC TTC AGA TGA TTA GCC ACA CTT TAC GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTC ACA GTT GGA GGA TTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA TTA AAA AGT CAA TTT TCT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGA CGT TAC TCA GAT TAT CCA GAT GCC TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTC TTT TTT TTC ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGT CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA

#R. lherminieri_AA36_FL_(Madison)_USA

TTA CTT CTA GTG AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCC AAT ATC
GCT CAT GGA GGG GCC TCT GTC GAT CTA GCT ATC TTT TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACT GCT TTA TTA CTA CTT TTA TCA TTA CCA GTT CTT GCC GGA GCT ATT ACT ATA CTA TTA ACT GAT CGA AAT

ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GCG TTT GGA ATA ATT TCT CAT ATT ATT AGC CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT GAT CCA GAT GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TFC ACC TCA GCA ACA ATA ATT ATT GCC GTA CCA ACA GGG ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT TAT GGA ACT CAA TAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTT GCA GGG TTT ACA GGA GTT GTT TTA GAT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TFC CAT TAT GTG TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAC AGC AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGG GTA AAT
TTA ACG TTC TTT CCC CAA CAT TTC TTA GGT CTT GCA GGT ATA CCC CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGG TCA ACA ATT TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAG CGA CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCT CCA

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#R_lherminieri_AN14_TN_(Jefferson)_USA
TTA CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATC
GCT CAT GGA GGG GCC TCT GTC GAT CTA GCC ATC TTT TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACT GCT TTA TTA CTA CTT TTA TCA TTA CCA GTT CTT GCC GGA GCT ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGC CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTC ACC TCA GCA ACA ATA ATT ATT GCC GTA CCA ACA GGG ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT -TA GGA TTT
GTA TTC TTA TTT ACA GTT GCA GGG TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTG TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAC AAC AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGG GTA AAT
TTA ACG TTC TTT CCT CAA CAT TTC TTA GGT CTT GCA GGT ATA CCC CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGG TCA ACA ATT TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCT CCA

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#R_lherminieri_AN15_TN_(Jefferson)_USA
-TA CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATC
GCT CAT GGA GGG GCC TCT GTC GAT CTA GCC ATC TTT TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACT GCT TTA TTA CTA CTT TTA TCA TTA CCA GTT CTT GCC GGA GCT ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGC CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTC ACC TCA GCA ACA ATA ATT ATT GCC GTA CCA ACA GGG ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT C-T TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTT GGA GGG TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTG TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAC AAC AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACG TTC TTT CCT CAA CAT TTC TTA GGT CTT GCA GGT ATA CCC CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGG TCA ACA ATT TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA A-- --- --- --- --- --- --- ---

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#R_lherminieri_AN16_TN_(Jefferson)_USA
TTA CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATC
GCT CAT GGA GGG GCC TCT GTC GAT CTA GCC ATC TTT TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACT GCT TTA TTA CTA CTT TTA TCA TTA CCA GTT CTT GCC GGA GCT ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGC CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTC ACC TCA GCA ACA ATA ATT ATT GCC GTA CCA ACA GGG ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTT GGA GGG TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTG TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAC AAC AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACG TTC TTT CCT CAA CAT TTC TTA GGT CTT GCA GGT ATA CCC CGA CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGG TCA ACA ATT TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCT CCA

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#R_lherminieri_AN17_TN_(Jefferson)_USA
TTA CTT CTA GTG AGT AGT ATA GTA GAA AAT GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCC AAT ATC
GCT CAT GGA GGG GCC TCT GTC GAT CTA GCC ATC TTT TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGG GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATG CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACT GCT TTA TTA CTA CTT TTA TCA TTA CCA GTT CTT GCC GGA GCT ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGG ATA ATT TCT CAT ATT ATT AGC CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTC ACC TCA GCA ACA ATA ATT ATT GCC GTA CCA ACA GGG ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTT GGA GGG TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTG TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAC AGC AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGG GTA AAT
TTA ACG TTC TTT CCC CAA CAT TTC TTA GGT CTT GCA GGT ATA CCC CGG CGA TAC TCA GAC TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGG TCA ACA ATT TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAG CGA CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCT CCA

GC
#R_lherminieri_AZ44_WV_(Kanawha)_USA
TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT CTA GCT ATC TTC TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGT GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATC ACT GCT TTA TTA CTA CTT TTA TCA TTA CCA GTT CTT GCC GGA GCT ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTC GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCT TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTA CCC ACA GGA ATT
AAA ATC TTC AGA TGA TTA GCT ACA CTT TAC GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT

GTA TTC TTA TTC ACA GTT GGA GGA TTA ACA GGA GGT GTT CTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GAT TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT TCT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TCT CCT CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGA CGT TAC TCA GAT TAT CCA GAT GCC TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTC TTT TTT TTC ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGT CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA
GC

#R._lherminieri_AZ58_WV_(Kanawha)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCC TCT GTT GAT CTA GCT ATC TTC TCT CTA ACT TTA GCA GGA ATT TCA TCA ATT TTA GGT TCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT CAT TTC GAT CGA ATA TCT TTA TTT GTT TGA TCT GTA
GTA ATC ACT GCT TTA TTA CTT CTT TTA TCA TTA CCA GTT CTT GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCT TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT TGA CCC ACA GGA ATT
AAA ATC TTC AGA TGA TTA GCT ACA CTT TAC GCA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTC ACA GTT GGA GGA TTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT TCT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TCT CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGA CGT TAC TCA GAT TAT CCA GAT GCC TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTC TTT TTT TTC ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGT CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA
GC

#R._planifrons_AE45_OR_(Klamath)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA CTT TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GAC CCT ATT TTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGA TTC ATT GTT TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCA TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCT CCT GCT ACT TTA TGA GCT TTA GGA TTC
GTA TTC TTA TTT ACT TTA GGA GGT CTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAC ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA CTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TCT CAA CAT TTC TTA GGT CTT GCA GGA ATA CCT CGA CGA TAC TCT GAC TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT CTG TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTG TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC

#R._planifrons_AF05_CA_(Siskiyou)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA CTT TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GGA GGA GAC CCT ATT TTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGA TTC ATT GTT TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCA TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCT CCT GCT ACT TTA TGA GCT TTA GGA TTC
GTA TTC TTA TTT GCT GTA GGA GGT CTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAC ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA CTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTC ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TCT CAA CAT TTC TTA GGT CTT GCA GGA ATA CCT CGA CGA TAC TCT GAC TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT CTG TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTG TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC

#R._planifrons_AF06_CA_(Siskiyou)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA CTT TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GGA GGA GAC CCT ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGA TTC ATT GTT TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCA TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCT CCT GCT ACT TTA TGA GCT TTA GGA TTC
GTA TTC TTA TTT ACT GTA GGA GGT CTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAC ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA CTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TCT CAA CAT TTC TTA GGT CTT GCG GGA ATA CCT CGA CGA TAC TCT GAC TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT CTG TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTG TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC

#R._planifrons_AK08_OR_(Jefferson)_USA

--- --- --- --- --- --- --- -AA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAC TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA CTT TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCT ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GAC CCT ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGA TTC ATT GTT TGA GCC CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCA TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCT CCT GCT ACT TTA TGA GCT TTA GGA TTC
GTA TTC TTA TTT ACT GTA GGA GGT CTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAC ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCC CAT TTC CAT TAT GTA CTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TCT CAA CAT TTC TTA GGT CTT GCG GGA ATA CCT CGA CGA TAC TCT GAC TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT CTG TTA GGA ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTG TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC

AGT TTA GCA TCA CAA CGA CAA GTG TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA GC

#R_pusiola_AX63_NM_(Grant)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA
AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTA TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCA CTA GGA ATA ATT TAC GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAC TTT ACT TCA GCC ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAC ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA TTA TTC CCC GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA GC

#R_pusiola_AZ05_NM_(Grant)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA
AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTA TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCA CTA GGA ATA ATT TAC GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAC TTT ACT TCA GCC ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAC ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA TTA TTC CCC GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA GC

#R_pusiola_BA18_NM_(Grant)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA
AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTA TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCA CTA GGA ATA ATT TAC GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAC TTT ACT TCA GCC ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GWA GGA GGA TTA ACT GTT TTT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAC ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA TTA TTC CCC GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA GC

#R_pusiola_AX61_NM_(Grant)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTG TTA CTT TTA TCT TTA CCA GTT CTT GCA GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAT TTT ACT TCA GCA ACA ATA ATT ATT GCG GTT CCT ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCC ACA CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAT ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTA TCT ATG GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAC TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT A-- --- ---

#R_pusiola_AW64_NM_(Grant)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA
AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTA TTA CTT TTA TCT TTA CCA GTT CTT GCG GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCA CTA GGA ATA ATT TAC GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAT TTT ACT TCA GCC ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAT ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA TTA TTC CCC GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA GC

#R_pusiola_AW73_NM_(Grant)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA

AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTA TTA CTT TTA TCA TCT TTA CCA GTT CTT GCA GGA ATT ACT TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC CGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCA CTA GGA ATA ATT TAC GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GGT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAC TTT ACT TCA GCC ACA ATA ATT ATT GCT GGT CCT ACA GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACA CTT TAT GGA ATT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA AAT GGA GTT GGT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GCA ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ACT GAT TTA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATT CCA CGA TAT TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA TTA TTC CCC GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA A-- --- --- ---

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#R_pusiola_AX62_NM_(Grant)_USA

TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTG TTA CTT TTA TCA TCT TTA CCA GTT CTT GCA GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAA GAA
ACT TTT GGT TCT CTA GGA ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GGT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAC TTT ACT TCA GCC ACA ATA ATT ATT GCT GGT CCT ACA GGA ATT
AAA ATT TTT AGT TGA TTA GCC ACA CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GGT TTA TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAT ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAC TCT GAT TAT CCA GAT GCC TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGT ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTA TTA TTC CCT GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT A-- --- --- ---

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#R_pusiola_AY36_NM_(Grant)_USA

TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA
AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTA TTA CTT TTA TCT TTA CCA GTT CTT GCG GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
CCT GAA GTT TAT ATT TTA GAT TTA CCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC GGA CAT
ACT TTT GGT TCA CTA GGA ATA ATT TAC GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GGT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAC TTT ACT TCA GCC ACA ATA ATT ATT GCT GGT CCT ACA GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GGT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAC ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA TTA TTC CCC GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA

GC
#R_pusiola_BA19_NM_(Grant)_USA

TTA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCC GTA
AAT TTT ATT ACT ACA GTT ATT AAC ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCC GTA
GTA ATT ACT GCT TTA TTA TTA CTT TTA TCT TTA CCA GTT CTT GCG GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCA GCA GGA GGA GGT GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTT GGT TCA CTA GGA ATA ATT TAC GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GGT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCA TAC TTT ACT TCA GCC ACA ATA ATT ATT GCT GGT CCT ACA GGA ATT
AAA ATT TTC AGT TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GGT TTA TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAC ACT
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTT CTT TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTA CAT TGA
TAT CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTC TTT TTT TAT ATT ATC TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA TTA TTC CCC GTA CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA

GC
#R_lherminieri_A22_OH_(Hocking)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
GCT CAT GGA GGA GGC TCT GTT GAT TTA GCT ATC TTC TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGT GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATC ACT GCT TTA TTA CTT CTT TTA TCA TTA CCA GTT CTT GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GGA GAC CCA ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT T?? ??? ??? ?TA YCA GGA TTC GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCT TTA GGA ATT TAT TTA GCT ATA TTT GCA GCT ATT GGT CTT TTA GGA TTT ATT TTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTA CCC ACA GGA ATT
AAA ATC TTC AGA TGA TTA GCT ACA CTT TAC GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTT TTA TTC ACA GTT GGA GGA TTA ACA GGA GTT GGT TTA GCT AAT TCT TCT ATT GAT ATT ATC CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACT CTA AAT AGA AAG ATA CTA AAA AGT CAA TTT TCT GAT TAT CCA GAT GCT TAC ACA
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGA CGT TAC TCA GAT TAT CCA GAT GCC TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTC TTT TTT TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGT CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA

GC
#R_lherminieri_AQ58_NY_(Saratoga)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACT GGA TGA ACT GTT TAC CCA CCA TTA TCT TCT AAT ATT
GCT CAT GGA GGA GGC TCT GTT GAT TTA GCT ATC TTC TCT CTA CAT TTA GCA GGA ATT TCA TCA ATT TTA GGT GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATC ACT GCT TTA TTA CTT CTT TTA TCA TTA CCA GTT CTT GCT GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GGT GAT CCA ATT TTA TAT TAC CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTC GGA ATA ATC TCT CAT ATT ATT AGT CAA GAA TCT GGT AAA AAG GAA
ACT TTT GGA TCT TTA GGA ATA ATT TAT GCT ATA CTA GCT ATT GGT CTT TTA GGA TTT ATT TTA TGA GCT CAT CAT ATA

TTT ACA GTA GGA ATA GAT GTT GAT ACT CGA GCA TAT TTT ACC TCA GCA ACA ATA ATT ATT GCT GTA CCC ACA GGA ATT
AAA ATC TTC AGA TGA TTA GCT ACA CTT TAC GGA ACT CAA TTA AAT TAC TCC CCA GCT ACT CTT TGA GCT TTA GGA TTT
GTA TTC TTA TTC ACA GTT GCA GGA TTA ACA GGA GTT GTT CTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTT GCT CAT TTC CAT TAT GTA TTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT TCT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGA CGT TAC TCA GAT TAT CCA GAT GCC TAT ACA
ACT TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTC TTT TTT TTC ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGT CAA GTT TTA TTC CCT ATT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACC CCA CCA
GC
#R_ querula_AW30_NM_(Grant)_USA
--- TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCC TCT GTT GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCT TCA ATT TTA GGG GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA GCA GGA ATT ACT TTT GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTG TTA TTA CTT TTA TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GGA GAT CCT ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTT GGA TCA CTA GGT ATA ATT TAT GCT ATA TTT GAT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCT TAC TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAC TAT TCT CCA GCT ACT TTA TGG GCT TTA GGG TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAC ACT
TAT TAT GTA GTA GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACA ATA AAT AGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGT CTT GCA GGA ATA CCT CGA CGA TAC TCT GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAC GTT ATT TCT ACA ATT GGA TCA ACA ATC TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTA TTA TTT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA A-- --- --- ---
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#R_ querula_AZ61_VA_(Bland)_USA
TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATC
GCT CAT GGA GGA GCC TCT GTT GAT TTA GCT ATT TTT TCT CTA CAT TTA GCA GGA ATT TCT TCA ATT TTA GGG GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA GCA GGA ATT ACT TTT GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTG TTA TTA CTT TTA TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GGA GGA GGA GAT CCT ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTC GGA TCA CTA GGT ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCT TAC TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT TAT GGA ACT CAA TTA AAC TAT TCT CCA GCT ACT TTA TGG GCT TTA GGG TTT
GTA TTT TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAC ACT
TAT TAT GTA GTA GCC CAT TTC CAC TAT GTA TTA TCT ATA GGA GCA GTA TTT GCA ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACA ATA AAT AGT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGT CTT GCA GGA ATA CCT CGA CGA TAC TCT GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAC GTT ATT TCT ACA ATT GGA TCA ACA ATC TCT TTA CTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTA TTA TTT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACT CCA CCA
GC
#R_ querula_BA16_NM_(Grant)_USA
TTA TTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTA TAC CCT CCC TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA GCA GGA ATT ACT TTT GAT CGA ATA CCT CTA TTT GTT TGA TCT GTA
GTA ATT ACT GCC TTG TTA TTA CTT TTA TCT TTA CCA GTT CTT GCT GGA GCA ATT ACT ATA CTT TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GGA GGA GGA GAT CCT ATT TTA TAC CAA CAT TTA TTT TGA TTT TTT GGG CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACT TTC GGA TCA TTA GGT ATA ATT TAT GCT ATA TTA GCT ATT GGT CTT TTA GGT TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCT TAC TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTC AGA TGA TTA GCT ACA CTT CAT GAT TAC TCT CCA GCT ACT TTA TGA GCT TTA TGA GCT TTA GGG TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCT TCT ATT GAT ATT ATT CTT CAT GAC ACT
TAT ---

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#R_ stimulans_E7_OH_(Hamilton)_USA
CTT CTT CTA GTT AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCC TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACT TTC CAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA CTT CTT TCT TTA CCA GTT CTT GCA GGA GAT ATC ATA CTT TTA ACA GAT CGA AAT
GTT AAT ACT TCA TTT TTT GAT CCT GCT GGA GGA GGA GAT CCA ATT TTA TAC CAA CAT TT? ??? ??? ??? ??? ???
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ACC TTT GGA TCA TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCC TAT TTT ACT TCA GCT ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA CTA GCT ACA CTT TAT GGA ACT CAA TAT AAT TAT TCC CCT GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCT GGA TTT GTA CAC TGA
TAC CCT TTA TTC ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTC CCA CAA CAC TTC CTT GGT TTA GCA GGA ATA CCT CGA CGT TAT TCT GAC TAC CCA GAT GCC TAC ACA
ACA TGA AAT GTT ATC TCA ACA ATT GGA TCA ACA ATT TCA TTA TTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCA GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACC CCT CCA
GC
#R_ stimulans_AR33_NY_(Saratoga)_USA
CTT CTT CTA GTT AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCC TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACG GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCA GGA GCA ATC ACT ATA CTT TTA ACA GAT CGA AAT
GTT AAT ACT TCA TTT TTT GAT CCT GCT GGA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATC TCC CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTT GGA TCA TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCC TAT TTT ACT TCA GCT ACA ATA ATT ATT GCT GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA CTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCT GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCT GGA TTT GTA CAC TGA
TAT CCT TTA TTC ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT

-TT CTT CTA GTT AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCC TTA TCT TCT AAT ATC
GCT CAT GGA GCA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACT GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCA GGA GCA ATC ACT ATA CTT TTA ACA GAT CGA AAT
GTT AAT ACT TCA TTT TTT GAT CCT GCT GGA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTC GCA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA TAC TCC CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTT GGA TCA TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCC TAT TTT ACT TCA GCT ACA ATA ATT ATT GCC GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA CTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCT GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT A-- --- --- --- ---

#R._stimulans_AZ60_VA_(Balnd)_USA

--- --- CTA GTT AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCC TTA TCT TCT AAT ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTC TCT CTT CAT TTA GCA GGA ATT TCT TCA ATT TTA GGA GCA GTT
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACT GGA ATT ACT TTC GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCA GGA GCA ATC ACT ATA CTT TTA ACA GAT CGA AAT
GTT AAT ACT TCA TTT TTT GAT CCT GCT GGA GGA GGA GAT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTT TTC GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA TAC TCC CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACC TTT GGA TCA TTA GGA ATA ATT TAT GCA ATA TTA GCT ATT GGT CTT TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACT CGA GCC TAT TTT ACT TCA GCT ACA ATA ATT ATT GCC GTT CCT ACA GGT ATT
AAA ATT TTT AGT TGA CTA GCT ACA CTT TAT GGA ACT CAA TTA AAT TAT TCC CCC GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACA GGA GTA GTA TTA GCT AAT TCA TCA ATT GAT ATT ATT CTT CAT GAT ACT
TAC TAT GTA GTA GCT CAC TTC CAT TAT GTT CTA TCT ATA GGA GCT GTA TTT GCA ATT ATA GCT GGA TTT GTA CAC TGA
TAT CCT TTA TTC ACA GGA TTA ACA TTA AAT AGA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TCC CAA CAC TTC CTT GGT TTA GCG GGA ATA CCT CGA CGT TAT TCT GAC TAC CCA GAT GCC TAC ACA
ACA TGA AAT GTC ATC TCA ACA ATT GGA TCG ACA ATT TCA TTA TTA GGT ATT TTA TTC TTT TTT TAT ATT ATT TGA GAA
AGT TTA GCT TCA CAA CGA CAA GTA TTA TTC CCA GTA CAA TTA AAT TCA TCA ATT GAA TGA CTA CAA AAT ACC CCC CCA
GC

#Sx_lambens_BG79_FL_(Broward)_USA

TTA CTT CTT GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT CTA TCT TCT AAT ATT
GCT CAT GGT GGA GCA TCT GTT GAT TTA GCT ATC TTT TCT CTT CAT TTA GCT GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTT ATC ACT ACA GTA ATT AAT ATA CGA TCT ACT GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCA GGA GCA ATC ACT ATA CTT TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT CTT TAT CAA CAT CTA TTT TGA TTT TTT GGG CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCG GGT AAA AAG GAA
ACA TTT GGA TCT CTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA CTT TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAC GTA GAT ACA CGA GCT TAT TTT ACA TCT GCT ACA ATA ATT ATT GCT GTC CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACT CTA TAC GGA ACA CAA CTA AAT TAT TCA CCA GCT ACA CTT TGA GCT CTT GGT TTT
GTA TTT TTA TTC ACA GTA GGA TTA ACA GGA TTA GTA TTA GCT AAT TCT TCA ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTA GCA CAT TTT CAT TAT GTT CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGG TTT GTA CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT ATA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TCC CCT CAA CAC TTC TTA GGA CTT GCA GGA ATA CCT CGT CGA TAC TCA GAT TAT CCA GAT GCT TAT ACA
GCA TGA AAT GTA ATT TCA ACT ATC GGT TCA ACA ATT TCT TTA CTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGA TTA GTA TCC CAA CGA CAA GTA ATA TTT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT AC- --- ---

#Sx._lambens_BG80_FL_(Broward)_USA

TTA CTT CTT GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAT CCA CCT CTA TCT TCT AAT ATT
GCT CAT GGT GGA GCA TCT GTT GAT TTA GCT ATC TTT TCT CTT CAT TTA GCT GGA ATT TCA TCA ATT TTA GGA GCA GTA
AAT TTT ATC ACT ACA GTA ATT AAT ATA CGA TCT ACT GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACT GCT TTA TTA TTA CTT CTT TCT TTA CCA GTT CTT GCA GGA GCT ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GCA GGA GGA GAT CCT ATT CTT TAT CAA CAT CTA TTT TGA TTT TTT GGG CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCG GGT AAA AAG GAA
ACA TTT GGA TCT CTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA CTT TTA GGA TTT ATT GTT TGA GCT CAT CAT ATA
TTT ACT GTA GGA ATG GAC GTA GAT ACA CGA GCT TAT TTT ACA TCT GCT ACA ATA ATT ATT GCT GTC CCA ACT GGA ATT
AAA ATT TTT AGT TGA TTA GCT ACT CTA TAC GGA ACA CAA CTA AAT TAT TCA CCA GCT ACA CTT TGA GCT CTT GGT TTT
GTA TTT TTA TTC ACA GTA GGA TTA ACA GGA TTA GTA TTA GCT AAT TCT TCA ATT GAT ATT ATT CTT CAT GAT ACT
TAT TAT GTA GTA GCA CAT TTT CAT TAT GTT CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGG TTT GTA CAC TGA
TAT CCT TTA TTT ACA GGA TTA ACT ATA AAT ACA AAG ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TCC CCT CAA CAC TTC TTA GGA CTT GCA GGA ATA CCT CGT CGA TAC TCA GAT TAT CCA GAT GCT TAT ACA
GCA TGA AAT GTA ATT TCA ACT ATC GGT TCA ACA ATT TCT TTA CTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGA TTA GTA TCC CAA CGA CAA GTA ATA TTT CCT GTT CAA TTA AAT TCA TCA ATT GAA TGA CTT CAA AAT ACA CCT CCT
GC

#S._acrophila_AL34_GA_(Rabun)_USA

TTA CTT CTA GTA AGC AGT ATA GTA GAA AAC GGA GCC GGG ACA GGA TGA ACT GTT TAC CCC CCA TTA TCA TCT AAT ATT
GCC CAT GGA GGA GCC TCT GTT GAT TTA GCT ATC TTT TCT CTT CAT TTA GCT GGA ATT TCT TCA ATT TTA GGG GCC GTA
AAT TTT ATT ACT ACA GTA ATC AAT ATA CGA TCT ACA GGA ATT ACC TTT GAC CGA ATA CCT CTA TTT GTA TGG TCA GTA
GTA ATT ACT GCC CTT CTT CTA TTA CTT TCT TTA CCT GTT CTT GCC GGA GCT ATT ACT ATA TTA CTA ACA GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GGA GGA GAT CCC ATT TTA TAC CAA CAT TTG TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAA GAA
ACA TTC GGG TCA TTA GGC ATA ATT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGG TTC ATT GCT TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACA CGA CCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACT CCA GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCG GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA CTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAC ATT ATT TTA CAT GAT ACA
TAC TAC GTA GTA GCT CAT TTT CAT TAT ACT CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGA CTA ACT TTA AAC ACA AAA ATA TTA AAA AGT CAA TTC ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCC CAA CAT TTC TTA GGA CTT GCA GGT ATA CCT CGC CGA TAC TCT GAT TAT CCA GAT GCT TAT ACA
GCT TGA AAT GTA ATT TCA ACT ATT TGA TGA ATT TCT TTT TTA GGA ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA ACA TCC CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACC CCA CCA
GC

#S._aldrichi_E23_OH_(Hocking)_USA

TTA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTT CAT TTA GCT GGA ATT TCT TCA ATT TTA GGG GCC GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA CTT TTA TTG CTT TCT TTA CCT GTA CTT GCA GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCG GCA GGA GGA GAC CCT ATT TTA TAT CAA CAC TTA TTT TGA TTT TTT GGA CAT

CCT GAA GTT TAT ATT TTA TTT TTA CCG GGA TTT GGA ATA ATT TCT CAC ATT ATT AGA CAA GAA TCA GGT AAA AAG GAA
ACA TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA CTA GCA ATT GCA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA ATT ACT GCA GCT TAT TTT ACA TGA GCA ACA ATA TTT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACC CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT CTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCA CAT TTC CAT TAT GTA CTT TCC ATA GGA GCT GTA TTT GCC ATT ATA GCA GGA TTT GTC CAT TGA
TAC CCT TTA TTT ACC GGA TTA ACA TTA AAT ACA AAA ATA CTG AAA AGC CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCA CAA CAC TTC TTA GGA TTA GCA GGT ATA CCA CGG CGA TAT TCT GAT TAT CCA GAT GCT TAT ACT
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT CTA CTA GGA ATT CTT TTT TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTT ATA TTT CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC

#S._aldrichi_NY_(Onondoga)_AV03

-TA TTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTT CAC TTA GCT GGA ATT TCT TCA ATT TTA GGG GCA GTA
AAC TTT ATC ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT TTA CTT TTA TTG CTT TCT TTA CCT GTA CTT GCA GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT TTT GAT CCG GCA GGA GGA GCA CCT ATT TTA TAT CAA CAC ATA TTT TGA TTT TTT GCA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGA CAA GAA TCA GGT AAA AAG GAA
ACA TTC GGA TCA TTA GGA APTA ATT TAT GCT ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA ACT GAT ACT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACC CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT CTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCA CAT TTC CAT TAT GTA CTT TCC ATA GGA GCT GTA TTT GCC ATT ATA GCA GGA TTT GTC CAT TGA
TAC CCT TTA TTT ACC GGA TTA ACA TTA AAT ACA AAA ATA CTG AAA AGC CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCA CAA CAC TTC TTA GGA TTA GCA GGT ATA CCA CGG CGA TAT TCT GAT TAT CCA GAT GCT TAT ACT
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT CTA CTA GGA ATT CTT TTT TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTT ATA TTT CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTA CAA AAT ACT CCA CCA
GC

#S._africa_AJ12_OR_(Coos)_USA

TTG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTA TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT CTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATC ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATC ACA GCC CTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCC GGA GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCA GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGG CAT
CCT GAA GTT TAT ATT ??? ??? ?A CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA TTA TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAC TCT CCA GCT ACT CTA TGA GCT TTA GGG TTT
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCG TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCA GGT TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGG TTA ACA TTA AAT GCT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC CTA GGA CTT GCA GGT ATG CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACA ATT GGA TCA ACA ATT TCT CTA TTG GGA ATT TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA ATA TTC CCG GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
GC

#S._africa_AJ13_OR_(Coos)_USA

TTG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTA TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT CTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATC ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATC ACA GCC CTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCC GGA GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCA GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGG CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA APTA ATT TAT GCT ATA TTA GCA ATT GCA TTA TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAC TCT CCA GCT ACT CTA TGA GCT TTA GGG TTT
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCG TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCA GGT TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGG TTA ACA TTA AAT GCT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC CTA GGA CTT GCA GGT ATG CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACA ATT GGA TCA ACA ATT TCT CTA TTG GGA ATT TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA ATA TTC CCG GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
GC

#S._africa_AL31_GA_(Rabun County)_USA

TTG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTA TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT CTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATC ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATC ACA GCC CTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCC GGA GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCA GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC ??? ???
??? GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA TTA TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAC TCT CCA GCT ACT CTA TGA GCT TTA GGG TTT
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCG TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCA GGT TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGG TTA ACA TTA AAT GCT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTC CTA GGA CTT GCA GGT ATG CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACA ATT GGA TCA ACA ATT TCT CTA TTG GGA ATT TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA ATA TTC CCG GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
GC

#S._africa_AW42_NM_(Grant)_USA

TTG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTA TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT CTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATC ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATC ACA GCC CTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCC GGA GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCA GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGG CAT
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA APTA ATT TAT GCT ATA TTA GCA ATT GGA TTA TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAC TCT CCA GCT ACT CTA TGA GCT TTA GGG TTT
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCG TCA ATT GAT ATT ATT CTT CAT GAT ACA

TAC TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GAA GTA TTT GCT ATT ATA GCA GGT TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGG TTA ACA CTA AAT GCT AAA ATA GAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTC CTA GCA CTT GCA GGT ATG CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACA ATT GGA TCA ACA ATT TCT CTA TTG GGA ATT TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA ATA TTC CCG GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
GC
#S._africa_BA20_NM_(Grant)_USA
TTG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTA TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT CTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATC ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATC ACA GCC CTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCC GGA GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCA GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGG CAT
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA TTA TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAC TCT CCA GCT ACT CTA TGA GCT TTA GGG TTT
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCG TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCA GGT TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGG TTA ACA TTA AAT GCT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTC CTA GGA CTT GCA GGT ATG CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACA ATT GGA TCA ACA ATT TCT CTA TTG GGA ATT TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA ATA TTC CCG GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
GC
#S._africa_AL31_GA_(Raburn)_USA
-TG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTA TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT CTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATC ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATC ACA GCC CTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCC GGA GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCA GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGG CAT
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA TTA TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAC TCT CCA GCT ACT CTA TGA GCT TTA GGG TTT
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCG TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCA GGT TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGG TTA ACA TTA AAT GCT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTC CTA GGA CTT GCA GGT ATG CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACA ATT GGA TCA ACA ATT TCT CTA TTG GGA ATT TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA ATA TTC CCG GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
GC
#S._africa_AW43_NM_(Grant)_USA
TTG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTA TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT CTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCA ACA GGA ATC ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCT GTA
GTA ATC ACA GCC CTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCC GGA GCT ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCA GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGG CAT
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGA AAA AAG GAA
ACT TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA TTA TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAT ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTC AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAC TCT CCA GCT ACT CTA TGA GCT TTA GGG TTT
GTA TTC TTA TTC ACA GTA GGA GGA TTA ACA GGA GTT GTT TTA GCT AAT TCG TCA ATT GAT ATT ATT CTT CAT GAT ACA
TAC TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCA GTA TTT GCT ATT ATA GCA GGT TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGG TTA ACA TTA AAT GCT AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTC CTA GGA CTT GCA GGT ATG CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACA ATT GGA TCA ACA ATT TCT CTA TTG GGA ATT TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GCA TCT CAA CGA CAA GTA ATA TTC CCG GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACA CCA CCA
--
#S._argyrostoma_AE73_CA_(Shasta)_USA
--A CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT CTA GCT ATT TTT TCT CTT CAC TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCC CTT TTT GTT TGA TCA GTA
GTA ATT ACC GCT TTA CTT CTT CTT CTA TCC CTA CCC GTA CTT GCA GGA GCA ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GGA GAT CCA ATT CTA TAT CAA CAC TTA TTT TGA TTT TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGT TCA CTA GGA ATA ATC TAT GCT ATG TTA GCA ATT GGA CTT TTA GGA TTC ATT GTA TGA GCC CAT CAC ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCC TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTT TAT GGA ACT CAA CTA AAT TAT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAT TAC TCA CTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAT CCC CTA TTC ACT GGA TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ACT ATT TTA TTA GGA TTA AAC
TTA ACT TTC TTT CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCA GAT TAT CCA GAT GCT TAC ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATC TCA TTA TTA GGA ATC TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT ATA TTC CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAC ACC CCA CCA
GC
#S._argyrostoma_AE74_CA_(Shasta)_USA
-TA CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT CTA GCT ATT TTT TCT CTT CAC TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCC CTT TTT GTT TGA TCA GTA
GTA ATT ACC GCT TTA CTT CTT CTT CTA TCC CTA CCC GTA CTT GCA GGA GCA ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GGA GAT CCA ATT CTA TAT CAA CAC TTA TTT TGA TTT TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAA GAA
ACA TTT GGT TCA CTA GGA ATA ATC TAT GCT ATA TTA GCA ATT GGA CTT TTA GGA TTC ATT GTA TGA GCC CAT CAC ATA
TTT ACA GTA GGA ATA GAC GTA GAT ACA CGA GCC TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTT TAT GGA ACT CAA CTA AAT TAT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTC
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAT TAC TCA CTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAT CCC CTA TTC ACT GGA TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ACT ATT TTA TTA GGA TTA AAC
TTA ACT TTC TTT CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCA GAT TAT CCA GAT GCT TAC ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATC TCA TTA TTA GGA ATC TTA TTT TTC TTT TAT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT ATG TTC CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAC ACC CCA CCA

GTA ATT ACC GCT TTA CTT CTC CTT CTA TCC CTA CCC GTA CTT GCA GGA GCA ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT TTT CCA GCA GGA GGA GAT CCA ATT CTA TAT CAA CAC TTA TTT TGA TTT TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAA GAA
ACA TTT GGT TCA CTA GGA ATA ATC TAT GCT ATA CTA GCA ATT GGA CTT TTA GGA TTC ATT GTA TGA GCC CAT CAC ATA
TTT ACA GTA GCA ATA GAC GAT ACA GAT ACA GCC TAT TTT ACT TCA GCA ACA ATA ATT GCT GTT CCA ACA GCA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTT TAT GGA ACT CAA CTA AAT TAT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAT TAC GAT CTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAT CCC CTA TTC ACT GGA TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAC
GTA ACT TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCA GAT TAT CCA GAT GCT TAC ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATC TCA TTA TTA GGA ATC TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT ATG TTC CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAC ACC CCA CCA
GC

#S._argyrostoma_BE18_KY_(Kenton)_USA

--A CTA CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT CTA GCT ATT TTT TCT CTT CAC TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCT CTT TTG GTT TGA TCA GTA
GTA ATT ACC GCT TTA CTT CTC CTT CTA TCC CTA CCC GTA CTT GCA GGA GCA ATT ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCA GCA GGA GGA GGA GAT CCA ATT CTA TAT CAA CAC TTA TTT TGA TTT TTT GGT CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGT TCA CTA GGA ATA ATC TAT GCT ATA CTA GCA ATT GGA CTT TTA GGA TTC ATT GTA TGA GCC CAT CAC ATA
TTT ACA GTA GGA ATA GAC GAT ACA GAT ACA GCC TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTT TAT GGA ACT CAA CTA AAT TAT TCC CCA GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACT GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAT TAC GTA CTA TCA ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAT CCC CTA TTC ACT GGA TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAC
TTA ACT TTC TTC CCT CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCA GAT TAT CCA GAT GCT TAC ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATC TCA TTA TTA GGA ATC TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GTA TCT CAA CGA CAA GTT ATG TTC CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAC ACC CCA CCA
GC

#S._bullata_AA02_FL_(Duval)_USA

TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAC GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CGA GCC TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTC CCA ACA GGG ATT
AAA ATT TTT AGC TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GGA GGT CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTT CTA GGA CTT GCA GGA ATA CCT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC

#S._bullata_AA45_FL_(Walton)_USA

TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAC GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CGA GCC TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTC CCA ACA GGG ATT
AAA ATT TTT AGC TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGT CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTT CTA GGA CTT GCA GGA ATA CCT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC

#S._bullata_AB07_FL_(Washington)_USA

TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAC GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CGA GCC TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTC CCA ACA GGG ATT
AAA ATT TTT AGC TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGT CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT TAT GTA GTA GCT CAT TTC MAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCT CAA CAT TTT CTA GGA CTT GCA GGA ATA CCT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC

#S._bullata_AD70_SC_(Newberry)_USA

TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAC GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CGA GCC TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTC CCA ACA GGG ATT

AAA ATT TTT AGC TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGT CTA ACT GGA TTA GTC CTA TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC CTT CCA CAT TTT CTA GGA CTT GCA GGA ATT CCA GGA TTA CTT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC
#S_bullata_AE39_SC_(Newberry)_USA
TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCC CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CTA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC CTT CCA CAT TTT CTA GGA CTT GCA GGA ATA CTT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC
#S_bullata_AF37_CA_(Yolo)_USA
TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
TTT ACA GTA GGA ATA GAT GTT GAC ACT CTA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
GTA TTT TTA TTT ACT GTA GGA GGT CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC CTT CCA CAT TTT CTA GGA CTT GCA GGA ATA CTT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC
#S_bullata_AD09_FL_(Highland)_USA
TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CTA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGT CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT -AT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC CTT CCA CAT TTT CTA GGA CTT GCA GGA ATA CTT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC
#S_bullata_BA69_MN_(Winona)_USA
TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CTA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GGA GGT CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT -AT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC CTT CCA CAT TTT CTA GGA CTT GCA GGA ATA CTT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC
#S_bullata_BC10_OH_(Hamilton)_USA
TTA CTT CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCC TTA TCT TCT AAC ATC
GCC CAC GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC TTA CAT TTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTG ACT GAT CGA AAT
ATT AAT ACC TCA TTC TTT GAC CCT GCA GGA GGA GGA GAT CCA ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAC ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAT CAC ATA
TTC ACA GTA GGA ATA GAT GTT GAC ACT CTA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTA CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGT CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT TAT GTA GTA GCT CAT TTC CAC TAT GTA CTA TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCC CTA TTT ACT GGA CTA ACT TTA AAC TCG AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC CTT CCA CAT TTT CTA GGA CTT GCA GGA ATA CTT CGA CGT TAC TCT GAC TAT CCT GAT GCT TAC ACA
TCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTT CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTG CAA AAT ACT CCA CCA
GC

#S._crassipalpis_AQ25_NY_(Rockland)_USA

-TG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACG GGG TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT CTA CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACC TTT GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCC CTA CTT TTA CTT TTA TCT TTA CCG GTA CTT GCA GGA GCT ATT ACT ATA TTA TTA ACT GAC GCA AAT
ATT AAT ACC TCT TTT TTC GAC CCA GCA GGA GGA GAT CCT ATT TTA TAC CAA CAC CTA TTT TGA TTT TTC GGT CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTC GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA CTT CTA GGA TTC ATT GTA TGA GCT CAC CAT GTA
TTC ACA GTA GGA ATA GAC GTA GAC ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAC GGA ACT CAA TTA AAT TAC TCT CCT GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TAT ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAC ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCT GTA TTT GCC ATT ATA GCA GGA TTT GTT CAC TGG
TAC CCT TTA TTT ACC GGA TTA ACA TTA AAT GCA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCG CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
GCT TGA AAT GTA ATT TCA ACA ATC GGA TCA ACA ATT TCA TTA TTA GGA ATC TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTT ATA TTC CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAC A-- --- ---

#S._crassipalpis_AR61_WI_(Waukesha)_USA

-TG CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACG GGG TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT CTA CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTA ATT AAT ATA CGA TCT ACA GGA ATT ACC TTT GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCC CTA CTT TTA CTT TTA TCT TTA CCG GTA CTT GCA GGA GCT ATT ACT ATA TTA TTA ACT GAC GCA AAT
ATT AAT ACC TCT TTT TTC GAC CCA GGA GGA GAT CCT ATT TTA TAC CAA CAC CTA TTT TGA TTT TTC GGT CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTC GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTC GGA TCA TTA GGA ATA ATT TAT GCT ATA TTA GCA ATT GGA CTT CTA GGA TTC ATT GTA TGA GCT CAC CAT ATA
TTC ACA GTA GGA ATA GAC GTA GAC ACA CGA GCT TAT TTT ACT TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAC GGA ACT CAA TTA AAT TAC TCT CCT GCT ACT TTA TGA GCC TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TAT ACT GGA GTT GTT TTA GCT AAT TCA TCA ATT GAC ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAC TTC CAC TAT GTA CTT TCA ATA GGA GCT GTA TTT GCC ATT ATA GCA GGA TTT GTT CAC TGG
TAC CCT TTA TTT ACC GGA TTA ACA TTA AAT GCA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTC TTC CCG CAA CAT TTC TTA GGA CTT GCA GGA ATA CCT CGA CGA TAT TCA GAT TAC CCA GAT GCT TAT ACA
GCT TGA AAT GTA ATT TCA ACA ATC GGA TCA ACA ATT TCA TTA TTA GGA ATC TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GTA TCA CAA CGA CAA GTT ATA TTC CCA GTT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAC ACT CCG CCA

#S._georgiana_AB10_FL_(Washington)_USA

TTA CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAT CCT CCT TTA TCA TCT AAT ATT
GCT CAT GGG GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT CTT CAC TTA GCT GGA ATT TCT TCT ATT TTA GGG GCA GTA
AAC TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCT CTA TTT GTT TGA TCC GTA
GTA ATT ACA GCT CTA CTT TTA CTT CTT TCT TTA CCA GTA CTT GCT GGA GCA ATT ACT ATA TTA CTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCA GGA GGA GAC CCA ATT TTA TAC CAA CAC TTA TTC TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATC AGA CAA GAA TCA GGT AAG AAG GAA
ACA TTT GGA TCC TTA GGA ATA ATT TAT GCA ATA CTA GCA ATT GGA TTA TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTT GGT ATA GAC GTA GAT ACT CTA GAT GCT TAT TTT ACA TCA GCA ACT ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAC GCT ACT CAA TAT TAT TCC CCA GCT ACA CTA TGA GCT CTA GGA TTT
GTA TTT TTA TTT ACA GTT GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT TAT GTA GTA GCT CAC TTT CAT TAT GTT CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACT GGA TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAG TTT ACT ATT ATA TTT ATA GGA GTA AAC
TTA ACA TTC TTC CCT CAA CAT TTT TTA GGT CTT GCA GGA ATA CCT CGA CGT TAT TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATT TCA ACA ATT GGG TCA ACT ATT TCT CTA CTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA GTG TCA CAA CGA CAA GTA ATA TTT CCA GTA CAA TTA AAT TCA TCT ATT GAA TGA CTC CAA AAT ACT CCT CCA

#S._hinei_AV61_NY_(Niagara)_USA

TTA CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCA CCC CTA TCA TCT AAC ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT TTA CAT CTA GCC GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTC ATT ACA ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACC TTT GAC CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTG ATT ACA GCT TTA CTT TTA TTA CTT TCT TTA COT GTT CTT GCT GGA GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCG GCG GGA GGA GAC CCT ATT CTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCA GAA GTT TAT ATC TTA ATT TTA CCA GGG TTT GGA ATA ATC TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGA ATA ATT TAT GCA ATA CTA GCA ATT GGA CTT TTA GGT TTC ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGG ATA GAT GTA GAC ACT CGA GCT TAT TTC ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGC TGA CTT GCT ACT CTT TAC GGA ACT CAA TTA AAT TAT TCC CCC GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAT TAT GTA GTA GCT CAT TTT CAT TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGG TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGA TTG ACT TTA GAC ACA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACC TTT TTT CCT CAA CAT TTC TTG GGA CTT GCA GGT ATG CCA CGA CGA TAC TCT GAT TAC CCA GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCA ACA ATT GGT TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TTT ATT ATT TGA GAA
AGC TTA ACA TCC CAT CGA CAA GTT ATA TTC CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTC CAA AAT ACC CCA CCT

#S._houghi_AO50_OH_(Hamilton)_USA

TTA CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT CTA TCA TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCA ATT TTT TCT CTA CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACA ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACC TTT GAT CGA ATA CCT TTA TTT GTT TGA TCA GTA
GTA ATT ACA GCT CTA CTT TTA CTA CTT TCT TTA COT GTT CTT GCT GGA GCA ATT ACT ATA TTA TTA ACT GAC GCA AAT
ATT AAT ACT TCA TTT TTT GAT CCG GCA GGA GGA GGA GAT CCT ATT TTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA --- --A CCA GGA TTT GGA ATA ATC TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGA ATA ATT TAT GCA ATA TTA GCA ATT TGA CTT TTA GGT TTC ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACT CGA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTC CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTT TAT GGA ACT CAA TTA AAT TAT TCT CCA GCT ACT CTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
TAC TAT GTA GTA GCT CAT TTT CAT TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAC CCT TTA TTT ACT GGA TTA ACT TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ATT ATT ATA TTT ATA GGA GTA AAT
TTA ACT TTT TTT CCT CAA CAT TTC TTA GGA CTT GCA GGT ATG CCA CGA CGA TAT TCT GAC TAT CCA GAT GCT TAC ACA
ACT TGA AAT GTA ATT TCA ACA ATT GGT TCA ACA ATT TCT TTA TTA GGA ATT CTA TTT TTC TTC TTT ATT ATT TGA GAA
AGT TTA TCG TCT CAA CGA CAA GTT ATA TTT CCA ATT CAA TTA AAC TCA TCT ATT GAA TGA CTT CAA AAT ACT CCA CCT

#S._idonea_AT30_MN_(Winona)_USA

TTA CTT CTA GTA AGC AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCC CCA TTA TCA TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCT ATC TTC TCT CTT CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGA ATT ACC TTT GAT CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCT CTT TTA CTA CTT TCT TTA COT GTT CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT

ATT AAT ACT TCA TTT TTT GAT CCA GCT GGG GGA GGA GAC CCT ATT TTA TAC CAA CAT CTA TTT TGA TTT TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAC ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATT TAT CGA GCA ATA TTA GCA ATT GGA CTT TTA GGA TTC ATT GTA TGA GCT CAC CAT ATA
TTC ACA GTA GGA ATA GAT GAT ACA CGA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAT GGA ACT CAA TAT TCT CCG GCC ACT TTA TGA GCT TTA GGT TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAC ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTT CAT TAT GTT CTA TCT ATA GGA GCC GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAT CCC CTA TTT ACT GGA CTA ACT TTA AAT GTA AAA ATA CTA AAA AGT CAA TTC ATT ATT ATA TTT ACA GGA CAT AAT
TTA ACC TTC TTT CCT CAA CAT TTC TTA GGA CTT GAA GGA ATA CCT CGT CGA TAT TCT GAT TAT CCA GAC GCT TAC ACA
GCT TGA AAT GTA ATA TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT TTA TTT TTC TTC TAC ATC ATT TGA GAA
AGT TTA ATA TCC CAA CGA CAA GTA TTA TTT CCA ATT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACC CCC CCA
GC

#S._johnsoni_AQ40_NY_(Richmond)_USA
TTA TTA CTA GTG AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCA CAT GGG GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTT CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACT TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCT TTG CCF GTG CTT GCC GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAC ACT TCT TTC TTT GAC CCA GCA GGA GGA GGA GAT CCT ATT TTA TAC CAA CAT CTA T-T TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCT CAT ATT ATT AGA CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCC TTA GGA ATA ATT TAT GCA ATA TTA GCA ATT GGA TTA TTA GGA TTT ATT GTA TGA GCA CAT CAT ATA
TTT ACA GTT GGA ATA GAC GTA GAT ACT CGA GCA TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCT ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACC CTT TAC GGA ACT CAA TTA AAT TAT TCC CCA GCT ACA TTA TGA GCC TTA GGA TTT
GTA TTT TTA TTC ACA GTA GAT GGT GGA TTA ACT GGA ATT TTA GCA AAT TCA TCT CTT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAC TTT CAT TAT GTT CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAC TGA
TAC CCT TTA TTT ACT GGA TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAA TTC ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTT TTT CCT CAA CAT TTT TTA GGT CTT GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCC TAT ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGT ATT TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GTG TCA CAA CGA CAA GTA ATA TTC CCT GTA CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCA CCT
GC

#S._mimoris_E39_OH_(Adams)_USA
TTA CTT CTA GTA AGC AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCC CCT TTA TCA TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT CTT CAT TTA GCT GGA ATT TCT TCA ATT TTA GGT GCA GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGA ATT ACT TTT GAT CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACT GCT CTT CTA TTA TTA CTT TCT TTA CCT GTT CTT GCT GGA GCA ATT ACT ATA TTA TTA ACA GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAT CCT GCT GGA GGA GGA GAT CCT ATT TTA TAT GAA CAT TTA TTT TGA TTT TTT GGA CAT
CCT GAA GTT TAT ATT TT? ??? ??? ??? ??? ?TA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCA TTA GGT ATA ATC TAT GCA ATA TTA GCA ATT GGA CTT TTA GGA TTC ATT GTA TGA GCT CAT CAT ATA
TTT ACA GTA GGA ATA GAT GAT ACA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACT GGA ATT
AAA ATT TTT AGT TGA CTT GCC ACT CTT TAT GGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAC TAT GTA GTA GCT CAT TTT CAT TAC GTC CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTT CAT TGA
TAT CCT TTA TTT ACT GGA TTA ACT TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ACA GGA GTA AAT
TTA ACT TTC TTT CCT CAA CAT TTC TTA GGT CTT GCA GGT ATA CCT CGC CGA TAC TCT GAT TAT CCA GAT GCT TAT ACA
GCT TGA AAT GTA TGC TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGG ATC TTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GCA TCC CAA CGA CAA GTA TTA TTT CCT ATT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCG CCA
GC

#S._nearctica_AS38_WI_(Marathon)_USA
CTA CTT CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCC GGA ATC TCA TCA ATT TTA GGA GCC GTA
AAT TTC ATT ACT ACA GTG ATT AAT ATA CGA TCA ACA GGT ATT CAA TTT GAC CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCC TTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAC CCT GCC GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GAT ACA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TGA ACT CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTT CAC TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAT CCC CTA TTT ACT GGA TTA ACA TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGT GTA AAT
TTA ACT TTT TTT CCA CAA CAT TTC TTA GGA CTT GCC GGT ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACT ATT GGG TCT ACA ATT TCT CTT TTA GGA ATT TTA TTT TTT TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTC CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCA CCT
GC

#S._nearctica_BB67_MN_(Kittson)_USA
CTA CTT CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCC GGA ATC TCA TCA ATT TTA GGA GCC GTA
AAT TTC ATT ACT ACA GTG ATT AAT ATA CGA TCA ACA GGT ATT CAA TTT GAC CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCC TTA CTT TTA TTA CTT TCT TTA COT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAC CCT GCC GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GAT ACA GCA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAT GGA ACC CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT? GAT ATT ATT TTA CAT GAT ACA
TAT ?AT GTA GTA GCT CAT TTT CAC TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAT CCC CTA TTT ACT GGA TTA ACA TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGT GTA AAT
TTA ACT TTT TTT CCA CAA CAT TTC TTA GGA CTT GCC GGT ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACT ATT GGG TCT ACA ATT TCT CTT TTA GGA ATT TTA TTT TTT TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTC CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCA CCT
G-

#S._nearctica_AQ52_NY_(Richmond)_USA
CTA CTT CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCC GGA ATC TCA TCA ATT TTA GGA GCC GTA
AAT TTC ATT ACT ACA GTG ATT AAT ATA CGA TCA ACA GGT ATT CAA TTT GAC CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCC TTA CTT TTA TTA CTT TCT TTA COT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTC TTT GAT CCT GCC GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GAT ACA CGA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAT GGA ACC CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT


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AGT TTA GCA TCA CAA CGA CAA GTT ATA TTC CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCA CCT
GC
#S._nearctica_AU24_MN_(Koochiching)_USA
CTA CTT CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCC GGA ATC TCA TCA ATT TTA GGA GCC GTA
AAT TTC ATT ACT ACA GTG ATT AAT ATA CGA TCA ACA GGT ATT ACA TTT GAC CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCC TTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCC GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACA CGA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAT GGA ACC CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTT CAC TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAT CCC CTA TTT ACT GGA TTA ACA TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGT GTA AAT
TTA ACT TTT TTT CCA CAA CAT TTC TTA GGA CTT GCC GGT ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACT ATT GGG TCT ACA ATT TCT CTT TTA GGA ATT TTA TTT TTT TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTC CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCA CCT
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#S._nearctica_AU70_NY_(Onandaga)_USA
CTA CTT CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCC GGA ATC TCA TCA ATT TTA GGA GCC GTA
AAT TTC ATT ACT ACA GTG ATT AAT ATA CGA TCA ACA GGT ATT ACA TTT GAC CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCC TTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCC GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCT TTA GGA ATA ATT TAT GCA ATA TTA GCA ATT GGA CTT TTA GGA TTT ATT GTA TGA GCT CAC CAT ATA
TTT ACA GTA GGA ATA GAT GTA GAT ACA CGA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAT GGA ACC CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
TAT TAT GTA GTA GCT CAT TTT CAC TAT GTA CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCT GGA TTT GTT CAT TGA
TAT CCC CTA TTT ACT GGA TTA ACA TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGT GTA AAT
TTA ACT TTT TTT CCA CAA CAT TTC TTA GGA CTT GCC GGT ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACT ATT GGG TCT ACA ATT TCT CTT TTA GGA ATT TTA TTT TTT TTC TTT ATT ATT TGA GAA
AGT TTA GCA TCA CAA CGA CAA GTT ATA TTC CCA ATT CAA CTA AAT TCA TCT ATT GAA TGA CTT CAA A-- --- --- ---
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#S._nearctica_AH21_OR_(Douglas)_USA
CTA CTT CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCC GGA ATC TCA TCA ATT TTA GGA GCC GTA
AAT TTC ATT ACT ACA GTG ATT AAT ATA CGA TCA ACA GGT ATT ACA TTT GAC CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCC TTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
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CCT GAA GTT TAT ATT TTA ATT TTA CCT GGA TTT GGA ATA ATT TCT CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
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TTT ACA GTA GGA ATA GAT GTA GAT ACA CGA GCT TAT TTT ACA TCA GCA ACA ATA ATT ATT GCT GTT CCA ACA GGA ATT
AAA ATT TTT AGT TGA CTT GCT ACT CTA TAT GGA ACC CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
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TAT CCC CTA TTT ACT GGA TTA ACA TTA AAT ACA AAA ATA CTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGT GTA AAT
TTA ACT TTT TTT CCA CAA CAT TTC TTA GGA CTT GCC GGT ATA CCT CGA CGA TAT TCT GAT TAT CCA GAT GCT TAT ACA
ACT TGA AAT GTA ATC TCA ACT ATT GGG TCT ACA ATT TCT CTT TTA GGA ATT TTA TTT TTT TTC TTT ATT ATT TGA GAA
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#S._nearctica_AS62_WI_(Barron)_USA
CTA CTT CTA GTA AGC AGC ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCT TTA CAT TTA GCC GGA ATC TCA TCA ATT TTA GGA GCC GTA
AAT TTC ATT ACT ACA GTG ATT AAT ATA CGA TCA ACA GGT ATT ACA TTT GAC CGA ATA CCT TTA TTT GTA TGA TCA GTA
GTA ATT ACA GCC TTA CTT TTA TTA CTT TCT TTA CCT GTA CTT GCT GGA GCA ATT ACT ATA CTA TTA ACT GAT CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCC GGA GGA GGA GAT CCT ATT CTA TAT CAA CAT TTA TTT TGA TTC TTT GGA CAC
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AAA ATT TTT AGT TGA CTT GCT ACT CTA TAT GGA ACC CAA TTA AAT TAT TCC CCA GCT ACT TTA TGA GCT TTA GGA TTT
GTA TTC TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAT ACA
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ACT TGA AAT GTA ATC TCA ACT ATT GGG TCT ACA ATT TCT CTT TTA GGA ATT TTA TTT TTT TTC TTT ATT ATT TGA GAA
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#S._polistensis_E43_OH_(Adams)_USA
CTA CTT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAC ATC
GCT CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTT TCC CTA CAC TTA GCT GGA ATT TCT TCA ATT TTA GGA GCT GTA
AAT TTT ATT ACT ACA GTT ATT AAT ATA CGA TCT ACA GGT ATT ACA TTT GAT CGA ATA CCT TTA TTT GTT TGA TCT GTA
GTA ATT ACA GCT TTA CTT TTA CTT CTT TCC CTG CCT GTA CTT GCT GGA GCA ATT ACT ATA TTA TTA ACT GAT CGA AAT
ATT AAC ACT TCA TTT TTT GAC CCT GCA GGA GGA GGT CCA ATT TTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAT
CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATA ATT TCC CAT ATT ATT AGT CAA GAA TCA GGT AAA AAG GAA
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AAA ATT TTT AGC TGA CTT GCC ACT CTT TAT GGA ACC CAA TTA AAT TAT TCC CCG GCC ACA CTA TGA GCT CTT GGA TTT
GTA TTT TTA TTT ACT GTA GGA GGA CTA ACT GGA GTA GTC TTA GCT AAT TCA TCT ATT GAC ATT ATT TTA CAT GAT ACA
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#S._saraceniodes_BC36_NY_(Suffolk)_USA
TTA CTA CTA GTA AGC AGT ATA GTA GAA AAT GGG GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCT TCT AAT ATT
GCT CAT GGG GGA GCT TCT GTT GAC TTA GCT ATT TTT TCT CTA CAT TTA GCT GGA ATC TCT TCA ATT TTA GGA GCA GTA

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TTA ACT TTC GTC CCA CAA CAT TTC CTG GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCT GAC GCT TAC ACA
GCT TGA AAT GTC ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
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#S._subvicina_AP06_NY_(Chatauqua)_USA
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GCC CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
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GC
#S._subvicina_AP72_NY_(Schuyler)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATC AAT ATA CGA TCT ACA GGT ATT ACT TTT GAC CGA ATA CCT TTA TTC GTG TGA TCA GTA
GTA ATT ACA GCT TTA CTT TTA TTA CTT TCT CTG CCT GTA CTT GCC GGA GCA ATC ACT ATA TTA TTA ACT GAC CGA AAT
ATT AAT ACT TCA TTT TTT GAC CCT GCC GGA GGT GGA GAC CCA ATC TTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAT
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TTA ACT TTC TTC CCA CAA CAT TTC CTG GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCT GAC GCT TAC ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
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GC
#S._subvicina_AS68_WI_(Waukesha)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATC AAT ATA CGA TCT ACA GGT ATT ACT TTT GAC CGA ATA CCT TTA TTC GTG TGA TCA GTA
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CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGC CAA GAA TCA GGT AAA AAG GAA
ACA TTC GGA TCA CTA GGT ATA ATT TAT GCA ATG CTA GCA ATT GGA CTT TTA GGG TTT ATT GTA TGA GCT CAT CAT ATA
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AGT TTA GCA TCC CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCC CCA
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#S._subvicina_AV23_NY_(Onondaga)_USA
TTA CTA CTA GTA AGT AGT ATA GTA GAA AAC GGA GCT GGA ACA GGA TGA ACT GTT TAC CCT CCT TTA TCA TCT AAT ATT
GCC CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
AAT TTT ATT ACT ACA GTT ATC AAT ATA CGA TCT ACA GGT ATT ACT TTT GAC CGA ATA CCT TTA TTC GTG TGA TCA GTA
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ATT AAT ACT TCA TTT TTT GAC CCT GCC GGA GGT GGA GAC CCA ATC TTA TAC CAA CAT TTA TTT TGA TTC TTT GGA CAT
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ACA TTC GGA TCA CTA GGT ATA ATT TAT GCA ATG CTA GCA ATT GGA CTT TTA GGG TTT ATT GTA TGA GCT CAT CAT ATA
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GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
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GC
#S._subvicina_AW02_NY_(Saratoga)_USA
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GCC CAT GGA GGA GCT TCT GTT GAT TTA GCT ATT TTC TCT CTC CAT TTA GCT GGA ATT TCT TCA ATT TTA GGA GCA GTA
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CCT GAA GTT TAT ATT TTA ATT TTA CCA GGA TTT GGA ATG ATT TCT CAT ATT ATT AGC CAA GAA TCA GGT AAA AAG GAA
ACA TTC GGA TCA CTA GGT ATA ATT TAT GCA ATG CTA GCA ATT GGA CTT TTA GGG TTT ATT GTA TGA GCT CAT CAT ATA
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AAA ATT TTT AGT TGA CTT GCT ACT CTT TAC GGA ACT CAA CTA AAT TAT TCT CCA GCT ACT TTA TGA GCT TTA GGA TTC
GTA TTT TTA TTT ACA GTA GGA GGA TTA ACT GGA GTT GTT TTA GCT AAT TCA TCT ATT GAT ATT ATT TTA CAT GAC ACA
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TTA ACT TTC TTC CCA CAA CAT TTC CTG GGA CTT GCA GGA ATA CCT CGA CGA TAT TCT GAT TAT CCT GAC GCT TAC ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA TTA GGA ATT CTA TTT TTC TTT TTT ATT ATT TGA GAA
AGT TTA GCA TCC CAA CGA CAA GTA TTA TTC CCA GTT CAA TTA AAT TCA TCT ATT GAA TGA CTT CAA AAT ACT CCC CCA
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#S._triplasia_A23_OH_(Hocking)_USA

TAT TAT GTA GTA GCT CAT TTT CAT TAT GTC CTT TCT ATA GGA GCT GTA TTT GCT ATT ATA GCA GGA TTT GTA CAC TGA
TAC CCT TTA TTT ACA GGA TTA ACA TTA AAT ACA AAA ATA TTA AAA AGT CAA TTT ACT ATT ATA TTT ATA GGA GTA AAT
TTA ACA TTC TTT CCT CAA CAT TTT TTA GGA CTT GCA GGA ATA CCT CGA CGT TAT TCT GAT TAC CCA GAT GCT TAC ACA
GCT TGA AAT GTA ATT TCA ACA ATT GGA TCA ACA ATT TCT TTA CTA GGA ATT TTA TTT TTC TTT TTC ATT ATC TGA GAA
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#S. utilis AE22_SC (Newberry)_USA
--- -TT CTA GTA AGT AGT ATA GTA GAA AAT GGA GCT GGA ACA GGT TGA ACT GTT TAT CCT CCT TTA TCA TCT AAT ATT
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GTA ATT ACA GCT CTA CTT TTA CTC CTT TCT TTA CCG GTA CTT GCT GGA GCT ATT ACT ATA TTA CTA ACA GAT CGA AAT
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CCC GAA GTT TAT ATT TTA ATT TTA CCG GGA TTT GGA ATA ATT TCC CAT ATT ATT AGA CAA GAA TCA GGT AAA AAG GAA
ACA TTT GGA TCC TTA GGA ATA ATT TAT GCA ATA CTA GCA ATT GGA TTA TTA GGA TTT ATT GTA TGA GCC CAC CAT ATA
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#Sb. flavipalpis AV21_NY (Onondaga)_USA
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GC

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Table 11. Average Tamura-Nei-corrected distances within and between the groups indicated.
Average Intra-specific Distances

<i>L. cluvia</i>	0.9%
<i>L. coeruleiviridis</i>	0.4%
<i>L. cuprina</i>	0.1%
<i>L. illustris</i>	0.5%
<i>L. mexicana</i> (NM and TX)	0.9%
<i>L. mexicana</i> (CA)	0.2%
<i>L. sericata</i>	0.1%
<i>L. silvarum</i>	0.3%

Average Inter-group Distances

<i>L. mexicana</i> : CA vs. NM+TX	5.0%
<i>L. cuprina</i> vs. <i>L. sericata</i>	1.1%

3. **Figures:** The figure(s) is clearly designed and accurately describes a relevant aspect of the results. Figures should include a title as well as a legend that concisely conveys the content displayed in the figure.

Fig. 1. Bootstrap support for ME analysis for *L. cuprina* and *L. sericata*, based on 600bp windows of the COI-COII region.

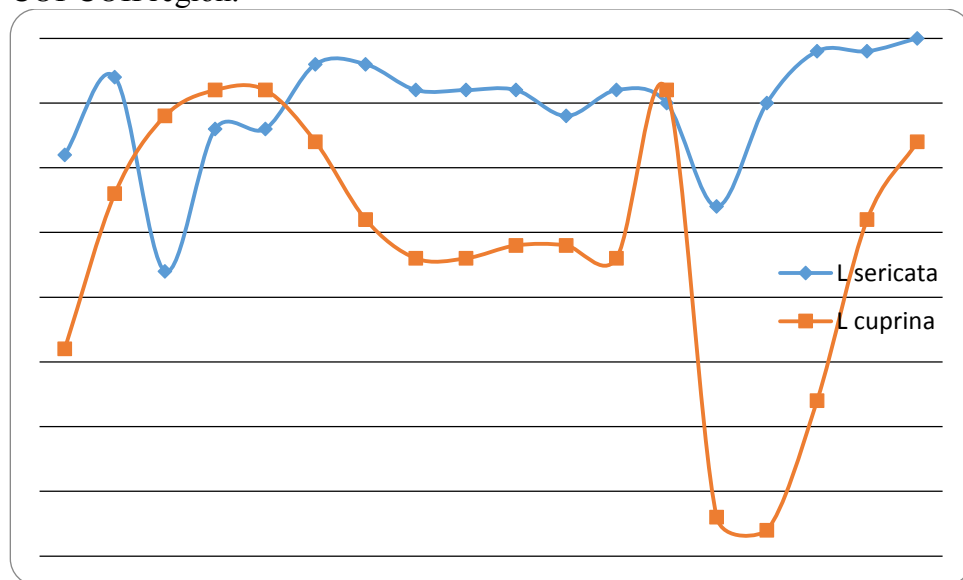
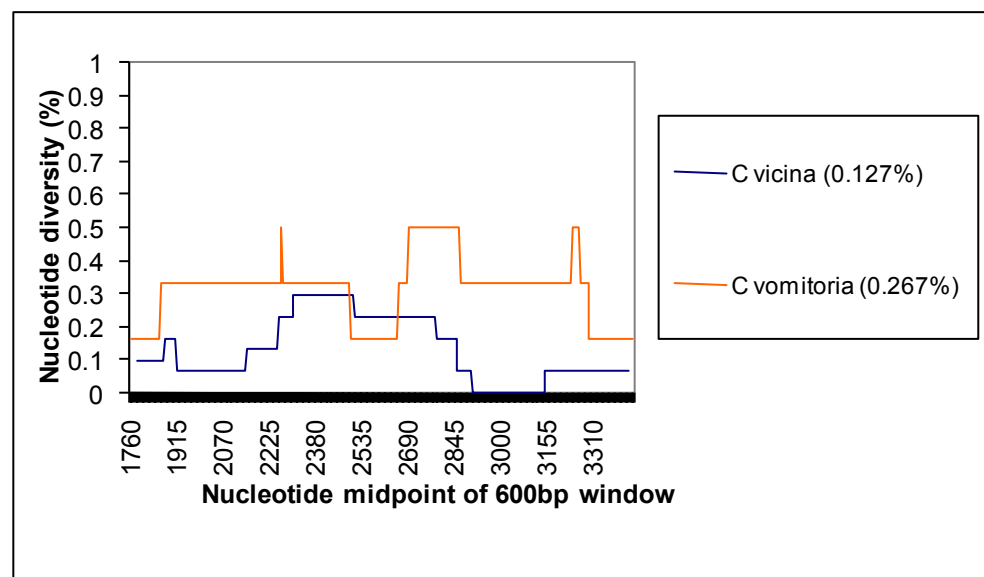


Fig 2. Sliding window profiles (600bp window) of COI-COII intraspecific nucleotide diversity.



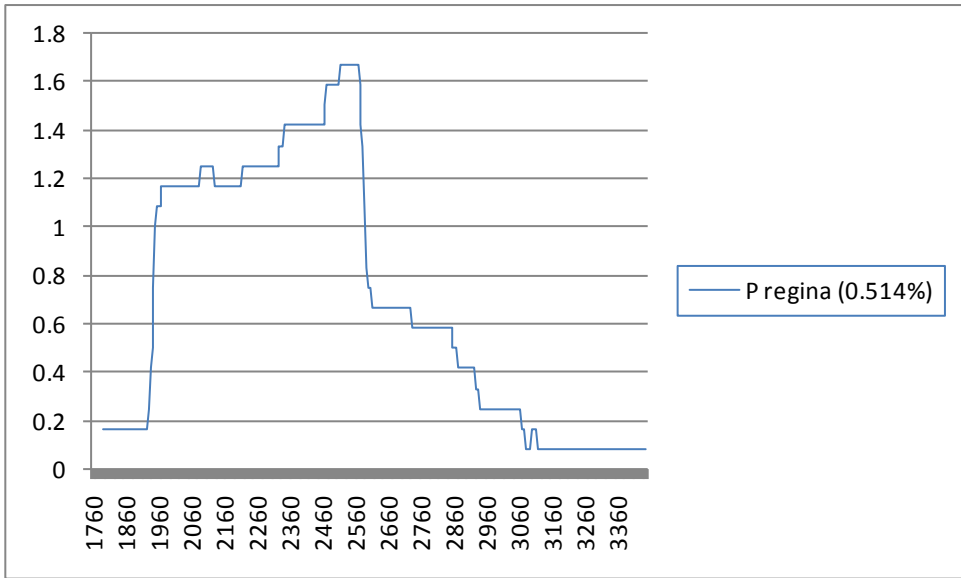
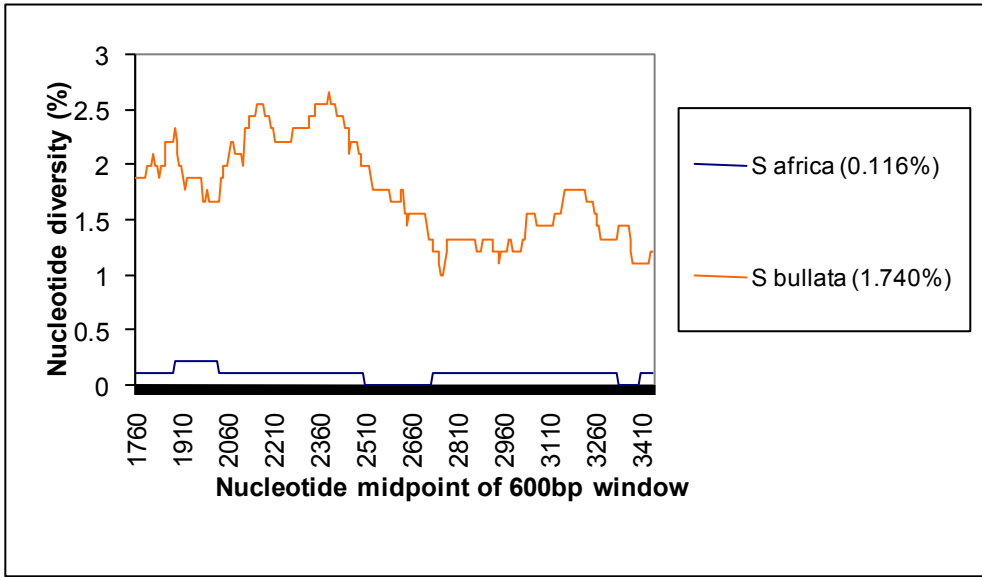
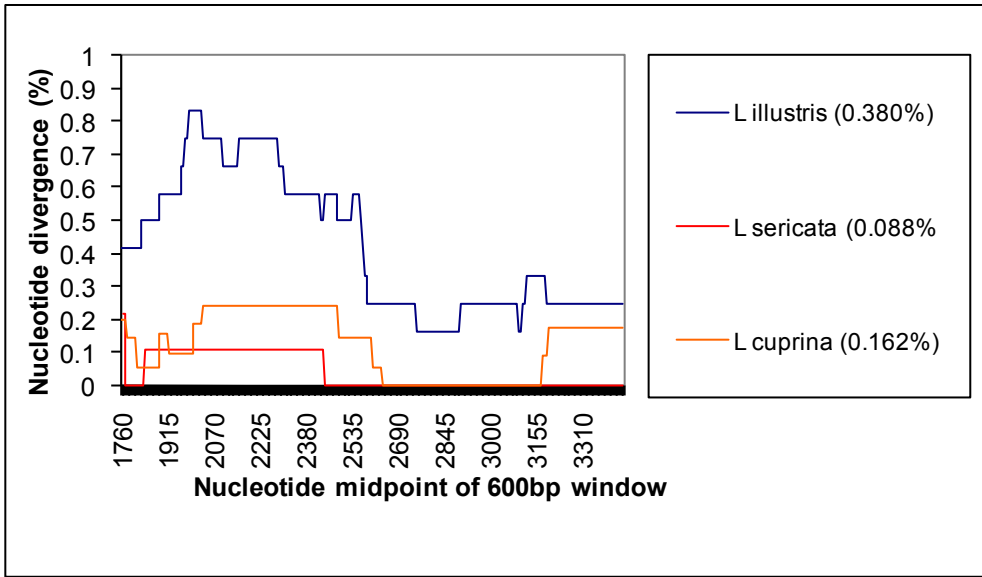


Fig. 3. Sliding window (600bp) analysis for five species pairs, based on analysis of the COI-COII region.

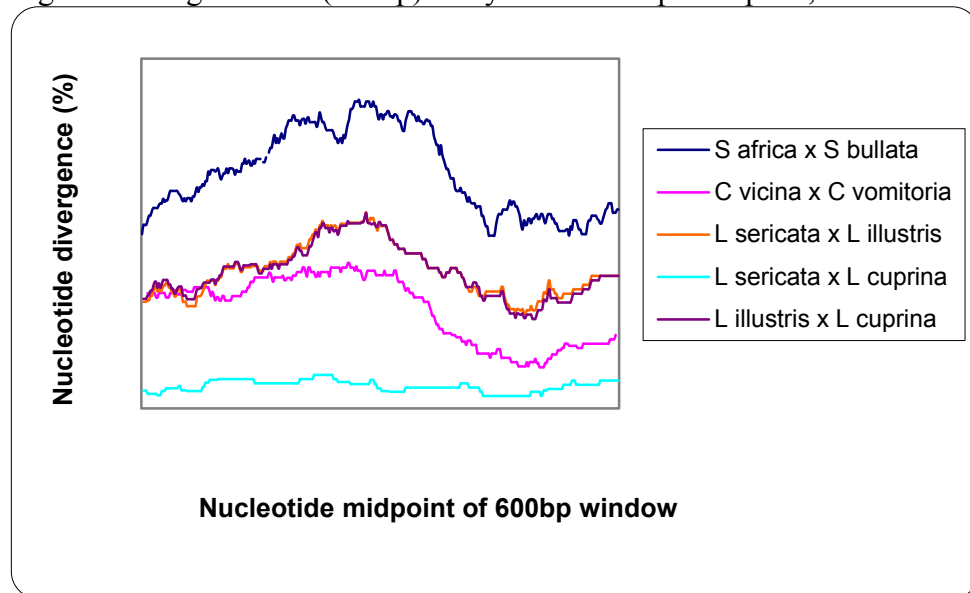


Fig. 4. Sliding window profile of mean relative divergence and $\pm 1SD$ for all 5 sister pairs for the COI-II region.

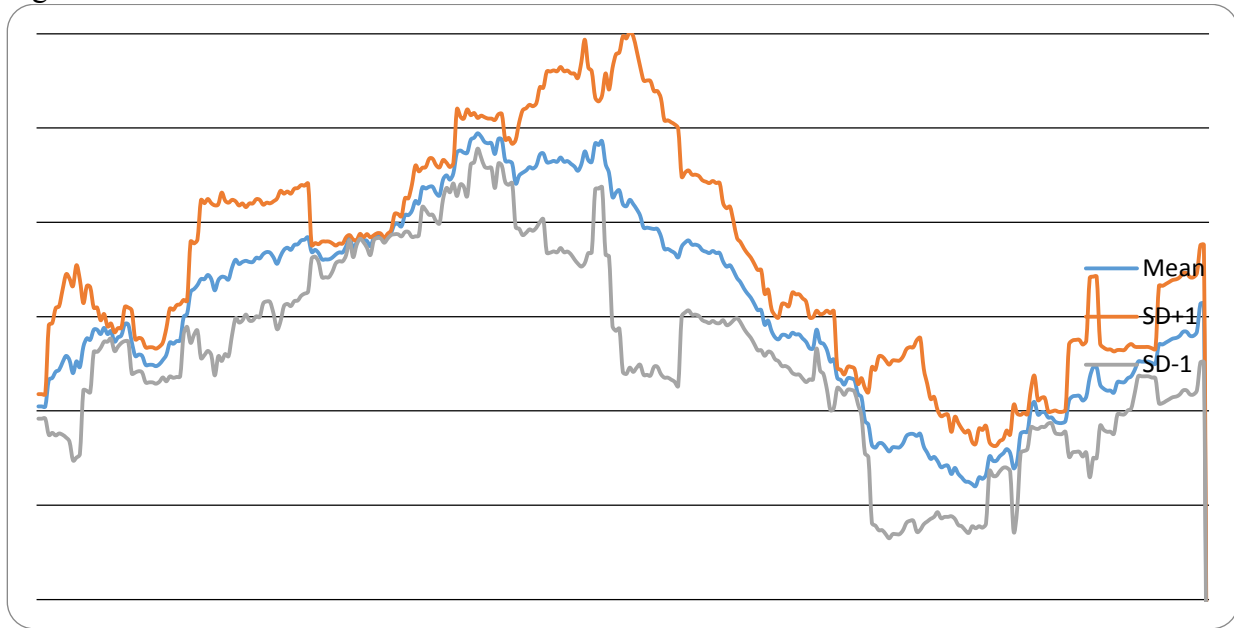


Fig. 5. ME tree based on 3 concatenated DNA sequences of the COI, COII and ND4 genes.

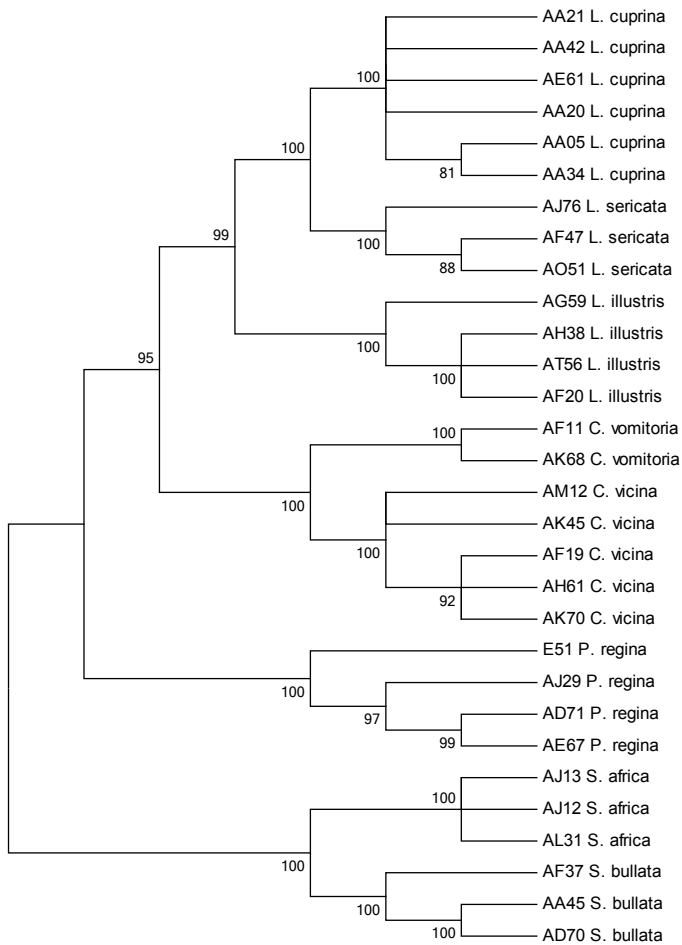


Fig. 6. Number of informative characters in the COI-COII region calculated for each adjacent 200bp window.

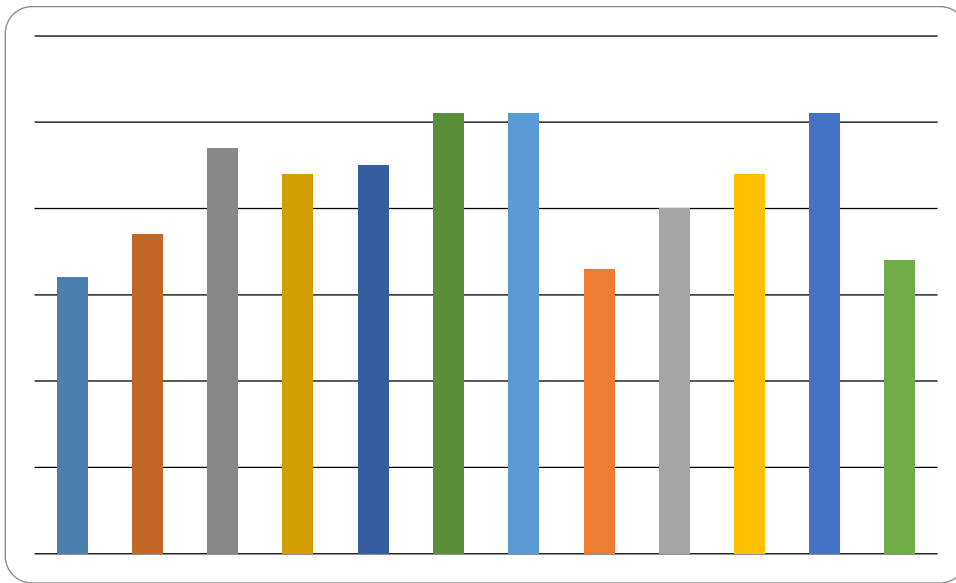


Fig. 7. Mean relative divergence for fragments of increasing length for concatenated DNA sequences of the COI, COII and ND4 genes starting at either the 5' end of COI, the midpoint of COI, COII and ND4 or the 5' end of ND4.

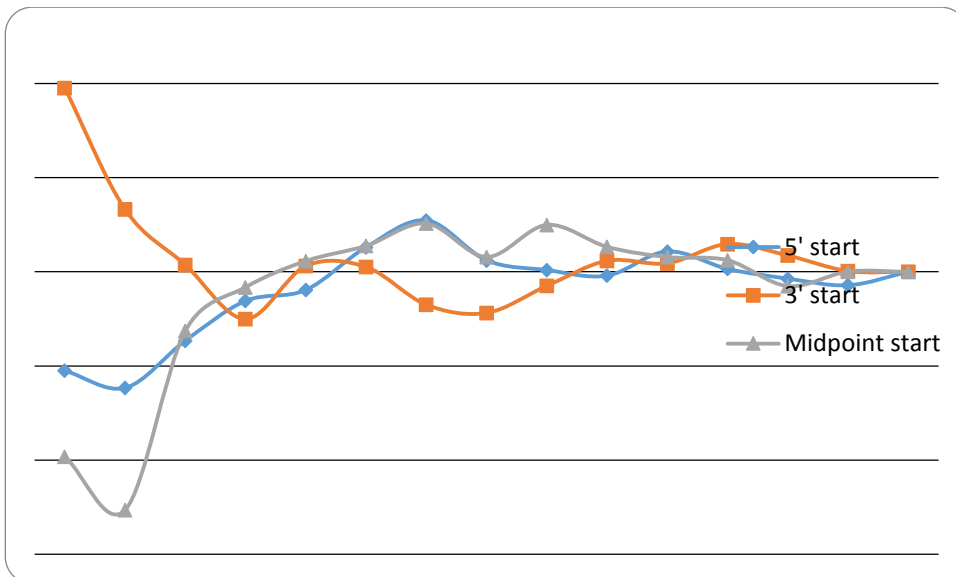


Fig. 8. ME tree based on a 200bp fragment at the 5' end of the COI gene.

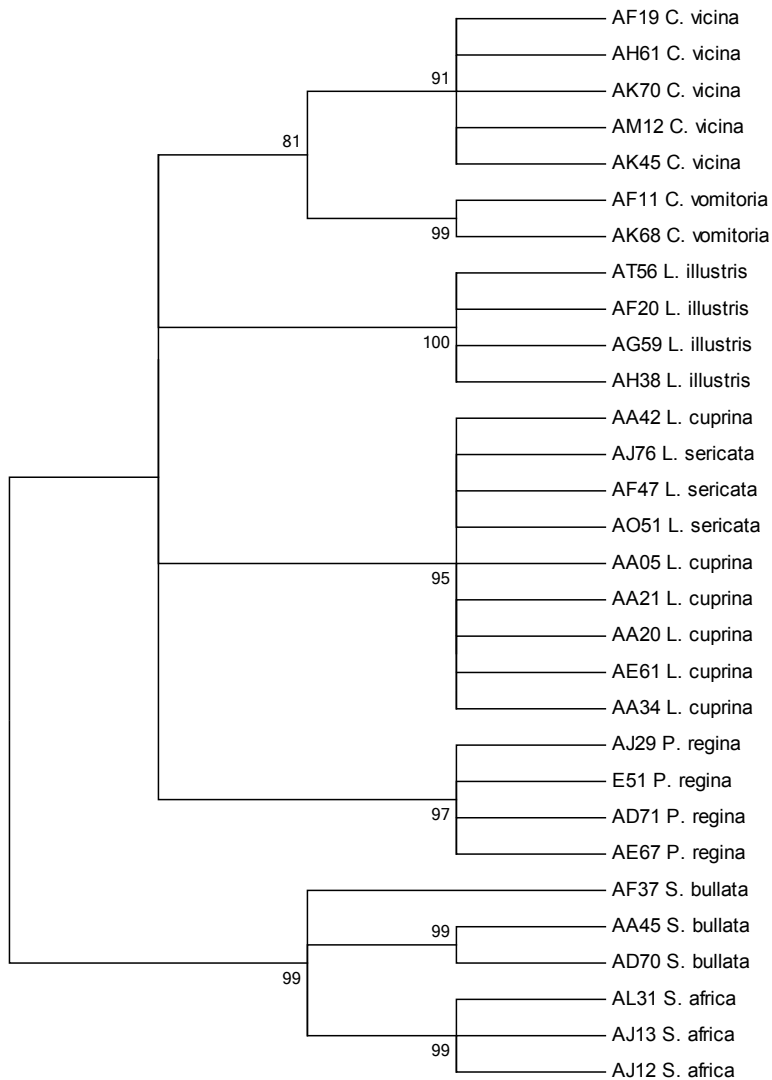


Fig. 9. Bootstrap support for increasing fragment length of *L. cuprina*.

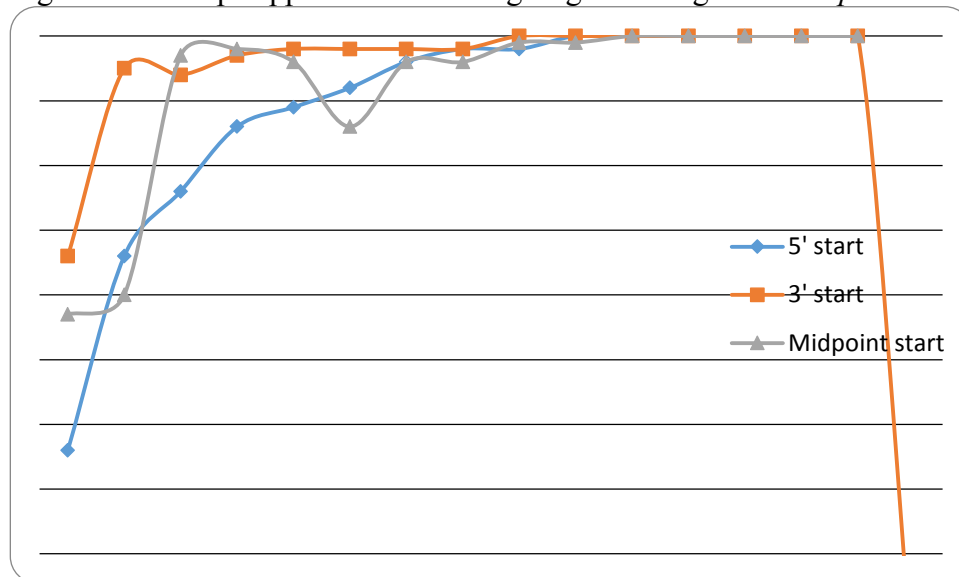
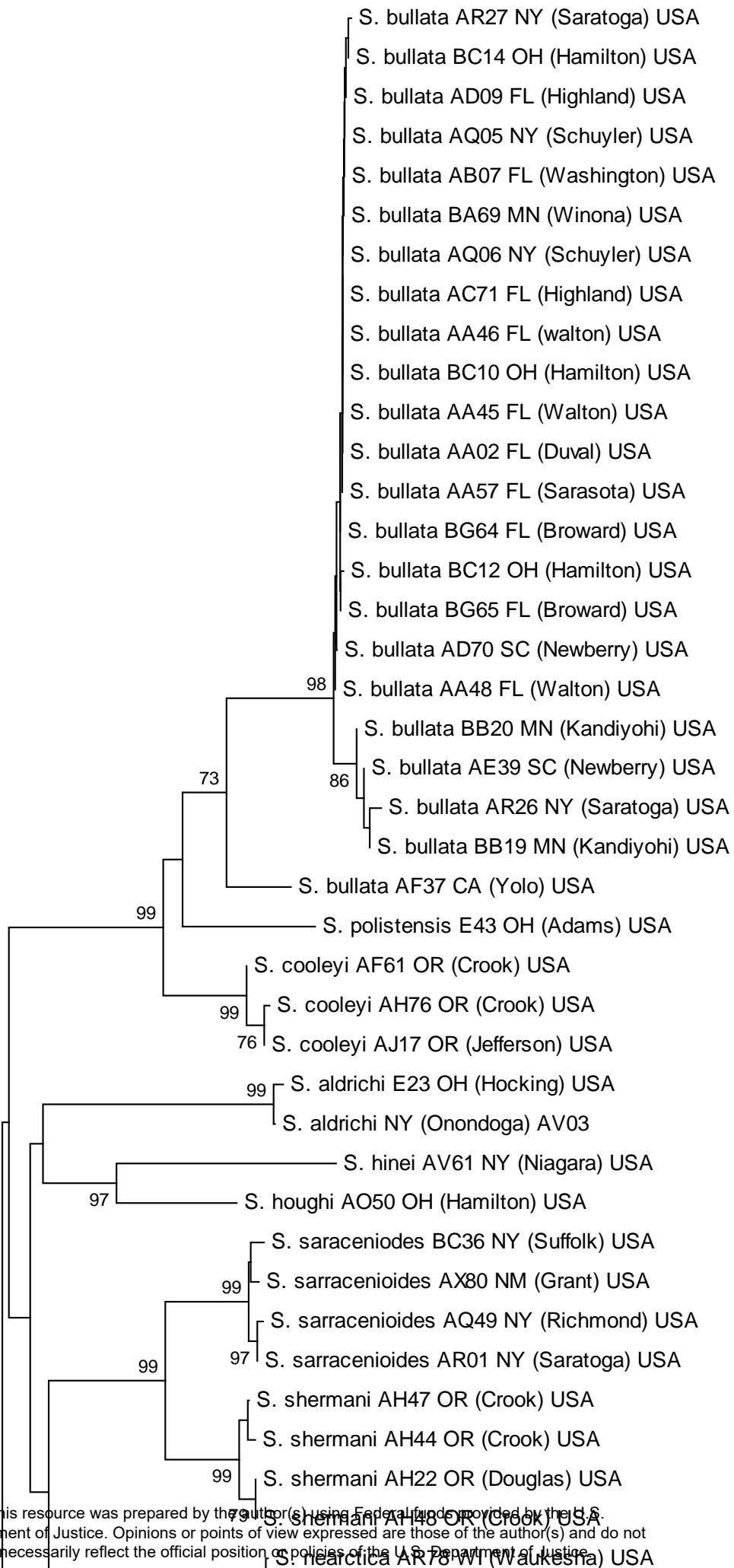
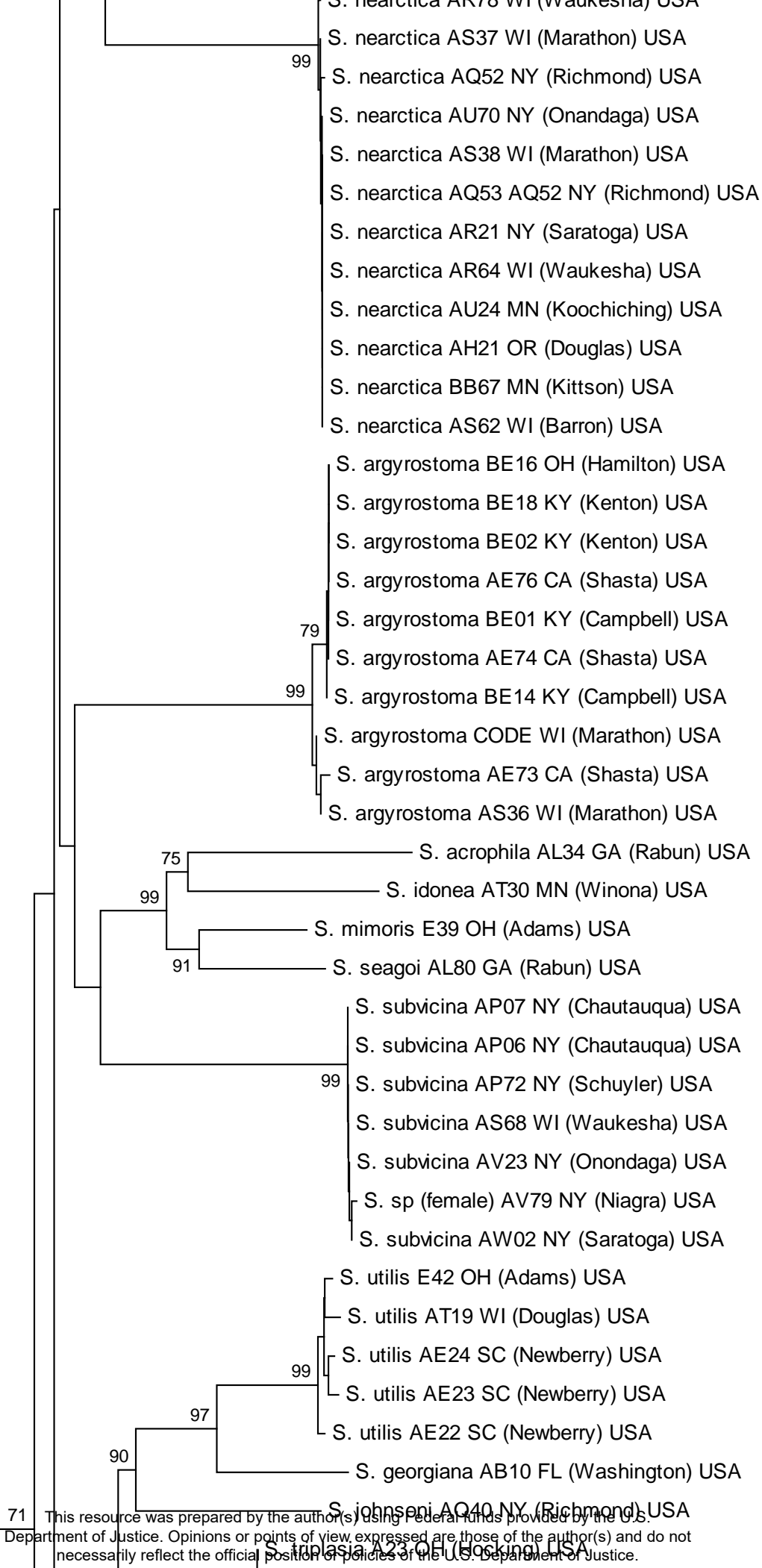
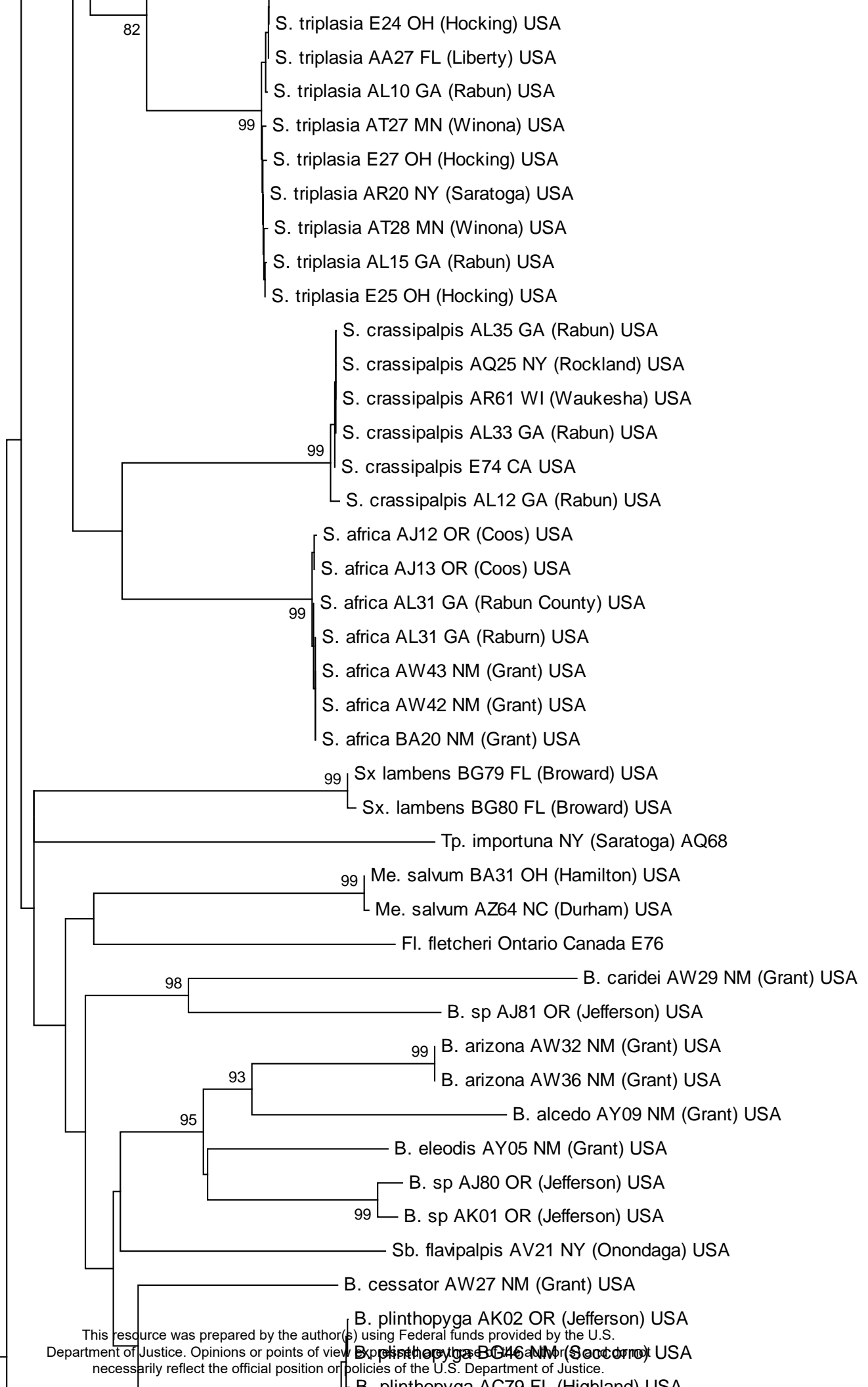


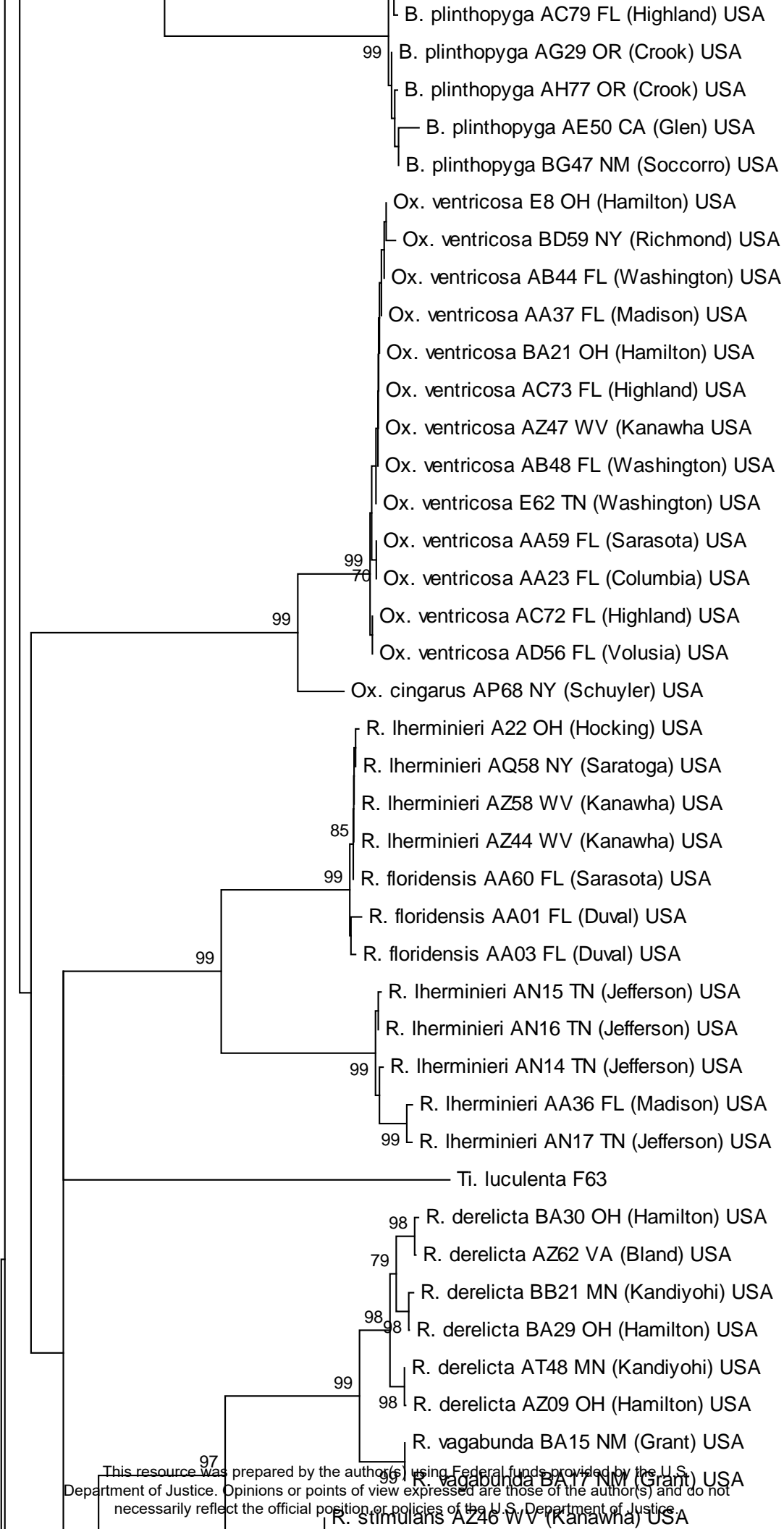
Fig. 10 ME phylogenetic analysis of the reference database. (Note: this is the largest that Word will allow for the figure. A larger sized PDF version will be uploaded as an Appendix)



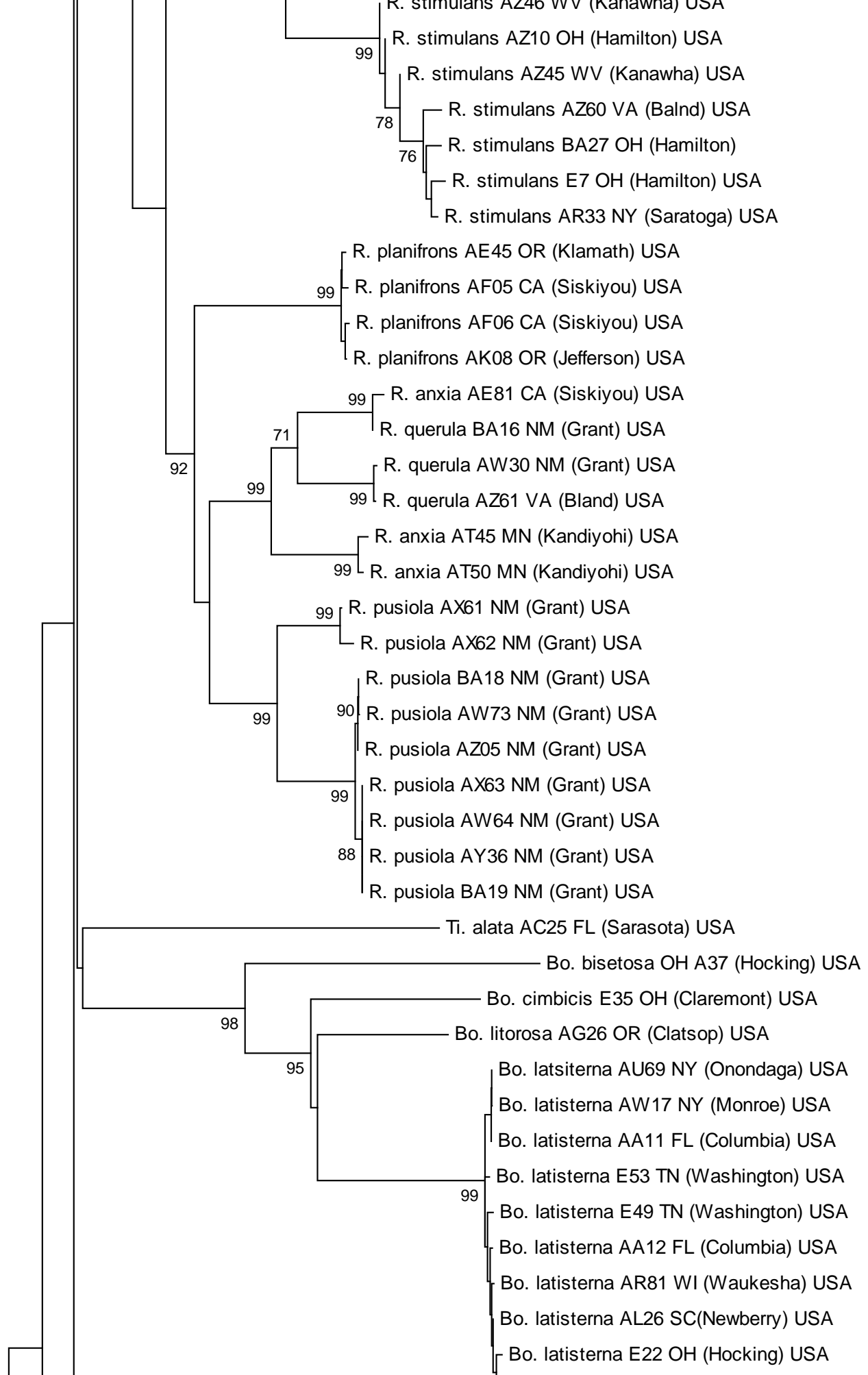
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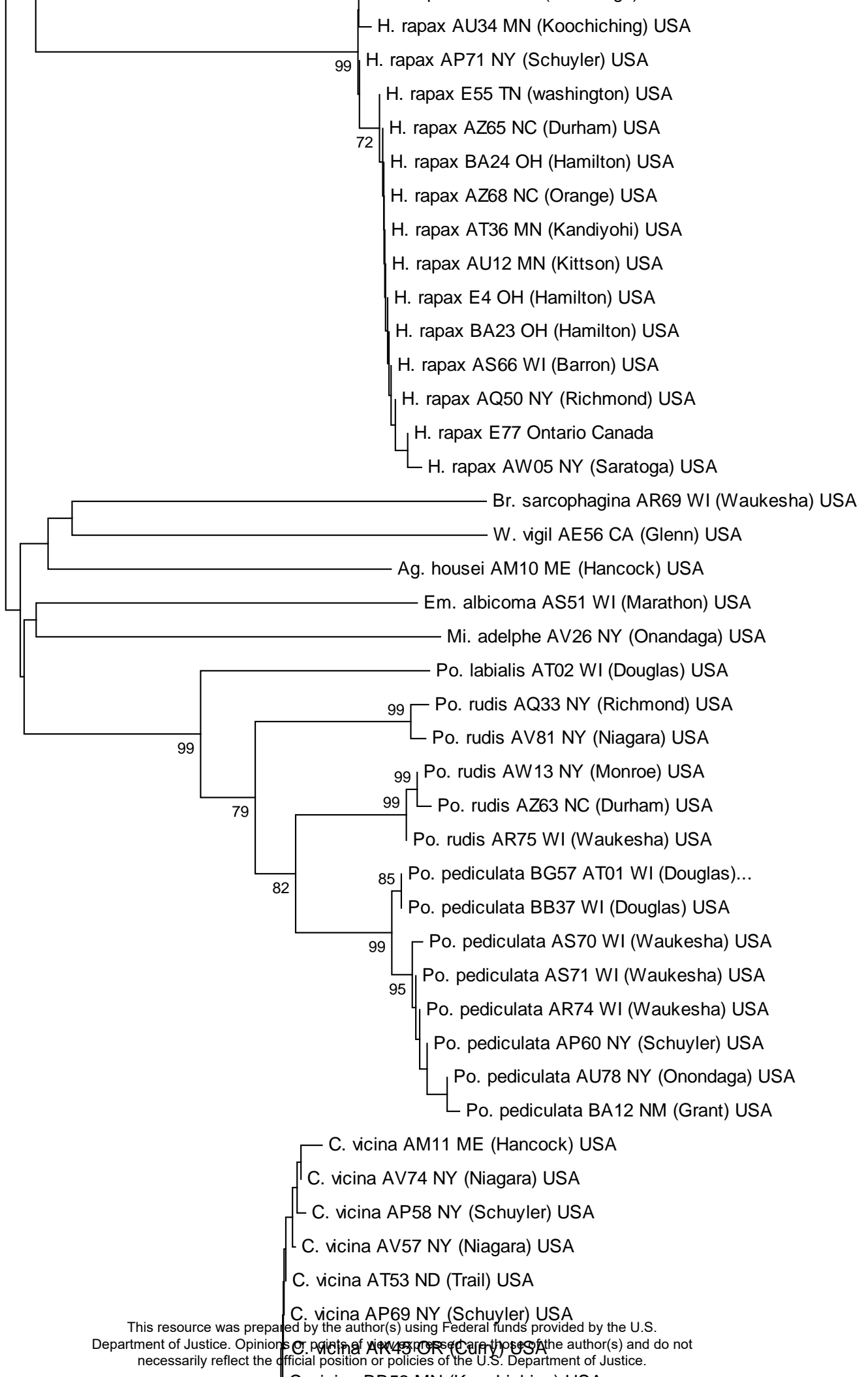




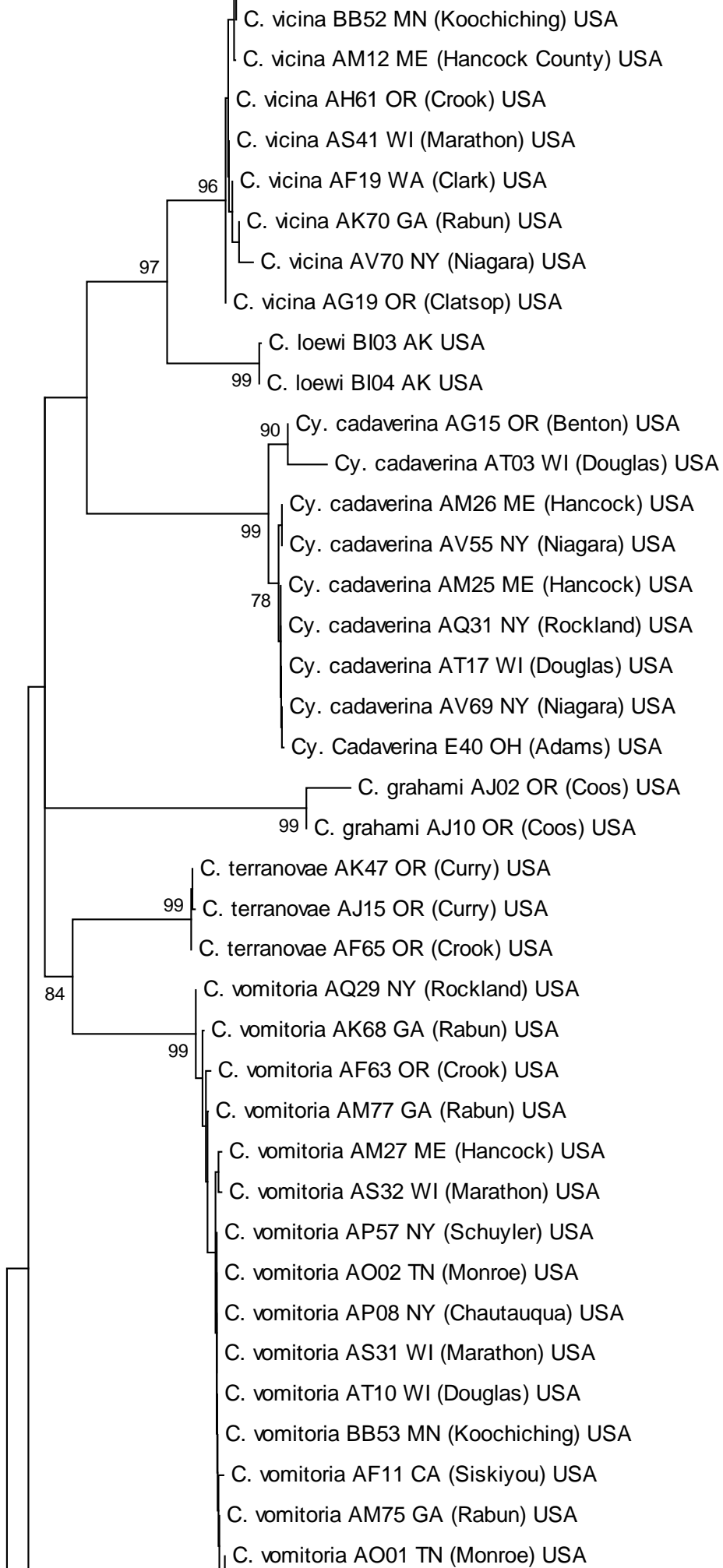


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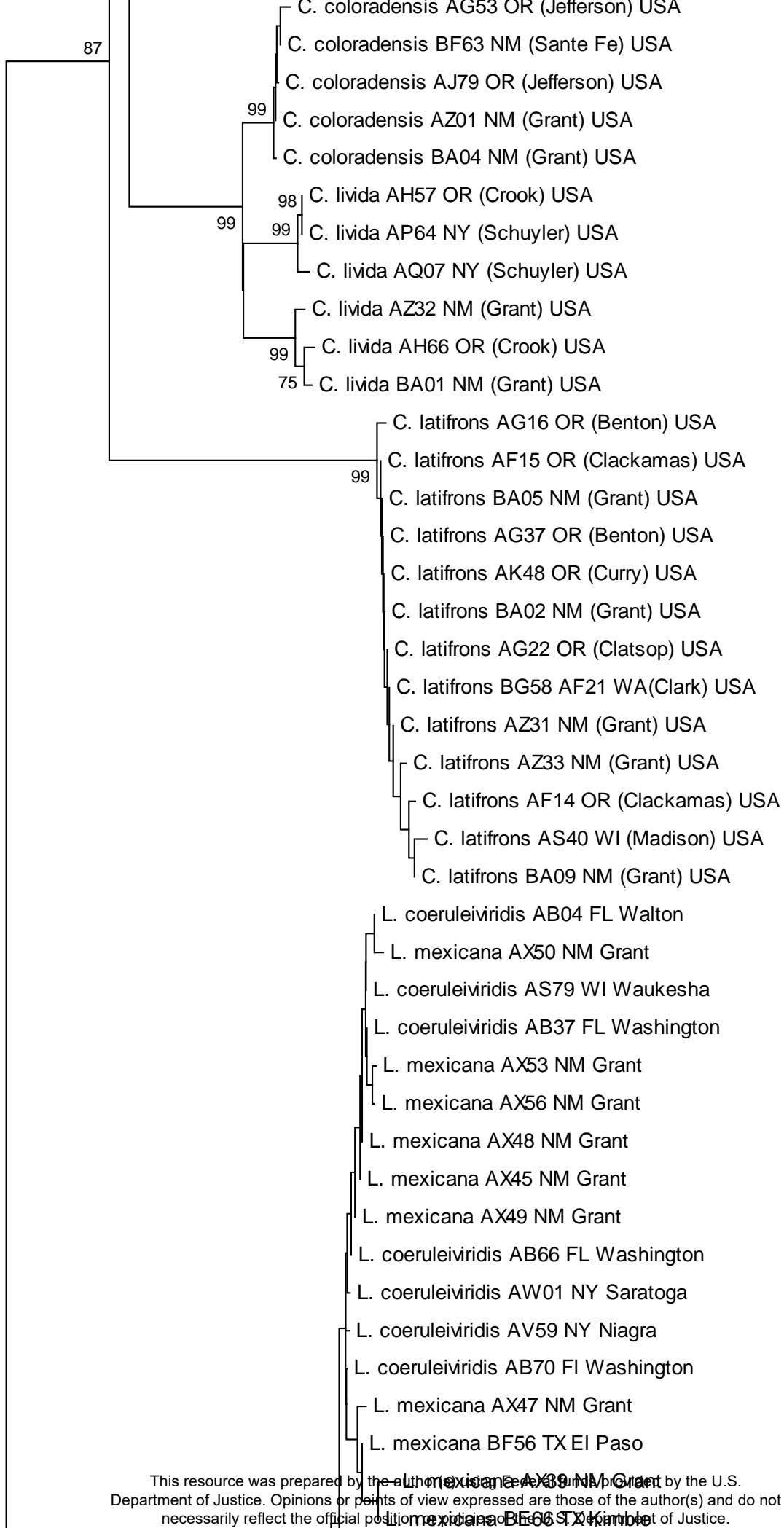




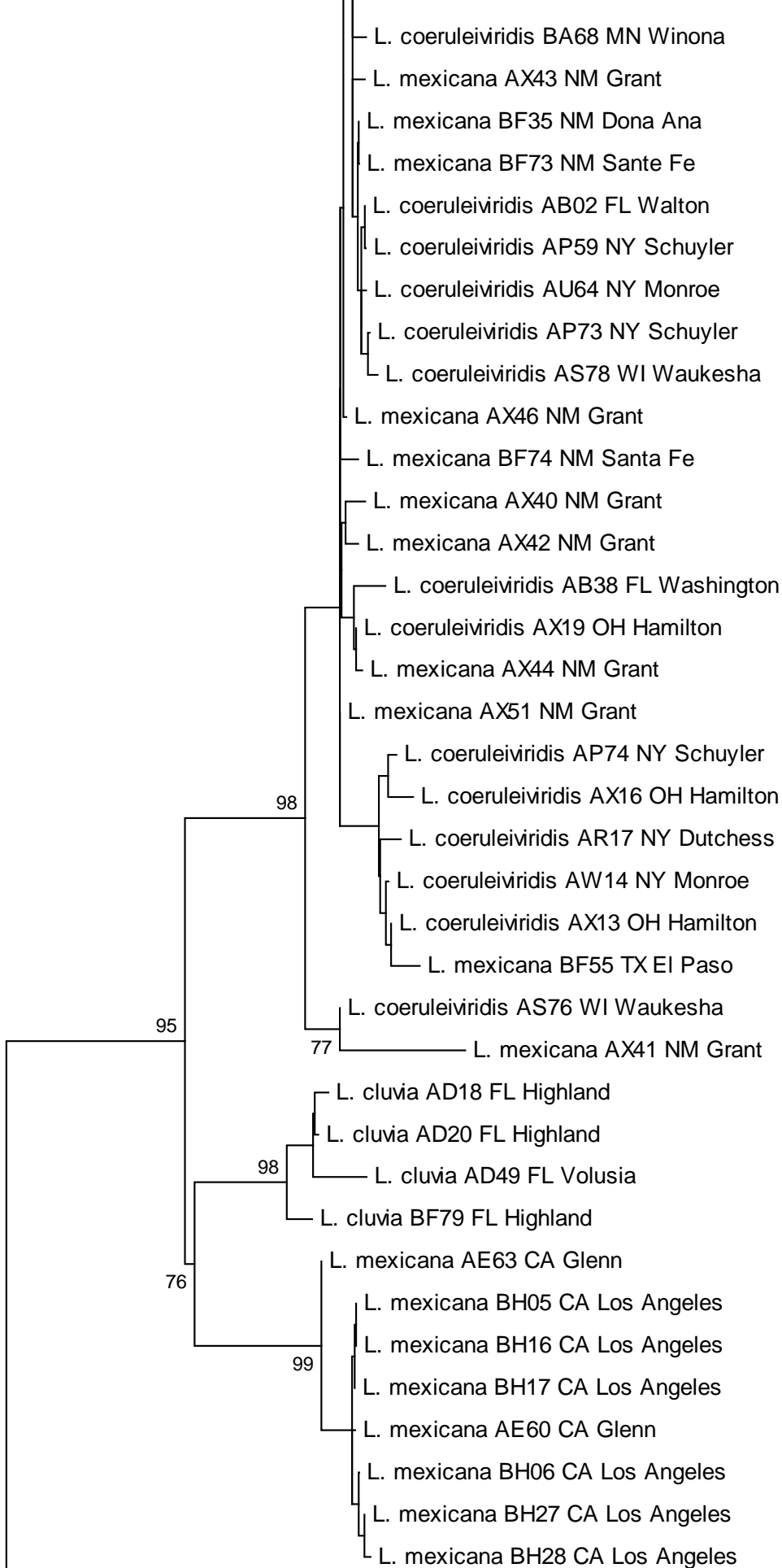
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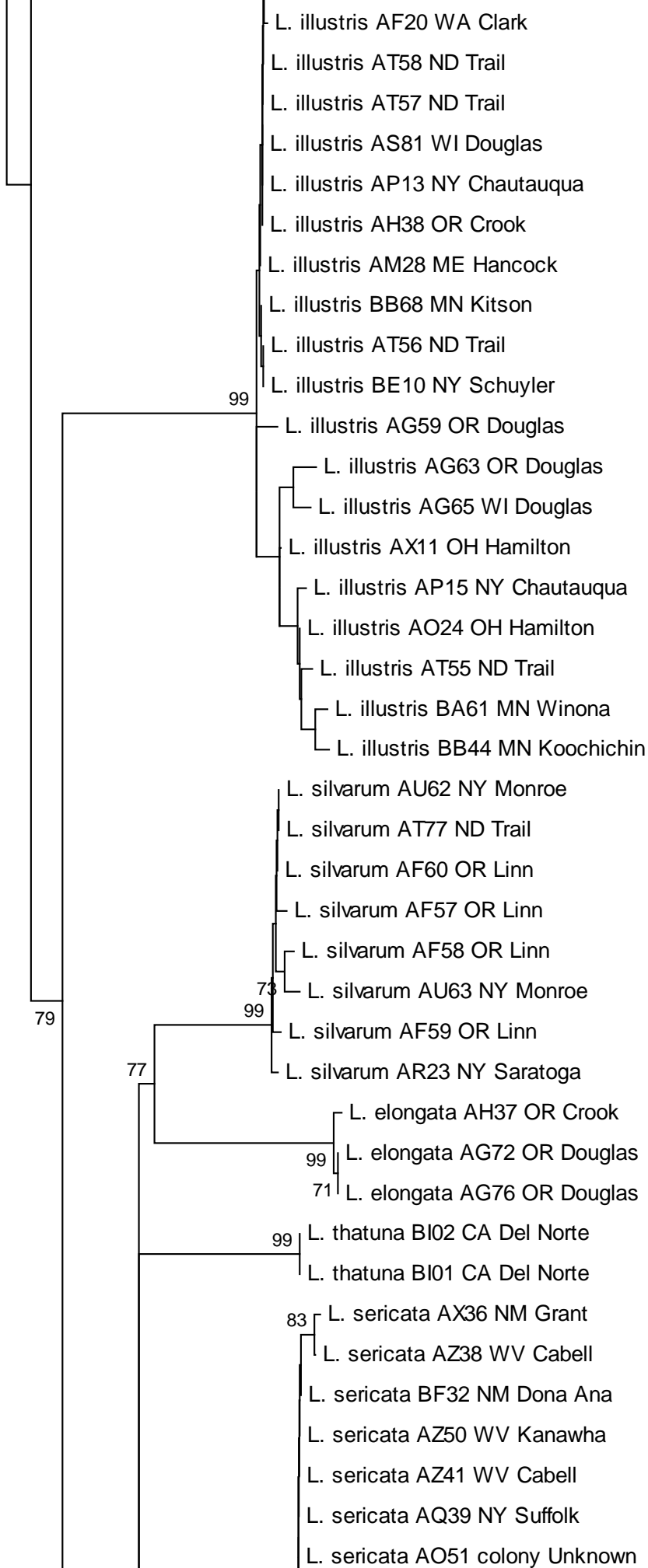


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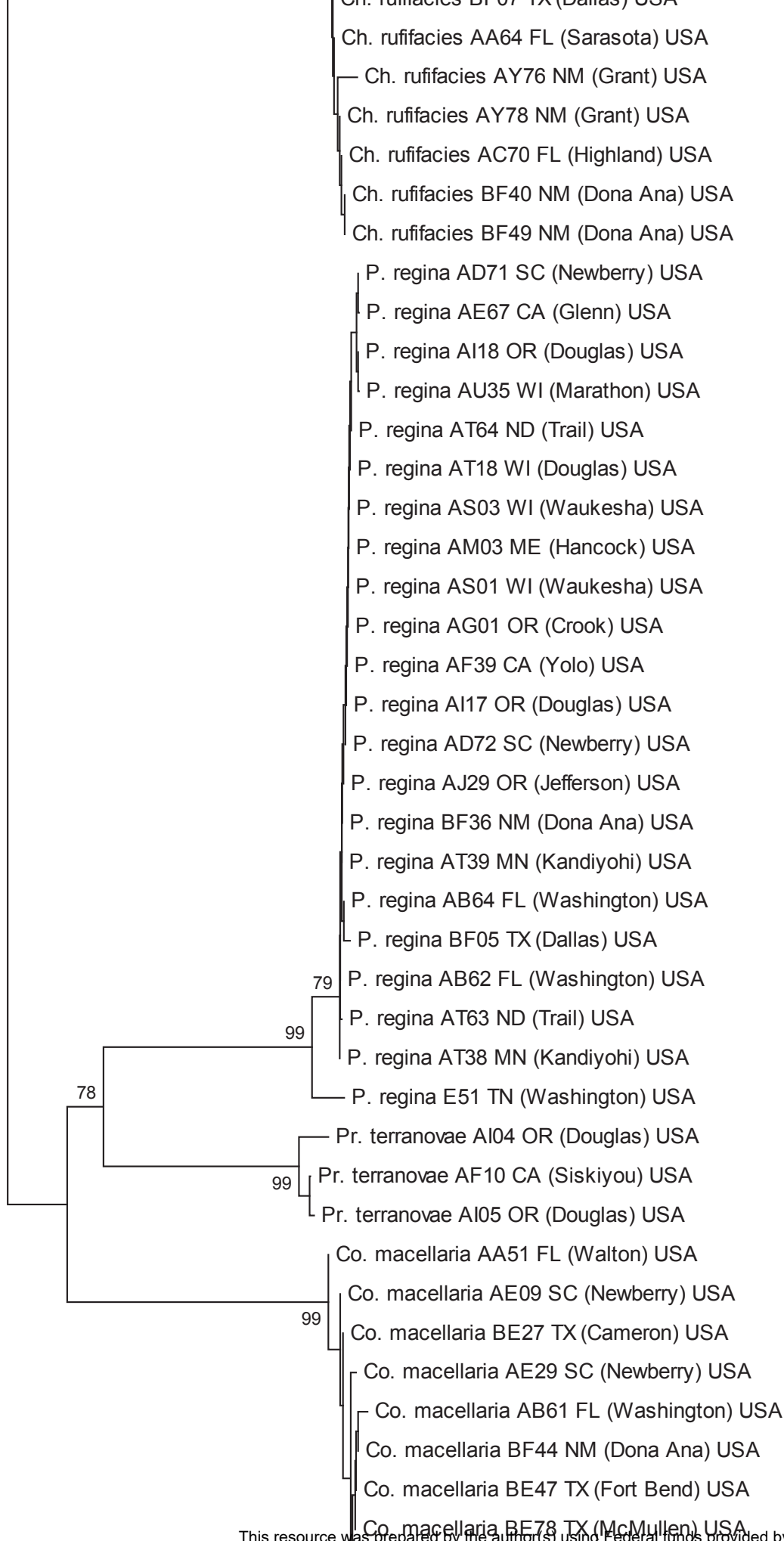
This resource was prepared by the author(s) using Federal funds provided by the U.S. Department of Justice. Opinions and conclusions are those of the author(s) and do not necessarily reflect the official position or policies of the U.S. Department of Justice.

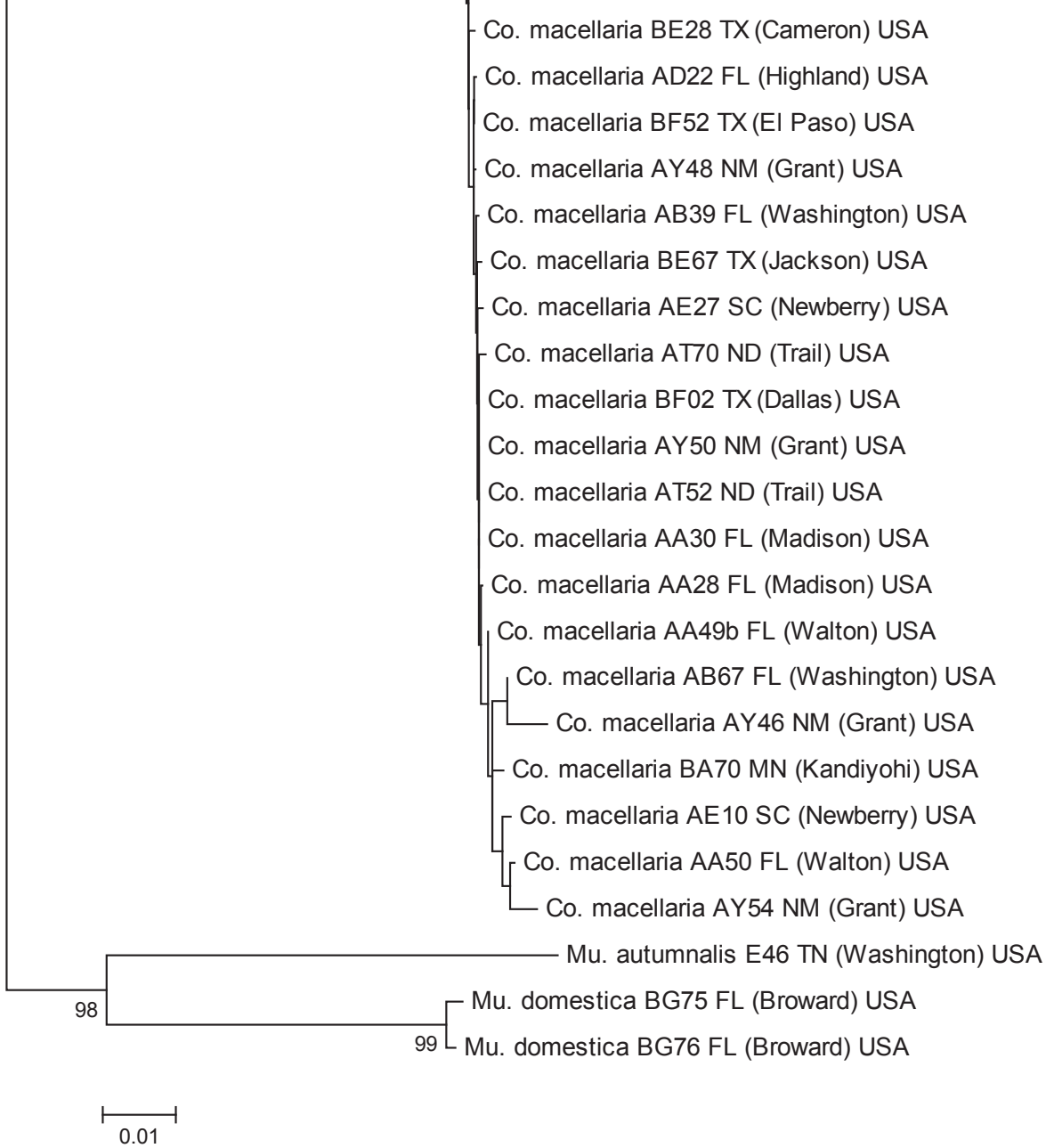
L. sericata AF47 CA Yolo

L. sericata AF47 CA Yolo
 L. sericata AZ12 OH Hamilton
 L. sericata AZ70 NC Davie
 L. sericata BF60 TX El Paso
 L. sericata AJ76 OR Jefferson
 L. sericata BF62 TX El Paso
 L. sericata BG37 NM Los Alamos
 L. cuprina AE61 CA Glenn
 L. cuprina BF11 TX Dallas
 L. cuprina AA42 FL Bay
 L. cuprina AA40 FL Bay
 L. cuprina AA21 FL Columbia
 L. cuprina AA18 FL Columbia
 L. cuprina BG72 FL Broward
 L. cuprina BF12 TX Dallas
 L. cuprina AA05 FL Duval
 L. cuprina BG02 FL Monroe
 L. cuprina BF13 TX Dallas
 L. cuprina AA34 FL Madison
 L. cuprina BE33 TX Cameron
 L. cuprina AA20 FL Columbia
 L. cuprina BE34 TX Cameron
 L. cuprina BE35 TX Cameron
 L. cuprina BF10 TX Dallas
 L. cuprina BF09 TX Dallas

Ch. megacephala BG60 FL (Monroe) USA
 Ch. megacephala BE20 TX (Cameron) USA
 Ch. megacephala BG59 FL (Monroe) USA
 Ch. megacephala BG24 FL (Monroe) USA
 Ch. megacephala BG23 FL (Monroe) USA
 Ch. megacephala BF21 TX (Cameron) USA
 Ch. megacephala BE19 TX (Cameron) USA

Ch. rufifacies AD43 FL (Volusia) USA
 Ch. rufifacies BE65 TX (Kimball) USA
 Ch. rufifacies AD19 FL (Highland) USA
 Ch. rufifacies AC78 FL (Highland) USA
 Ch. rufifacies AC77 FL (Highland) USA
 Ch. rufifacies BE50 TX (Fort Bend) USA





IV. Conclusions

1. Discussion of findings:

The specific goals of this project were:

1. Identify a locus within the mitochondrial genome (mtDNA) that is maximally informative for species identification using phylogenetic analysis.

We identified a locus within the mitochondrial genome that is highly informative with respect to DNA-based species identification of forensically important Diptera. The region we selected for Phase II (part of the 3' region of COI) is highly efficient for discriminating among the vast majority of forensically important Diptera found in the contiguous United States. It is able, for example, to discriminate between the particularly difficult species pair *Lucilia cuprina* and *L. sericata*.

2. Produce a comprehensive database for this locus for all carrion fly species in the families Calliphoridae and Sarcophagidae that are found in the contiguous U.S. and that have literature reports of association with large-animal carrion or human remains.

We produced a database comprising approximately 1,200 bp from the 3' end of the mitochondrially encoded gene COI. The database includes data for 36 of the 39 species of calliphorid and sarcophagid flies, plus two species of muscid flies, that were identified earlier as being of probable forensic importance. The database also includes data for 68 additional species of Sarcophagidae and Calliphoridae that, while not likely to deposit larvae on human remains, were collected at chicken meat bait traps and, therefore, may well be encountered as adults at a human death scene.

For the species of greatest forensic importance, one very closely related species pair (*L. mexicana* and *L. coeruleiviridis*) cannot be distinguished. This reflects the fact that no single short segment of DNA is likely to be 100% successful at discriminating among all species. In the particular case of *L. mexicana* and *L. coeruleiviridis*, we examined additional regions of the mitochondrial genome without finding a region (or, indeed, any combination of regions) that was capable of discriminating between that pair. It is apparent that discrimination between those two species by DNA data will require use of a nuclear locus (or loci). Even then, the prospects for success will depend on the (unknown) nature of the biological process that has led the two species to share a single mitochondrial lineage. It is possible, for example, the different nuclear loci will show different patterns of genealogical relationships among individuals.

For the remaining 16 species of primary forensic importance and 16 of the 20 species of limited forensic importance in the database, however, the level of discrimination is excellent. We see no reason to believe that it will be necessary to develop additional mitochondrially encoded loci for species identification of forensically important Diptera in the contiguous United States.

The reference database produced by this project is far more comprehensive than any similar database, in terms of species coverage, geographic diversity within the 48 contiguous United States, and depth of sampling within forensically important species.

2. Implications for policy and practice:

The project can safely be described as a success. The database can be used for DNA-based identification of forensically important Diptera in the contiguous United States. Nearly all of the calliphorid and sarcophagid species of the greatest forensic importance and a very large majority of species with marginal forensic relevance are included in the database. In addition, nearly all of these species (with the exceptions noted in detail above) have COI haplotype sequences that are sufficiently distinct from other forensically relevant species so as to allow robust species determination.

Availability of this database will, hopefully, lead to an expansion in the use of entomological evidence in death-scene investigations. The use of DNA instead of adult morphology to identify dipteran

larvae recovered at a death scene can allow for such identifications to be performed by crime lab personnel, without the need to send crucial evidence to a specially trained outside consulting entomologist. The expertise of a forensic entomologist will always be necessary, but if species identity is established within the crime lab, the forensic entomologist can concentrate on determining the developmental age of the larvae recovered and on converting those ages into a best estimate for PMI.

The ability to use DNA sequence data to identify these important forensic indicator species will also, hopefully, spur additional research into species-specific developmental trajectories. Such data are required for PMI estimation, regardless of the species recovered from the death scene, and regardless of the method used to obtain species identification.

The remaining problems will be expanded upon in the next section, as implications for further research. Briefly, they include: 1) one particular pair of species with high forensic importance (*Lucilia coeruleiviridis* and *L. mexicana*) cannot be distinguished using COI sequence data (and indications from additional mtDNA data suggest that no mitochondrial locus is likely to be successful in that case); 2) one additional pair of species, also in the genus *Lucilia* (*L. cuprina* and *L. sericata*) are genetically distinct, yet sufficiently similar in DNA sequence so that it would be wise to expand the reference collection for both species; 3) despite the inclusion of approximately 500 individual reference specimens, the utility and reliability of the database can still be increased by a further expansion of the reference collection; 4) the utility of the database for identification of unknown larvae needs to be validated.

3. **Implications for further research:** Recommendations for further research in this area should be described.
 - a. **The reference database is never “complete”** Forensic entomology must deal with natural, evolving species composed of multiple populations distributed across large geographic ranges. Consequently, a reference database for species identification can never really be considered “complete”. The database is always subject to expansion in terms of geographic coverage and depth of coverage (the total number of individuals of any particular species included). The variability inherent in natural populations was the original rationale for taking a phylogenetic approach to DNA-based species identification. Approaches that rely on a singular “signature” – including such technologies as microarrays of SNPs, or DNA fingerprinting based on length-variable-length microsatellite alleles – cannot as easily account for the high likelihood that an individual sampled from nature will be genetically unique and, therefore, not directly represented in the database. We have endeavored to sample each species in our reference database over a wide geographic area, but there will always remain some as-yet-un-discovered haplotypes. Therefore the reference database should, properly, be continually growing, with new reference specimens from new locations added.

Examination of the results presented in Fig. 10 leads to speculation that not every species carries the same need for additional specimens. For example, within the genus *Lucilia*, one species (*L. illustris*) is both well separated from any other species in the phylogenetic analysis and highly homogeneous, with very low levels of intra-species diversity and no hint of any geographic pattern to the intra-species diversity that is present. It is extremely unlikely that a member of any other species will, by chance, have a COI haplotype that could group with those of *L. illustris* or that, conversely, that there exists a yet-unsampled *L. illustris* haplotype that would group with a different species in the phylogenetic analysis (note, however, that this inference is not based on an explicit probabilistic model). Contrast that situation to the one presented by the closely related pair of species *L. cuprina* and *L. sericata*. Although levels of intra-species diversity are rather low, COI haplotypes from the two species are very similar to each other. This similarity means that there will always be a much higher possibility that there exists somewhere within one of those species a genotype that is more similar to the “wrong” species, compared to the situation with *L. illustris*. There might be no such genotype, of course, but it would still be prudent to

continue sampling additional *L. cuprina* and *L. sericata* to obtain an even more detailed picture of the scope of variation within each species.

- b. Additional theory needs to be developed to justify conclusions of positive identification based on reciprocal monophyly in phylogenetic analysis.** As alluded to in **a.**, there is currently no probabilistic framework to directly answer the question “What is the probability that a newly sampled individual of species X will, by chance, fall within the clade defining species Y in the reference database?” The answer, while undoubtedly related to sample size, is not likely to be a simple function of only sample size – the degree of observed genetic differentiation in the reference database will, no doubt, also be a critical factor. Such theory would be useful beyond the realm of forensic entomology in human death cases, as similar data and methods are applied to many forms of DNA-based species identification systems.
- c. Two kinds of validation studies will be required**
- i. *The ability to correctly identify unknown samples must be documented.* This critical step had been included in the original proposal submitted to NIJ. However, the reviewers of the original proposal suggested that we would develop a much better database if we postponed any validation studies and focused instead on increasing the number and geographic scope of field collecting activities and the total number of individual specimens incorporated into the database. We agreed with the review commentary, and implemented those changes via a change-of-scope GAN. That change in focus proved to be quite valuable, as the addition of several extra collecting trips enabled the reference database to have a much greater depth and breadth of coverage. This form of validation study should be accomplished in at least two phases. The first phase would be to challenge the database in a blind study with reared, known-species larvae. The second phase would be to test the reference database on non-probative casework samples. Ideally, these samples would be processed in parallel with specimens (larvae) that were collected live and reared to adulthood for traditional morphological identification.
 - ii. *The laboratory protocols will need to be validated.* There is a need to establish the range around the standard PCR conditions for which our primers will give adequate data. The laboratory procedures needed to obtain DNA sequence for an unknown sample are routine but will still require validation. The specific primer pair developed in this project to amplify the region of interest should be tested on a wide range of templates, including damaged or degraded template DNA. The database should be tested using query sequences that are shorter than the full length. In addition, the primers should be tested under conditions in which multiple potential targets of amplification are present. For example, the primers should be tested when the target is present in a mixture with other (e.g., human) genomic DNA.

V. References: Citations are in a format similar to that used by professional journals, and are appropriate for the study performed.

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VI. Dissemination of Research Findings: Publications and presentations resulting from this award should be listed, including the full journal citation, scientific conference and location, etc.

Presentations based on this project have been given at:

NIJ DNA Grantees Workshop / NIJ Conference, Arlington, VA: 2005-2011

Entomological Society of America, Indianapolis, IN, 2009

Entomological Society of America, San Diego, CA, 2010

North American Forensic Entomology Association

American Association for Forensic Sciences

Publications:

DeBry, RW, AE Timm, GA Dahlem, T Stamper. 2010. mtDNA-based identification of *Lucilia cuprina* (Wiedemann) and *Lucilia sericata* (Meigen) (Diptera: Calliphoridae) in the continental United States. *Forensic Sci Int* 202:102-109.

In Press:

DeBry, RW, AE Timm, ES Wong, T Stamper, C. Cookman, GA Dahlem. 2012. DNA-based identification of forensically important *Lucilia* (Diptera: Calliphoridae) in the continental United States. *Forensic Science International*.

At least 3 additional journal submissions are in preparation based on this work.