



**73<sup>rd</sup> Annual Meeting of the  
Lepidopterists' Society**

**Sierra Vista, Arizona  
July 30<sup>th</sup> to August 3<sup>rd</sup> 2025**

**73<sup>rd</sup> Annual Meeting of the Lepidopterists' Society**  
Sierra Vista, Arizona 🦋 July 30<sup>th</sup> — August 3<sup>rd</sup> 2025

## Hosts

Kelly Richers  
Chuck Harp  
Christopher Grinter  
Todd Gilligan

## Volunteer Assistants

Laura Franzel  
Olivia Lucas and Nathan Orwick  
Christina Nimmo  
Chase Coiner

## Meeting Location

Clarion Inn Sierra Suites Boutique Hotel  
391 East Fry Boulevard, Sierra Vista, AZ, 85635

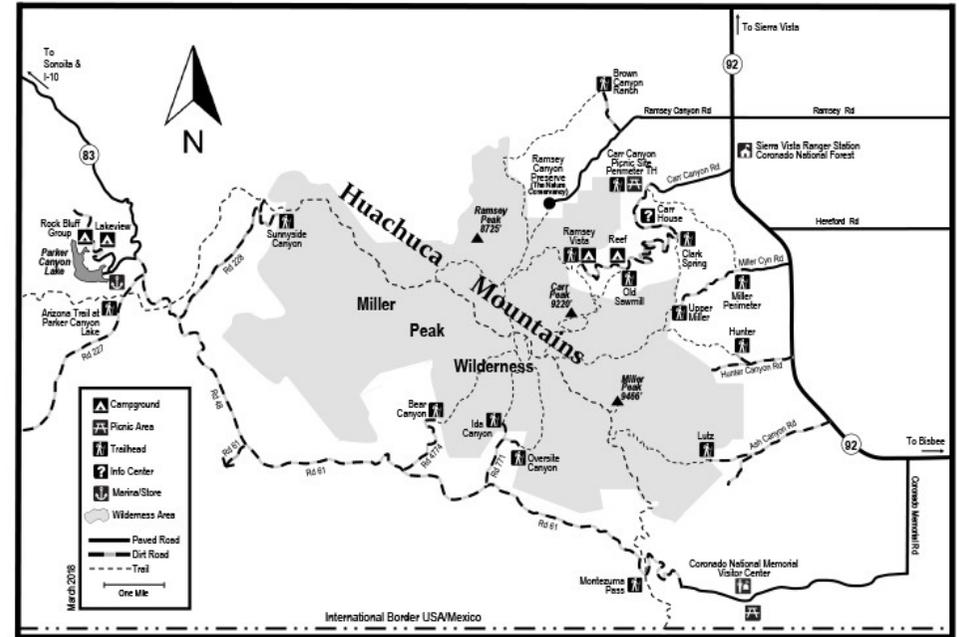
## Barbeque

5:30 p.m. to 8:00 p.m.  
Bam Bam BBQ  
Veterans Memorial Park  
3105 East Fry Boulevard, Sierra Vista, AZ, 85635

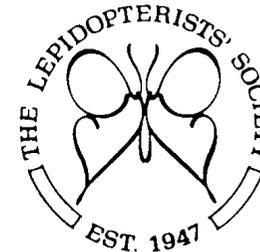
## Banquet

Orient Express Restaurant  
391 East Fry Boulevard, Sierra Vista, AZ, 85635

## Coronado National Forest Sierra Vista Ranger District



Thanks to the California Academy of Science, Colorado State University, and the Lepidopterists' Society for sponsoring and facilitating this meeting!



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ACADEMY OF  
SCIENCES

## Vendors either attending or donating merchandise

JMS Plastics medium density Plastrozote and signage/cards/pricing for a place at our vending tables.

H-H Elements (Harris Haynie) a CalAcad and Cornell drawer with signage/cards/pricing also for the table and raffle.

[millermoth.com](http://millermoth.com) samples of their new LED bucket light traps for a spot and for giveaway.

Here are some GPS coordinates for collecting locations:

N31°24.14' W111°05.35' Pena Blanca

N31°21.69' W110°18.05' Copper Canyon

N31°42.76' W110°52.42' Madera Canyon

N31°47.95' W110°47.95' Box Canyon

N31°22.79' W110°21.82' Bear Creek

N31°23.64' W111°08.55' Sycamore Canyon

N31°27.259' W111°14.822' California Gulch

N31°21.535' W110°17.587' W Montezuma Pass

These are approximate so don't set your GPS and blindly drive or walk to these spots!

## Schedule of Events:

### Tuesday, July 29

10:00am-3:00pm

Wedge Foundation Board meeting

8:00pm-late

Moth field trip

### Wednesday, July 30

9:00am-3:00pm

Executive Council meeting

3:00pm-8:00pm

Registration

5:00pm-7:30pm

Informal Opening Reception

### Thursday, July 31

9:00am-8:00pm

Registration

10:00am-Noon

Talks (Session 1)

Noon-1:30pm

Lunch (on own)

1:30pm-4:00pm

Talks (Session 2)

4:30pm

Dinner (on own)

8:30pm-late

Moth field trip

### Friday, August 1

9:00am-Noon

Registration

10:00am-Noon

Talks (Session 3)

Noon-4:00pm

Field Trip to Carr Canyon

5:00pm

Group Picture at Clarion Sierra Suites

5:30pm-8:00pm

Barbeque at Veterans Memorial Park

### Saturday, August 2

9:00am-noon

Registration

10:00-11:30am

Talks (Session 4)

11:30am-Noon

Student poster judging

5:30pm-9:00pm

Banquet by Orient Express

## **Tuesday, July 29**

10:00am-3:00pm

Wedge Entomological Research  
Foundation Board meeting  
Vernadero Room, Clarion Sierra Suites,  
39 E. Fry Boulevard  
Sierra Vista AZ (520) 459-4221

8:00pm

Moth Collecting Field Trip

## **Wednesday, July 30**

9:00am-3:00pm

Lepidopterists' Society Executive Council meeting  
Vernadero Room, Clarion Sierra Suites,  
39 E. Fry Boulevard  
Sierra Vista, AZ (520) 459-4221

3:00pm-8:00pm

Registration

5:00pm-7:30pm

Informal Opening Reception  
La Casita Mexican Restaurant & Cantina  
465 E. Fry Boulevard  
Sierra Vista, AZ

## **Saturday, August 2, continued**

**11:30 to Noon**

### **Student Poster Viewing, continued**

Chase Coiner

student at Colorado State University and student intern at the C.P. Gillette Museum.

### **A New Mountain Research Center for the Native American Cultural Center at Colorado State University**

A property near Livermore, Colorado will potentially become a new mountain research center for the Native American Cultural Center at Colorado State University. To begin biological research at the site, baseline data for species diversity is needed to document the important players in the ecosystems. The biodiversity of macro-moths was assessed through field collections using light traps during the summer of 2024. Moths collected were preserved and identified to species or morpho-species based on appearance. This poster will help illustrate the means and measures taken to start this project.

**5:30 to 9:00pm**

### **Banquet**

**Presentation of the Mix Family Award  
Presentation of the Karl Jordan Award  
Presidential Address  
Announcement of Next Year's Meeting**

## Saturday, August 2, continued

11:30 to Noon

### Student Poster Viewing

Yair Ben Dor, Linyi Zhang, and John Lill

Address: George Washington University, Department of Biological Sciences, Washington, DC 20052 [yairbendor@gwmail.gwu.edu](mailto:yairbendor@gwmail.gwu.edu)

### Evidence for Host-Associated Differentiation in the Oak Dagger Moth (Noctuidae: *Acronicta*)

We investigated whether the moth *Acronicta increta*, traditionally associated with white oak, is undergoing host-associated differentiation (HAD) following a shift to American beech. HAD was assessed with no-choice feeding experiments measuring larval survival, growth, and behavior on natal vs. novel hosts. Larval coloration was also quantified to assess morphological divergence. Lastly, genomic differences between host races were examined with ddRAD seq. Early instar caterpillars reared on their natal host exhibited higher survival and growth rates than those switched to novel hosts, indicating strong host-specific adaptation. Notably, oak-associated larvae reared on beech experienced complete mortality, whereas the converse exhibited marginal survival. Additionally, larval ground coloration differed significantly by host, potentially reflecting divergence in morphology associated with habitat specialization. Despite these ecological and morphological differences, genomic analyses revealed no genetic differentiation among host races. Therefore, while *A. increta* exhibits features consistent with local adaptation, it remains genetically intermixed across most of the genome, a characteristic of incipient HAD.

## Thursday, July 31

9:00am-8:00pm

Registration

Clarion Sierra Suites,  
39 E. Fry Boulevard  
Sierra Vista AZ (520) 459-4221

9:30 a.m. Vernadero Room

### Introductory Remarks

Richers, Harp, Grinter

### Session 1

(Arizona and New Mexico presentations)

10:00 to 10:30am

Christopher C. Grinter

California Academy of Sciences, 55 Music Concourse Dr  
[cgrinter@calacademy.org](mailto:cgrinter@calacademy.org)

### 14 years of the LepCourse

A look into the past and future of the 10 day field course held at the Southwestern Research Station in Portal, AZ. Founded in 2009, the course has evolved into a popular resource for students, professionals, and naturalists. How can the LepCourse continue to grow and turn into a resource for everyone? What tools, techniques, or topics would you like to see in future courses?

## Thursday, July 31, continued

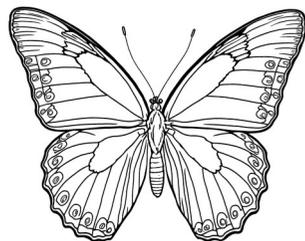
**10:30 to 11:00am**

Jing Zhang, Qian Cong, Jinhui Shen, Leina Song, and Nick V. Grishin

University of Texas Southwestern, 5323 Harry Hines Blvd., Dallas, TX, USA. 75390  
[grishin@chop.swmed.edu](mailto:grishin@chop.swmed.edu)

### **Southeastern Arizona as a hotspot for new butterfly species**

Southeastern Arizona is a hotspot of Lepidoptera diversity. The whole genome analysis provides unprecedented insight into the classification of butterflies and reveals several new species in the region, both residents and strays. From Mark Walker's and Roever's Cloudywing, Arizona streaky-skipper, and Brown-banded Skipper to Elfoid and Faria Metalmark, these examples will be discussed and the steps leading to the discovery of these new species shown. Some of these findings were rooted in previous misidentifications revealed by the analysis of primary type specimens, while other species are split from their known relatives hinted by genetic differentiation and supported by phenotypic considerations.



## Saturday, August 2, continued

**11:00 to 11:30am**

**(Student Presentation)**

Andrea Murillo Vazquez, Ivonne J. Garzón Orduña,  
Jorge Leonel León Cortéz

Colegio de la Frontera Sur, San Cristobal de las Casas, Chiapas; Universidad Nacional Autónoma de México, CDMX, México

[andrea.murillo@posgrado.ecosur.mx](mailto:andrea.murillo@posgrado.ecosur.mx); [ivonne.garzon@ib.unam.mx](mailto:ivonne.garzon@ib.unam.mx); [jleon@ecosur.mx](mailto:jleon@ecosur.mx)

### **An annotated checklist of Ennominae (Lepidoptera: Geometridae) from Chiapas, Mexico**

Ennominae is the most diverse clade of geometrids with over 11,100 named species. The identities and number of Mexican geometrids are unknown, even for regions well sampled for butterflies, like Chiapas.

We present an annotated list of Ennominae of Chiapas, based on records from scientific collections, field work, and online sources. 213 taxa were recorded: 197 species in 98 genera, including 177 new records and 16 taxa that likely represent undescribed lineages. *Pero*, *Opisthoxia*, *Macaria*, and *Hygrochroma* have the most species in Chiapas. Of 12 documented tribes, the most diverse was by far Ennomini, followed by Palyadini, Boarmiini, Macariini, and Odontoperini. Tropical rainforests were dominated by Ennomini and Macariini, dry deciduous forests displayed a balanced representation of Boarmiini and Ennomini. This checklist provides a baseline for future research on Ennominae in Chiapas and is expected to be expanded and refined as taxonomic work in the region progresses.

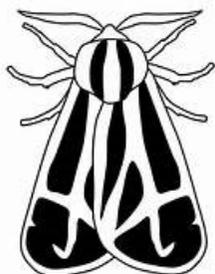
## Saturday, August 2, continued

10:30 to 11:00am

Garzón-Orduña, Ivonne J., Sihvonen, Pasi, Murillo-Ramos, Leidys & Matson, Tanner  
Universidad Nacional Autónoma de México, CDMX, México  
[ivonne.garzon@ib.unam.mx](mailto:ivonne.garzon@ib.unam.mx)

### Phylogenetic relationships among ennomine looper moths (Geometridae: Ennominae): A cladistic total evidence approach

This talk reports on the results of a phylogenetic analysis that concatenates a previously published molecular supermatrix of 10 DNA markers with a novel morphological matrix for the largest subfamily of geometrid moths, the Ennominae under a cladistic approach. At the center of this study is the phylogenetic position of Mexican taxa included in a phylogenetic analysis for the first time. The patterns of character change of various characters traditionally used in loopers' taxonomy are explored to test their value as potential synapomorphies.



## Thursday, July 31, continued

11:00 to 11:20am

Steven J. Cary & Michael E. Toliver  
Cary: 202 Solana Drive, Santa Fe, NM 87501 Toliver: 613 E Center, Eureka, IL 61530  
Cary: [sjcary@outlook.com](mailto:sjcary@outlook.com) Toliver: [tolivermichael1@gmail.com](mailto:tolivermichael1@gmail.com)

### Butterflies of New Mexico: A web-based book

On 2 June, 2020, Steve Cary initiated an on-line book (<https://peecnature.org/butterflies-of-new-mexico/>) detailing information on all butterfly species known from the state of New Mexico, generously hosted by the Pajarito Environmental Education Center. Mike Toliver joined this effort in 2021. Together, they have developed a resource which includes descriptions, range and habitat, life histories, flight periods and additional comments of all species known or suspected from New Mexico, illustrated whenever possible with live photos. The site includes distribution maps and comparison plates for difficult groups. The advantages of an on-line book revolves around the ability to incorporate new information and provide easy accessibility to a wide range of users. Many individuals have contributed photos or information to this effort.



## Thursday, July 31, continued

11:25am to Noon

### (Student Presentation)

Simon Doneski, Quinlyn Baine, David Lightfoot, Esteban Muldavin, Jade McLaughlin, Anna Walker, Richard Norwood

1013 Summit Drive NE, Albuquerque, New Mexico 87106  
[sdoneski@unm.edu](mailto:sdoneski@unm.edu)

### Introducing the New Mexico Rare Arthropods Resource

The New Mexico Rare Arthropods Resource (NM-RARe) is a one stop shop to learn about the rare, threatened, and endemic arthropods that call New Mexico home.

We have been compiling information on these unique, poorly known species, for the last two years so that we can better understand and protect them. There are currently 70 lepidopterans listed with full write ups on [nmrare.org](http://nmrare.org) and we have plans to add many more soon. In this talk I plan to discuss the resource and open discussions about how best to improve and add to the resource in the coming years.

Noon to 1:30pm

Lunch on your own

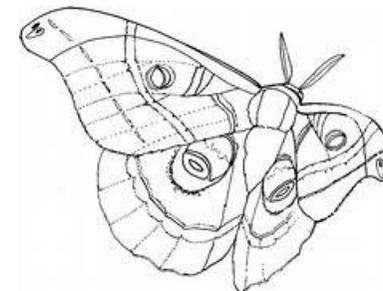
## Saturday, August 2

10:00 to 10:30am

Jeffrey M. Marcus and Nataliia Kopchak  
Dept. Biological Sciences, Univ. Manitoba, Winnipeg MB R3T 2N2  
[jeffrey.marcus@umanitoba.ca](mailto:jeffrey.marcus@umanitoba.ca)

### Above It All? Genetic Variation in the Andean buckeye, *Junonia vestina*, and Lowland Congeners: Is there Gene Flow?

In 1928, Forbes proposed that all New World *Junonia* belonged to a ring species, except for *Junonia vestina*, which occurs at moderate to high elevations in the Andes. The question of the distinctiveness of *J. vestina* has never been evaluated using modern genetic tools. Here we report the results of our survey of mitochondrial haplotypes from throughout the range of *J. vestina* and compare them with haplotypes from lowland *Junonia* species in South America. We conclude that *J. vestina* is genetically distinct from other *Junonia* species, but there is likely also some gene flow between *J. vestina* and other species.



## Friday, August 1, continued

3:30 to 4:00pm

Jing Zhang, Qian Cong, Jinhui Shen, Leina Song, and Nick V. Grishin

University of Texas Southwestern, 5323 Harry Hines Blvd., Dallas, TX, USA. 75390  
[grishin@chop.swmed.edu](mailto:grishin@chop.swmed.edu)

### Surprises from butterfly genomics

While nature is full of deceit, the true colors of Lepidoptera are revealed by their genes. Whole-genome sequencing of the worldwide butterfly fauna yields numerous surprises. From unexpected evolutionary relationships to extreme phenotypic variation and misidentifications, large-scale genomic analysis revolutionizes our understanding of butterflies on all levels. Stemming from nomenclature puzzles solved by DNA sequencing of primary type specimens, some up to 250 years old, we address questions of systematics and speciation. Selected examples to showcase these surprises will be discussed, focusing on American species, including their higher classification along with species delimitation in cryptic species complexes.

5:00pm

**Group Picture at the Hotel**

**Barbeque**

**5:30 p.m. to 8:00 p.m.**

Bam Bam Barbeque, Veterans Memorial Park  
3105 East Fry Blvd., Sierra Vista

## Thursday, July 31, continued

1:30 to 2:00pm

### Session 2

(Western Continent Presentations)

Felix Sperling

Dept. Biological Sciences, University of Alberta, Edmonton, AB Canada  
[felix.sperling@ualberta.ca](mailto:felix.sperling@ualberta.ca)

### Six Decades of Butterfly Surveys at Bragg Creek, Alberta

In 1966 my family moved to a newly cleared farm site on the south slope of Fish Butte, near Bragg Creek, in the Rocky Mountain foothills west of Calgary Alberta. I enthusiastically launched into collecting, recording and observing the wonderful diversity of skippers and butterflies on my doorstep. By 1993 I had published an article that documented 67 species at this location, almost half of the known butterfly fauna for Alberta at that time, and pointed out some of the long-term faunal changes that had occurred during the previous 27 years. I have continued to survey butterflies at that location each year for 60 years, documenting a variety of changes that relate to current issues on the evolution, conservation and taxonomy of Lepidoptera.



## Thursday, July 31, continued

2:00 to 2:30pm

Colin Chiu, Oksana Vernygora, Felix Sperling

Dept. Biological Sciences, University of Alberta, Edmonton, AB Canada

[houwaico@alberta.ca](mailto:houwaico@alberta.ca)

### **Genetic mixing, adaptation and speciation in the spruce budworm species complex (Lepidoptera: Tortricidae: Choristoneura)**

Introgression describes gene flow between previously diverged species, a process which can enhance genetic variation that enables adaptation and even speciation. Such introgression may have shaped the evolution of the spruce budworm species complex (Lepidoptera: Tortricidae: Choristoneura) which is Canada's most important defoliator of conifer trees. Using whole-genome sequences, we are assessing the role of introgression in adaptive divergence, focusing on the origin of the two-year-cycle spruce budworm (*Choristoneura occidentalis biennis*). We are also comparing the genomes of interacting species in the complex to find signatures of adaptive introgression and reproductive barriers using a sliding window approach. In addition to providing better understanding of evolutionary processes, this work aims to inform predictive modeling of these forest pests.

## Friday, August 1, continued

3:00 to 3:30pm

Akito Y. Kawahara

McGuire Center for Lepidoptera and Biodiversity, Florida Museum, University of Florida

[kawahara@flmnh.ufl.edu](mailto:kawahara@flmnh.ufl.edu)

### **20 Years of research and education at the McGuire Center for Lepidoptera and Biodiversity**

The McGuire Center for Lepidoptera and Biodiversity, located at the University of Florida in Gainesville, is a global leader in butterfly and moth research and education. Over the past 20 years, it has expanded to house millions of specimens from around the world, placing it among the largest Lepidoptera collections. The Center supports cutting-edge research in behavior, ecology, conservation, genomics, and systematics. As environmental changes accelerate, understanding the historical and ecological roles of butterflies and moths becomes increasingly urgent. The McGuire Center plays a vital role in uncovering how these insects have adapted over time and in forecasting their future. With world-class expertise and resources, the Center is committed to protecting Lepidoptera biodiversity through research, training, and public engagement. In this talk, I will highlight key milestones from the past two decades and share the McGuire Center's vision for the future.

## Friday, August 1, continued

2:30 to 3:00pm

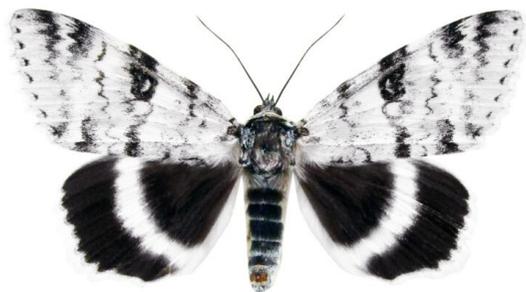
James K. Adams

Dept. of Life Science, Dalton State College, Dalton, GA 30720

[jadams@daltonstate.edu](mailto:jadams@daltonstate.edu)

### Updates on the *Catocala* (Erebidae) of Taylor's Ridge, Walker Co., Georgia

I presented a brief, pre-recorded talk on the *Catocala* of Taylor's Ridge in the Covidyear (2020) at the online 2021 Lep Soc meeting. I have since compiled a more complete picture of the *Catocala* species that have been recorded there (47 species verified, with some others possible). Bob Borth wrote up a discussion of this location for the Southern Lep Soc News, and I thought I would share not only my observations but some of his as well. Come and enjoy!



## Thursday, July 31, continued

2:30 to 3:00pm

Andrew B. Overton and Donald G. Miller

Department of Biological Sciences, CSU Chico

### Post-wildfire recovery of macrolepidopteran communities in the northern California chaparral.

Succession of chaparral plant communities following wild-fire has been well-documented. Lepidoptera can be biological indicators of plant community health based on four criteria: i) their well-defined taxonomy, ii) the ease of surveying them, iii) their wide distribution and host-plant specificity, iv) the concordance of lepidopteran diversity with that of other taxa. Here we evaluated whether changes in moth communities parallel those in plant communities following fire. We used UV traps to sample macrolepidoptera in burned chaparral and two unburned reference sites in northern California. We sampled 2204 moths in eight families in the burned (treatment) area and 3647 in the reference (control) sites. A Bray-Curtis similarity index indicated moderately dissimilar species composition (estimate: 0.614) between treatment and control sites. We recorded significant seasonal changes in species richness and abundance from January-December 2019, and we identified several candidates potentially useful as indicator species in response to fire.

**Friday, August 1**

**Session 3**

(Western presentations and parts East)

**10:00 to 10:30am**

Kelly Richers

139 Liberty Drive, Bolivar, PA 15923  
[kerichers74@gmail.com](mailto:kerichers74@gmail.com)

**Producing a State List of Lepidoptera by County**

Producing a state list of Lepidoptera by county is a process requiring significant research and assistance from collectors, museums and universities. Categories of information will be discussed, based on work done to this point. The presenter has worked on California, Arizona and Nevada, and now has a significant Arizona checklist. I will have the AZ information on a flash drive for interested moth students. There will also be a hard copy of many of the moths on the list with photos of many AZ species for reference available during the meeting.



**Friday, August 1, continued**

**2:00 to 2:30pm**

Paul Masonick [1], Pamela Rivera [2], Jesse Barber [3], and Akito Kawahara

- 1] McGuire Center, Florida Museum of Natural History, University of Florida, Gainesville, FL 32611, USA; [2] Boise State University, Boise, ID 83725, USA; [3] American Museum of Natural History, New York City, NY 10024, USA- Phone: 847-650-2132 (cell) Email: [p.masonick@ufl.edu](mailto:p.masonick@ufl.edu)

**Convergent evolution of antipredatory bioacoustic traits in bombycoid moths**

Moths exhibit remarkable antipredatory traits. Many Saturniidae and Sphingidae employ disparate bioacoustic tactics to thwart bat attacks. Saturniidae have evolved elongated hind wings with twisted tails that lure bat strikes away from vital body parts. Sphingidae disrupt bat attacks by producing ultrasound via genital-based stridulation where modified valvular scales rub against specialized scales, spines, or microsculpturing on the 8th abdominal tergite. Surprisingly, detailed investigations into the bioacoustics, three-dimensional morphology, and microsculpturing of these traits have been lacking. Using ensonification experiments, microtomography, and scanning electron microscopy, we examine the evolution of saturniid tails and their complex morphologies that create decoy echoes to confuse bats. Similarly, we survey the acoustics, diversity, and evolution of stridulatory systems in male Sphingidae. In both groups, we investigate traits in a phylogenetic context and demonstrate their tendencies to coalesce on adaptive optima. Our findings ultimately provide novel insights into the iconic bat-moth arms race.

## Friday, August 1, continued

1:30 to 2:00pm

James K. Adams

Dept. of Life Science, Dalton State College, Dalton, GA 30720

[jadams@daltonstate.edu](mailto:jadams@daltonstate.edu)

### **Dinumma deponens (Erebidae: Scoliopteryginae): the continuing saga of an introduced moth**

*Dinumma deponens*, native to southeast Asia, was first recorded in the U.S. in northern Georgia in 2012. It has since spread throughout much of the range of its introduced foodplant, Silktree (*Albizia julibrissin*). I have presented updates on its progress across the southeastern U.S. at two past Lep Soc meetings, with a lot of the information being collected by citizen scientists and shared on social media. This is the latest update and nearly complete story of its spread and current range in the U.S., and includes a discussion of the possible origins of the U.S. populations.

## Friday, August 1, continued

10:30 to 11:00am

Kyhl Austin, Michael San Jose, Camiel Doorenweerd, Dan Rubinoff

3050 Maile Way, Room 310, Honolulu, HI 96822

[kaaustin@hawaii.edu](mailto:kaaustin@hawaii.edu)

### **Phylogenetics of *Spheterista* (Tortricidae: Tortricinae: Archipini): leaf-rolling, fruit-feeding, stem-boring, and 100 other ways to make a living in Hawaiian rainforests**

*Spheterista* Meyrick is an endemic Hawaiian tortricid genus with 17 described species entirely restricted to the family Araliaceae as host plants. Despite several species being presumed extinct, others can be among the most abundant native moths in Hawaiian wet forests. To better understand relationships in *Spheterista*, we assembled and annotated a 332 Mb reference genome using PacBio HiFi long-read sequencing for an undescribed species from O'ahu. We then sequenced 31 specimens representing 13 described and 8 undescribed species using Illumina whole genome resequencing. Using more than 500 orthologous genes, we reconstructed a robust phylogeny for *Spheterista* using Maximum Likelihood and Bayesian approaches. This allowed us to identify three clades within *Spheterista*: one as leaf-tiers on *Cheirodendron*, one as leaf-tiers on *Polyscias*, and one exhibiting diverse feeding strategies on both plant genera. Niche partitioning in this third clade appears to have facilitated higher rates of speciation than in the other two.

## Friday, August 1, continued

11:00 to 11:30am

Lucas Micheels

750 N. Glebe Road, Arlington, VA 22203  
[lbmicheels@gwmail.gwu.edu](mailto:lbmicheels@gwmail.gwu.edu)

### **One Way, but not Another: Hybridization and Reproductive Isolation in two species of *Euclea* (Lepidoptera: Limacodidae)**

*Euclea delphinii* Boisduval [1832] and *Euclea incisa* Harvey [1876] are two species of North American slug caterpillar moths (Lepidoptera: Limacodidae) that have partially overlapping ranges in Eastern Texas north through Western Missouri. Across this zone there have been many recorded intermediate adult phenotypes, suggesting potential hybridization. To test this hypothesis, I placed pairs of *E. delphinii* and *E. incisa* in mating chambers to record copulation and oviposition. Thus far, I have mated male *E. delphinii* with female *E. incisa* without the assistance of female *E. delphinii* pheromones. All crosses (N=3) produced viable offspring exhibiting a diverse suite of intermediate phenotypes. Curiously, the reverse cross of male *E. incisa* with female *E. delphinii* produced no eggs, suggesting unidirectional interspecific hybridization, a phenomenon indicative of asymmetric reproductive isolation. Analyses of late larval and adult phenotypes are forthcoming and may provide important reference material for interpreting the biogeographic patterns in adult wing patterning.

## Friday, August 1, continued

11:30 to Noon

Chuck Harp

Research Associate II/Collections Manager  
C.P. Gillette Museum of Arthropod Diversity  
Colorado State University, Department of Agricultural Biology, 1177H Campus  
Delivery, Fort Collins, CO 80523

### **The Montana Moth Project: Growing Beyond Our Wildest Dreams**

After completing five seasons collecting moths across the state of Montana, great interest has developed in 2024, with articles in three regional magazines, a segment in *All Things Considered*, an NPR radio broadcast, and a flurry of new volunteer groups helping across the state. The talk will highlight the recent volunteer groups and will feature some of the discoveries from this past season, including two new species, published in 2024.

Noon

Lunch on your own

