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**IDENTIFICATION OF FIREFLIES (COLEOPTERA: LAMPYRIDAE) AT THAYER  
FARM, OTSEGO COUNTY, NY**

BY

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THESIS

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## PREFACE

32  
33 Organismal bioluminescence is an intriguing phenomenon. The ability for something  
34 living to glow and produce light seems unnatural. The process of converting chemical energy  
35 into light is a popular field of research (e.g., Day et al. 2004). Fireflies (Coleoptera:  
36 Lampyridae) are a family of insects that can emit light via unique biochemical pathways. The  
37 presence and interaction of the enzyme luciferase with its substrate luciferin in the abdominal fat  
38 bodies of fireflies catalyzes the chemical reaction that produces visible light. In the process,  
39 luciferase also converts luciferin into oxyluciferin, which is considered the “light emitter”  
40 because this transformation aids in the production of bioluminescence (Marques and Esteves da  
41 Silva 2009). Oxyluciferin then interacts with luciferin-regenerating enzyme (LRE) to create  
42 additional luciferin so the reaction can start over again (Day and Bailey 2003). The repetition of  
43 this chemical reaction is necessary for fireflies, as each species uses a different flash pattern in  
44 mate recognition: females lay flashing in the grass while males are airborne, signaling to females  
45 using species-specific flashing patterns. These patterns differ in flash duration, interval, and  
46 color, which are influenced by the amount of luciferin and luciferase produced by the firefly  
47 (Demary et al. 2006). Several firefly species are active during the day and do not require  
48 bioluminescence. Instead, these diurnal fliers use cuticular pheromones, or chemical signals, to  
49 find a mate (Shibue et al. 2004).

50 Fireflies are a notable group of organisms not only in their varied communication  
51 abilities, but in their taxonomic breadth. More than 100 genera and 2,000 species of Lampyridae  
52 have been recorded worldwide, and these numbers are expected to increase, as many species are  
53 yet to be described (Stanger-Hall et al. 2007). More than 75 of these species are found in the  
54 eastern United States and Canada (Faust 2017); however, there is a clear lack of organized  
55 knowledge regarding the identity and specific distribution of fireflies in New York. There are

56 keys available to identify fireflies, but a local, comprehensive, key does not exist, making it  
57 difficult to properly classify individuals. Luk et al. (2011) produced the only geographically  
58 proximal key; however, their work focuses on the Canadian province of Ontario. Taxonomic  
59 identifications are further complicated by species boundaries based on bioluminescence. In the  
60 genus *Photuris*, individuals are nearly indistinguishable physically, requiring a known flash  
61 pattern for identification. It is difficult to identify species based on their flash patterns alone.  
62 For example, species can differ based on subtle changes in the wavelength or frequency of the  
63 light emissions. Further confounding identification attempts, female *Photuris*, commonly  
64 referred to as “femmes fatales,” can mimic the flash patterns of other species to lure in and  
65 devour males (Faust 2017, Lloyd 2018).

66         During the last half century, molecular science has transformed the field of biology.  
67 Ground-breaking techniques have allowed us to understand the world on a molecular level,  
68 letting us discover information about the structure and function of molecules, including DNA,  
69 RNA, and proteins (Davis 2012). Molecular data can be used to elucidate relationships between  
70 organisms that may not be discernable by morphological systematics alone. Molecular  
71 phylogenies use information such as DNA sequences to help hypothesize relationships and  
72 describe evolutionary patterns of related species. DNA barcoding, the mass sequencing of a  
73 single gene from multiple species to create a molecular standard, is a tool used for distinguishing  
74 species that are physically indistinct (Hebert et al. 2003). Use of phylogenetics, specifically in  
75 comparative research, has increased as phylogenies are helpful in resolving species identities and  
76 evolutionary uncertainties (Soltis and Soltis 2003).

77         With increased use of DNA sequence data for a variety of purposes, global online  
78 repositories and databases have also become widely used. The genetic sequence database

79 GenBank contains over 150 million DNA sequences from many organisms but should be used  
80 with caution when trying to determine species identifications. Comparing a sequence to one in  
81 the database can help confirm a species if they match closely enough, but only if the sequence  
82 submitted to GenBank is associated with the correct species. Revisions to the database can be  
83 made by the original submitter, but this process tends to be problematic, as it requires the person  
84 to recognize and admit fault. Mis-classified sequences can lead to unreliable results (Fritz et al.  
85 2012). Critiques of GenBank and other databases including the Barcode of Life Data systems  
86 (BoLD) are common in taxonomic literature. Many discuss the ambiguity of identifications and  
87 ways to fix some common issues (e.g., Chang and James 2011, Fritz et al. 2012, Sonet et al.  
88 2013).

89 Morphology and molecular methods started out as separate analysis techniques, but the  
90 integration of phylogenetic results with traditional morphological classifications has gained  
91 traction with many biologists. The integrative taxonomy approach, combining both  
92 morphological and molecular techniques, is a valuable tool in determining the identity of an  
93 organism. It has become prominent in recent literature (e.g., Chang and James 2011, Darienko et  
94 al. 2015, Kirichenko et al. 2015, Goulding et al. 2018) and will most likely continue to assist in  
95 taxonomic classifications into the future. The two methods that are typically employed by  
96 taxonomists are “integration by congruence”, which uses mitochondrial DNA and morphology in  
97 species identifications, and “integration by cumulation”, which looks at only mitochondrial DNA  
98 or morphological characteristics to identify an individual (Padiál et al. 2010). “Integration by  
99 cumulation” tends to increase type I error, over-estimating the number of distinct species  
100 identified. On the other hand, because molecular databases may not be up to date with recently  
101 discovered and/or cryptic species, “integration by congruence” may under-estimate the number

102 of species that are actually present (increased type II error). It is essential that we focus on the  
103 big picture when dealing with taxonomy, keeping in mind that species are not only defined by  
104 the visual patterns we observe, but also by the genetic diversity responsible for functional  
105 differences (Padial et al. 2010).

106         Understanding what species are present in an area is imperative for both scientific  
107 research and conservation efforts, and species distribution data are necessary for unbiased and  
108 realistic conservation policies. In order to plan conservation efforts, biological surveys of  
109 species distribution patterns should be performed (Margules et al. 1994). Without understanding  
110 the range across which a population exists, it is almost impossible to assess which populations  
111 are at risk. Peripheral populations, or populations at the geographic edge of a species range, are  
112 known for their lack of genetic diversity. These populations experience less gene flow since they  
113 are typically established through founders' effects (Channell 2004). On the other hand, Lesica  
114 and Allendorf (1995) made a case for conserving peripheral populations. Central populations  
115 adapt to different environmental conditions compared to peripheral populations. Since these  
116 populations are genetically distinct from central populations, their ability to adapt to  
117 circumstances outside of the norm in our changing environment make them essential in  
118 protecting the evolutionary future of populations.

119         Examining how large-scale environmental changes impact species over time requires first  
120 identifying what organisms are currently present and where they exist. For example, we cannot  
121 understand the full impacts of increased artificial light and other anthropogenic alterations on  
122 fireflies without first knowing the range of species found in specific areas. Ineichen and  
123 Rüttimann (2012) found that male fireflies do not have the ability to locate females that establish  
124 mating sites under street lamps or other light sources, or males simply avoid illuminated areas

125 altogether, both lowering the chances of reproduction (Ineichen and Rüttimann, 2012). There is  
126 evidence based on species distribution data taken in Florida that the species *Micronaspis*  
127 *floridana* is threatened due to artificial light pollution, increased development, and other human  
128 induced modifications (Faust 2017); however, it is unclear which populations are likely to be  
129 affected by artificial light in New York.

130 Published literature on fireflies has not been consistent over time. In the late 1800s, John  
131 LeConte was the first to describe many species of fireflies that we see in the United States  
132 (LeConte, 1881). About a century later, James Lloyd discovered new lampyrid species and  
133 published on the flash behavior of fireflies (Lloyd 1966, 1969). Marc Branham and Michael  
134 Greenfield (1996) published a study on female attraction to flash patterns, being the first to  
135 examine the natural selection of fireflies. Lloyd (2018) recently published the first extensive  
136 guide to North American *Photuris*, introducing new species and helping “fireflyers” identify  
137 species by creating elaborate graphics of flash patterns. Research on firefly bioluminescence at  
138 the organismal level has been overshadowed by studies performed with bioluminescence at the  
139 chemical and cellular levels. The chemical reaction that produces bioluminescence is commonly  
140 used in medical research, specifically bioluminescent imaging and tagging in tumor detection  
141 and cancer treatment assessment (Rehemtulla et al. 2000). Not only is the literature lacking, but  
142 the habitats of bioluminescent organisms, typically dark forests and oceans, are challenging to  
143 explore (McInnis 2015).

144 In order to expand upon our knowledge of fireflies and understand what species are in the  
145 area, this study focused on the identification of fireflies at Thayer Farm, a 104-hectare property  
146 of the much larger 607-hectare holdings of the State University of New York, College at  
147 Oneonta, Biological Field Station (BFS). The history of Thayer Farm goes back to the 1800s. It



148 was home to many generations of the Thayer family who used the property for agricultural  
149 purposes and built structures to support the production of grains, hops, dairy, and wool (Staley  
150 2010). The property was left to the Oneonta Foundation to be used by the State University of  
151 New York, College at Oneonta, as a Biological Field Station in 1999. Most structures were  
152 remodeled and are currently used for research and education purposes. The fields of the property  
153 are partially old-field/hay and are currently surrounded by mixed forests. A group of chain  
154 ponds can be found among the fields, which are still maintained.

155         The spacious property supports a diverse community of fireflies in a very accessible  
156 location in Otsego County. Lampyrids were collected and morphologically identified to species  
157 based on regional keys developed for Ontario, Canada. To confirm identifications, genetic  
158 techniques were implemented and the highly-conserved mitochondrial gene Cytochrome  
159 Oxidase I (COI) was used as a barcoding gene for species identifications (Folmer et al. 1994). A  
160 Bayesian phylogenetic tree was generated using the Bayesian clustering program MrBayes,  
161 giving insight into the evolutionary relationships between the species of fireflies discovered.  
162 Species-accumulation curves were made to determine species coverage of the survey. A  
163 complete taxon list was then compiled and new species records for Otsego County and New  
164 York were documented. In addition, a comprehensive list of fireflies that can be found in New  
165 York was created by combining the results of the current study with publicly available species  
166 distributions, and a Lampyridae genus and species key for the county was established.

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1 **SPECIES SURVEY OF LAMPYRIDAE AT SUNY ONEONTA BIOLOGICAL FIELD**  
2 **STATION; INCLUDING NEW STATE AND COUNTY RECORDS FOR OTSEGO**  
3 **COUNTY, NY**

4 SAMANTHA CASSATA, JEFFREY HEILVEIL

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6  
7 *Abstract*—

8 Fireflies are a family of insects that are understudied, especially in their species distribution.  
9 Minimal information is documented for the state of New York, and there are no species records  
10 for Otsego County. A lack of distributional data makes it difficult for other research to be  
11 completed on fireflies, and impedes understanding about which firefly species, if any, require  
12 protection. In addition, there are currently no firefly identification keys for the state. The most  
13 appropriate key to use was from the Canadian province of Ontario. Fireflies from Thayer Farm,  
14 a property of the State University of New York College at Oneonta (SUNY Oneonta) Biological  
15 Field Station, were collected and initially identified based on morphology. Molecular techniques  
16 were also applied and the DNA barcoding gene, Cytochrome Oxidase I (COI), was used for  
17 molecular species identifications. A total of 13 species from 6 different genera were discovered.  
18 As no previous county record exists in the literature, each species recovered was documented as  
19 a new species record for Otsego County. *Pyractomena palustris* is recorded for the first time as  
20 occurring in New York. A collective taxa list for New York was created based on current  
21 published literature on fireflies and a local-specific species key was constructed. (Lampyridae,  
22 Otsego County, Distribution).

26 INTRODUCTION

27 Almost 200 years after lampyrid species were first described, Marc Branham, a pioneer  
28 on the interaction of selection and firefly flash patterns reminisced in an interview, “I remember  
29 saying: ‘[Fireflies] are so common and so charismatic, I bet everything is known about them’ ...  
30 I was flabbergasted to see that there was far less than I thought” (McInnis 2015). As humans, we  
31 cannot help but be attracted to the glow of fireflies (Coleoptera: Lampyridae) and everyone is  
32 bound to have a memory of catching fireflies, but these creatures are still surprisingly  
33 understudied.

34 There are more than 2,000 different species of fireflies found globally (Stanger-Hall et al.  
35 2007) and most species produce a bioluminescent flash pattern that is unique in flash duration,  
36 interval, and wavelength (Demary et al. 2006). The patterns assist in species-specific mate  
37 recognition. Females typically flash from a stationary position on the ground or in vegetation  
38 while males fly searching for the appropriate response signal. Not all species use  
39 bioluminescence to guide mate selection. Several diurnal firefly species lack an abdominal  
40 lantern and rely on the use of cuticular pheromones to find a mate (South et al. 2008).

41 Sometimes mate selection can end poorly. Female fireflies, especially in the genus  
42 *Photuris*, can mimic the flash pattern of other species to bait fellow fireflies. Males expecting to  
43 mate with a female of the same species risk being eaten by these “femmes fatales” (Eisner et al.  
44 1997). The flash of fireflies can also serve as a response to threats. When distressed, some  
45 species flash erratically to emphasize toxicity or cease flashing to evade detection, while others  
46 induce thanatosis, or produce droplets of hemolymph from their elytra as a defense (Day 2003,  
47 Faust 2017, Eisner et al. 1997).

48           Despite general life history knowledge at the family-level and more in-depth  
49 comprehension of select species, other information, including basic topics like species  
50 distribution, is often unknown. Distributional knowledge tends to get overlooked, which is  
51 problematic. It is necessary to understand the range of a species in order to appropriately protect  
52 populations and establish informative conservation policies. Determining where species are  
53 located also establishes a basis for species-specific research that relies on presence/absence  
54 information, such as how environmental changes impact populations over time. Species-specific  
55 distributions of fireflies are rare. Low-resolution ranges are more common but are of little help  
56 when trying to perform local research. A consolidated lampyrid taxa list for New York does not  
57 exist and there are currently no documented records of firefly species specifically in Otsego  
58 County. This work remediates some of those gaps by using integrative taxonomy to confidently  
59 identify fireflies present on Thayer Farm at the Biological Field Station in Cooperstown, NY,  
60 providing a taxa list for New York fireflies, a local identification key, and new records for New  
61 York and Otsego County.

62

63

## MATERIALS AND METHODS

### *Study Site and Species Collection*

65           Thayer Farm in Otsego County was the study site for this survey. It is part of the SUNY  
66 Oneonta Biological Field Station (BFS) located in Cooperstown, NY (42.73°N, 79.91°W). The  
67 104-hectare property is a part of the of the much larger 607-hectare holdings of the BFS. The  
68 farm is partially old-field and partially operational hay fields. Sampling occurred over ~ 6.9 ha,  
69 which spanned across the main road of the property, several grassy fields, and along the mixed  
70 hardwood forest lines. Lampyrids were haphazardly collected via aerial net on Thayer Farm in

71 the summers of 2017 and 2018. Individuals were caught in the air and in the grass between 2100  
72 and 0130 hours. To preserve the DNA of each sample, lampyrids were frozen at -80°C.

73

#### 74 *Morphological and Molecular Identifications*

75 All individuals caught were morphologically identified according to Luk et al. (2011).  
76 Fireflies of the genus *Photinus* were identified to species. A subset of individuals from each  
77 genus found were used for molecular species identifications. Genomic DNA was extracted from  
78 the prothoracic tissue of each sample selected for molecular identification using the DNeasy  
79 Blood and Tissue Kit (Qiagen, Valencia, CA). The gene Cytochrome Oxidase I (COI) was  
80 amplified in 25 µL reactions containing DNase- and RNase-free water, 1x Hot Master Buffer,  
81 0.5 µM forward and reverse primers (LCO1490 & HCO2198 of Folmer et al. 1994), 0.2 mM  
82 dNTPs, 0.04 units Hot Master Taq polymerase (Quantabio), and approximately 30 ng template.  
83 Hebert's (2003) thermal cycler protocol was followed. Upon successful amplification, the DNA  
84 was purified using AMPure XP Beads (Beckman Coulter Genomics, Brea, CA) and sent to  
85 Eurofins Genomics for sequencing (EG, Louisville, KY). Sequences were visualized using  
86 Chromas Lite (Technelysium Pty Ltd, Australia), trimmed by hand, and examined for allelic  
87 differences. Sequences were then compared to those in GenBank using n-BLAST to provide a  
88 molecular species identification. All sequences were submitted to GenBank (accession numbers  
89 MK635154-MK635340).

90

#### 91 *Sampling Coverage, Taxa Lists, and New Records*

92 Species-accumulation curves were created for genera with 3 or more species to ensure the  
93 maximum number of species were recovered and identified using the “vegan” package in R



94 (Version 1.1.463). Molecular species identifications were randomly sampled to determine how  
95 many individuals should be collected in order to retrieve the maximum number of species  
96 present. A curve for all nocturnal species was created as well. A complete firefly taxon list for  
97 New York was compiled from the data from this study and a review of the literature. Most of the  
98 distributional data from the literature came from Faust (2017), Lloyd (1966, 1969), and Green  
99 (1961). New species records for Otsego County and New York were documented.

100

### 101 *Phylogenetic Analyses*

102 A Bayesian phylogenetic tree was created using MrBayes (v3.2.6, Huelsenbeck and  
103 Ronquist 2001, 2003) after jModelTest (version 2.1.3, Posada and Crandall 1998) was used to  
104 select the most appropriate mutational model (invgamma). The consensus tree was created using  
105 5 chains (4 hot, 1 cold) to evaluate 1,000,000 simulated trees each with every 100<sup>th</sup> tree sampled  
106 and a burn-in value of 2,000 generations. The burn-in value was chosen as a generation beyond  
107 the point at which posterior likelihoods reached stationarity (Ronquist et al. 2005). A species of  
108 net-wing beetles that belong to the family Lycidae was used as an outgroup (Accession number =  
109 MG060334.1), as it was the closest related family to Lampyridae within the subfamily  
110 Elateroidea. The phylogenetic tree viewer FigTree was used to create a tree of the inferred  
111 relationships based on the molecular data (Morariu et al. 2008).

112

113

## RESULTS

### 114 *Morphological and Molecular Identifications*

115 A total of 383 fireflies were caught and morphologically identified to genus (Table 1).  
116 We found 13 firefly species on the Thayer Farm property (Table 4). Most individuals belonged

117 to the genus *Photinus*, and a total of 6 genera were discovered (Table 1). Initial morphological  
118 identifications revealed 7 species of the genus *Photinus*. The number of species identified in that  
119 genus was reduced to 4 species after both molecular and morphological results were examined  
120 (Table 3). The genus *Pyropyga* was not used in molecular identifications, as we were unable to  
121 obtain usable DNA from any of the individuals despite attempting extractions from the  
122 prothorax, head, legs, elytra, head+legs, head+thorax, and whole body.

123

#### 124 *Sampling Coverage, Taxon Lists, and New Records*

125 The species-accumulation curve for the genus *Photuris* plateaued at 4 species after  
126 approximately 12 individuals (Figure 2). The curve for *Photinus* nearly, but did not completely  
127 plateau, after sampling 140 individuals (Figure 1). There was uncertainty in both simulations;  
128 however, the estimated number of species in the genus *Photinus* showed much more ambiguity.  
129 The accumulation curve for the 10 nocturnal species did not plateau after sampling 186  
130 individuals (Figure 3).

131 The review of the literature revealed a total of 30 documented species in New York  
132 (Table 2). The following species are documented here for the first time for Otsego County:  
133 *Photinus ignitus* Fall, 1927, *Photinus macdermotti* Lloyd, 1966, *Photinus carolinus* Green, 1956,  
134 *Photinus obscurellus* LeConte, 1851, *Photuris quadrifulgens* Barber, 1951, *Photuris lucicrescens*  
135 Barber, 1951, *Photuris tremulans* Barber, 1951, *Photuris pennsylvanica* (De Geer, 1774),  
136 *Pyractomena angulata* (Say, 1825), *Pyractomena palustris* Green, 1957, *Lucidota atra* (G.  
137 Oliver, 1790), *Ellychnia corrusca* (Linnaeus, 1767), and *Pyropyga sp(p)* (Harris, 1836), (Say,  
138 1823). *Pyractomena palustris* Green, 1957 had not been previously documented in New York.

139

140 *Phylogenetic Analyses*

141           The consensus Bayesian tree (Figure 5) was separated into two major clades (PP= 100).  
142 One clade included most of the genera: *Photinus*, *Ellychnia*, *Pyractomena*, and *Lucidota*, while  
143 the other contained species of the genus *Photuris*. *Photinus macdermotti* and *Photinus ignitus*  
144 were found to be sister taxa (PP= 100). Relationships between *Lucidota* haplotypes, *Ellychnia*  
145 haplotypes, and *Pyractomena* species were strong (PP= 100), while the weakest relationships  
146 included the group of all *Photuris* species except *Photuris pennsylvanica* (PP=50) and the  
147 grouping of *Pyractomena* with *Photinus* and *Ellychnia* with only 57% of trees supporting that  
148 lineage.

149

150

DISCUSSION

151           We present the first lampyrid species records for Otsego County, for at least 13 different  
152 species. The species *Pyractomena palustris* was also documented as a new species record for the  
153 state of New York, bringing the total number of species for the state to 31. Otsego County and  
154 New York now have firefly species taxa lists, as this information was previously lacking. Since  
155 there was no local species key, the number and identity of species discovered morphologically  
156 was much different than what was found molecularly. *Photinus macdermotti* was not  
157 morphologically identified because it was not present on the *Photinus* species key used, but it  
158 appeared in more than half of the COI species identifications that were “morphologically”  
159 *Photinus ignitus*. The lack of regional species keys and distributional data made it difficult to  
160 understand what species were present and made it nearly impossible to begin other research on  
161 lampyrids.

162           The combination of morphological and molecular taxonomy proved to be the most  
163 appropriate and accurate way to identify species, as we ran into issues using both methods  
164 separately. Visually determining different firefly genera is straightforward, but species  
165 morphology can be confusing, as there are many characteristics that vary only slightly and  
166 required significant practice to discern. If we used morphology alone, we would have  
167 overestimated the number of species present in the genus *Photinus* by 3 species, and entirely  
168 missed another common species (*Photinus macdermotti*). There were issues with our molecular  
169 identifications as well. GenBank was untrustworthy at times and gave us reason to believe that  
170 some sequences previously submitted to the database were incorrectly identified. Several times,  
171 we discovered two species that were 98-99% identical to our sequence, which was within the 2%  
172 sequence divergence used to delineate species boundaries in insects. One way to clarify this  
173 issue was re-examining morphological characters. For example, a few of our sequences were  
174 99% identical to both *Photinus ignitus* and *Photinus indictus*. The species descriptions revealed  
175 one major physical difference: *Photinus ignitus* has a lantern, while *Photinus indictus* does not.  
176 This discrepancy was enough evidence to prove that the individual in question was *Photinus*  
177 *ignitus*, and ultimately made us more skeptical of GenBank identifications. Most of the time,  
178 looking at morphological distinctions solved the problem; however, for species of the genus  
179 *Photuris*, we were not able to use this method because species cannot be delineated based on  
180 morphology.

181           Based on the species-accumulation curves, we appear to have sufficiently sampled  
182 *Photuris* species that are at Thayer Farm. It is likely that we sampled the common species of  
183 *Photinus* in the area, but there is a chance other rare species are present on the property, since the  
184 estimated species-accumulation did not reach a plateau. We were unable to produce plots for the

185 other genera, as we only observed one or two species on Thayer Farm. If we collected more  
186 individuals from these genera, we may have encountered more species; however, it is evident  
187 that these genera are much less common than *Photinus* and *Photuris* fireflies. The nocturnal  
188 species-accumulation curve resulted in a similar curve to *Photinus*; a phenomenon likely driven  
189 by the large number of *Photinus* in our samples.

190 The Bayesian phylogeny provided support for the monophyly of most genera, and  
191 *Photuris* was clearly distinct from the other taxa recovered from the BFS. The weak support for  
192 internal branches in the *Photuris* groupings (PP= 50, 52, 63, 75) was not unexpected, as the  
193 nucleotide diversity between species in this genus were sometimes below the 2% species  
194 boundary.

195 In the remainder of the tree, several polytomies arose, which signified that there was not  
196 enough information present to evolutionarily separate species in many of the hypothesized  
197 groupings. This occurred with several haplotypes of *Photuris* (*P. quadrifulgens* 01, 02, 03, *P.*  
198 *lucicrescens* 01, 02, 04, 07), *Pyrractomena* (*P. angulata* 01, 03, 04) and *Lucidota* (*L. atra* 01, 02,  
199 03). There are different biological reasons evolutionary divergences occur. Actual  
200 morphological differences may define lineages as two different species, or species could diverge  
201 because of strong selection pressures, geographic isolation, or a plethora of genetic diversity seen  
202 within that species. Since the highly conserved gene COI was used, it may fail to detect  
203 evolutionarily recent divergences, even when clear morphological or ecological differences exist.  
204 Integrating both molecular and morphological techniques is one way to resolve this ambiguity;  
205 however, it is likely that morphological traits (e.g., flash patterns) drove the divergence of  
206 lampyrid groups.

207 Our phylogenetic tree strongly supported the paraphyly of *Photinus*, first reported by  
208 Stanger et al. (2007). The diurnal genus *Ellychnia* appeared in the middle of the *Photinus*  
209 species group, suggesting that either one clade of *Photinus* needs assignment to a new genus or  
210 the *Ellychnia* group should be reassigned into *Photinus*. Our phylogeny included multiple  
211 species of *Photinus* that were not present in Stanger et al.'s (2007) study, thus providing stronger  
212 support for their finding. Similarly, corroborating Stanger et al. (2007), it is clear that based on  
213 this phylogeny, the diurnal habit arose multiple times in lampyrids.

214 Integrative taxonomy allowed us to properly identify almost all of the fireflies  
215 encountered in this survey. The species lists compiled for the county and the state should aid  
216 others who work with lampyrids. The distribution of existing firefly species and species that  
217 have yet to be described needs to be understood in order to facilitate and complete other research  
218 on lampyrid populations (e.g., impacts of artificial light, anthropogenic modifications, and  
219 climate change).

220

#### 221 ACKNOWLEDGEMENTS

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308 List of Figures

309 Figure 1. Estimated species-accumulation curve for the genus *Photinus* at the SUNY Oneonta  
310 Biological Field Station, Cooperstown, NY. The COI sequence identifications for each  
311 individual were randomly sampled and are represented by the black curve. The gray polygon  
312 around the curve represents uncertainty based on multiple runs of the model.

313 Figure 2. Estimated species-accumulation curve for the genus *Photuris* at the SUNY Oneonta  
314 Biological Field Station, Cooperstown, NY. The COI sequence identifications for each  
315 individual were randomly sampled and are represented by the black curve. The gray polygon  
316 around the curve represents uncertainty based on multiple runs of the model.

317 Figure 3. Estimated species-accumulation curve for all nocturnal lampyrid species present at the  
318 SUNY Oneonta Biological Field Station, Cooperstown, NY. These species belong to the genera  
319 *Photinus*, *Photuris*, and *Pyractomena*. The COI sequence identifications for each individual  
320 were randomly sampled and are represented by the gray line. The black curve is the estimated  
321 curve based on the molecular data.

322 Figure 4. The phylogenetic relationships of firefly species found at Thayer Farm with appropriate  
323 Bayesian posterior probabilities (based on a consensus of 10,000 trees). *Lycidae spp.* was the  
324 outgroup taxon. The following posterior probabilities are not listed: *Photuris quadrifulgens* 01,  
325 02, 03= 52; *Photuris lucicrescens* 01, 02, 03, 04, 06= 92; *Photuris lucicrescens* 01, 02, 03, 04,  
326 06, 05= 75; *Photuris tremulans* 01, 03= 63; *Photuris tremulans* 01, 03, 02, 04= 96; *Photinus*  
327 *carolinus* 02, 03= 57; *Photinus carolinus* 02, 03, *Photinus obscurellus* 01= 54; *Pyractomena*  
328 *angulata* 01, 03, 04= 85.

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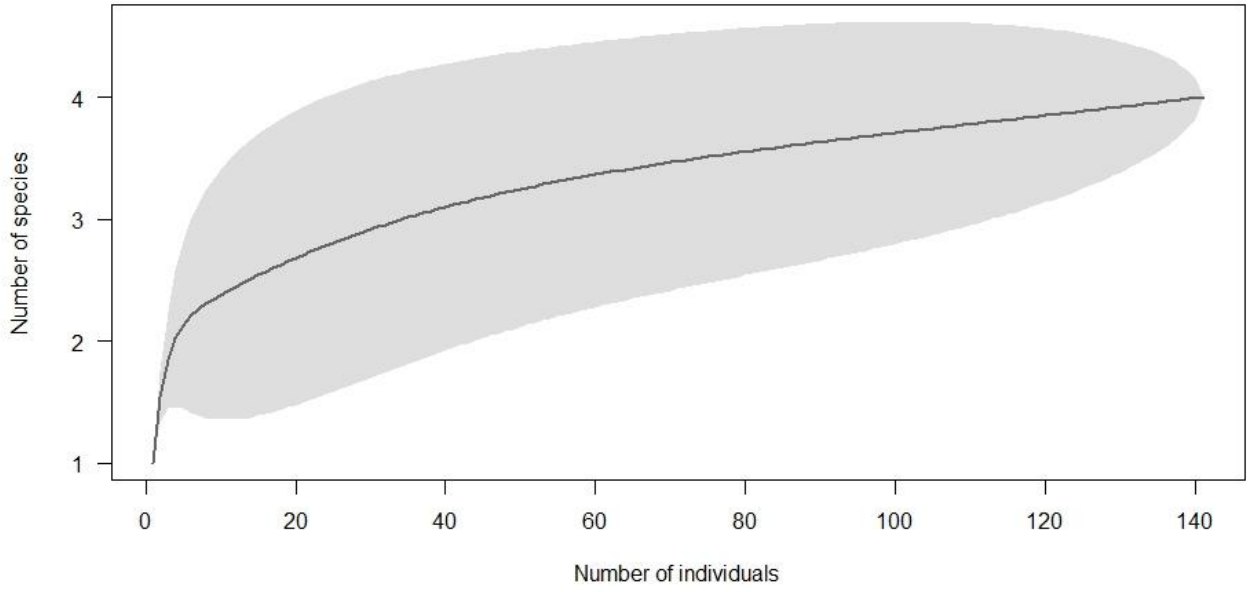
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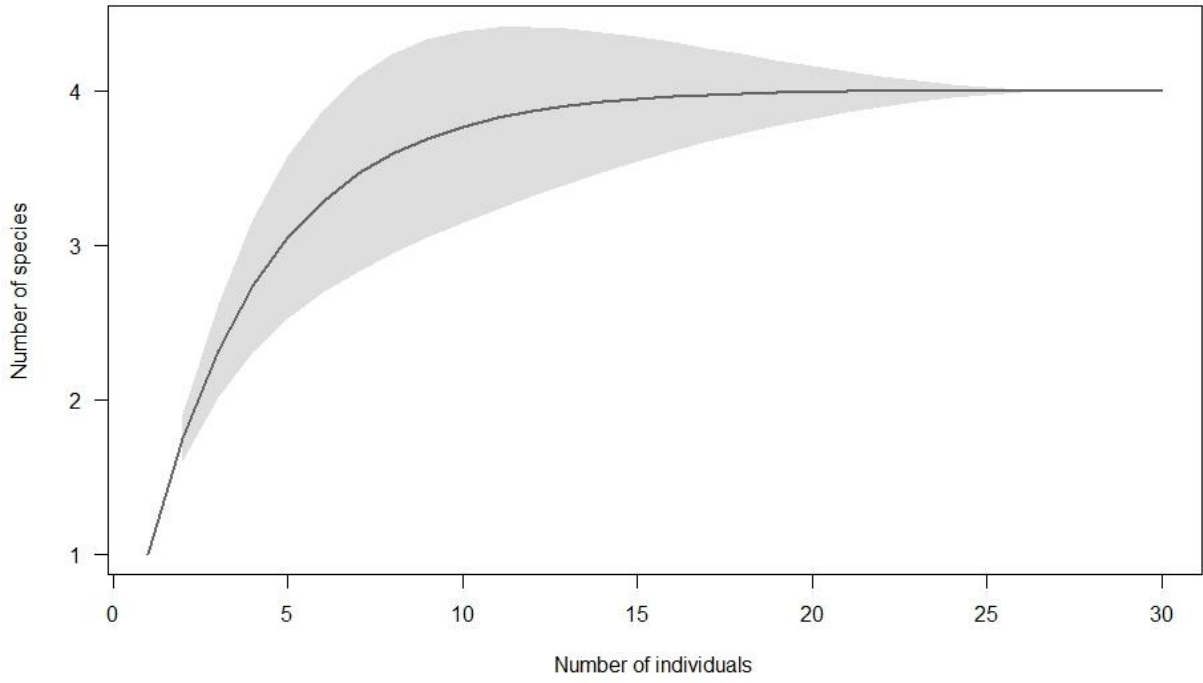
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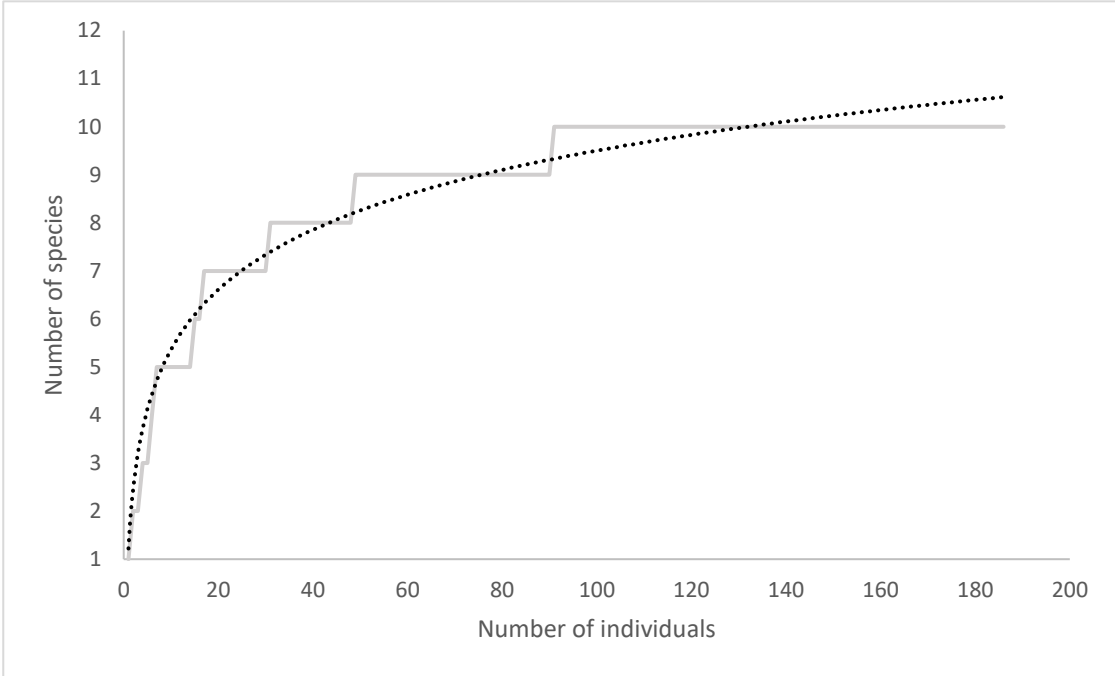
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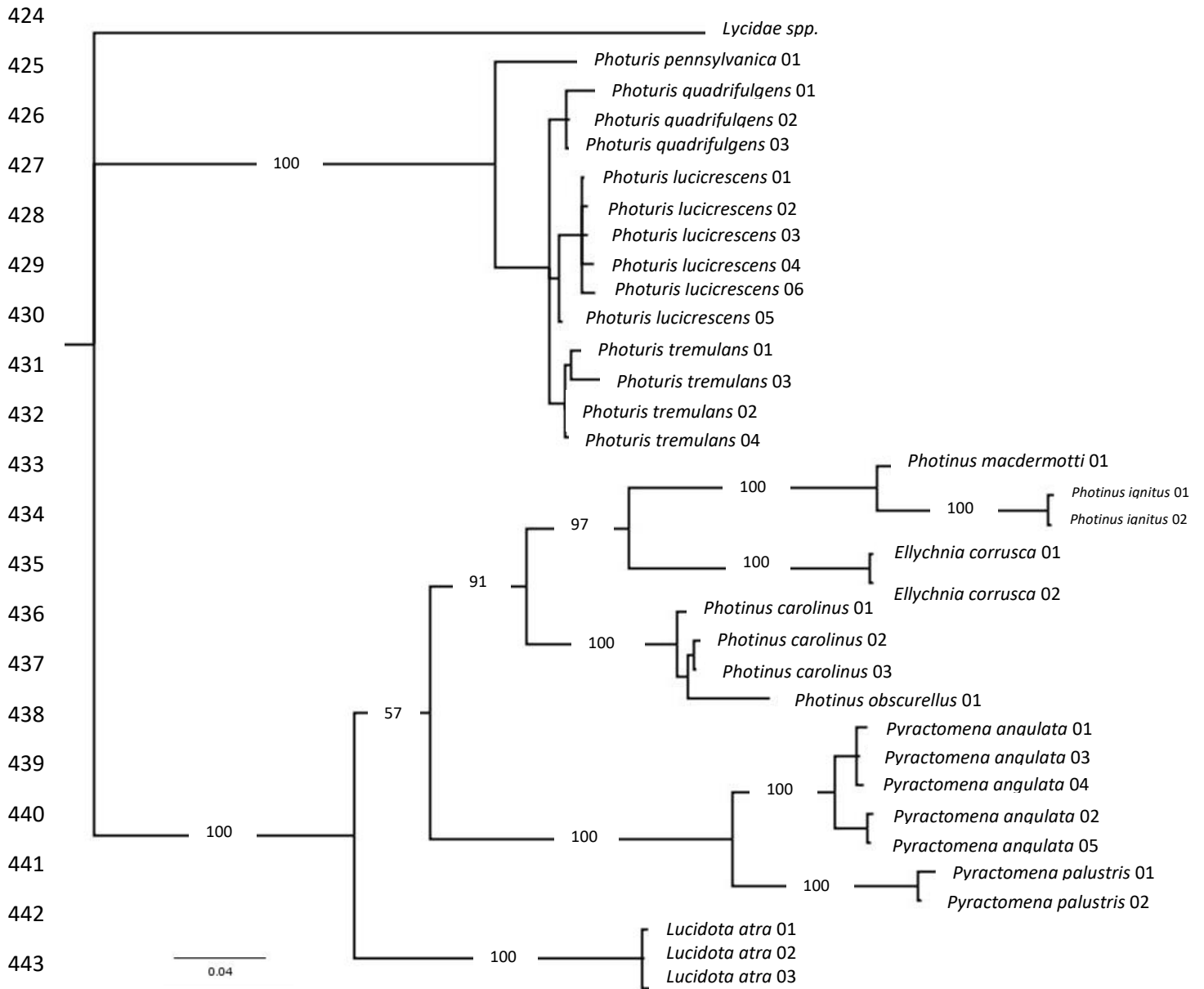


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451 Table 1. Lampyrid genera and number of individuals surveyed at Thayer Farm, Richfield  
452 Springs, NY. Individuals were separated to genus based on morphology. Individuals of the  
453 genus *Pyropyga* were collected in the summer of 2018.

Genus	Individuals
<i>Photinus</i>	205
<i>Photuris</i>	90
<i>Pyractomena</i>	14
<i>Ellychnia</i>	6
<i>Lucidota</i>	4
<i>Pyropyga</i>	64

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465 Table 2. List of firefly species found in New York. The general range of each species is  
 466 documented in relation to New York.

Species	Range	Reference
<i>Ellychnia corrusca</i>	Widespread across eastern U.S.	Faust 2017
<i>Lucidota atra</i>	Widespread across eastern U.S.	Faust 2017
<i>Lucidota punctata</i>	Widespread across eastern U.S.	Faust 2017
<i>Photinus ardens</i>	Oneida, NY	Lloyd 1966
<i>Photinus carolinus</i>	Allegheny region of NY	Faust 2017
<i>Photinus consimilis</i>	Eastern U.S.	Faust 2017
<i>Photinus consanguineus</i> complex ( <i>consanguineus</i> , <i>macdermotti</i> , <i>greeni</i> )	Southern NY	Lloyd 1969
<i>Photinus curtatus</i>	Central NY	Lloyd 1966
<i>Photinus ignitus</i>	Northeastern U.S.	Faust 2017
<i>Photinus indictus</i>	States bordering Great Lakes	Faust 2017
<i>Photinus macdermotti</i>	Eastern U.S.	Faust 2017
<i>Photinus marginellus</i>	Northeast states	Faust 2017
<i>Photinus obscurellus</i>	Northeast states	Faust 2017
<i>Photinus pyralis</i>	Widespread across eastern U.S.	Faust 2017
<i>Photinus sabulosus</i>	Eastern U.S.	Faust 2017
<i>Photinus scintillans</i>	Southeastern NY	Faust 2017
<i>Photuris</i> “Chinese Lanterns”	Southern NY	Faust 2017
<i>Photuris hebes</i>	Eastern U.S.	Faust 2017
<i>Photuris lucicrescens</i>	Eastern U.S.	Faust 2017
<i>Photuris pennsylvanica</i>	Mid-Atlantic states	Faust 2017
<i>Photuris</i> “Primitive Unnamed” -species of <i>versicolor</i> complex	Found where other <i>Photuris</i> are	Faust 2017
<i>Photuris tremulans</i>	Widespread across eastern U.S.	Faust 2017
<i>Photuris quadrifulgens</i>	Widespread across eastern U.S.	Faust 2017
<i>Pyrractomena angulata</i>	Every state east of the Mississippi River	Faust 2017

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<i>Pyractomena borealis</i>	Every state east of the Mississippi River	Faust 2017
<i>Pyractomena linearis</i>	States bordering Great Lakes	Faust 2017
<i>Pyractomena lucifera</i>	Eastern U.S.	Faust 2017
<i>Pyractomena marginalis</i>	Eastern U.S.	Faust 2017
<i>Pyractomena palustris</i>	Otsego County, NY	Current study
<i>Pyropyga decipiens</i>	Northern species, northeastern states	Faust 2017
<i>Pyropyga nigricans</i>	All but southeastern U.S., uncommon east of Mississippi River	Green 1961

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486 Table 3. List of New York lampyrid species that were morphologically and molecularly  
 487 discovered in Otsego County.

Species	Morphological ID	Molecular (N)	Molecular ID(Haplotype)	Accession numbers
<i>Ellychnia corrusca</i>	6	6	3(2)	MK635338-MK635340
<i>Lucidota atra</i>	4	4	4(3)	MK635334-MK635337
<i>Lucidota punctate</i>	0	0	0	
<i>Photinus ardens</i>	2	2	0	
<i>Photinus carolinus</i>	0	0	6(3)	MK635284-MK635289
<i>Photinus consimilis</i>	23	23	0	
<i>Photinus consanguineus</i> complex ( <i>consanguineus</i> , <i>macdermotti</i> , <i>greeni</i> )	12	12	0	
<i>Photinus curtatus</i>	1	1	0	
<i>Photinus ignitus</i>	164	54	63(2)	MK635224-MK635283
<i>Photinus indictus</i>	0	0	0	
<i>Photinus macdermotti</i>	0	0	71(1)	MK635154-MK635223
<i>Photinus marginellus</i>	0	0	0	
<i>Photinus obscurellus</i>	1	1	1(1)	MK635290
<i>Photinus pyralis</i>	1	1	0	
<i>Photinus sabulosus</i>	2	2	0	
<i>Photinus scintillans</i>	0	0	0	
<i>Photuris</i> “Chinese Lanterns”	0	0	0	
<i>Photuris hebes</i>	0	0	0	
<i>Photuris lucicrescens</i>	0	0	6(6)	MK635295 MK635300 MK635302 MK635308 MK635316 MK635320
<i>Photuris pennsylvanica</i>	0	0	9(1)	MK635293 MK635297 MK635304-MK635306 MK635310 MK635312 MK635315 MK635319
<i>Photuris</i> “Primitive Unnamed” -species of <i>versicolor</i> complex	0	0	0	

<i>Photuris tremulans</i>	0	0	10(4)	MK635294 MK635298 MK635299 MK635301 MK635303 MK635307 MK635309 MK635311 MK635313 MK635317
<i>Photuris quadrifulgens</i>	0	0	5(3)	MK635291 MK635292 MK635296 MK635314 MK635318
<i>Pyractomena angulata</i>	0	0	6(5)	MK635321-MK635323 MK635326 MK635329 MK635332
<i>Pyractomena borealis</i>	0	0	0	
<i>Pyractomena linearis</i>	0	0	0	
<i>Pyractomena lucifera</i>	0	0	0	
<i>Pyractomena marginalis</i>	0	0	0	
<i>Pyractomena palustris*</i>	0	0	7(2)	MK635324 MK635325 MK635327 MK635328 MK635330 MK635331 MK635333
<i>Pyropyga decipiens</i>	**	0	0	
<i>Pyropyga nigricans</i>	**	0	0	

488 \*Species not previously documented in New York but was recovered in Otsego County.

489 \*\*Genus morphologically identified but was unable to be molecularly verified to species.

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500 Table 4. List of Lampyridae on Thayer Farm Property; SUNY Oneonta Biological Field Station.  
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<b>Species</b>	<b>Authority</b>
<i>Photinus ignitus</i>	Fall, 1927
<i>Photinus macdermotti</i>	Lloyd, 1966
<i>Photinus carolinus</i>	Green, 1956
<i>Photinus obscurellus</i>	LeConte, 1851
<i>Photuris quadrifulgens</i>	Barber 1951
<i>Photuris lucicrescens</i>	Barber, 1951
<i>Photuris tremulans</i>	Barber, 1951
<i>Photuris pennsylvanica</i>	(De Geer, 1774)
<i>Pyractomena angulata</i>	(Say, 1825)
<i>Pyractomena palustris</i>	Green, 1957
<i>Lucidota atra</i>	(G. Olivier, 1790)
<i>Ellychnia corrusca</i>	(Linnaeus, 1767)
<i>Pyropyga spp.</i>	<i>Pyropyga decipiens</i> (Harris, 1836) <i>Pyropyga nigricans</i> (Say, 1823)

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APPENDIX 1

Species key with morphology and flash patterns for Otsego County

GENERA

- 1. Lantern present; or if absent, yellow stripe on inner margin of wings ..... 2
- 1'. Lantern absent (right) ..... 4



- 2. Large in size (+13 mm) with median stripe on each elytron ..... *Photuris*



- 2'. Small to medium in size (<10 mm) ..... 3

- 3. Dark mark on pronotum is prominent in the center ..... *Photinus*



- 3'. Pronotum center and edges are dark; clear median keel ..... *Pyraclomena*



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4. Tiny in size (5-8 mm) ..... *Pyropyga*



4'. Medium to large (>8 mm) ..... 5

5. Distinctively large and serrated antennae ..... *Lucidota*



5'. Normal antennae; yellow and pink parentheses on dark pronotum ..... *Ellychnia*



SPECIES

*Photinus*

1. Light/pale scutellum; narrow body (~3x longer than wide) ..... *P. ignitus*  
(yellow flash once every 5 seconds)



1'. Dark scutellum ..... 2

575 2. Broad body (~2.4x longer than wide) ..... *P. macdermotti*  
576 (two yellow flashes that are two seconds apart, then 4 seconds of dark)

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582 2'. Different pronotum pattern (typical dark patch down middle with additional  
583 characteristic) ..... 3

584 3. Small dark patches on edge of pronotum ..... *P. carolinus*  
585 (Repeating yellow flash, ~6 every half second, then 6-9 seconds of dark)

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590 3'. Maroon, rather than pink, side patches on pronotum ..... *P. obscurellus*  
591 (3 quick yellow flashes every 5-7 seconds)

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595 *Pyractomena*

596 1. Wide pale margins on elytra ..... *P. angulata*  
597 (orange diagonal sputtering flash every 2-4 seconds)

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605 1'. Pale brown elytra with 2 paler ribs down middle ..... *P. palustris*  
606 (orange crescendo dive every 3 seconds)

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611 *Photuris* (Identification of species requires flash pattern)

- 612 - 2 different flashes; 2 to 5 green flashes every 4 seconds (more common), or half a second  
613 green sparkler flash once every 4 seconds ..... *P. quadrifulgens*
- 614 - Green upward crescendo flash (2 seconds long) every 5 seconds; females aggressively  
615 mimic others ..... *P. lucicrescens*
- 616 - Green, vibrating, 1 second flash every 4 seconds ..... *P. tremulans*
- 617 - Dot-dash flash lasting 1-3 seconds, yellow-green (sometimes dot without the dash, and vice  
618 versa) ..... *P. pennsylvanica*

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620 *Lucidota*

621 Long, flatly serrated antennae ..... *L. atra*

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627 *Ellychnia*

628 Yellow/pink pronotum parentheses; elytra covered in yellowish hairs ..... *E. corrusca*

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635 *Pyropyga*

636 1. Dark border usually present on pronotum ..... *P. nigricans*

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643 1'. Pronotum edges are paler ..... *P. decipiens*

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