Evolution 2017 June 23-27



Portland, Oregon

GENERAL CONFERENCE INFORMATION









The 2017 annual meeting of the <u>Society for the Study of Evolution</u> (SSE), the <u>Society of Systematic Biologists</u> (SSB), and the <u>American Society of Naturalists</u> (ASN).

Jun. 22	Fri, June 23	Sat., June 24	Sun., June 25	Mon., June 26	Tues., June 27	Jun. 28
8:30		8:30		8:00 Undergrad Diversity bkfst social (B110)	i	
JoeFest: a meeting to honor Joe Felsenstein	iEvoBio Pre-conference workshop (B113) ASN Science Communication workshop (B115) Professional Development for Undergraduate Educators workshop (B112) SSB Workshop: Species in the age of discordance (A109) SSE/BEACON Undergrad Diversity: Professional Development Workshop (B111) Joint council meeting (B119)	Regular contributed talks - SSB Mayr symposium (Ball 202) - ASN VP symposium: The causes & consequences of evolutionary conflicts (Ball 204) - SSE Education symposium (C123)	Regular contributed talks - SSB symp: Conceptual issues surrounding genetic-based species delimitation in the genomic era (Ball 202) - Spotlight: Evolutionary mechanisms underlying population-level differentiation (Ball 204)	Joint council exit meeting (C126) Regular contributed talks - Spotlight: New approaches & examples in the iden. of signal, noise & conflict in phylogenetic data (Ball 202) - SSE symp: Evolution in urban ecosystems (Ball 203)	SSB council exit meeting (B110); ASN council exit mtg (C126) Regular contributed talks Spotlight: Genetic transmission at the population level (Ball 203)	9:00
		Regular contributed talks - SSB Mayr symposium (Ball 202) - SSE Hamilton symposium (Ball 203) - ASN VP symposium: The causes and consequences of evolutionary conflicts (Ball 204) - SSE Education symposium (C123)	Regular contributed talks - SSB symp: Conceptual issues surrounding genetic-based spp delimitation in the genomic era (Ball 202) - Spotlight: Speciation with & without sex chromosomes (Ball 203)	Regular contributed talks - SSE symp: Evolution in urban ecosystems (Ball 203) - Spotlight: Eco-evolutionary dynamics (Ball 204)	Regular contributed talks • Spotlight: Sex in the wild (Ball 203)	American Institute of Biological Sciences workshop: Informing and Engaging Decision-makers (B110; lunch 'own your own')
12:10	LUNCH	11:30 SSE student-faculty & postdoc networking lunches	(Ball 201)	NSF Information session (B113)	LUNCH	rkshop:I
		Taylor & Francis publishing wkshp (B113) 13:30	wksp (B113)	SSE Council exit meeting (C126)		'nformir ır own')
JoeFest: a meeting to honor Joe Felsenstein 49 (B113-114)	iEvoBio Pi ASN Science SSB Workshop: Trees, traits and SSB Workshop: Sj SSE Diverse ca Professional Development Council meetings: SSE (Regular contributed talks - SSE Hamilton symposium (Ball 203) - ASN symp: Quant. & conceptual advances in the study of niche breadth (Ball 204) - ASN Education symposium (C123)	Regular contributed talks - SSB symp: Phylogenetic approaches to connecting genotypes to phenotypes (Ball 202) - Spotlight: Sexual selection via male competition & the process of speciation (Ball 203)	Regular contributed talks Spotlight: Impact of gene flow and reticulation in phylogenetics (Ball 202) SSE symp: The impact of stress on genetic variation (Ball 203)	Awards symposium (Ball 203)	3 and Engaging Decision-makers
	iEvoBio Pre-conference workshop (B113) ASN Science Communication workshop (B115) SSB Workshop: Trees, traits and functions: semantics for comparative biology (B114) SSB Workshop: Species in the age of discordance (A109) SSE Diverse careers committee workshop (B116) Professional Development for Undergraduate Educators workshop (B112) Council meetings: SSE (B119); ASN (C126); SSB (B110)	Regular contributed talks SSE Hamilton symposium (Ball 203) ASN symp: Quant. & conceptual advances in the study of niche breadth (Ball 204) ASN Education symposium (C123)	Regular contributed talks regular contributed talks - SSB symp: Phylogenetic approaches to connecting genotypes to phenotypes (Ball 202) - Spotlight: Evolution of hybrid incompatibilities (Ball 203)	Meet the Editors (Ball 201) Regular contributed talks - SSE symp: The impact of stress on genetic variation (Ball 203) - Spotlight: '25 years of sensory drive' (Ball 204)	Awards symposium (Ball 202-204)	
16:50		16:30 ASN mixer	SSB mixer	SSE mixer		
		ASN awards announcements & Pres.	SSB awards announcements & Pres.	SSE awards announcements & Pres. address (Ball 202-		
	Registration (Pre-fun	address (Ball 202-204) 18:30 Poster session 1 & (AU business (Exhibit Hall A)	Poster session 2 & reception (Exhibit Hall A)	Poster session 3 & reception (Exibit Hall A)	18:00	
	19:30 Gould lecture (Ball 202-204) Exhibit Hall A	Free time - dinner & drinks	19:30 Evolution film festival (A107-9) Free time - dinner & drinks	Free time - dinner & drinks	Super social (Oregon Zoo)	

Welcome

We are delighted to welcome you to *Evolution 2017*, the joint annual meeting of the Society for the Study of Evolution, the Society of Systematic Biologists, and the American Society of Naturalists. This year's meeting will be held from June 23-27 at the Oregon Convention Center in Portland, Oregon. Portland is an incredible destination with a convention center in the heart of town that is surrounded by great restaurants and several famous entertainment districts.

The conference is the premier annual opportunity for sharing scientific research in the broad field of evolutionary biology and we anticipate approximately 1,700 attendees. The meeting features multiple workshops and symposia, three new Spotlight sessions, 800+contributed talks, several plenary and outreach talks, and three poster sessions. Product and service providers are also contributing through their sponsorship and exhibits.

Social activities will include an opening reception and SSE Public outreach lecture, mixers during each of three evening poster sessions, and a Super Social with open bar at the newly expanded Oregon Zoo on the final night (no speeches). With the exception of only a small additional charge for the Super Social, all of this is included with your registration.

Thanks for coming and we look forward to seeing you in Portland!

Howard Rundle – Chief Meeting Officer Mitch Cruzan - Assistant Meeting Officer Danielle Wiggins – Conference Manager

Acknowledgements

The meeting has been made possible through the hard work of many people. These include current and past members of the Joint Meeting Committee including Aneil Agrawal, Butch Brodie, Andrea Case, Becky Fuller, Lacey Knowles, Matthew Walsh, and Kelly Zamudio, who served as valuable consultants throughout. We thank Candace Brown, Stephanie Risbon, and Jory Weintraub for helping onsite. Arman Bilge coded our poster invitation and talk timing systems, Emily Behrman and Mohamed Noor organized the recording/posting of talks and archiving of posters, Stephen Christy organized our bike and hiking tours, and Alf Fink who was tasked with picking up and dropping off all the wine we will consume during the conference. Numerous local volunteers also helped in various capacities and Jory Weintraub graciously coordinated them all. To all the organizers of the various symposia, spotlight sessions, workshops, and other special events – a heartfelt thank you for your time and effort!

Table of Contents

Welcome	
Acknowledgements	1
General Information	3
Venues	3
Registration & Information desk	3
Childcare & nursing	4
Practice room	4
Message board	4
Internet access & electronic devices	4
First aid & emergencies	5
Conference policies	5
Oral presentations	6
Speaker Ready Room hours	7
Poster sessions	8
Super Social	8
TriMet's MAX light rail	8
Sponsors & Exhibitors	9
Sponsors	9
Exhibitors	10
Dining Guide – OCC Area	11
Field Trips	12
Registration Fee	13
Travel Support	14
Special Talks	15
Presidential addresses	15
Spotlight sessions	15
Symposia	18
Special Events	21
Pre-conference workshops & satellite events	21
Social events	26
Council, business and other meetings	27
Special events during the meeting	
Post-Conference Workshop	30
Oregon Convention Center floorplan	32

General Information

Venues

The <u>Oregon Convention Center (OCC)</u> is the largest convention center in the Pacific Northwest, making it an ideal venue for conventions, industry trade shows, meetings and banquets. As a <u>LEED – Platinum certified building</u> that is located in beautiful and lively downtown Portland, the OCC is the preferred destination for many groups because of its commitment to customer service and leadership in sustainable practices.

All Evolution 2017 events will be held in the north-east portion of the building (room blocks A-C, Exhibit Hall A, and Ballrooms 201-204), close to the MAX light rail stop outside on Holladay St.

Registration & Information desk

The Registration & Information desk will be located in Pre-function A on Level 1 of the OCC and will be attended during the times listed below. Staff and volunteers will be identified by distinctive blue conference t-shirts; don't hesitate to ask any of them for help. Your conference registration covers attendance at most scientific events over the four days (June 23-27) including concurrent sessions, symposia, spotlight sessions, plenary addresses, and three evening poster sessions. It also includes morning and afternoon coffee breaks (June 24-27), the June 23 evening opening reception, the three evening poster session mixers with light food, and wireless access throughout the OCC. There is a small, additional charge for the conference-ending Super Social and some workshops have separate registration fees. With the exception of the Super Social, no meals are provided. The Oregon Convention Center will have a concession stand in the building on Saturday – Tuesday with coffee, drinks, and lunch available for purchase. See the Dining in Portland section later in this program for information about local restaurants.

In the interest of reducing waste, no conference bag will be provided. Your name badge is your entrance ticket to all of these events so <u>please wear it</u>.

Registration & information desk hours:

Friday, June 23	4:00 pm - 8:00 pm
Saturday, June 24	7:30 am - 5:30 pm
Sunday, June 25	8:30 am - 4:30 pm
Monday, June 26	8:30 am - 4:30 pm
Tuesday, June 27	8:30 am - 3:00 pm

Childcare & nursing

Onsite childcare services for children ages 6 months – 12 years are located in Rm A103/104 on Level 1 of the OCC. Bookings had to be made prior



to June 23, 2017 (i.e. no unscheduled drop-ins). Childcare services are provided by <u>Little Vikings Child Care</u>, a professional organization on the campus of their sponsor, Portland State University, which normally serves PSU students, staff, and faculty. Their goal is to provide your children with a comfortable, safe, and happy experience. Snacks and beverages are provided but parents lunch is not (parents/guardians must pick up their child for lunch each day). There will be a fridge to store food/drinks but no way to heat them. Little Vikings enforces a strict NUT FREE policy. All arrangements and payments were handled directly by Little Vikings.

A quiet, private room with comfortable seating and a mini-fridge is available for nursing mothers in Rm A102 on Level 1, right near the childcare rooms.

Practice room

Between 8 am – 5 pm daily (Sat-Tues), B110 will often be available as a quiet room in which to practice your talk. It will be configured with boardroom-style seating and will have a laptop, projector and screen, but you will have to bring your presentation on a USB key (uploaded presentations will not be accessible in this room). The room is occasionally reserved for special meetings so please check the signage outside prior to entering and if it is busy, try again later.

Message board

A message board will be located near the Registration & Information desk in Pre-function A on Level 1 of the OCC. Registrants are welcome to post notices about events, jobs, announcements, and messages for other attendees.

Internet access & electronic devices

Wireless access is provided to participants free of charge throughout the OCC. No password is required.

Please remember to turn off or mute all electronic devices during all concurrent sessions.

First aid & emergencies

A pharmacy (Lloyd Center Pharmacy, 438 E Burnside St, Portland, OR 97214) is a 9 min. walk away from the Oregon Convention Center.

Call 911 immediately for life threatening situations. Life threatening situations include an apparent heart attack, stroke, serious accident or other major injury.

For non-life threatening first aid occurrences, please locate a staff member wearing a blue conference t-shirt and they will contact the Security, the on-site Medical Technician and/or other Agent on duty and the Event Manager. Non-threatening occurrences include minor cuts or abrasions, minor illnesses, etc.

Conference policies

Code of Conduct for Evolution Meetings

The Evolution Meetings are intended to foster the exchange of scientific ideas, providing participants with an opportunity to present research findings, establish/renew collaborations, recruit people to their laboratories, and learn, teach, and network with an international community of evolutionary biologists. The Society for the Study of Evolution (SSE), the American Society of Naturalists (ASN), and the Society for the Study of Systematic Biology (SSB) are committed to creating an environment where everyone can participate without harassment, discrimination, or violence of any kind. All meeting participants must be treated with respect and consideration. Registration for the meeting is considered an agreement to abide by this code of conduct.

Harassment of any participant (attendee, speaker, volunteer, exhibitor, staff member, service provider, organizer, or meeting guest) will not be tolerated. Unacceptable behavior includes (but is not limited to) unwanted verbal attention, unwanted touching, intimidation, stalking, shaming, or bullying. Discrimination on the basis of gender or gender identity, sexual orientation, age, disability, physical appearance, race, religion, national origin, or ethnicity will not be tolerated. Harassment presented in a joking manner is still harassment and constitutes unacceptable behavior. Retaliation for reporting harassment is also unacceptable, as is reporting an incident in bad faith.

People wishing to report a violation of this code of conduct should immediately contact a member of the <u>Joint Meeting Committee</u> or any member of the executive council of the <u>SSE</u>, <u>ASN</u>, or <u>SSB</u>. Incidents of harassment and discrimination are taken extremely seriously. Confidentiality will be maintained unless disclosure is legally required.

The meeting organizers, members of the JMC, and Society executive officers reserve the right to enforce this code of conduct in any manner deemed appropriate. Anyone violating the code of conduct may be: (a) asked to stop, (b) expelled from the meeting (without refund), and/or (c) prohibited from attending future meetings. Establishing this code of conduct is intended to prevent incidents of harassment, discrimination, and violence, and to maintain the high quality of scientific discourse that our members have come to expect from the Evolution meetings.

Policy on liability

The Societies shall not be responsible for any defamatory, offensive, or illegal conduct of all Evolution Meeting participants, and shall not be held liable for personal injury, property damage, theft or damage of any kind suffered by the participants at or in connection with the Evolution meeting. By registering for and attending the annual Evolution Meeting, each participant acknowledges that they have read this Disclaimer, and expressly releases the Societies and its board members, directors, officers, employees, or agents from any and all liability in connection with such Meeting as provided herein.

Weapons policy

For the safety of all attendees, the Evolution Meetings are 'weapons free' and all conference participants, including staff, volunteers, and attendees, are banned from possessing any object or substance intended to cause injury to others, including but not limited to firearms.

Broadcasting policy

The Evolution Meetings support the communication and discussion of science. Information presented at the Meeting (in oral or poster format) may be reported and discussed by attendees and science writers via blogs, Twitter, or other formats. However, we require that this be done respectfully and without direct reproduction of visual materials (e.g., no posting photos of slides or posters) unless permission is obtained from the presenter or they have already made this information freely available in an open-source forum. If a presenter does not want information from his/her presentation to be broadcast, they should make this clear in their talk/poster and we ask that attendees respect this.

If you have questions or concerns about this policy, or would like to report an abuse of it, please contact any member of the Joint Meeting Committee.

Policy on multiple presentations

The Evolution Meetings have grown dramatically of late, increasing the demand for both oral and poster presentations. To reduce competition and give more people an opportunity to present, attendees are limited to presenting one talk, no matter what type (i.e. contributed or invited of any length) and/or one poster*. Please note that this only applies to the presenting author; you can be a co-author on multiple presentations given by others. *There is an exception for those giving a 5 min. talk in a Spotlight Session.

Oral presentations

Regular contributed talks are 14 min in duration INCLUDING QUESTIONS, leaving 1 min setup/movement time between them. This will be rigorously enforced by an automatic timing system that will sound a chime at 12 min (indicating two 2 min left), then at 14 min to denote the end of the talk and the start of 1 min of movement time, then finally at 15 min to denote the start of the next talk. Note that the 14 min includes time for questions, so we suggest 12-13 min for the talk and 1-2 min for questions. Special talks of other duration (e.g. '30' min. symposium talks) will likewise include 1 min of movement time, meaning a

30 min talk is actually 29 min including questions, and the warning chime will similarly sound at 2 min remaining, at the end of the talk (start of 1 min movement time), and at the start of the next talk. The chime system is controlled by the internet browser on the presentation laptop. DO NOT EXIT THE BROWSER.

The 5 min 'lightning' talks during the Spotlight sessions receive the full 5 min, so the next speaker has to be 'waiting in the wings' ready to begin as soon as the previous speaker is done. In this case a warning chime will sound at 4 min indicating one min left, and then a second chime will sound at 5 min denoting the end of that talk and the start of the next.

Should the automated timing system fail, or if there is any other AV-related or other problem in the room, click the Call for Help button in the presentation software and a technician will be immediately sent to your room. There will be a clock radio in each room and should the chimes fail, designate an audience member as a Session Chair and have them enforce the above timing using this clock.

The AV system is PC-based and will support both PowerPoint (all versions) and PDF files (unfortunately Keynote and Prezi are NOT supported). **Presentations should be formatted using a standard (4:3) slide size, NOT widescreen (16:9).** If your presentation was prepared on a Mac or other non-Windows based system, it is STRONGLY recommended that you review it in the Speaker Ready Room prior to giving your talk (see below). Additional details about audio and video support can be found here.

Talks can be uploaded online before the conference, or at the conference via the Speaker Ready Room (Rm C125 on Level 1 of the OCC) at the conference. **This must be done a minimum of 2 h before your presentation**. Instructions on how to upload presentations online will be emailed to every speaker ahead of the meeting. You can also review your talk in the Speaker Ready Room and AV technicians will be present to help resolve any compatibility or formatting issues and can also explain the in-room setup. The computers in the Speaker Ready Room will be configured with hardware and software exactly like the ones in the meeting room.

Speaker Ready Room hours

Hours for the Speaker Ready Room (Rm C125 in OCC) are as follows:

Friday, June 23 3:00 pm – 8:00 pm

Saturday - Monday, June 24-26 8:00 am - 5:00 pm

Tuesday, June 27 8:30 am – noon

Speaker Ready Room and Presentation Software are sponsored in part by Oregon State University.



Poster sessions

Posters are assigned to one of the three poster sessions that will be held in conjunction with social mixers that include light food, beer, wine and non-alcoholic beverages. Presenters should attend their poster during their assigned session (see the daily schedule here). The location (i.e. poster board) on which you display your poster can be found in the online program. Note that it is NOT the ID number you received when submitted your poster. All poster sessions will be held in Exhibit Hall A (EH-A).

Poster session #1: Saturday, June 24, 6:30 - 8:30 pm

Posters should be displayed by noon on Saturday and removed by noon on Sunday.

Poster session #2: Sunday, June 25, 6:30 - 8:30 pm

Posters should be displayed by 1:30 pm on Sunday and removed by noon on Monday.

Poster session #3: Monday, June 26, 6:30 - 8:30 pm

Posters should be displayed by 1:30 pm on Monday and removed by 10 am on Tuesday.

Posters can be a maximum of 1.2 m x 1.2 m (4 ft x 4 ft) and pushpins will be provided. Poster boards will be numbered and presenters should use the space assigned to them in the online program (note that is NOT the poster ID number you were emailed at the time of submission). If the previous poster is still displayed when you arrive to put up your own, please carefully remove it, roll it up, and place it next to your poster board. We will collect unclaimed posters prior to the next poster session and relocate them to a side table.

Super Social

The conference-ending Super Social will be held in the indoor/outdoor Oregon Zoo starting at 6:30 pm on Tuesday, June 27. The zoo is located at 4001 SW Canyon Rd. and is accessible via the MAX light rail pass. Advance ticket purchase is required for the Super Social. Tickets will not be available for purchase at the registration desk.

Participants are responsible for all tickets. Lost tickets will not be replaced.

TriMet's MAX light rail

All registered attendees will receive a complimentary MAX light rail pass at the Evolution registration desk located in Pre-function A at the OCC. The pass will be valid through the dates of the conference. You will only need to purchase a pass should you wish to ride the light rail prior to registering on Friday afternoon (e.g., to travel from the airport into Portland when you arrive).

MAX light rail, buses, and streetcars are generally within a few blocks of any place you might want to go in the Portland Metro area. In addition to downtown area museums, sights, and shopping, a number of nature parks are easily accessible on MAX light rail.

Sponsors & Exhibitors

Sponsors

We are grateful to our sponsors whose financial contributions have helped keep registration costs more affordable. Each sponsor has generously selected to sponsor the items or events listed above the logo below.

Gold

Name Badge

ecology & evolution

Presenation Management



Silver

Daycare



Bronze

Opening Reception



Society Social Mixers

YXLON

X-ray & Computed Tomography

Saturday Morning Break



Saturday Afternoon Break Sponsor



Sunday Morning Break Sponsor



Sunday Afternoon Break Sponsor

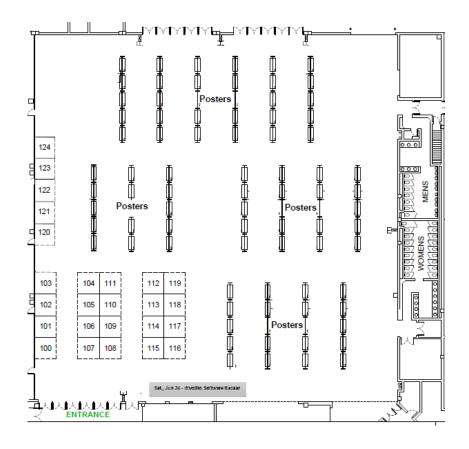


Exhibitors

A wide range of exhibitors have booths in Exhibit Hall A. We encourage you to visit them and learn about the publications, products, and services they have to share. Exhibitor hours are:

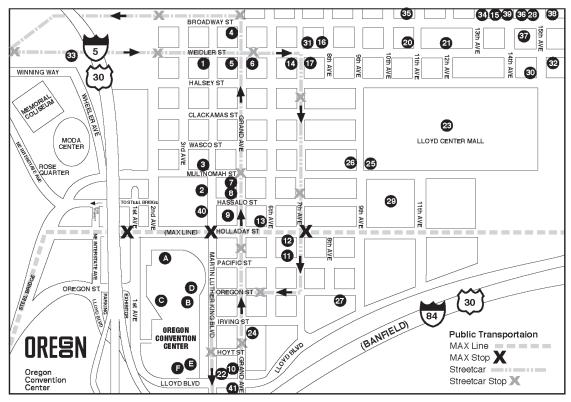
Friday, June 23 7:00 pm - 9:30 pmSaturday - Monday, June 24-26 9:45 am - 8:30 pmTuesday, June 27 9:00 am - 12 noon

Exhibitor	Booth	Exhibitors	Booth
Oxford University Press	100, 101	FSU Center for Anchored Phylogenomics	112
Polymorphic Games	102	National Science Foundation	113
Springer Nature	103	University of Chicago Press	114, 115
RTLGenomics	104	W. W. Norton	116
Penguin Random House	105	British Ecological Society	117
Sinauer Associates, Inc., Publishers	106	Taylor & Francis Group	118, 119
Princeton University Press	107	SSE GSAC	120
YXLON	108	SSE Diversity Careers Committee	120
University of Southern Maine	109	Harvard University Press	121
SimBio	110	BEACON Center for the Study of Evolution in Action	122
Wiley	111	Reconyx Inc.	123
		Macmillian Learning	124



Dining Guide - OCC Area

The Oregon Convention Center will have a concession stand in the building on Sat-Tues with coffee, drinks, and lunch available for purchase. See below for information on local restaurants near the OCC, or here for an online guide.



Dining Guide - Oregon Convention Center Area

- A **Orbit Cafe** (open depending on building activity)
 1st Floor OR CONVENTION CENTER, Deli Food
- B Dragon Cafe (open depending on building activity) 1st Floor OR CONVENTION CENTER, Deli Food
- C **Skyview Terrace** (open depending on building activity) 4th Floor OR CONVENTION CENTER, Restaurant/Bar
- D **Portland Roasting Coffee I** (open depending on building activity) 1st Floor OR CONVENTION CENTER
- E **Cucina Rosso** (open depending on building activity) 1st Floor OR CONVENTION CENTER
- F Portland Roasting Coffee II (open depending on building activity) 1st Floor OR CONVENTION CENTER
- 1 Starbucks 1503 NE MLK Jr Blvd
- Burgerville 1135 NE MLK Jr Blvd Fast Food, Sit-down, take out
- 3 **Subway** 1211 NE MLK Jr Blvd Sit-down, take out
- 4 Starbucks 1621 NE Grand Ave
- 5 Wendy's 1421 NE Grand Fast Food
- 6 McDonald's 1520 NE Grand Fast Food
- 7 **Denny's** 425 NE Hassalo St Sit-down restaurant
- 8 **Red Robin** 422 NE Multnomah St Sit-down restaurant
- 9 Hotel Eastlund 1021 NE Grand Ave Altabira City Tavern- Sit-down (6th floor) Citizen Bakery (Lobby level)

- 10 Starbucks 525 NE Grand Ave Variety of coffees, pastries
 - Cafe Today 650 NE Holladay St
- Sit-down/take-out restaurant

 City Coffee 650 NE Holladay

 Variety of coffees, pastries
- 13 **J Cafe** 533 NE Holiday Sit-down cafe
- 14 **Burger King** 632 NE Weidler Fast food
- 15 **Yuki Sushi** 1337 NE Broadway Sit-down/take-out
- Taco Bell 725 NE Weidler Opening Soon
 Chipotle Mexican Grill 7th & NE Weidler
- Sit-down restaurant
- 20 **Village Inn** 11th & Weidler Sit-down restaurant
- 21 Prime Rib & Chocolate Cake
- 12th & NE Broadway Sit-down restaurant
 Spirit of '77 500 NE MLK Jr. Blvd.
- 23 Food Court 3rd Floor -Lloyd Center Mall
- 24 **Table 6**622 NE Grand Ave 7:30 am 2 pm
 Sit-down/take-out
- 25 Stanfords Lloyd Center Mall, SW Corner, Sit-down dining
- 26 **Lloyd Tower** 825 NE Multnomah 2nd floor cafeteria & deli
- 27 Oregon Building 800 NE Oregon Cafeteria

- Petisco 1411 NE Broadway Sit-down
- **DoubleTree Lloyd Center** Entry on 11th St. Multiple restaurants, sit-down dining
- 30 Applebee's 15th & NE Halsey Sit-down restaurant

29

- 31 Muchos Grasias 707 NE Weidler Sit-down
 - Koji 15th & NE Weidler Japanese Fare - Sit-down restaurant
- Mucho Grande 10 NE Weidler
 Sit-down restaurant
 Milos City Cafe 1325 NE Broadway
- Sit-down restaurant restaurant (open event days only)
- 35 Chens Dynasty II 1025 NE Broadway Hunan & Szechuan, sit-down dining, takeout, delivery
- 36 Chai Yo Thai 1411 NE Broadway Sit-down
- 37 Pastini Pastaria 1426 NE Broadway Sit-down
- McMenamin's Broadway Pub 1504 NE Broadway - Sit-down
- 39 Namaste Indian Cuisine 1403 NE Weidler - Sit-down 40 Starbucks - 1035 NE MLK Blvd
- 41 Dutch Brothers Coffee 430 NE Lloyd Blvd

rev. 2/10/16

Field Trips

Daily Lunchtime Bike Tours

Need to get outside? Come explore Portland on a bike tour! Portland ranks near the top of the list of bicycle friendly cities, so it is relatively easy to get around the city by bike. Biking is a great way to see a little bit of Portland while getting some exercise! June 24th, 25th, and 26th (Sat - Mon) 11:35 am - 1:30 pm. Sign up here or at the conference registration table. Renting a bike is easy (see our Getting Around Portland page) and all skill levels are welcome. We'll do a shorter 8-mile ride on Saturday, and longer 12-mile rides on Sunday and Monday. Bring a brown bag or cash for lunch; we'll stop at Flying Elephants on our way down to Oaks Bottom Wildlife Refuge. We will follow a relatively car-free route, with great views of the city, and an amateur history lesson along the way! Email Stephen if you have any questions: schristy@pdx.edu.

Cost: Bike rental, food

Time and location: Meet at 11:35 am at the orange BIKETOWN bike racks near the northeast

corner of the Oregon Convention Center grounds (NE Holladay & MLK).

Bring: Bike, water, and brown bag lunch or cash for food.

Pre-Super Social Hiking Tour

A trip to Oregon just isn't complete without a hike! Luckily, there are great trails within the city limits. A serene and memorizing maze of trails zigzags though the <u>Hoyt Arboretum</u>, a roughly 190-acre tree sanctuary nestled just above downtown. We highly recommend you check it out! The trails are well maintained—but hilly—and a world apart from the hustle and bustle of downtown. Come join us for a tour of the park right before the Super Social on Tuesday, June 27. Sign up here or at the conference registration table. Email Stephen if you have any questions: schristy@pdx.edu.

Trees of interest: Redwood, sequoia, larch, madrone, magnolias, California bay, firs, pines, oaks, cottonwoods, and many more!

Cost: Free, though the park always appreciates donations no matter how small.

Time and location: Meet at 4:30 pm outside the Washington Park MAX Station (take the elevators to the top of the deepest subway station in the U.S.). We will finish at the Oregon Zoo for the Super Social.

Distance: approximately 2 miles of rolling hills, but individuals/groups are welcome to split off at any time.

Bring: Water, shoes that can get muddy, and snacks.

We'll have: Knowledge of all the best routes and trees, and a car trunk that you can store non-valuable things in during the hike (space-dependent).

Registration Fees

Fees cover attendance at all scientific events over the four days (June 24-27) including concurrent sessions, symposia, spotlight sessions, plenary addresses, outreach talk(s), and three evening poster sessions with light food. They also include morning and afternoon coffee breaks (June 24-27) and the June 23 opening reception. There is a small additional charge for the conference-ending Super Social (which includes dinner and a host bar).

REGISTRATION	Early (by Apr 15)	Late (Apr 16-June 22)	On-site
Professional (member)	\$465	\$565	\$615
Professional (non-member)	\$565	\$665	\$715
Postdoctoral (member)	\$390	\$490	\$540
Postdoctoral (non-member)	\$490	\$590	\$640
Graduate student (member)	\$315	\$415	\$465
Graduate student (non-member)	\$415	\$515	\$565
Undergraduate	\$300	\$400	\$450
Guest*	\$150	\$200	\$250

^{*}A guest is a person accompanying someone with a standard (i.e. non-guest) registration and who wishes to attend the social functions only (i.e. opening reception, coffee breaks, evening poster sessions). Anyone wanting to attend talks, symposia, spotlight sessions, or other academic events must independently register as a regular attendee.

SUPER SOCIAL	Early (by Apr 15)	Late (Apr 16-June 22)	On-site
Professional/faculty	\$25	\$45	Not available
Postdoctoral / student	\$15	\$35	Not available
Registered guest	\$25	\$45	Not available
Accompanying person not registered as a guest	\$70	\$90	Not available

Travel Support

Financial support to attend the meeting is available from various sources, as detailed below. Travel grants funds are normally provided at, or after, the conference.

Volunteer at the conference

In return for volunteering for approximately two 4-hour shifts during the meeting, graduate students can receive a rebate of their early registration fee. Applications are made during main conference registration and will be accepted until the end of early registration (i.e. midnight on Apr. 15). Recipients will be randomly selected from the list of applicants and will be informed by May 15th at the latest. Registration fees must be paid in advance and the rebate will be processed after the meeting, subject to satisfactory completion of assigned volunteer duties.

ASN Graduate Student travel grants

Graduate student members of the ASN can request to be considered for a \$500 US travel award. To be eligible, the student must present a talk or poster and must not have received the travel award in the previous year. Applications are made during main conference registration and will be accepted until the end of early registration (i.e. midnight on Apr. 15). Recipients will be randomly selected from the list of applicants and will be informed by the ASN treasurer by email.

SSE/BEACON Undergraduate Diversity at Evolution travel award

This travel award brings talented and diverse undergraduates (domestic and international) to the meeting to present a poster, receive mentoring and participate in a career-oriented 'Undergraduate Futures in Evolutionary Biology' panel and discussion. The program covers travel, registration, and room and board. For more information and to submit an application visit the organizer's website. Applications will be accepted until 11:59 PM (EST) on March 31, 2017 and awardees will be notified by April 6.

SSE Graduate Student and Postdoc travel supplements

Graduate student and postdoc members of the SSE can request to be considered for a \$500 US travel award. To be eligible, they must present a talk or poster and must provide a written explanation of why they need the funds (150 words maximum). They must also not have received the travel award in the previous year, and postdocs must be fewer than 6 years post-PhD without a permanent position. Applications are made during main conference registration and will be accepted until the end of early registration (i.e. midnight on Apr. 15). Applicants will be informed of the outcome by May 15th at the latest.

SSE International travel stipends

These stipends support attendance at the conference for scientists at various stages of their professional career (e.g., Masters/PhD students, postdocs, and lecturers). You must be a member of the SSE. Those working in countries with high GDP are not eligible. For additional information and the application procedure, consult the SSE's website. Deadline for application is March 31, 2017.

ESEB International travel stipends

These stipends support attendance at the conference for scientists at various stages of their professional career (e.g., Masters/PhD students, postdocs, and lecturers). You must be a member of ESEB. NOTE THAT THESE STIPENDS ARE GIVEN IN CONJUNCTION WITH ANALOGOUS INTERNATIONAL TRAVEL STIPENDS OFFERED BY THE SSE ABOVE, SO THERE IS NO NEED TO APPLY FOR BOTH. Additional details can be found here. Deadline for application is March 31, 2017.

Special Talks

Presidential addresses

ASN Presidential address & award announcements
 Saturday, June 24 from 5:30 - 6:30 pm, Ballroom 202-204 (Level 2)
 Kathleen Donohue (Duke U.), ASN Presidential Address

SSB Presidential address & award announcements
 Sunday, June 25 from 5:30 - 6:30 pm, Ballroom 202-204 (Level 2)
 Anne Yoder (Duke U.), One Tree to Rule Them All

SSE Presidential address & award announcements
 Monday, June 26 from 5:30 - 6:30 pm, Ballroom 202-204 (Level 2)
 Sally Otto (U. British Columbia), Linking Levels of Selection With Genetic Modifiers

Spotlight sessions

See the online program or mobile app for a listing of the talks in each session.

American Society of Naturalists

1) Processes underlying pattern: considering the evolutionary mechanisms underlying population-level differentiation

Sun. June 25, 8:30 - 9:45 am (Oregon ballroom 204)

Organizer: Rebecca Safran (U. Colorado)

Evolutionary mechanisms lead to changes in the phenotypic and genomic features of populations; population genetics and patterns of phenotype differentiation are often used to infer which of these mechanisms are at work. Here, we highlight the need to more directly study evolutionary process as a way to understand the mechanistic basis of population-level patterns. That is: what adaptive or random processes are at work that contribute to population differences? In this session, we hope to touch on processes (e.g., local adaptation, migration, mate-selection) that influence evolution within populations and contribute to larger scale patterns of differentiation, with a consideration of analytical and experimental approaches.

2) Eco-evolutionary dynamics

Mon. June 26, 10:15 - 11:30 am (Oregon ballroom 204)

Organizer: Andrew Hendry (McGill U.)

It is now widely recognized that substantial evolutionary change can occur on contemporary (or "ecological") time scales. This is the phenomenon of contemporary (or "rapid") evolution. What we now need to know is the extent to which contemporary evolution shapes ecological dynamics at the population, community, and ecosystem levels. This Spotlight Session will feature talks on how ecological change influences contemporary evolution and how contemporary evolution influences ecological change.

3) '25 Years of sensory drive'

Mon. June 26, 3:15 - 4:30 pm (Oregon ballroom 204)

Organizer: Becky Fuller (U. Illinois)

In 1992, the American Naturalist published a supplemental issue on the topic of 'Sensory Drive' organized by John Endler. This issue featured a number of influential papers that emphasized the role of the environment, phylogenetic history, and development on the evolution and functioning of sensory systems. This supplement introduced many ecologists and evolutionary biologists to the intriguing and well-developed topic of sensory ecology. In this spotlight session, we will consider the role of sensory drive as a guiding framework in ecological and evolutionary studies of signals/sensory systems over the past 25 years, and the extent to which it serves as a useful framework going forward.

Society for the Study of Evolution

1) Speciation with and without sex chromosomes

Sun. June 25, 10:15 - 11:30 am (Oregon ballroom 203)

Organizer: <u>Suzanne Edmands</u> (U. Southern California)

Studies of the genetics of speciation frequently focus on sex chromosomes. In fact, the "two rules of speciation" (Coyne & Orr 1989) are both based on sex chromosomes. In this session we address how sex chromosomes affect speciation and explore how speciation mechanisms may be different in the wide diversity of taxa lacking sex chromosomes.

2) Sexual selection via male competition and the process of speciation

Sun. June 25, 1:30 - 2:45 pm (Oregon ballroom 203)

Organizer: Janette Boughman (Michigan State U)

The role of sexual selection in speciation remains controversial, despite the last two decades of intensive work. The importance of male competition has been almost entirely ignored, as nearly all the work focuses exclusively on how female preferences generate sexual selection in ways that hinder gene flow. This is despite that ecological competition over resources plays a central role in speciation via natural selection, and that male competition can generate very strong selection. However, recent work suggests that incorporating both male competition and female choice into research on speciation can help resolve the controversy over whether and when sexual selection causes speciation, prevents it, or has no real effect. Male competition sometimes complements and other times opposes female choice, which can either accelerate or put a brake on the speciation process, or lead to asymmetries in gene flow. Another complexity is that males compete both before and after copulation through sperm competition or the analogous pollen competition in plants; including this component can draw interesting parallels between animal and plant speciation.

3) Evolution of hybrid incompatibilities

Sun. June 25, 3:15 - 4:30 pm (Oregon ballroom 203)

Organizer: Andrea Sweigart (U. Georgia)

As a common cause of reproductive isolation in diverse taxa, hybrid incompatibilities are fundamentally important for genomic divergence and speciation. This session explores new insights into the molecular genetic mechanisms and evolutionary origins of hybrid incompatibilities.

4) Genetic transmission at the population level

Tues. June 27, 8:30 - 9:45 am (Oregon ballroom 203)

Organizer: <u>Jeremiah Busch</u> (Washington State U.)

The transmission of genes between parents and offspring is a critical feature of biological populations. In this spotlight session, talks will explore the modes whereby genes are transmitted across generations and their evolutionary relevance. Topics include, but are not limited to, meiotic drive systems, organelle and nuclear genomes, ploidy, mating systems, and genetic conflict. We hope you will join us to learn more about the intriguing diversity, evolution, and macro-evolutionary consequences of genetic transmission systems.

5) Sex in the wild

Tues. June 27, 10:15 - 11:30 am (Oregon ballroom 203)

Organizer: Maurine Neiman (U. Iowa)

A session focusing on exciting work relevant to the evolution/maintenance of reproductive mode polymorphism in non-model organisms and/or using field-based approaches to study sex (and its alternatives) in natural populations.

Society of Systematic Biology

1) Uncertainty in the face of strong support: new approaches and examples in the identification of signal, noise, and conflict in phylogenetic datasets

Mon. June 26, 8:30 - 9:45 am (Oregon ballroom 202)

Organizers: <u>David Weisrock</u> (U. Kentucky) and <u>Emily Jane McTavish</u> (U. California Merced)

Genome-scale data sets provide large quantities of information with which to resolve challenging phylogenetic relationships. However, large data sets can yield strongly supported trees, even when there is conflict in the data. Consequently, it is important to understand what exactly underlies the resolution of a particular topological result. While we are often looking for phylogenetic signal, it is important to investigate other factors affecting reconstruction. These factors may include biological processes such as hybridization and horizontal gene transfer, and incomplete lineage sorting, or methodological biases, such as insufficient models of evolution, choice of reference genome, or bias in genomic or taxonomic sampling. While the idea of examining sources of conflict is not a new idea in systematics, its importance is elevated in the study of genome-scale data. We have developed this spotlight session to highlight new

approaches for discerning how individual loci or sets of loci impact a phylogenetic hypothesis, and provide examples of how these approaches have been implemented with contemporary datasets.

2) Impact of gene flow and reticulation in phylogenetics

Mon. June 26, 1:30 - 2:45 pm (Oregon ballroom 202)

Organizers: <u>Cécile Ané</u> (U. Wisconsin-Madison) and <u>Frank Burbrink</u> (American Museum of Natural History)

Genomic data hold the power to robustly infer phylogeny, yet they also present new challenges for resolving the tree of life due to conflicting signals across loci. Currently, coalescent-based methods depend on incomplete lineage sorting to account for phylogenetic conflict among loci, while mostly ignoring the confounding effects of gene flow. In this spotlight session, we will consider the following questions: In what ways do gene flow and other biological processes (e.g., introgression, hybridization, and selection) alter our ability to study the evolutionary history of a group? Should we represent this history as a species or population network? What do empirical studies say about the prevalence of gene flow and hybridization across phylogeny? How does this affect our ability to perform downstream comparative phylogenetic analyses?

Symposia

See the daily schedule listed online for a detailed listing of the talks in each symposium.

1) ASN VP Symposium: The causes and consequences of evolutionary conflicts

Sat. June 24, morning (Oregon ballroom 204)

Organizer: Locke Rowe (U. Toronto)

Sponsor: ASN

In this symposium we explore conflict at several levels of biological organization, with the aim of highlighting some unifying principles of evolutionary conflicts.

2) Across the Nth dimension: Quantitative and conceptual advances in the study of niche breadth

Sat. June 24, afternoon (Oregon ballroom 204)

Organizers: Kelly Anne Carscadden (U. Colorado-Boulder) and Nancy Emery (U.

Colorado-Boulder);

Sponsor: ASN

The niche is central to many of the most enduring ecological and evolutionary questions; however, fundamental challenges remain in defining and quantifying the niche and interpreting niche breadth data. This symposium will highlight key limitations and advances in niche concepts, explore network- and trait-based measures of niche breadth, and propose a general classification and guide to niche breadth metrics. Then, speakers will provide an updated view on niche architecture based on the idea that species-level niche breadth may arise through a combination of mechanisms at different nested levels of organization. Because niche breadth is a concept with not only deep historical roots, but

also relevance to many of today's environmental challenges, we collectively aim to emphasize how niche breadth relates to conservation and environmental change.

3) A debate of conceptual issues surrounding genetic-based species delimitation in the genomic era

Sun. June 25, morning (Oregon ballroom 202)

Organizers: David M. Hillis (U. Texas-Austin) and L. Lacey Knowles (U. Michigan)

Sponsor: SSB

With recently developed coalescent-based analytical methods, coupled with technological advances for collecting genomic datasets across many individuals per putative species, we no longer have to worry about missing recently diverged taxa because of limited power. With a diversity of approaches to handle different genomic data types, much of the focus to date regarding genetic-based species delimitation has been either on (1) performance differences among methods and data types, and (2) inference for empirical application. What have not received much attention are the conceptual issues surrounding genetic-based species delimitation in the genomic era. This is what our symposium will address.

The symposium will be organized in a debate style format to foster discussion about issues that have broad implications beyond the endeavor of inferring species boundaries, including understanding the underlying macroevolution dynamics, as well as conservation of, biodiversity, and the study of the speciation process itself. The themes the symposium will focus upon include:

Does the multispecies coalescent delimit species?
Is species delimitation necessary for assessing the drivers of diversification?
Should we seek a "standardized protocol" for species delimitation?
What is the role of morphology in species delimitation in the genomic era?

4) Phylogenetic approaches to connecting genotypes to phenotypes

Sun. June 25, afternoon (Oregon ballroom 202)

Organizer: Stacey D. Smith (U. Colorado-Boulder)

Sponsor: SSB

A central challenge for evolutionary biologists is to understand the relationship between genotype and phenotype, the so-called 'rules of life.' Making this connection is difficult beyond the species-level, where traditional population genetic methods (e.g., QTL mapping) often cannot be used. This symposium will explore new approaches that aim to dissect the mechanisms of phenotypic evolution at macroevolutionary scales and to place underlying changes along branches of the phylogeny.

5) Evolution in urban ecosystems

Mon. June 26, morning (Oregon ballroom 203)

Organizers: Emily Puckett (Fordham U.) and Jason Munshi-South (Fordham U.)

Sponsor: SSE

The rapid growth of existing cities and creation of entirely new cities, have created unique

environments and novel selection pressures for microbes, animals, and plants. Aspects of urban environments that may act as selection pressures include: increased exposure to noise, exposure to pollution and other synthetic materials, constant artificial light, availability of calorie dense food, linearized environments, novel substrates, novel predators or release from predation, exposure to high-density human populations, and interactions within novel ecological communities. Our symposium will focus on how species from distant taxa are evolving to these pressures. Further, we will explore how urban-rural gradients and the urban matrix serve as either barriers or corridors for gene flow in different species, and implications for genetic drift, local adaptation, the evolution of mating systems, and inbreeding tolerance.

6) The impact of stress on genetic variation

Mon. June 26, afternoon (Oregon ballroom 203)

Organizer: Laurie Stevison (Auburn U.)

Sponsor: SSE

Understanding the influence of environmental stress on genetic variation typically requires very large sample sizes. Many previous studies in this area have been limited to model organisms such as *Drosophila* and have used mutant phenotypic markers as a proxy for genetic variation. As sequencing costs have continued to decline, the possibility of addressing these questions in other systems has become easier. Here, we include a broad swath of taxonomic diversity to represent the increased possibility of addressing this important evolutionary question outside of traditional model organisms. Further, we present both theoretical and empirical work at the interface of evolution and ecology to understand more broadly the impact of stress on natural variation.

7) SSE Education symposium – Communicating (the relevance of) human evolution Sat. June 24, 8:30 - 11:30 am (Rm C123)

Organizers: Jim Smith (Michigan State U. - Lyman Briggs College) and Norman Johnson (U. Massachusetts-Amherst)

In recent years, educators and researchers have made substantial progress in addressing the challenges of evolution education. Despite this excellent work, a large fraction of students and the general public still refuses to accept or struggles with the concept of humans as an evolved and evolving species. Moreover, many of those who accept evolution do not fully appreciate the relevance to society and/or hold misconceptions about how evolution works. The three speakers in this SSE-sponsored symposium describe their work in human evolution and how they have overcome obstacles to communicating human evolution and the relevance of evolution to humans. We'll cap off our session with a Panel discussion in which we explore avenues for effective action.

8) ASN Education symposium – Natural history as the inspiration for scientific inquiry: Stories and tools for teaching

Sat. June 24, 1:30 - 4:30 pm (Rm C123)

Organizers: George Gilchrist (NSF) and Becky Fuller (U. Illinois)

The American Society of Naturalists has promoted the unification of the biological sciences and the importance of natural history in inspiring research. In this anniversary symposium

celebrating the 150th anniversary of The American Naturalist, life scientists will reflect on the role of natural history in directing their research, providing examples and materials that could be incorporated into teaching in modern conceptually based courses.

Outreach talk

SSE Public Outreach Lecture

Friday, June 23, 7:30 - 8:30 pm, Oregon Ballroom 202-204

Speaker: Ann Reid, Executive Director of the National Center for Science Education

Title: TBD

Special Events

Pre-conference workshops & satellite events

All of these require registration and some have an additional fee. Sign-up and payment (if applicable) for most, but not all, are part of main conference registration. All events, except #1, are in the Oregon Convention Center.

1) Active LENS Train-the-trainer satellite workshop: Avida-ED Digital Evolution Education Software

Wed. June 21 - Friday, June 23, U. Washington, Seattle WA; Cost: free, application required (separate from main conference registration; see below).

Organizers: Robert Pennock, Jim Smith, Louise Mead

Topic: The purpose of this workshop is to train instructors in the use of the Avida-ED software package, developed to help students learn about evolution and the nature of science, so that workshop participants can both implement classroom interventions using this software and also train other educators. Teams of two will learn to use Avida-ED and how to best incorporate it into courses that they teach. Travel and expenses related to the workshop will be covered for the 20 participants for each workshop as part of an NSF-funded IUSE grant. Avida is a digital evolution software platform used to study evolutionary processes, and harness evolution to solve engineering problems. Avida-ED is a free, user-friendly version of Avida developed specifically for educational purposes, with a graphical user interface and visualizations that allow the user to observe evolution in action. (See http://avida-ed.msu.edu for more information.) Workshop participants will join a growing community of educators using digital evolution to let their students directly observe evolutionary processes through inquiry-based exercises that advance reformoriented active learning. The team application form for the Active LENS Workshop must be completed online on this page. Applications should be submitted no later than March 17, 2017.

Speakers: Ben Kerr, Richard Lenski, Louise Mead, Robert Pennock, Jim Smith, and Mike Wiser.

2) JoeFest: a symposium to honor Joe Felsenstein

Thursday, June 22, 9 am - 5 pm with optional evening reception, Rm. B113-114;

Cost: \$20 (+\$30 optional reception; reception includes two drinks and appetizers), registration required (during conference registration). Open to everyone. Maximum enrollment: 300 (first-come, first-served).

Organizer: Michael Turelli (UC Davis), Monty Slatkin (UC Berkeley), Ray Huey (U. Washington) If you want to attend JoeFest only (i.e. not the main Evolution conference), then please <a href="mailto:emailto:

JoeFest Schedule:

9:00 Lee Felsenstein, The Fonly Institute. *The early years: an appreciation of Joey Felsenstein* 9:20 Michael Turelli, UC Davis. *A star is born*

9:40 Bill Hill, University of Edinburgh. *Joe Felsenstein's Hill-Robertson Effect* 10:00 Ray Huey, University of Washington. *Comparative biology: before, during, and after Felsenstein 1985*

10:20 — COFFEE BREAK

10:50 Wayne Maddison, University of British Columbia. "Ask Joe"

11:10 Jeff Thorne, North Carolina State. Phylogenetic inference of interlocus gene conversion

11:30 Peter Beerli, Florida State University. Everything you always wanted to know about frogs/computers/population genetics but were afraid to ask

11:50 Mary Kuhner/Jon Yamato, University of Washington. The insane geometry of tree space: early development of genealogy samplers

12:10 — LUNCH

1:40 Mark Kirkpatrick, University of Texas, Austin. *The war between the sexes: genome evolution under sexual antagonism*

2:00 Fred Allendorf, University of Montana. Population genomics: Do we really need theory?

2:20 Steve Arnold, Oregon State. Quantitative genetics on evolutionary timescales

2:40 Scott Edwards, Harvard. The tree thinker and the long way to amphioxus

3:00 — COFFEE BREAK

3:30 Shozo Yokoyama, Emory. Evolution by molecular interaction

3:50 Ted Garland, UC Riverside. Zen and the art of phylogenetically independent contrasts

4:10 Ruth Shaw, University of Minnesota. Evaluating the capacity for ongoing adaptation

4:30 Monty Slatkin, UC Berkeley. Isolation by distance and time--another pain in the torus

5:30-7:30 Reception (separate registration \$30, private room at Altabira, across the street from the OCC, includes two drinks and appetizers)

3) SSB workshop – Trees, traits and functions: semantics for comparative biology Friday, June 23, 1:30 - 5 pm, Rm. B114; Cost: free, registration required (during conference registration). Open to everyone. Maximum enrollment: 130 (first-come, first-served) Organizers: Todd Vision (UNC Chapel Hill) and Hilmar Lapp (Duke U)

Topic: A rich set of comparative phylogenetic methods has been developed in recent decades for the analysis of traits, functions, and other macroevolutionary patterns. A similarly rich set of tools,

such as ontologies, data annotation models, and machine reasoners, have also recently been introduced allowing researchers to represent the semantics of phenotypes, traits, functions, and environmental properties in such a way that computers can access what had previously been the implicit knowledge of biological experts. Although these two efforts have remained largely separate from each other, a few studies at their intersection have shown that their synthesis has the potential to enable new classes of comparative analyses and provide an interdisciplinary bridge between different phylogenetics user communities. This symposium brings together research at the nexus of comparative phylogenetics and computable semantics of biological knowledge. The aim is to stimulate discussion and awareness of the opportunities that are emerging for the comparative biology community.

Speakers:

Carrine Blank (U. Montana) Use of natural language processing algorithms, ontologies, and phylogenomic trees to automate and accelerate phenotypic trait evolutionary studies for prokaryotes

Paula Mabee (U. South Dakota) Progress in comparative biology can be accelerated through semantic annotation

Paul Thomas (U. Southern California) Modeling the evolution of gene function using phylogenies and the Gene Ontology

Martin Ramirez (Museo Argentino de Ciencias Naturales) *Tracing the evolution of complex traits:* the integration of atomized morphological data on phylogenies

Plus additional speakers (TBD)

4) SSB workshop - Species in the age of discordance

Friday, June 23, 8:45 am - 5:00 pm, Rm A109; Cost: \$20, registration required (during conference registration). Open to everyone. Maximum enrollment: 70 (first-come, first-served). Organizer: Matt Haber (U. Utah)

Topic: Biological lineages move through time, space, and each other. As they do, they diversify, diverge, and grade away from and into one another. One result of this is genealogical discordance, i.e., the lineages of a biological entity may have different histories. We see this on numerous levels, from population-level lineages, to microbial networks, to holobionts. Recent work in systematics, in particular genomics and phylogenomics, is providing an ever more refined view of this (and other sorts of) discordance. The goal of this workshop is to pause to consider how and whether our better understanding of discordance impacts how we think about species. The invited speakers represent a diverse group along several dimensions from multiple career stages, from graduate student to full professor. The workshop will include structured, interactive discussions in addition to panel presentations by invited speakers. The central question will be addressed from a variety of theoretical perspectives, including phylogenetic systematics, population genetics, microbiology, and philosophy of biology. This workshop is a follow-up session to an NSF supported interdisciplinary conference on the same topic. The intention is to use this workshop to help maintain and build on the cross-disciplinary connections made between researchers at that meeting, and to encourage broader disciplinary participation in systematics.

Speakers:

David Baum (U. Wisconsin, Madison) Genealogical individuation in the face of discordant gene trees: Does exclusivity still work?

Matt Haber (U. Utah) The species problem problem and the no solution solution

Ariadna Morales (Ohio State U.) Species delimitation with gene flow in Myotis bats

Leonie Moyle (Indiana University) Still 'good' species or: How I learned to stop worrying and love the incongruence

Jay Odenbaugh (Lewis & Clark College) Discordance, explanation, and levels of lineages

Makmiller Pedroso (Towson U.) Species and the assembly of microbial communities

Nitin Phadnis (U. Utah) Selfish genetic elements and speciation in Drosophila

Mike Shapiro (U. Utah) The pigeon's checkered past: pigment patterning and the hybrid origin of an adaptive trait

Olga Zhaxybayeva (Dartmouth U.) Of species and clusters: Could we distinguish the observed patterns of bacterial microdiversity from those produced by a neutral process of lineage birth and death?

5) ASN Science communication workshop: Effective communication of science and its impacts

Friday, June 23, Time 9 am - 4 pm, Rm B111; Cost: \$25 (Includes box lunch and morning/afternoon coffee breaks), registration required (during conference registration). Open to all conference attendees. Maximum enrollment: 40 (first-come, first-served).

Organizer: <u>Jory Weintraub</u> (Science Communication Director, Duke Initiative for Science & Society)

Topic: Sure, your work on [fill-in-the-topic-of-your-research-here] is the most exciting and important science out there! But to have an impact, it must be communicated in ways that enable people to understand and compel them to care. And, in order to be funded it must have a well-designed, easily-implemented and assessable broader impacts plan. This workshop will provide strategies and approaches to help you communicate your work effectively to diverse audiences, and provide insight into what the NSF really wants when they ask you to communicate the broader impacts of your work.

Topics covered will include:

Defining your message, identifying your audience and crafting the perfect elevator pitch Ready for Your Close-Up? – Strategies for nailing your media interview Social Media – The three W's (What?Why?Will using it take over your entire life?) Broader Impacts 101 – What does this have to do with science communication?(Answer: EVERYTHING!)

This will be a "hands-on" workshop, so come prepared to practice your skills with your colleagues!

This workshop will be co-facilitated by Jory Weintraub, PhD and Briana Pobiner, PhD. Jory is the Science Communication Director for the Duke Initiative for Science & Society, and Director of the Duke Broader Impacts Resource Center, and serves on the Steering Committee of the National

Alliance for Broader Impacts (NABI). He has over 20 years of experience in science education, outreach and communication, including 10+ years leading the education and outreach efforts at NESCent (the National Evolutionary Synthesis Center). Briana is a paleoanthropologist and educator at the Smithsonian's National Museum of Natural History (NMNH). While she continues to conduct research on human evolution, her primary responsibilities since she joined the NMNH's Human Origins Program in 2005 focus on onsite and online education, outreach, and public programming. She has led several science communication training workshops for NMNH staff, postdocs, and volunteers.

6) SSE Diverse careers committee workshop

Friday, June 23, 1 - 5:15 pm, Rm. B116; Cost: \$20, registration required (during conference registration). Restricted to graduate students/PDFs who are members of the SSE, ASN, and/ or SSB. Maximum enrollment: 70 (first-come, first-served).

Organizers: Mohamed Noor (Duke U) and Lex Flagel

Topic:The key goals of the workshop are to guide students and postdocs through the first steps of personal career exploration (academic and non-academic) and necessary actions for active planning of post-graduate school or post-postdoctoral stages. In addition to small-group conversations with professionals from various careers, there will be presentations on how to market/brand oneself, networking, informational interviews, etc. Based on our past workshop, we expect that attendees will leave the workshop with more confidence regarding career options, and with actionable items and strategies for successive efforts.

7) SSE Education Committee workshop: Professional Development for Undergraduate Educators

Friday, June 23; Time: 9 am - 5pm (approximately), Rm B112; Cost: \$25 (includes a boxed lunch), registration required (during conference registration). Maximum enrollment: 40 (first-come, first-served).

Organizer: Louise Mead

Topic: National reports such as <u>Vision and Change</u> call for engaging students in authentic scientific experiences, such as using models and working with real data. Teaching with data and models can deepen students understanding of key biological concepts, develop a greater appreciation of the process of science, and engage students in quantitative and computational thinking. In this workshop, we will explore some of the challenges and successful approaches to using models and data in the classroom. More information including a workshop schedule can be found here.

8) SSE/BEACON Undergraduate Diversity Program: Professional Development Workshop Friday, June 23; Time: 8:30 am - 1:00 pm, Rm B115; Cost: free, registration required (during conference registration). Restricted to undergraduates who are attending the meeting. Maximum enrollment: 40 (first-come, first-served).

Organizer: Alexa Warwick

Topic: This workshop will cover topics including networking, mentoring, work/life balance, and alternative careers (8:30-11:15 am), with presentations by Scott Edwards, Lex Flagel, Joanne Kamens, Stacy Farina, and April Wright. It will also feature a keynote speaker (Pamela Yeh) for the annual Undergraduate Futures Lecture followed by lunch and a panel discussion with all the

speakers (11:15am-1:00pm).

9) iEvoBio satellite conference

Organizer: Daisie Huang (U. British Columbia)

iEvoBio is a forum bringing together biologists working in evolution, systematics, and biodiversity, with software developers and mathematicians. This year, we'd like to focus on the computational challenges facing empirical biologists. Are you an empirical biologist? Is there some part of your research that you think software can help with? Are you a software developer who wants to find a new challenging project that will help scientists with their research? iEvoBio is the place to discuss these challenges and make some new connections.

iEvobio will host two events in association with the meeting:

- 1) Pre-conference workshop, Friday June 23, 9 am 5 pm, Rm. B113. Keynote speaker: Tracy Teal (Data Carpentry).
- 2) A 'Software Demo' during the Saturday poster session (6:30 8:30, Exhibit Hall A)

There was a separate registration fee for attending iEvoBio (during main conference registration). For more information about the schedule and agenda, please visit their website.

Social events

Friday, June 23

• Opening Reception, 6:30 - 10:00 pm, Exhibit Hall A

Saturday, June 24

• ASN Grad Student Mixer, 4:30 – 5:30 pm, Ballroom 201 & Ballroom Lobby (level 2)

Sunday, June 25

- SSB Society Mixer, 4:30 5:30 pm, Ballroom 201 & Ballroom Lobby (level 2)
- 7th Annual Evolution Film Festival & Video Contest, 7:30 9:30 pm, Room A107-109 One of the highlight of the conference that should not be missed! Join us for an evening of terrifically informative and entertaining videos, while enjoying popcorn and beer/wine/soda (courtesy of HHMI). During the first hour, event co-sponsor HHMI Biointeractive will screen some of the wonderful evolution videos they offer, and you will have a chance to ask questions of scientists and filmmakers involved in producing their videos. During the second hour, we will screen the top entrants to this year's contest. We asked the science and science education communities to tell a story about evolution in 3 min. or less, using animation, art, music, dance, etc. We have some terrific entries and you will be able to see them all and vote on your favorite. (To learn more and see past years' entries, go to evolutionfilmfestival.org and follow #evofilmfest on Twitter.)

Sponsored by the Duke Initiative for Science & Society, the Howard Hughes Medical Institute (HHMI), the Society for the Study of Evolution, and the BEACON Center for the Study of Evolution in Action.

Monday, June 26

- SSE/BEACON Undergraduate Diversity Program:
 - Undergraduate breakfast social, 8:00 9:30 am, Oregon Ballroom 201.
 Attendees should register by entering their information on this spreadsheet. Open to undergraduates and their advisers and mentors.
 Organizer: Alexa Warwick
 Come to meet and mingle with other undergrads, their advisors/mentors, and alumni of the Undergrad Diversity Program. This social is a great opportunity to discuss your research, career plans, etc. and expand your network in an informal setting while enjoying free continental breakfast!
 - SSE Student-Faculty Mixer, 4:30 5:30 pm, Ballroom 201 & Ballroom Lobby (level 2)

Tuesday, June 27

• Super Social, 6:30 pm – midnight, Oregon Zoo. Ticket required.

Council, business and other meetings

1) Executive joint council (ASN/SSB/SSE)

- Opening meeting: Friday, June 23, 8:30 am; Rm B119.
- Exit meeting: Monday, June 26, 8:30 am; Rm C126.

2) ASN

- Executive council opening meeting: Friday, June 23, 1:00 pm; Rm C126
- Business Members (open to all ASN members): Saturday, June 24, 6:30 pm;
 Rm A105
- American Naturalist journal editorial board meeting: Sunday, June 25, 8:30 am; Rm C126.
- Executive council exit meeting: Tuesday, June 27, 8:30 am; Rm C126.

3) SSB

- Executive council opening meeting: Friday, June 23, 1:00 pm; Rm B110
- Business meeting (open to all SSB members): Sunday, June 25, 6:30-7:30 pm;
 Room A105

- Systematic Biology journal editorial board meeting: Monday, June 26, 11:30 am -1:30 pm; Rm B110.
- Executive council exit meeting: Tuesday, June 27, 8:30 am; Rm B110.

4) SSE

- Executive council opening meeting: Friday, June 23, 1:00 pm; Rm B119
- Evolution journal editorial board meeting: Saturday, June 24, 11:30 am, Rm. B110.
- Education & Outreach committee: Sunday, June 25, 11:30 am; Rm. C126
- Hamilton award judges meeting: Sunday, June 25, 4:30 pm; Rm B110
- Executive council exit meeting: Monday, June 26, 11:30 am; Rm C126.
- Business meeting (open to all SSE members): Monday, June 26, 6:30 7:30 pm; Rm A105

5) Other meetings

- Taylor & Francis editorial board meeting: Sunday, June 25, 8:30 am; Rm B110
- Ecology and Evolution editorial board meeting: Monday, June 26, 11:30 am 1:30 pm; Rm B114-115

Special events during the meeting

Some of these require registration and additional fees. Sign-up and payment (if applicable) are done during main conference registration for most or via a separate registration process for others, as indicated below. All events are in the Oregon Convention Center.

1) Diversity in Science luncheon: Essentials of a successful diversity and inclusion program

Sunday, June 25 from 11:30 am - 1:30 pm, Oregon Ballroom 201. Cost: \$15 (includes a boxed lunch), registration required (during conference registration). Maximum enrollment: 300 (first-come, first-served). Open to conference attendees of all genders, ethnic groups, and backgrounds who are members of the SSE, ASN, and/or SSB.

Organizers: <u>SSE Graduate Student Advisory Council</u> with input from SSB and ASN Topic: Join us for a discussion on identifying bias and increasing diversity across all dimensions. Keynote speaker: <u>Lee Bitsoi</u> (Rush U.)

2) Evo101 Workshop for Science Educators

Saturday, June 24; Time: 8:30 am - 4 pm, Rm. C126; registration required (separate from conference registration - see below). Intended for K-12 science educators. Questions? Contact Kristin Jenkins

Topic: Evolution can often seem very distant from students' lives, making it hard to engage them in studying this core biological concept. Human evolution offers compelling and relevant examples of evolutionary processes for students. At this hands-on workshop we will share free resources based on human evolution you can use in your classroom. The workshop is designed for middle and high school teachers. We encourage you to bring a colleague! Lunch and certificates of participation will be provided for participants.

This workshop has its own registration process separate from the main conference. You don't need to register or attend the main conference to sign-up for this event.

Schedule, registration and other information can be found here. Registration deadline: June 1, 2017

Supported by the <u>Society for the Study of Evolution</u> Education Committee and the <u>Howard Hughes</u> <u>Medical Institute</u>

3) SSE Student-Faculty networking lunch

Saturday, June 24, 11:30 am -1:30 pm, Location: various offsite lunch venues. Cost: your own lunch; Registration: indicate interest during main conference registration.

Organizer: Megan Kobiela and the SSE Graduate Student Advisory Council

The SSE Graduate Student Advisory Council is organizing small group networking lunches to facilitate interactions between students and faculty. You can indicate your interest in participating during main conference registration, and we will send out surveys to match students with faculty in late May. Groups will be responsible for finding their own lunch venues and will be capped at 5 students per faculty member, so student attendance may be limited to a first-come, first-serve basis once the surveys go out.

4) SSE Postdoc networking lunch

Saturday, June 24, 11:30 am - 1:30 pm, Location: various offsite lunch venues. Cost: your own lunch; Registration: indicate interest during main conference registration.

Organizer: SSE Graduate Student Advisory Council

The SSE Graduate Student Advisory Council is organizing small group networking lunches to facilitate interactions among postdocs. You can indicate your interest in participating during main conference registration and we will send out surveys to set up groups in May. Groups will be responsible for finding their own lunch venues.

5) Taylor & Francis

Early career researcher's workshop (1 of 2): Publishing your research: tips for success Saturday, June 24, 12:15 -1:15 pm, Rm. B113. Cost: free, no registration required. Bring your own lunch.

Organizer: Ashlynne Merrifield, Ali Paskins, Andrew Polaszek, and Jeffrey Streicher This talk will guide new authors through the publishing process from selecting a journal, to preparing for submission, to navigating the peer review process. Andrew Polaszek (Editor-in-Chief, JNH) and Jeffrey Streicher (Associate Editor, JNH) will give you a valuable insight into the editorial process for Journal of Natural History, including advice for successful publication. We will also cover essential tips for writing a review article, how to respond to reviewer comments, acting as a reviewer, plus more, giving a thorough understanding of the steps involved, the key information sources to be aware of, and what authors can be doing to help get their paper published.

6) British Ecological Society

Early career researcher's workshop (2 of 2): Maximizing the impact of your research Sunday, June 25, 12:15 -1:15 pm, Rm. B113. Cost: free, no registration required. Bring your own lunch.

Organizers: Chris Grieves (Methods in Ecology and Evolution)

In recent years there has been a significant increase in the number of academic articles published. At the same time, readers are changing how they find content – searching for individual articles

rather than looking through journals. This interactive workshop will provide new authors with the necessary skills to maximize the impact of their research in the digital age. It will include guidance on optimising your articles for search engines, identifying the best forums to promote your work, pitching your paper to different audiences, and using social media.

7) 7th annual Evolution Film Festival/Video contest (Call for video submission)

Sunday, June 25, 7:30 - 9:30 pm, Rm. A107-109. Cost: free, no registration required. One of the highlight of the conference that should not be missed! Join us for an evening of terrifically informative and entertaining videos, while enjoying popcorn and beer/wine/soda (courtesy of HHMI). During the first hour, event co-sponsor HHMI Biointeractive will screen some of the wonderful evolution videos they offer, and you will have a chance to ask questions of scientists and filmmakers involved in producing their videos. During the second hour, we will screen the top entrants to this year's contest. We asked the science and science education communities to tell a story about evolution in 3 min. or less, using animation, art, music, dance, etc. We have some terrific entries and you will be able to see them all and vote on your favorite. (To learn more and see past years' entries, go to evolutionfilmfestival.org and follow #evofilmfest on Twitter.)

Sponsored by the Duke Initiative for Science & Society, the Howard Hughes Medical Institute (HHMI), the Society for the Study of Evolution, and the BEACON Center for the Study of Evolution in Action.

8) NSF Information session

Monday, June 26, 12:15 - 1:15 pm, Rm. B113. Cost: free, no registration required.

An information session on NSF funding opportunities and program changes. Lunch is not provided (you can bring your own).

Host: Samuel Scheiner & George Gilchrist (NSF Program Directors)

9) Meet the Editors: A question and answer session

Monday, June 26, 3:15 - 4:30 pm, Oregon Ballroom 201. Cost: free, no registration required. Open to all conference attendees.

An informal question and answer session with a panel of editors/associate editors from the leading journals in the field including:

Judith Bronstein, American Naturalist
Allen Moore, Ecology and Evolution
Mohamed Noor, Evolution
Zach Gompert, Evolution Letters (Associate editor)
Barbara Mable, Heredity
Thomas Near, Systematic Biology

Post-Conference Workshop

1) American Institute of Biological Sciences workshop: Informing and engaging decisionmakers

Wednesday, June 28, 9 am - 5pm, Rm B110; Cost: \$35 (lunch is 'own your own' and is not included in registration cost), application required (during conference registration). Open to everyone. Maximum enrollment: 30 (first-come, first-served).

Sponsor: SSE

Organizer: <u>Janette Boughman</u> (Michigan State U.)

Topic: Rarely has the need for effective and influential communication about science been more important than it is today. Politicians and political interests are redefining and reinterpreting science—with great persistence and impact. The majority of the public still respect scientists and value science, but they often find it challenging to discern who and what is legitimate.

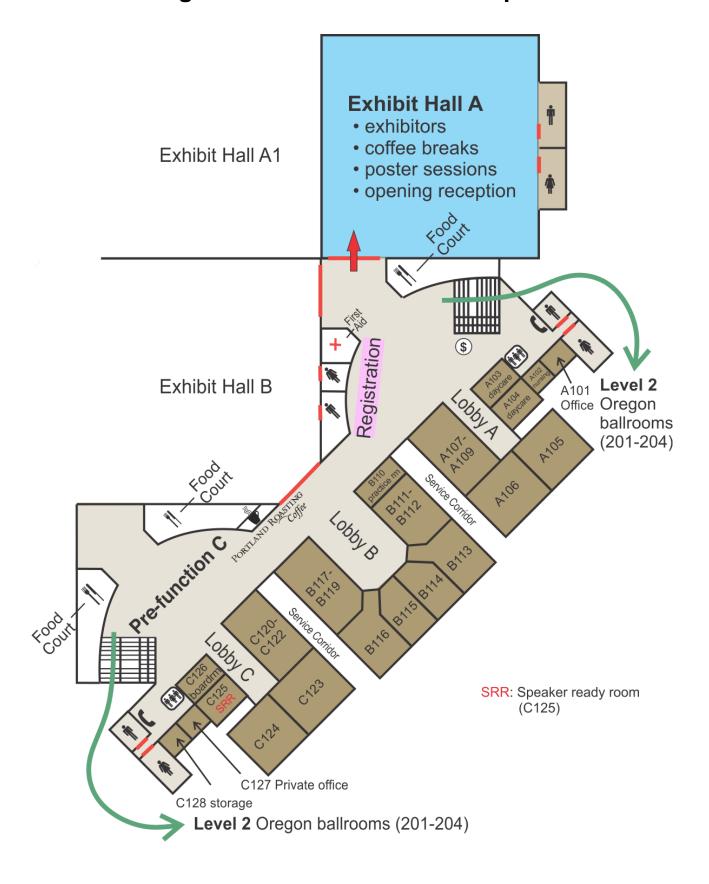
Simply adding to the noise will not provide individuals with the understanding they need to inform deliberations and decisions. Scientists must become more skilled at communicating with policymakers in impactful ways.

The American Institute of Biological Sciences (AIBS) will deliver this program. AIBS has a long and successful track record of engaging, informing, and influencing the public and science policy decision-makers. AIBS routinely work with include members of Congress, federal agency heads and program managers, state officials, and university administrators. AIBS has played important roles, for example, in helping to defeat state-level initiatives to introduce creationism into state curricula. AIBS has also been a leader in national efforts that have led to new federal investments in biology.

Participants will learn:

How to tell a resonant story that informs decision-makers
How to prepare for and engage in a meeting with a decision-maker
How to protect your scientific reputation
How to identify and define the audience you need to reach
What policymakers want and need to know from a scientist

Oregon Convention Center floorplan



Evolution 2017 Program

Session Thu-B113-B114 JoeFest: a symposium to honor Joe Felsenstein

Thursday - 9:00 - 17:00

Room B113-114

Session Executive joint council (ASN/SSB/SSE) Friday - 8:30 - 12:00 **Room B119**

Session SSE/BEACON Undergraduate Diversity Program: **Professional Development Workshop**

Friday - 8:30 - 13:00

Room B115

This is an optional pre-conference event. Pre-registration is required (during conference registration) and is r estricted to undergraduates who are attending the meeting.

This workshop will cover topics including networking, mentoring, work/life balance, and alternative careers (8:30-11:15 am), with presentations by Scott Edwards, Lex Flagel, Joanne Kamens, Stacy Farina, and April Wright. It will also feature a keynote speaker (Pamela Yeh) for the annual Undergraduate Futures Lecture followed by lunch and a panel discussion with all the speakers (11:15am-1:00pm).

Session SSB workshop - Species in the age of discordance Friday - 8:45 - 17:00 **Room A109**

An optional pre-confernce event. Prior registration required (during conference regisration).

Topic: Biological lineages move through time, space, and each other. As they do, they diversify, diverge, and grade away from and into one another. One result of this is genealogical discordance, i.e., the lineages of a biological entity may have different histories. We see this on numerous levels, from population-level lineages, to microbial networks, to holobionts. Recent work in systematics, in particular genomics and phylogenomics, is providing an ever more refined view of this (and other sorts of) discordance. The goal of this workshop is to pause to consider how and whether our better understanding of discordance impacts how we think about species. The invited speakers represent a diverse group along several dimensions from multiple career stages, from graduate student to full professor. The workshop will include structured, interactive discussions in addition to panel presentations by invited speakers. The central question will be addressed from a variety of theoretical perspectives, including phylogenetic systematics, population genetics, microbiology, and philosophy of biology. This workshop is a follow-up session to an NSF supported interdisciplinary conference on the same topic. The intention is to use this workshop to help maintain and build on the cross-disciplinary connections made between researchers at that meeting, and to encourage broader disciplinary participation in systematics.

Speakers:

• David Baum (U. Wisconsin, Madison) Genealogical individuation in the face of discordant gene trees: Does exclusivity still work?

• Matt Habér (U. Utah) The species problem problem and the no solution solution

- Ariadna Morales (Ohio State U.) Species delimitation with gene flow in Myotis bats
 Leonie Moyle (Indiana University) Still 'good' species or: How I learned to stop worrying and love the incongruence
- Jay Odenbaugh (Lewis & Clark College) Discordance, explanation, and levels of lineages

 Makmiller Pedroso (Towson U.) Species and the assembly of microbial communities
 Nitin Phadnis (U. Utah) Selfish genetic elements and speciation in Drosophila
 Mike Shapiro (U. Utah) The pigeon's checkered past: pigment patterning and the hybrid origin of an adaptive trait

 Olga Zhaxybayeva (Dartmouth U.) Of species and clusters: Could we distinguish the observed patterns of bacterial microdiversity from those produced by a neutral process of lineage birth and death?

Session ASN Science communication workshop: Effective communication of science and its impacts Friday - 9:00 - 16:00 **Room B111**

This is an optional pre-conference event. Prior registration required (during conference registration).

Topic: Sure, your work on [fill-in-the-topic-of-your-research-here] is the most exciting and important science out there! But to have an impact, it must be communicated in ways that enable people to understand and compel them to care. And, in order to be funded it must have a well-designed, easily-implemented and assessable broader impacts plan. This workshop will provide strategies and approaches to help you communicate your work effectively to diverse audiences, and provide insight into what the NSF really wants when they ask you to communicate the broader impacts of your work.

Topics covered will include:

- Defining your message, identifying your audience and crafting the perfect elevator pitch

- Ready for Your Close-Up? Strategies for nailing your media interview
 Social Media The three W's (What?Why?Will using it take over your entire life?)
 Broader Impacts 101 What does this have to do with science communication?(Answer: EVERYTHING!)

This will be a "hands-on" workshop, so come prepared to practice your skills with your colleagues!

This workshop will be co-facilitated by Jory Weintraub, PhD and Briana Pobiner, PhD. Jory is the Science Communication Director for the Duke Initiative for Science & Society, and Director of the Duke Broader Impacts Resource Center, and serves on the Steering Committee of the National Alliance for Broader Impacts (NABI). He has over 20 years of experience in science education, outreach and communication, including 10+ years leading the education and outreach efforts at NESCent (the National Evolutionary Synthesis Center). Briana is a paleoanthropologist and educator at the Smithsonian's National Museum of Natural History (NMNH). While she continues to conduct research on human evolution, her primary responsibilities since she joined the NMNH's Human Origins Program in 2005 focus on onsite and online education, outreach, and public programming. She has led several science communication training workshops for NMNH staff, postdocs, and volunteers.

Session SSE Education Committee workshop: Professional **Development for Undergraduate Educators** Friday - 9:00 - 17:00 **Room B112**

This is an optional pre-conference event. Prior registration required (during conference registration).

Topic: National reports such as <u>Vision and Change</u> call for engaging students in authentic scientific experiences, such as using models and working with real data. Teaching with data and models can deepen students understanding of key biological concepts, develop a greater appreciation of the process of science, and engage students in quantitative and computational thinking. In this workshop, we will explore some of the challenges and successful approaches to using models and data in the classroom.

More information including a workshop schedule can be found here.

Session ASN Executive council opening meeting Friday - 13:00 - 17:00 **Room C126**

Session SSB Executive council opening meeting Friday - 13:00 - 17:00 **Room B110**

Session SSE Executive council opening meeting Friday - 13:00 - 17:00 **Room B119**

Session SSE Diverse careers committee workshop Friday - 13:00 - 17:15 **Room B116** This is an optional pre-conference event. Prior registration required (during conference registration).

Topic:The key goals of the workshop are to guide students and postdocs through the first steps of personal career exploration (academic and non-academic) and necessary actions for active planning of post-graduate school or post-postdoctoral stages. In addition to small-group conversations with professionals from various careers, there will be presentations on how to market/ brand oneself, networking, informational interviews, etc. Based on our past workshop, we expect that attendees will leave the workshop with more confidence regarding career options, and with actionable items and strategies for successive efforts.

Session SSB workshop – Trees, traits and functions: semantics for comparative biology

Friday - 13:30 - 17:00

Room B114

An optinonal pre-conference event. Prior registration required (during conference registration).

Topic: A rich set of comparative phylogenetic methods has been developed in recent decades for the analysis of traits, functions, and other macroevolutionary patterns. A similarly rich set of tools, such as ontologies, data annotation models, and machine reasoners, have also recently been introduced allowing researchers to represent the semantics of phenotypes, traits, functions, and environmental properties in such a way that computers can access what had previously been the implicit knowledge of biological experts. Although these two efforts have remained largely separate from each other, a few studies at their intersection have shown that their synthesis has the potential to enable new classes of comparative analyses and provide an interdisciplinary bridge between different phylogenetics user communities. This symposium brings together research at the nexus of comparative phylogenetics and computable semantics of biological knowledge. The aim is to stimulate discussion and awareness of the opportunities that are emerging for the comparative biology community.

Speakers:

- Carrine Blank (U. Montana) Use of natural language processing algorithms, ontologies, and phylogenomic trees to automate and accelerate phenotypic trait evolutionary studies for prokaryotes
- Paula Mabee (U. South Dakota) Progress in comparative biology can be accelerated through semantic annotation
- Paul Thomas (U. Southern California) Modeling the evolution of gene function using phylogenies and the Gene Ontology
- Martin Ramirez (Museo Argentino de Ciencias Naturales) *Tracing the evolution of complex traits: the integration of atomized morphological data on phylogenies*
- Sergei Tarasov (NIMBioS, U. Tennessee, Knoxville) *Anatomy Ontology, Bayesian Networks and Evo-Devo suggest a new framework for modeling discrete morphological characters*
- Josef Uyeda (Virginia Tech U.) Can ontologies help understand trait evolution (and vice versa)?

Session Opening reception

Friday - 18:30 - 22:00

Room Exhibit Hall A

Session SSE Public Outreach Lecture Friday - 19:30 - 20:30 Room Oregon ballroom 202-204

Speaker: Ann Reid, Executive Director of the National Center for Science Education **Abstract:** Surveys consistently show that about 40% of American adults believe that humans were created in their current form in the last 10,000 years. One-third of American adults do not believe that human activities are changing Earth's climate. Lack of societal consensus around topics that are scientifically well-established echoes in our nation's science classrooms, where at least half of middle and high school science teachers report that they equivocate when they teach evolution and climate change. Ann Reid, Executive Director of the National Center for Science Education (NCSE), will discuss the more-timely-than-ever phenomenon of science denial and why she finds that label neither accurate nor particularly helpful. She will also share tools and lessons for communicating effectively about topics that are societally, but not scientifically, controversial.

Session Sat1-A105 Computational biology Saturday - 8:30 - 9:45 Room A105

- 8:45 The Effect of Evolution and Spatial Structure on Diversity in Biological Reserves
 Michael Wiser, Charles Ofria, Emily Dolson
- 9:00 Effects of Novel P Transposable Element Invasion in Drosophila yakuba Antonio Serrato, Stephania Zhang, Daniel Matute
- 9:15 Phylogenetic inference of tumor progression via copy number variation Russell Schwartz
- 9:30 Defining, understanding and exploiting eco-evolutionary dynamics in bone cancer David Basanta

Session Sat1-A106 Phylogeography / molecular ecology Saturday - 8:30 - 9:45 Room A106

- 8:30 Genetics and evolutionary relationships of Hemphillia slug populations Andrew Rankin, Anahí Espíndola, Michael Lucid, Jack Sullivan
- 8:45 Cryptic divergence on Florida's sand ridges: a new trap-jaw ant species Daniela Magdalena Sorger, Warren Booth, Adrian Smith, Robert Dunn
- 9:00 Evolution of a Subterranean Lifestyle in an Arboreal Frog Lineage Limits Dispersal and Enhances Genetic Diversification

 Jacobo Reyes Velasco, Joseph Manthey, Stephane Boissinot, Xenia Freilich
- 9:15 Severed connections and isolation of relictual Amazonian savannas inferred from genomic data

 Luciana C. Resende-Moreira, Andréa Thomaz, Joyce Prado, Andrea P. Souto, José P. Lemos-Filho, Maria Bernadete Lovato, L. Lacey Knowles
- 9:30 Geographic variation in the genetic structure of a mussel hybrid zone David Innes, Norah Saarman, Raymond Thompson

Session Sat1-A107-109 Species delimination / systematics Saturday - 8:30 - 9:45 Room A107-109

- 8:30 Revisiting species delimitation in the Lampropeltis triangulum complex *E. Anne Chambers*, Harry Greene, David Hillis
- 8:45 Genomic divergence in a widespread sea anemone complex: Implications for species delimitation

 Carlos Spano, Vreni Häussermann, Fabián Acuña, Charles Griffiths, Lisa Seeb, Daniel Gomez-Uchida
- 9:00 Species Delimitation in a recent and rapid radiation Sarah Jacobs, David Tank
- 9:15 Elucidating the evolutionary history of Mexican whiptail lizards

 Anthony Barley, Robert Thomson
- 9:30 Quantifying discordance in single locus species delimitation methods in horned lizards (Phrynosomatidae: Phrynosoma)

 Christopher Blair, Robert Bryson

Session Sat1-B111-112 Phenotypic plasticity / GxE 1 Saturday - 8:30 - 9:45 Room B111-112

- 8:30 The persistence of temperature effects through multi-generations Who Seung Lee, Stephan Munch, Youngrog Lee, Marc Mangel
- 8:45 Like grandfather like grandson: the trans-generational effects of dopamine manipulation on behavioural traits in the guppy, Poecilia reticulata

 Alex De Serrano, Helen Rodd, Kimberly Hughes
- 9:00 Complex interactions among individual traits affect behavioral plasticity expressed by female green swordtails (Xiphophorus hellerii) in two social contexts

 Rachael DiSciullo, Alexandra Basolo
- 9:15 The evolution of cooperation: Interacting phenotypes among social partners

 Bronwyn Bleakley, Mat Edenbrow, Safi Darden, Charles Tyler, Indar Ramnarine, Darren Croft
- 9:30 Social plasticity and the maintenance of alternative life histories: a modelling approach *Elizabeth Lange*, *Leithen M'Gonigle*

Session Sat1-B113 Evolutionary theory 1 Saturday - 8:30 - 9:45 Room B113

- 8:30 Transmission mode evolution in response to spatial heterogeneity in symbiont quality Alexandra Brown, Erol Akcay
- 8:45 Stags, hawks, and doves: Social evolution theory and individual variation in cooperation Jeremy Van Cleve
- 9:00 Genetic variation in legume sanctions and the maintenance of ineffective rhizobia Camille Wendlandt, Joel Sachs, Kelsey Gano-Cohen
- 9:15 Hidden Benefits Aid the Evolution of Altruism in Small Populations of Unrelated Individuals Sarah Fumagalli, Sean Rice
- 9:30 A cybernetic evolution theory: evolutionary duality of environmental stochasticity and natural selection

 Kurt Heininger

Session Sat1-B114-115 Conservation biology 1 Saturday - 8:30 - 9:45 Room B114-115

- 8:30 Large-scale genomic data change our understanding of the hybridization dynamics of the endangered California tiger salamander

 Evan McCartney-Melstad, Janet Vu, Jarrett Johnson, Bradley Shaffer
- 8:45 Novel solution to an old problem? Adaptation to transmissible cancer in the Tasmanian Devil Amanda Stahlke, Paul Hohenlohe
- 9:00 Hawaiian tree snails in a hot, dry world Melissa Price, Rob J. Toonen, Zac Forsman, Ingrid Knapp, Michael Hadfield
- 9:15 Investigating genome-wide patterns of deleterious variation in Channel Island foxes and Isle Royale wolves Jacqueline Robinson, Diego Ortega Del Vecchyo, Zhenxin Fan, Bernard Kim, Clare Marsden, Bridgett vonHoldt, Kirk Lohmueller, Robert Wayne
- 9:30 The effects of severe population crashes on genome-wide diversity in the Island Fox, Urocyon littoralis
 Nicole Adams, Xiaoming Wang, Suzanne Edmands

Session Sat1-B116 Speciation 1

Saturday - 8:30 - 9:45

Room B116

- 8:30 Climate-driven diversification in a clade of Peruvian poison frogs Connor French, Jason Brown
- 8:45 Amphibian and reptile speciation along elevational gradients in Ecuador Juan Guayasamin, Alejandro Arteaga
- 9:00 Insipient speciation in the high Andes? The tale of the Ecuadorian Hillstar, Oreotrochilus chimborazo

Elisa Bonaccorso, Carlos Rodríguez-Saltos, Nicolás Peñafiel, Nora Oleas

- 9:15 A hotspot for marine speciation: the Cape Verde Islands Regina Lopes da Cunha, Rita Castilho
- 9:30 Speciation in plethodontid salamanders of the Trans-Mexican Volcanic Belt Maria Guadalupe Segovia Ramírez, Sean Rovito

Session Sat1-B117-119 Adaptation 1 Saturday - 8:30 - 9:45 Room B117-119

8:30 Desiccation and the effect on diapause in two species of Rhagoletis fruit flies

Peter Meyers, Meredith Doellman, Lauren Ashley Miller, Christa Kohnert, Dietmar Schwarz, Gregory
Ragland, Jeffrey Feder

- 8:45 Is climate change disrupting local adaptation? A meta-analysis of field transplants Amy Angert, Megan Bontrager, Devin Gamble, Rachel Germain, Anna Hargreaves, Liz Kleynhans, Catriona Leven, Thomas Richards, Ken Thompson
- 9:00 Plant adaptation along drought frequency gradients Grey Monroe
- 9:15 Assessing genomic divergence in American pika along elevational transects reveals insights into thermal adaptation

 Matthew Waterhouse, Liesl Erb, Erik Beever, Michael Russello
- 9:30 To guard or to grow? The role of pleiotropy and life history tradeoffs in the evolution of chemical defense clines in monkeyflowers.

 Nicholas Kooyers, Benjamin Blackman, Liza Holeski

Session Sat1-Ball202 SSB Mayr Award Symposium 1 Saturday - 8:30 - 9:45 Room Oregon ballroom 202

- 8:30 Late Miocene biome turnover constrained body size evolution across Australian vertebrates Ian Brennan
- 8:45 Birth-death dynamics reveal how phylogeny and ecology shape mammalian vomerolfaction Laurel Yohe, Liliana Davalos
- 9:00 Multiple sequence alignment averaging improves phylogeny reconstruction Haim Ashkenazy, Itamar Sela, Eli Levy Karin, Giddy Landan, Tal Pupko
- 9:15 The evolution of biomechanical complexity during the global radiation of Strumigenys Douglas Booher, Evan Economo
- 9:30 Stochastic character mapping of state-dependent diversification reveals the tempo of evolutionary decline in self-compatible lineages

 Will Freyman

Session Sat1-Ball203 Social systems Saturday - 8:30 - 9:45 Room Oregon ballroom 203

- 8:30 Social norm choreography with tags Bryce Morsky, Erol Akcay
- 8:45 Social network position shapes nonlinear selection on weaponry in natural populations of forked fungus beetles

 Vincent Formica, David Tian, Butch Brodie
- 9:00 The foundress's dilemma: group selection for cooperation among queens of the harvester ant, Pogonomyrmex californicus

 Zachary Shaffer, Takao Sasaki, Brian Haney, Marco Janssen, Stephen Pratt, Jennifer Fewell
- 9:15 Coevolution of social behaviors and dynamic network structure *Erol Akcay*
- 9:30 What can social Isolation in mammals and insects teach us?

 Nadia Tsvetkov

Session Sat1-Ball204 ASN VP Symposium: The causes and consequences of evolutionary conflicts 1 Saturday - 8:30 - 9:45 Room Oregon ballroom 204

- 8:30 Introduction and some central ideas in evolutionary conflict Locke Rowe
- 8:45 Sexual conflicts: mechanisms and emerging themes Tracey Chapman
- 9:15 The role of phenotype plasticity in moderating evolutionary conflict Troy Day

Session Sat1-C120-122 Sensory systems

- 8:30 Color Preferences and Visual Tuning in Guppies: From Individuals to Populations Ben Sandkam, Mitchel Daniel, Karen Carleton, Helen Rodd, Felix Breden
- 8:45 Analyses of New World Vulture Olfactory Receptor Repertoires Reveals Dynamic Patterns of Evolution

 Joni E. Wright, Rebecca Kimball, J. Gordon Burleigh, Ed Braun
- 9:00 Evolutionary and Ecological Implications of Chromophore Usage in Rhodopsin Nihar Bhattacharya, Frances Hauser, James Morrow, Akimori Wada, Belinda Chang
- 9:15 What big eyes you have: Eye scaling and visual range in cephalopods Kate Thomas, Michael Vecchione, Sönke Johnsen
- 9:30 Spectral sensitivity via chromatic aberration-an alternative evolutionary path to color vision?

 Alexander Stubbs

Session Sat1-C123 SSE Education symposium Communicating (the relevance of) human evolution 1 Saturday - 8:30 - 9:45 Room C123

- 8:30 Why Communicating (The Relevance of) Human Evolution Still Matters James Smith
- 8:45 Using human examples to teach evolution in high school classrooms increases understanding and decreases cognitive biases and misconceptions

 Briana Pobiner, Bill Watson, Paul Beardsley, Constance Bertka
- 9:15 Evolution is not racist, sexist or anti-theist, and does not replace religion Holly Dunsworth

Session Sat1-C124 Expression studies / transcriptomics 1 Saturday - 8:30 - 9:45 Room C124

- 8:30 Regulatory evolution and the Island Rule Mark Nolte, Bret Payseur, Colin Dewey
- 8:45 Gene co-expression network connectivity is an important determinant of selective constraint
 Niklas Mähler, Jing Wang, Barbara K Terebieniec, Pär K Ingvarsson, Nathaniel R Street, Torgeir R
 Hvidsten
- 9:00 Using genetic variance of fitness and transcriptomics to estimate evolutionary potential in new environmental conditions

 Eva Koch, Frédéric Guillaume
- 9:15 Genes involved in musculature, metabolism, and the immune system are up-regulated in yellow-bellied marmots prior to dispersal

 Tiffany Armenta, Steven Cole, Robert Wayne, Daniel Blumstein
- 9:30 Contrasting gene expression programs correspond with predator-induced phenotypic plasticity within and across-generations in Daphnia.

 Nicole Hales, Drew Schield, Daren Card, Audra Andrew, Matthew Walsh, Todd Castoe

Session Evo101 Workshop for Science Educators Saturday - 8:30 - 16:00 Room C126

Prior registration required (separate from conference registration - see below). Intended for K-12 science educators. Questions? Contact Kristin Jenkins

Topic: Evolution can often seem very distant from students' lives, making it hard to engage them in studying this core biological concept. Human evolution offers compelling and relevant examples of evolutionary processes for students. At this hands-on workshop we will share free resources based on human evolution you can use in your classroom. The workshop is designed for middle and high school teachers. We encourage you to bring a colleague! Lunch and certificates of participation will be provided for participants.

- This workshop has its own registration process separate from the main conference. You don't need to register or attend the main conference to sign-up for this event.
- Schedule, registration and other information can be found here. Registration deadline: June 1, 2017

Supported by the <u>Society for the Study of Evolution</u> Education Committee and the <u>Howard Hughes Medical Institute</u>

Session Coffee break

Saturday - 9:45 - 10:15

Room Exhibit Hall A

Sponsored by:



Session Sat2-A105 Diversification

Saturday - 10:15 - 11:30

Room A105

10:15 Genomic analysis of a transposable element explosion in woodpeckers and allies (Aves: Piciformes)

Joseph Manthey, Stephane Boissinot

- 10:30 Structural genomic variation and the evolution of sugar accumulation in sweet sorghum Elizabeth Cooper, Stephen Kresovich, Christopher Saski
- 10:45 Sperm size evolution in passerine birds Jan T. Lifjeld
- 11:00 Population structure and hybrid zones in the tsetse fly Glossina fuscipes fuscipes (Diptera: Glossinidae) in Uganda: Implications for vector control
 Norah Saarman, Robert Opiro, Mary Burak, Serap Aksoy, Adalgisa Caccone
- 11:15 Visual signaling and morph discrimination in the rock-paper-scissors game Justin Yeager, Kinsey Brock, Danielle Edwards, Marie-Claire Chelini

Session Sat2-A106 Comparative phylogeography Saturday - 10:15 - 11:30 Room A106

- 10:15 Comparative phylogeography, accounting for lineage fusion. Ryan Garrick
- 10:30 Mitochondrial capture at the range edges of Enyalius lizards in the Atlantic Forests Roberta Damasceno, Miquel Rodriques
- 10:45 Comparative phylogeography of multiple Viburnum species based on RADseq data Elizabeth Spriggs, Deren Eaton, Yoshihisa Suyama, Tetsukazu Yahara, Erika Edwards, Michael Donoghue
- 11:00 Spatial and temporal congruence in regional genomic structure of Brazilian coastal fish communities

Andréa Thomaz, L. Lacey Knowles

11:15 Comparative adaptive genomics in the Brazilian Atlantic Forest: environmental and genomic clines across multiple species

Laura Bertola, Kyle McDonald, Mariana Vasconcellos, Nick Steiner, Ashfaq Khan, Roberta Damasceno, Ivan Prates, Natalia Quinteros, Ana Carnaval, Michael Hickerson

Session Sat2-A107-109 Species delimination Saturday - 10:15 - 11:30 Room A107-109

10:15 What the hellbender? Exploring species boundaries in North America's giant salamander.

Paul Hime, Steven Price, Shem Unger, Jeffrey Briggler, Amy McMillan, Michael Freake, Andrea Drayer,
Mary Foley, Emily Lemmon, David Weisrock

- 10:30 Integrated analyses of molecular, morphological, and bioacoustic data for species delimitation of an insular amphibian.

 Jim Labisko, Jeffrey Streicher
- 10:45 Species limits and relationships within tail-dropper slugs (genus Prophysaon) from the Pacific Northwest

 Megan Smith, Bryan Carstens, David Tank, Jack Sullivan, Megan Ruffley, Anahí Espíndola
- 11:00 Species Delimitation under the Multispecies Coalescent: Conflating Populations with Species in the Grey Zone

 Jeet Sukumaran, L. Lacey Knowles
- 11:15 Lineages vs species in macroevolution: an example from phylogenomics of lizards withdrawn 2017-06-23 00:01:05.0

 Craig Moritz

Session Sat2-B111-112 Phenotypic plasticity / GxE 2 Saturday - 10:15 - 11:30 Room B111-112

- 10:15 Transgenerational methylation variation induced by maternal vernalization in A. thaliana Mariano Alvarez, Gabriela Auge, Kathleen Donohue
- 10:30 Epigenetic signatures of parental effects in soil mites Stuart Piertney, Tim Benton, Marius Wenzel, Marianne Mugabo
- 10:45 Climatic gradients shape intraspecific divergence in plasticity for functional traits and gene expression in a South African Protea

 Melis Akman, Andrew M. Latimer, Jane E. Carlson
- 11:00 Transgenerational effects alter plant defense and resistance in nature Jack Colicchio
- 11:15 Genetic assimilation in the wild: evolution of gene expression plasticity during character displacement
 Nicholas Levis, Antonio Serrato, David Pfennig

Session Sat2-B113 Evolutionary theory 2 Saturday - 10:15 - 11:30 Room B113

- 10:15 Valley crossing in rapidly adapting sexual populations Taylor Kessinger, Jeremy Van Cleve
- 10:30 The effect of migration and environmental heterogeneity on the maintenance of standing genetic variation: a simulation study Tegan McDonald
- 10:45 Evolution of Pleiotropic Modules Jobran Chebib, Frédéric Guillaume
- 11:00 Fisher's geometrical model emerges as a property of complex integrated phenotypic networks GUILLAUME MARTIN

Session Sat2-B114-115 Conservation biology 2 Saturday - 10:15 - 11:30 Room B114-115

- 10:15 Global patterns of amphibian phylogenetic diversity across a gradient of human-modified environments
 - Dan Greenberg, Arne Mooers, Wendy Palen
- 10:30 Key factors influencing the reproductive success of natural- and hatchery-origin spring Chinook salmon

 Ilana Janowitz Koch, Maureen Hess, Craig Rabe, Shawn Narum
- 10:45 The importance of evolutionary significant lineages for conservation: lessons from a giant water bug

 Emily Hartfield Kirk
- 11:00 Importance of evolutionary conservation; lineage turnover and carbon storage in Madagascar Abagail Burrus, Sarah Federman, Alex Dornburg
- 11:15 Predictive methods for conservation prioritization

 Anahí Espíndola, Tara Pelletier, Bryan Carstens, Jack Sullivan, David Tank

Session Sat2-B116 Speciation 3

Saturday - 10:15 - 11:30

Room B116

- 10:15 Wax on wax off: tradeoffs between desiccation resistance and pheromone production in the Drosophila virilis clade

 Jeremy Davis, Leonie Moyle
- 10:30 Interspecific competition and temporal resource partitioning facilitate speciation and the formation of community biodiversity

 Glen Hood, Thomas Powell, Andrew Forbes, Scott Egan, Alice Harada, Jeffrey Feder
- 10:45 Reproductive isolation in an edaphic endemic monkeyflower species: Habitat selection in plants via manipulation of dormancy and dispersal James Sobel
- 11:00 Does ecological divergence accelerate premating reproductive isolation based on song in birds?

 Benjamin Freeman
- 11:15 A genomic and ecological view of (recent) speciation in the white stickleback Kieran Samuk, Hannah Visty, Dolph Schluter

Session Sat2-B117-119 Adaptation 2 Saturday - 10:15 - 11:30 Room B117-119

- 10:15 Genomic inference of selection on sperm proteins in the monarch butterfly Andrew Mongue, James Walters
- 10:30 Mutual selection pressures exerted by free-spawned gametes at fertilization Keyne Monro, Dustin Marshall
- 10:45 Transcriptomic evaluation of male reproductive adaptation in the cactus mouse (Peromyscus eremicus)

 Lauren Kordonowy, Matthew MacManes
- 11:00 Maternal Effects on Reproduction and Fitness in the Least Killifish, Heterandria formosa Samantha Levell, Joseph Travis, Samuel Bedgood
- 11:15 A local duplication of the melanocortin receptor 1 (Mc1r) locus in Astyanax fish Joshua Gross, James Weagley, Bethany Stahl, Li Ma, Luis Espinasa, Suzanne McGaugh

Session Sat2-Ball202 SSB Mayr Award Symposium 2 Saturday - 10:15 - 11:30 Room Oregon ballroom 202

- 10:15 Phylogeny, evolution, and the biogeographic history of Calandrinia (Montiaceae)

 Lillian Hancock, Abigail Moore, Jurriaan De Vos, Erika Edwards
- 10:30 TraitRateProp: An Integrated Model of Phenotypic Trait Changes and Site-Specific Sequence Evolution

 Eli Levy Karin, Susann Wicke, Tal Pupko, Itay Mayrose
- 10:45 Untangling the coevolutionary trees between New World ground-doves and their parasitic lice Andrew Sweet
- 11:00 Diagnosing Effects of Systematic Error and Phylogenetic Signal in Phylogenomics: A Case Study on the Order Cypriniformes

 Milton Tan, Jonathan Armbruster
- 11:15 Phenotypes linked to polyploidy: Revisiting Stebbins' 1938 hypothesis Rosana Zenil-Ferguson, Jose Miguel Ponciano, J. Gordon Burleigh

Session Sat2-Ball203 SSE Hamilton Award Symposium 1 Saturday - 10:15 - 11:30 Room Oregon ballroom 203

10:15 Rapid evolution of learning in natural populations of Drosophila melanogaster Emily Behrman, Tadeusz Kawecki, Paul Schmidt

- 10:30 The Reptile Gut Microbiome: Its Role in Host Evolution and Community Assembly Timothy Colston, Brice Noonan, Colin Jackson
- 10:45 The role of chromosomal inversions in life history divergence and reproductive isolation in Mimulus

 Jenn Coughlan, John Willis
- 11:00 Investigating performance costs incurred in exchange for adaptive resistance to tetrodotoxin in the Newt-Snake co-evolutionary arms Race

 Robert Eugene del Carlo, Jessica S. Reimche, Michael Hague, Edmund D. Brodie, Jr., Normand Leblanc, Chris Feldman
- 11:15 The evolution of parallel Arctic charr ecotypes via non-parallel evolutionary and genomic routes

Arne Jacobs, Kathryn Elmer, Natalia V. Gordeeva, Sergey S. Alekseyev, Colin E. Adams

Session Sat2-C120-122 Sensory systems / molecular evolution

Saturday - 10:15 - 11:30

Room C120-122

- 10:15 Photoreceptor transmutation in snakes and geckos Ryan Schott, Belinda Chang
- 10:30 The molecular evolution of sensory genes in neotropical bats Kalina Davies, Laurel Yohe, Elizabeth Dumont, Karen Sears, Liliana Davalos, Stephen Rossiter
- 10:45 Visual evolution in marine-derived Amazonian fishes
 Alexander Van Nynatten, Belinda Chang, Nathan Lovejoy
- 11:00 Convergent Cold-Adaptations of the Visual Pigment Rhodopsin at High Altitudes Gianni Castiglione, Nathan Lujan, Belinda Chang
- 11:15 Population differentiation and molecular evolution of pheromone communication in two orchid bee sibling species

 Philipp Brand, Thomas Eltz, Santiago Ramirez

Session Sat2-C123 SSE Education symposium Communicating (the relevance of) human evolution 2 Saturday - 10:15 - 11:30 Room C123

- 10:15 Teaching undergraduate life sciences majors Melissa Wilson Sayres
- 10:45 Panel discussion: Communicating (the relevance of) human evolution Norman Johnson

Session Sat2-C124 Expression studies / transcriptomics 2 Saturday - 10:15 - 11:30 Room C124

- 10:15 Evolutionary lability of vernalization responsiveness in Pooideae grasses

 Jinshun Zhong, Meghan Robbett, Siri Fjelheim, Jill Preston
- 10:30 Dosage Compensation in Yeast withdrawn 2017-06-09 10:32:10.0 Holly McQueary
- 10:45 Gene expression evolution in parallel cold-adapted Drosophila populations Yuheng Huang, John Pool
- 11:00 Differential responses under future ocean acidification between populations of Balanophyllia elegans corals from different upwelling environments

 Joanna Griffiths, Morgan Kelly
- 11:15 Extreme evolution: crustacean vision in a land-to-deep-sea gradient. Hernan Vazquez Miranda, Heather Bracken-Grissom

Session Sat2-Ball204 ASN VP Symposium: The causes and consequences of evolutionary conflicts 2 Saturday - 10:15 - 11:45 Room Oregon ballroom 204

- 10:15 The Causes and Consequences of Selection at the Gametic Level Simone Immler
- 10:45 War and Peace: The Invasion, Proliferation, and (Occasional) Domestication of Mobile DNA in Host Genomes

 Sarah Schaack, Michael Song
- 11:15 Socio-reproductive conflicts and the father's curse of hormones Mikael Mokkonen, Tanya Procyshyn, Bernard Crespi

Session Evolution editorial board meeting Saturday - 11:30 - 13:30 Room B110

Session Lunch / Student and PDF networking lunches Saturday - 11:30 - 13:30 Room On the town

Conference lunch is on your own.

The SSE Graduate Student Advisory Council organized small group networking lunches to facilitate interactions between i) students and faculty; and ii) postdocs and faculty. To participate, you had to indicate your interest during main conference registration. Groups are responsible for finding their own lunch venues.

Session Early career researchers workshop (1 of 2): Publishing your research: tips for success (Taylor & Francis) Saturday - 12:15 - 13:15 Room B113

Cost: free, no registration required. Bring your own lunch. Organizer: Ashlynne Merrifield, Ali Paskins, Andrew Polaszek, and Jeffrey Streicher

This talk will guide new authors through the publishing process from selecting a journal, to preparing for submission, to navigating the peer review process. Andrew Polaszek (Editor-in-Chief, JNH) and Jeffrey Streicher (Associate Editor, JNH) will give you a valuable insight into the editorial process for Journal of Natural History, including advice for successful publication. We will also cover essential tips for writing a review article, how to respond to reviewer comments, acting as a reviewer, plus more, giving a thorough understanding of the steps involved, the key information sources to be aware of, and what authors can be doing to help get their paper published.

Session Sat3-Ball204 ASN Symposium - Across the Nth dimension: Quantitative and conceptual advances in the study of niche breadth 1

Saturday - 13:20 - 14:45

Room Oregon ballroom 204

- 13:20 Introductory Remarks: Exploring the N Dimensions of Niche Breadth Nancy Emery
- 13:30 On the history and current status of niche concepts: Implications for niche breadth research Robert Holt
- 14:00 Mechanisms of niche breadth 1: Adaptive evolution and the nested niche. Copepod thermal tolerance and evolutionary potential in the face of warming waters.

 Morgan Kelly

14:15 Mechanisms of niche breadth 2: Plasticity and the nested niche. Ecological causes and consequences of individual-level specialization.

Richard Svanbäck

Session Sat3-A105 Phylogenetic comparative methods 1 Saturday - 13:30 - 14:45 Room A105

- 13:30 Morphological convergence among grassland-adapted clades of South American ghost spiders Nicolas Mongiardino Koch, F. Sara Ceccarelli, Eduardo Soto, Martin J. Ramirez
- 13:45 Continuously-varying adaptive landscapes over macroevolutionary timescales Josef Uyeda
- 14:00 Rapid adaptive shifts are a common mode of macroevolution Michael Landis, Joshua Schraiber
- 14:15 A Penalized Likelihood Framework For High Dimensional Phylogenetic Comparative Methods Julien Clavel, Hélène Morlon
- 14:30 A Bayesian Method for Inferring Discrete-trait-dependent Rates of Continuous-trait Evolution Michael May, Brian Moore

Session Sat3-A106 Comparative phylogeography / biogeography

Saturday - 13:30 - 14:45

Room A106

- 13:30 An Anthropocene map of genetic diversity
 - **Katharine Marske**, Andreia Miraldo, Sen Li, Michael Borregaard, Alexander Florez-Rodriquez, Shyam Gopalakrishnan, Mirnesa Rizvanovic, Zhiheng Wang, Carsten Rahbek, David Nogues-Bravo
- 13:45 Beyond topography: The role of rain asynchrony on the diversification of tropical frogs *Carlos Guarnizo*, Paola Montoya, Ignacio Quintero, Daniel Cadena
- 14:00 Biogeographic origins of the frog fauna on the semi-arid Caribbean Coast of South America withdrawn 2017-06-08 12:03:17.0

Andrew J. Crawford

- 14:15 Integrating island assembly models and comparative population genetics.

 Isaac Overcast, Michael Hickerson
- 14:30 Intraspecific genetic relationships of two coastal species (red and white mangroves) reveal broad phylogeographic patterns in the Caribbean Richard Hodel, Pamela Soltis, Douglas Soltis

Session Sat3-A107-109 Species delimination / speciation Saturday - 13:30 - 14:45 Room A107-109

- 13:30 Linking floral traits, pollination, and speciation in western Camassia: How important are geography and ecological contexts?
 - **Susan Kephart**, Theresa Culley, Jenny Archibald, Kathryn Theiss, Robert Tunison, Sebastian Mortimer, James Kephart
- 13:45 Using a classic speciation data set to test coalescent methods for species delimitation Luke Campillo, Robert Thomson, Anthony Barley
- 14:00 The mitonuclear compatibility species concept Geoffrey Hill
- 14:15 Caught in the act: incipient speciation in a fossorial mammal from Madagascar, the mole tenrec Oryzorictes hova (Tenrecidae)

 Kathryn Everson, Kyndall B.P. Hildebrandt, Link Olson
- 14:30 Testing the tri-trophic speciation patterns of parasitoid wasps in Peristenus pallipes complex (Hymenoptera: Braconidae)

 Miles Zhang, Amber Bass, Catalina Fernández, Barbara Sharanowski

Session Sat3-B111-112 Phenotypic plasticity / quantitative genetics

Saturday - 13:30 - 14:45

Room B111-112

- 13:30 Genetic basis of total energy budget in different nutrient environments Anna Perinchery, Elizabeth King
- 13:45 Testing the Importance of Phenotypic Plasticity in Facilitating Survival in a Novel Environment Shengpei Wang
- 14:00 Multiple forms of selection act on trait phenotypes and plasticities in nature: evidence from a manipulative field experiment.

 Freddy Herrera, Elizabeth Lacey
- 14:15 When three traits make a line: Evolution of phenotypic plasticity and genetic assimilation through linear reaction norms in stochastic environments

 Torbjørn Ergon, Rolf Ergon
- 14:30 The genetic basis of variation in phenotypic plasticity Karin van der Burg, Robert Reed

Session Sat3-B113 Evolutionary theory 3 Saturday - 13:30 - 14:45 Room B113

- 13:30 Why is cytoplasmic incompatibility so common?

 Michael Turelli
- 13:45 Gene network drift and speciation Joshua Schiffman, Peter Ralph
- 14:00 Excess of genomic defects in a woolly mammoth on Wrangel island Rebekah Rogers, Montgomery Slatkin
- 14:15 Fitness landscapes of drug interactions Bjørn Østman, Tina Kang, Van Savage, Pamela Yeh

Session Sat3-B114-115 Conservation biology 3 Saturday - 13:30 - 14:45 Room B114-115

13:30 Phylogeography of two species of minnow from the eastern Oregon desert withdrawn 2017-06-19 10:51:47.0

Christian Smith, Jennifer Von Bargen, Patrick DeHaan, Paul Scheerer

- 13:45 Phylogenetic Assessments for California's Highest Risk Mammals Christopher Jorgensen, Joshua Reece
- 14:00 Using RAD data to inform the population and conservation strategies of Melipona subnitida (Apidae, Meliponin), an endemic bee in dry forest of Brazil

 Vanessa Bonatti, Gregory Thom, Sonia Andrade, Maria Cristina Arias, Michael Hickerson, Tiago
 Francov
- 14:15 Climate drives adaptive genetic responses associated with survival in big sagebrush (Artemisia tridentata)

 Lindsay Chaney, Bryce Richarson, Mathew Germino
- 14:30 Biologically informed niche models predict slightly less dire future for mountaintop plethodontid salamanders.

 Marta Lyons, Ken Kozak

Session Sat3-B116 Speciation 2

Saturday - 13:30 - 14:45

Room B116

13:30 Geographic variation in New Guinean paradise kingfishers: the classic case of peripatric speciation

Gregory Mayer

- 13:45 Does size matter? Giant sperm in Caenorhabditis worms Rebecca Schalkowski
- 14:00 A model for coexistence and divergence in insect-vectored yeast populations Kelly Thomasson
- 14:15 "Fishing" for vertebrate fertilization genes: the evolution of threespine stickleback reproductive proteins

 Emily Killingbeck, Damien Wilburn, Gennifer Merrihew, Michael MacCoss, Catherine Peichel, Willie Swanson
- 14:30 Speciation in the Tropical Eastern Pacific? A Case Study with Prionurus Surgeonfishes William Ludt, Moises Bernal, Eva Salas, Prosanta Chakrabarty

Session Sat3-B117-119 Adaptation 3 Saturday - 13:30 - 14:45 Room B117-119

- 13:30 Effects of gene flow on performance at the northern range margin of Clarkia pulchella Megan Bontrager, Amy Angert
- 13:45 Functional divergence of the dim-light opsin accompanies cichlid invasion of Central America Frances Hauser, Katriina Ilves, Gianni Castiglione, Ryan Schott, Hernan Lopez-Fernandez, Belinda Chang
- 14:00 Molecular evolution of cold response in temperate grasses (Pooideae)

 Marian Schubert, Lars Grønvold, Simen Sandve, Torgeir R Hvidsten, Siri Fjelheim
- 14:15 Life at high latitudes: the role of photoperiodic flowering in diversification of the grass subfamily Pooideae in northern temperate regions

 Siri Fjelheim, Jill Preston
- 14:30 Non-adaptive plasticity promotes adaptation to high-elevation in *Peromyscus* mice *Jonathan Velotta*, *Zachary Cheviron*, *Graham Scott*

Session Sat3-Ball202 Host-parasite Saturday - 13:30 - 14:45 Room Oregon ballroom 202

- 13:30 Microbial-Host Dynamics and the Evolution of Nearshore Diversity Jimiane Ashe
- 13:45 Avian Haemosporidian Diversification: from hosts to habitats Johanna Harvey, Gary Voelker
- 14:00 Infrapopulation size and composition influence reproductive success of a parasitic plant *Kelsey Yule*, *Judith Bronstein*
- 14:15 Predictors of the correlation between sexual reproduction and infection: a meta-analysis of the Red Queen Hypothesis

 Meredith Krause, Marc Lajeunesse
- 14:30 Mating biology, cheating behavior, and the evolution of social parasitism in ants Romain Dahan, Ben Gerstner, Martin Bollazzi, Christian Rabeling

Session Sat3-Ball203 SSE Hamilton Award Symposium 2 Saturday - 13:30 - 14:45 Room Oregon ballroom 203

- 13:30 Immune and antioxidant performance of carotenoid-free canaries questions a longstanding hypothesis for the evolution and maintenance of colored signals

 Rebecca Koch, Geoffrey Hill*
- 13:45 Direct evidence that metabolic genes underlie temperate adaptation of Drosophila melanogaster life history Spencer Koury
- 14:00 The origin and evolution of polymorphic warning signals

 J.P. Lawrence, Bibiana Rojas, Antoine Fouquet, Johanna Mappes, Annelise Blanchette, Ralph Saporito,
 Renan Bosque, Elodie Courtois, Brice Noonan
- 14:15 Beyond topology: Coevolution of structure and flux in metabolic networks Erin Morrison, Alexander Badyaev

14:30 The extent of parallelism in independent lineages in shared environments: Migration phenology of even-year and odd-year pink salmon

Krista Oke, Curry Cunningham, Thomas Quinn, Gregor Rolshausen, Caroline LeBlond, Andrew Hendry

Session Sat3-C120-122 Sexual selection 1 Saturday - 13:30 - 14:45 Room C120-122

- 13:30 Are we wrong about territoriality in Anolis lizards?

 Ambika Kamath, Jonathan Losos
- 13:45 A sexual selection model of mate location: a foundational, but missing, mechanism of mate recognition evolution

 Kerry Shaw, H. Kern Reeve
- 14:00 A "sexy" co-evolutionary arms race: structural insights into the rapid evolution of egg-sperm interactions in Pacific abalone

 Damien Wilburn, Willie Swanson
- 14:15 Beyond resource trade-offs: An alternative mechanism for maintaining honesty from colorful ornaments used in mate choice

 Ryan Weaver, Rebecca Koch, Geoffrey Hill
- 14:30 Evolutionary ecology of sexual communication in Bahamian mosquitofish Sean Giery

Session Sat3-C123 ASN Education symposium Natural history as the inspiration for scientific inquiry: Stories and tools for teaching 1

Saturday - 13:30 - 14:45

Room C123

- 13:30 Natural History's place in Science and Society

 Josh Tewksbury
- 13:45 Killifish and sensory drive in the active classroom Becky Fuller
- 14:00 Using the extraordinary natural history of manakin courtship to teach fundamental concepts in evolution and behavior Emily DuVal, Kevin Dixon, Charles Eldermire
- 14:15 Born to run: inquiry-based learning about experimental evolution of voluntary exercise in mice
 Theodore Garland
- 14:30 "There are plenty of fish in the sea"—A scientific guide for finding the perfect match for each of our unique research interests *Molly Cummings*

Session Sat3-C124 Expression studies / molecular evolution Saturday - 13:30 - 14:45 Room C124

- 13:30 Cis-regulatory variation underlying population differences in HSPB1 expression and thermal tolerance in the copepod Tigriopus californicus

 Sumaetee Tangwancharoen, Ron Burton
- 13:45 Gene expression dynamics are modified at the protein level by natural genetic variation in the Saccharomyces cerevisiae population

 Daniel Pollard, Ciara Asamoto, Homa Rahnamoun, Austin Abendroth, Suzanne Lee, Scott Rifkin
- 14:00 Differential gene expression among developmental stages characterize genes implicated in sex determination of the cichlid fish Astatotilapia burtoni

 Athimed El Taher, Nicolas Lichilin, Geoffrey Fucile, Walter Salzburger, Astrid Boehne
- 14:15 The transcriptional response of microbes to long-term warming and short-term seasonal fluctuations in a terrestrial forest

 Priyanka Roy Chowdhury, Kristen DeAngelis

14:30 Copy number variation and a role in vision of CRAL-TRIO domain genes in Heliconius melpomene

Aide Macias-Muñoz, Kyle McCulloch, Adriana Briscoe

Session Coffee break

Saturday - 14:45 - 15:15

Room Exhibit Hall A

Sponsored by:



Session Sat4-A105 Phylogenetic theory Saturday - 15:15 - 16:30 Room A105

- 15:30 Climate niche variation of Hawaiian Psychotria in an endemic island diversification Elaine Zhang, John Paul
- 15:45 Are Bryophytes monophyletic? Evidence from Organellar and Nuclear Protein-Coding Genes Filipe Sousa, Cymon Cox
- 16:00 Assessing Combinability of Phylogenomic Data using Bayes Factors
 Suman Neupane, Karolina Fučíková, Louise A. Lewis, Lynn Kuo, Ming-Hui Chen, Paul Lewis
- 16:15 The probability of monophyly of a sample of gene lineages on a species tree. Rohan Mehta, David Bryant, Noah Rosenberg

Session Sat4-A106 Biogeography / phylogeography Saturday - 15:15 - 16:30 Room A106

- 15:15 Phylogeography, Biogeography, and the Origins of Indo-Pacific Reef Fishes Brian Bowen
- 15:30 A phylogeographic study of birds supports that the Andean-Atlantic forests connection occurred through the Cerrado

 Gustavo Sebastian Cabanne, Leonardo Campagna, Natalia Trujillo, Irby Lovette, Pablo Tubaro
- 15:45 A temperate armadillo: a genomic portrait of the nine-banded armadillo's northward migration

 Lucy Tran, Lydia Smith
- 16:00 Phylogenetic biogeography of fossil Canidae: combining models of imperfect detection, fossil databases as taphonomic controls, and Biogeographical Stochastic Mapping Nicholas Matzke, Laura Säilä
- 16:15 Model selection in historical biogeography: why DEC and DEC+J likelihoods are not statistically comparable Richard Ree, Isabel Sanmartín

Session Sat4-A107-109 Macroevolution 1
Saturday - 15:15 - 16:30 Room A107-109

- 15:15 The role of diversification rates on the assembly of frugivory networks.

 Gustavo Burin, Paulo Roberto Guimaraes Jr, Tiago Quental
- 15:30 Elaborating the macroevolutionary dynamics of ants-plant interactions Matthew Nelsen, Corrie Moreau, Richard Ree
- 15:45 Predictors of secondary sympatry across the global radiation of extant bats Jeff Shi, Erin Westeen, Elizabeth Dumont, Daniel Rabosky
- 16:00 Mutualism breakdown and ecological opportunity in leafflower-leafflower moth interactions David Hembry, Noah Whiteman, Katrina Dlugosch
- 16:15 Macroevolution of perfume signalling in orchid bees Marjorie Weber, Santiago Ramirez, Thomas Eltz

Session Sat4-B111-112 Quantitative genetics Saturday - 15:15 - 16:30 Room B111-112

- 15:15 Genetic costs of domestication and improvement Brook Moyers, Peter Morrell, John McKay
- 15:30 Predicting evolutionary responses when genetic variance and selection covary with the environment: a large-scale Open Access Data approach

 Jip Ramakers, Phillip Gienapp, Antica Culina, Marcel E Visser
- 15:45 Genetic constraints on wing pattern variation in Lycaeides butterflies Lauren Lucas, Chris Nice, Zachariah Gompert
- 16:00 Observational evidence of directional and stabilizing selection in a contemporary human population

 Jaleal Sanjak, Kevin Thornton, Matthew Robinson, Peter Visscher
- 16:15 Indirect genetic effects, resource competition, and a potential evolutionary constraint in North American red squirrels

 David Fisher, Andrew McAdam*, Alastair Wilson*

Session Sat4-B113 Evolutionary theory 4 Saturday - 15:15 - 16:30 Room B113

- 15:15 Evolution and the duration of a doomed population Richard Gomulkiewicz, Stephen Krone, Christopher Remien
- 15:30 Role of Sustainability in Evolution PANKAJ VALLABH
- 15:45 Modeling How Selection in One Trait Clonally Interferes with Adaptation in Another Kevin Gomez, Jason Bertram, Joanna Masel
- 16:00 When do fluctuating environments select for epigenetic memory? Stephen Proulx, Henrique Teotonio
- 16:15 Epigenetic change increases local adaptation in small, poorly adapted populations Mark Smithson, Mark Dybdahl, Scott Nuismer

Session Sat4-B114-115 Conservation biology / population genetics

Saturday - 15:15 - 16:30

Room B114-115

- 15:15 Population Genetics and Adaptive Variation within a Cryptic and Threatened Hawaiian Seabird withdrawn 2017-06-08 00:03:54.0

 Carmen Antaky, Melissa Price
- 15:30 Conservation genetics of a rare species of beardtongue: Penstemon caryi (Plantaginaceae)

 Benjamin Stone, Andrea Wolfe, Alexander Ward
- 15:45 Old school vs. new school: comparing inferences of population structure using RADseq and microsatellite datasets

 Justin Bohling

16:00 Population genomics of Oenothera primiveris, insights into plant conservation and role of inbreeding.

Anita Cisternas Fuentes, Rick Overson, Jeremie Fant

16:15 Genome-wide historical DNA analysis reveals insights into population genetic theory and conservation practice

Evelyn Jensen, Adalgisa Caccone, Michael Russello

Session Sat4-B116 Speciation / adaptation Saturday - 15:15 - 16:30 Room B116

- 15:15 Repeated dune adaptation and speciation in sunflowers Kate Ostevik, Rose Andrew, Loren Rieseberg
- 15:30 Speciation and adaptive diversification within the box-ironbark eucalypts

 Jasmine Janes, Kevin Murray, Norman Warthmann, Jamieson Gorrell, Timothy Collins, Rebecca

 Jordan, Jeremy Bruhl, Justin Borevitz, Rose Andrew
- 15:45 Across the tipping point? The transition from host race to species in the Rhagoletis pomonella species complex

 Meredith Doellman, Thomas Powell, Stewart Berlocher, Gregory Ragland, Scott Egan, Peter Meyers, Glen Hood, Jeffrey Feder
- 16:00 Repeated patterns of adaptation along the speciation continuum: insight from studies of North American rattlesnakes

 Drew Schield, Daren Card, Richard Adams, Giulia Irene Maria Pasquesi, Blair Perry, Nicole Hales, Audra Andrew, Andrew Corbin, Stephen Mackessy, Todd Castoe

16:15 Hybridization maintains of ecological divergence between Pinus strobiformis and Pinus

flexilis Mitra Menon, Chris Friedline, Anna Schoettle, Sam Cushman, Daniel McGarvey, Kristen Waring, Amy Whipple, Richard Sniezko, Lluvia Flores-Renteria, Andrew Eckert, Justin Bagley, Christian Wehenkel, Alejandro Sáenz

Session Sat4-B117-119 Adaptation / expression studies Saturday - 15:15 - 16:30 Room B117-119

- 15:15 Complexities of gene expression patterns in natural populations of an extremophile fish (Poecilia mexicana, Poeciliidae)

 Courtney Passow, Joanna Kelley, Michael Tobler
- 15:30 Environmental adaptation in House mice: the role of gene regulation along a latitudinal cline Katya Mack, Mallory Ballinger, Megan Phifer-Rixey, Michael Nachman
- 15:45 Ecological genomics of rapidly differentiating exotic horned beetle populations Melissa Pespeni, Armin Moczek
- 16:00 A master regulator of oxygen homeostasis contributes to adaptive hypoxia signaling in deer mice.

Zachary Cheviron, Jonathan Velotta, Rena Schweizer, Catie Ivy, Graham Scott

16:15 Transcriptome and whole genome sequencing promise to reveal the genomic substrate of a textbook adaptive radiation

Marylaure de La Harpe, Margot Paris, Hess Jaqueline, Michael Barfuss, Walter Till, Nicolas Salamin, Michael Kessler, Ray Ming, Christian Lexer

Session Sat4-Ball202 Host-parasite / pathogens Saturday - 15:15 - 16:30 Room Oregon ballroom 202

15:15 Balancing Selection on a Cis-Regulatory Region Maintains Disease-Associated Genetic Variation in a Wild Primate

Noah Simons, Geeta Eick, Maria Jose Ruiz Lopez, Colin Chapman, Tony Goldberg, Nelson Ting, Kirstin Sterner

- 15:30 Natural selection on immune function Otto Seppälä
- 15:45 Contagious Clam Cancer: Evolution at the intersection of cancer and infectious disease Michael Metzger, Antonio Villalba, María Carballal, David Iglesias, James Sherry, Carol Reinisch, Annette Muttray, Susan Baldwin, Stephen Goff

16:00 Evolution under competition: signatures of strain structure within the major blood-stage antigen of the malaria parasite P. falciparum

Mary Rorick, Yael Artzy-Randrup, Shazia Ruybal-Pesántez, Kathryn Tiedje, Thomas Rask, Abraham Oduro, Anita Ghansah, Kwadwo Koram, Karen Day, Mercedes Pascual

16:15 Does phylogeny predict what ails you? – A pollinator community study David Pascall. Lena Wilfert

Session Sat4-Ball203 SSE Hamilton Award Symposium 3 Saturday - 15:15 - 16:30 Room Oregon ballroom 203

15:15 Negative genetic interactions among adaptive mutations in yeast turn positive in a harsher environment

Jasmine Ono, Aleeza Gerstein, Sally Otto

- 15:30 Misaligned interests between plants and pollinators revealed through functional exploration of flower morphospace

 Foen Peng, Toby Bradshaw, Tom Daniel
- 15:45 Microsatellites as "tuning knobs" of adaptation in Helianthus annuus (Asteraceae)
 Chathurani Ranathunge, Gregory Wheeler, Melody Chimahusky, Andy Perkins, Mark Welch
- 16:00 Selection against LTR retrotransposons is balanced by locally adapted transposable element alleles in Arabidopsis thaliana

 Michelle Stitzer, Jeffrey Ross-Ibarra
- 16:15 There is no cure that does not cost: fitness trade-offs and putative mechanisms of resistance to bovine tuberculosis in a wild mammal Hannah Tavalire, Anna Jolles

Session Sat4-C120-122 Sexual selection 2 Saturday - 15:15 - 16:30 Room C120-122

15:15 Gene Expression and Reduced Sexual Dimorphism in Threespine Stickleback Throat Coloration

Jeffrey McKinnon, Chris Balakrishnan, Burns Newsome

- 15:30 The non-sexual origins of sexual preferences and ornaments in live-bearing fishes Michael Foisy, Jeremy Beaulieu, Luke Frishkoff, Mark Springer, David Reznick, Arcadio Gonzalez, Helen Rodd, D. Luke Mahler
- 15:45 The influence of resource limitation on male reproductive investment Josiah Ritchey
- 16:00 Consequences of inbreeding and environmental stress on a species with sexually selected weaponry

 Pablo Allen, Christine Miller

Session Sat4-C123 ASN Education symposium Natural history as the inspiration for scientific inquiry: Stories and tools for teaching 2

Saturday - 15:15 - 16:30

Room C123

- 15:15 Teaching natural history in the Anthropocene: some rules of engagement Harry Greene
- 15:30 Using electron-mediated mutualism to teach from molecule to community Kristina Hillesland
- 15:45 "Ordinary, extraordinary junco: bringing field and evolutionary biology to the classroom" Ellen Ketterson, Jonathan Atwell
- 16:00 Teaching animal physiology from physical principles Chris Cheng
- 16:15 The unnatural concept of extinction in species-area relationships and teaching conservation biology

 Liliana Davalos, M. Elise Lauterbur, Anusha Shankar

Session Sat4-C124 Expression studies / hybridization Saturday - 15:15 - 16:30 Room C124

- 15:15 Variation of mitochondrial function in hybrid Tree lizards: assessing the role of differential gene expression

 Greg Haenel
- 15:30 Do regulatory differences maintain the phenotypic boundary between sister species of Streamertail Hummingbirds (Trochilus polytmus & T. scitulus)?

 Caroline Judy, Robb Brumfield, Gary R. Graves
- 15:45 Sex differences in mouse hybrids: Differential expression and morphology in placenta Lena Arévalo, Polly Campbell
- 16:00 Introgressive hybridization among Virroconus Andrew Wood, Thomas Duda
- 16:15 RNA-Seq reveals unexpected patterns of endosperm imprinting in interploid Mimulus hybrids Taliesin Kinser, Ronald Smith, Mario Vallejo-Marin, Joshua Puzey

Session Sat4-Ball204 ASN Symposium - Across the Nth dimension: Quantitative and conceptual advances in the study of niche breadth 2

Saturday - 15:15 - 16:45

Room Oregon ballroom 204

- 15:15 Does niche breadth predict vulnerability to changing environments? From population-level traits and demography to diversification in deep time

 Seema Sheth
- 15:45 Defining niche breadth through statistical modelling A historical perspective Guillaume Blanchet
- 16:15 A guide to quantifying niche breadth: classifying and comparing niche breadth metrics Kelly Carscadden, Carlos Alberto Arnillas, Nancy Emery, Marc Cadotte, Dominique Gravel, Michelle Afkhami, J. Scott MacIvor

Session American Society of Naturalists 'Happy Hour' and graduate student networking

Saturday - 16:30 - 17:30 Room Oregon ballroom 201

Open to all ASN members - students, postdocs & faculty. Come chat with your ASN colleagues. Cash bar and ASN grad member free drink tickets accepted.

Session Sat5-Ball202 ASN Presidential address & award announcements

Saturday - 17:30 - 18:30 Room Oregon ballroom 202-204

ASN Presidential Address Kathleen Donohue

Session ASN Members business meeting

Saturday - 18:30 - 19:30

Room A105

Open to all ASN members.

Poster Session Sat-01 Adaptation Saturday - 18:30 - 20:30 Room EHA-01

- 1 Dynamics of seasonal adaptation in Drosophila melanogaster Emily Behrman, Dmitri Petrov, Paul Schmidt
- 2 The influence of lake productivity on life history evolution in Daphnia: natural vs. experimental tests

 Michelle Packer, Matthew Walsh
- **Feeling the Squeeze: Adaptation to High Hydrostatic Pressure in Ctenophore Proteomes** *Jacob Winnikoff, Warren Francis, Steven Haddock*
- 4 Local adaptation in leeches: Estimating the relative fitness of three northern Arizona Helobdella modesta populations in natal and non-natal water Rebecca Beresic-Perrins, James Boothroyd, Angela Rincon, Cassidy Klovanish, Stephen Shuster
- 5 Evaluating the role of growth defense trade-offs in the evolution of locally adapted ecotypes across the Mimulus guttatus species complex Damian Popovic

Topic Sat-02 Behavior

- 6 Feeding and neural mechanisms underlying the predatory behavior of a model aquatic invertebrate

 Elizabeth Preza, Elizabeth J. Walsh
- 7 Do you see what I see? The Evolution of Visual Signals Across Spotted Winged Drosophila Species Groups
 Paula Roy, Jennifer Gleason
- 8 Behavioral, neuroendocrine, and reproductive implications of exposure to an endocrine disrupting pesticide in Gambusia affinis.

 Isaac Ligocki, Amelia Munson, Victoria Farrar, Rebecca Calisi
- 9 Reproductive isolation driven by pheromones in mimetic and closely related butterflies Carolina Pardo-Diaz, Maria Fernanda Gonzalez, Camilo Salazar, Chris Jiggins, Kathy Darragh
- Foraging Rates of a Freshwater Drift-Feeding Salmonid, Salvelinus malma, Under a Dominance Hierarchy withdrawn 2017-06-24 10:26:28.0
 Jeff Cullen
- 11 Habitat Utilization Preferences in Potamopyrgus antipodarum, a Freshwater Snail from New Zealand
 Colette Ohotnicky, Deanna Soper

Topic Sat-03 Biogeography

- 12 Biodiversity and distribution of hydroids (Cnidaria, Hydrozoa) in southern South America and Antarctica: an approach using marine ecoregions and provinces

 Thais Miranda, Marina Fernandez, Antonio Marques
- 13 Dispersal, not vicariance, explains the biogeographic origin of Madagascar's endemic iguanas (Opluridae)

 Rachel Welt, Christopher Raxworthy
- 14 The temporal puzzle of North American disjunct plants Chloe Drummond, Kenneth Sytsma
- 15 Phylogenetic and Biogeographic History of the Conifer Family Araucariaceae inferred from Chloroplast Genome Sequences
 Linda Raubeson, Mei Wenbin, Markus Ruhsam, Sarah Mathews
- Panmixia across the range of the assortatively mating Lesser Snow Goose withdrawn 2017-06-19 22:24:09.0

 Tom Ouinn, Brent Horowitz

Topic Sat-04 Bioinformatics

- 17 De novo detection of transposable elements in high-throughput sequence data. Laura Helou, Joanna Kelley, Anna-Sophie Fiston-Lavier
- 18 High Quality Phylogenomic Data Through Accurate Sorting of Anchored MetaPrep Reads Sean Holland
- 19 A Pipeline for Rapid and Inexpensive Processing of Anchored MetaPrep Data Kirby Birch, Alan Lemmon

Topic Sat-05 Coevolution

- 20 Coevolutionary Forces Reveal an Alternative Origin for Dim-Light Vision Gianni Castiglione, Belinda Chang
- 21 High-throughput investigation of species-specific binding of abalone reproductive proteins Jolie Carlisle, Willie Swanson, Damien Wilburn
- 22 Gene-culture Price equation for the evolution of behavior. *Elliot Aguilar*, *Erol Akcay*
- 23 Apoptotic machinery: An ancient adaptation to aerobic conditions

 Joanna Klim, Arkadiusz Gladki, Roza Kucharczyk, Urszula Zielenkiewicz, Szymon Kaczanowski
- 24 Preference driven Co-evolution of Drosophila melanogaster and Saccharomyces cerevisiae

 Janine Fischer, Amanda Thompson, Kelly Thomasson, Stephen Proulx
- 25 Genetic analysis of host immunity and microbiota variation in threespine stickleback Emily Beck, Clayton Small, Mark Currey, William Cresko

Topic Sat-06 Comparative biology

- 26 Evolutionary origin and functional impact of the GHR deletion Hannah Calkins
- 27 Evolution of swimming in Neotropical tree frogs (Anura: Hylidae) *Itzue Caviedes-Solis*
- 28 Three-dimensional analysis of craniofacial traits across distinct cavefish populations using micro-computed tomography (micro-CT)

 Daniel Berning, Amanda Powers, Joshua Gross, Patricia Ornelas-García
- 29 Fundulus shape and osmoregulation efficiency Jason Pienaar, Ryan Earley, Joe Styga, Peter Scott
- 30 Comparision of skin in two european species of lizards (Lacerta agilis and Anguis fragilis) withdrawn 2017-06-16 06:15:20.0

 Paweł Kaczmarek
- 31 Incorporating the logic of anatomical homology in a gene-phenotype knowledgebase: Phenoscape KB

 Wasila Dahdul, James Balhoff, Hilmar Lapp, Todd Vision, Paula Mabee

Topic Sat-07 Comparative phylogeography

- 32 Phylogeographic comparisons of Leptasterias clades near estuarine outflow along the Pacific Coast

 Jeyna Perez, C. Sarah Cohen
- 33 Rivers as barriers across space, time and species: dissecting the role of river barriers using tests of concordant divergence in Amazonian species

 Renata Pirani, Fernanda Werneck, Andréa Thomaz, Mariah Kenney, Marcelo Sturaro, Teresa Avila-Pires, Miguel Rodrigues, L. Lacey Knowles

- 34 Drivers of Phylogenetic Endemism and the Conservation of Centres of Endemism in the Coastal Forests of Eastern Africa
 Simon Loader, Chris Barratt
- 35 Improving the predictive power of comparative phylogeographic tests using refined hypotheses

 L. Lacey Knowles

Topic Sat-08 Computational biology

- 36 The coexistence of competing mathematical and natural solutions to the 'paradox of the plankton': does the paradox still exist? withdrawn 2017-05-31 21:13:58.0

 Richard Knecht
- 37 genotypeR: an integrated R package for SNP genotype marker design and data analysis Stephen Sefick, Magdalena Castronova, Laurie Stevison
- 38 Between a rock and a hard place: climate change and limited distribution of the California native plant Arabis blepharophylla (Brassicaceae)

 Nila Le, John Paul
- 39 Evolutionary Conserved Alternative Splicing in the Rice Genus Oryza Nathan Catlin, Rod Wing, W. Brad Barbazuk

Topic Sat-09 Conservation biology

- 40 Population genetics and adaptive variation within a cryptic and threatened Hawaiian seabird Carmen Antaky, Melissa Price
- 41 Genetic structure of the water rat Nectomys squamipes in a fragmented habitat of the Brazilian Atlantic forest Hudson Lemos, Luana Maroja, Pablo Gonçalves
- 42 Does habitat specific seed sources result in differential success of Chamaecrista fasciculata in a restored tallgrass prairie?

 Robert Philips, Diane Byers*
- 43 Don't Cross the Streams: Population Structure Among Three Adjacent Steelhead Streams? Brice Adams, Christian Smith
- 44 How do Niches Evolve? Leveraging observational community data Luke Frishkoff
- 45 Genetic diversity in the endangered parakeet Pyrrhura griseipectus and implications for conservation

 Roberta Canton, Chrysoula Gubili, Tomas Hrbek, Camila Ribas, Fabio Nunes

Topic Sat-10 Development

- 46 Developmental Timing In Periodical Cicada (Hemiptera: Magicicada) Lifecycle Evolution Diler Haji
- 47 The Influence of Temperature on the Development, Mortality, and Spatial Behavior of Vernal Pool Odonate Larvae

 Nene Ugbah
- 48 Evo-Devo, metamorphosis and critical developmental periods for shaping the feeding biomechanics of fishes

 W. James Cooper, Sarah McMenamin, Demi Galindo, Mitchell Wagner, Casey Carter, Adrian Deleon
- 49 Role of Genetic Variation in the Ontogeny of Metabolism during Development of Drosophila Omera Matoo, Cole Julick, Kristi Montooth
- 50 The origin of muscle precursors in non-amniotic and amniotic species. withdrawn 2017-06-16 06:15:23.0

 Mateusz Hermyt

Topic Sat-11 Diversification

- 51 Diversity-dependent diversification: but do modern planktonic foraminifera actually compete? Marina Rillo, Michal Kucera, Thomas Ezard
- 52 Diversification of lineages and functional phenotypes in the Neotropical electric fish Apteronotidae

 Lesley Kim, Kory Evans, Maxwell Bernt, James Albert
- 53 Glacial cycles drive rapid genomic differentiation in the field vole Nicholas Fletcher

Topic Sat-12 Ecological genetics

- 54 Genetic variation at a CNE correlates with microhabitat-associated behavioral differentiation in Malawi African cichlid fishes

 Emily Moore, Lynea Bull, Reade Roberts
- 55 Reintroducing Western Pond Turtles to Yosemite Valley: A Conservation Genomics Approach Laura Van Vranken, Ninette Daniele, Molly Thompson, Rob Grasso, Danielle Edwards

Topic Sat-13 Evolutionary ecology

- The role of urbanization on the evolution of selfing and population genetic structure in Impatiens capensis Ruth Rivkin, Marc Johnson
- 57 Alternative survival strategies in vectored yeasts Kate McKee
- 58 Using ABC to Understand Trait-Mediated Community Assembly Megan Ruffley, Luke Harmon
- 59 Influence of temporal selection in evolutionary response to temperature in Daphnia Hamanda Badona Cavalheri, Celia Symons, Marika Schulhof, Natalie Jones, Jonathan Shurin
- 60 Multivariate selection and the range limit of alderia willowi: effects of salinity and trait covariance

 Jackson Powell, Patrick Krug, Dustin Marshall

Topic Sat-14 Evolutionary theory

- 61 Spontaneous-mutation spectra and fitness loss of Daphnia pulex mutation-accumulation lines exposed to chronic, sub-lethal concentrations of heavy metals withdrawn 2017-06-10 10:21:56.0
 - James Bull, Jullien Flynn, Frederic Chain, Melania Cristescu
- 62 Most colorful example of genetic assimilation? Exploring the evolutionary destiny of recurrent phenotypic accommodation

 Ahva Potticary, Erin Morrison, Alexander Badyaev
- 63 Self-deception in the Service of Deceit: An Adaptationist Account Shahab Zareyan, Christoph Hauert, Sally Otto
- 64 Comparing patterns of selection on Phospholipase A2 toxins within and between species of North American Pitvipers

 Michael Hogan, Darin Rokyta, Christopher Parkinson

Topic Sat-15 Experimental evolution

- 65 Experimental adaptation of a non-host-associated bacterium to the zebrafish digestive tract
 - Jarrett Lebov, Brendan Bohannan
- 66 Tempo, Mode, and Fitness Effects of Mutation in C. elegans over 400 Generations of Minimal Selection
 - Ayush Saxena, Matthew Salomon, Charles Baer, Chikako Matsuba, Shudan Yeh

Topic Sat-16 Gene flow

- **67** From spawning to settlement: Identifying fine-scale connectivity in the Convict Tang, Acanthurus triostegus, across Oʻahu
 - Richard Coleman, Derek Kraft, Rob J. Toonen, Brian Bowen
- The role of gene flow on local adaptation in a field cricket contact zone Emma Collosi, Jeanne Robertson, David Gray
- 69 Jumping through the desert: population genetics and structure of a kangaroo rat in northern Mexico
 - Alejandro Flores-Manzanero, Ella Vázquez-Domínguez, Rodney Dyer
- 70 Genetic distinctiveness of Atlantic Spotted dolphin along coastal Brazil

 Karina do Amaral, Valentina Franco-Trecu, Dalia Barrera, Nohelia Curtidor, Susana Caballero, Ana
 Carolina de Meirelles, Vidal Martin, Manuel Carrillo, Paula Mendez-Fernandez, Marcos Oliveira Santos,
 Salvatore Siciliano, L. Lacey Knowles, Nelson Fagundes, Ignacio Moreno, Ana Rita Amaral
- 71 The Evolution of Dispersal in Trinidadian Guppies Robert Prather, David Reznick

Topic Sat-17 Genomics

- 72 3D genome structure and the splicing of the mysterious minor spliceosomal introns Swadha Singh, Graham Larue, Scott Roy
- 73 Variation in fluorescent protein gene family cause expression differences among individuals in a stony coral, Acropora digitifera Shiho Takahashi-Kariyazono
- 74 Sexually Dimorphic Exon Splicing and Gene Expression in the Rock Dove HPG Axis Andrew Lang, Suzanne Austin, Rebecca Calisi, Matthew MacManes

Topic Sat-18 Geographic variation

- 75 Maintenance of the geographic variation of flower color in Leavenworthia stylosa: Role of local adaptation in different life history stages

 Mahaguruge Fernando, Chris Herlihy, Jeffrey Walck
- 76 Genetic underpinnings of transgressive segregation: Increased thermal tolerance in experimental interpopulation hybrid crosses of Tigriopus californicus. withdrawn 2017-06-20 15:32:11.0
 - Reginald Blackwell, Ron Burton
- 77 Evidence of introgression in a geographically widespread venomous marine snail species complex

 Peter Cerda, Thomas Duda
- Teter cerua, momas bada
- 78 Mitochondrial evolution in Tigriopus californicus is marked by geographic variation in tRNA structure and codon usage bias *Eric Watson*, Heath Blackmon, Suzanne Edmands

Topic Sat-19 Host-parasite

79 Testing for selection in chipmunk sucking lice Kayce Bell, Anna Phillips, Julie Allen, Kevin Johnson, John Demboski, Joseph Cook

- 80 Landscape impacts on white footed mouse movement in southern New England Kerry Barrett, Jonathan Richardson, Georgi Silveira
- 81 How does endoparasitism evolve? Insights from Venom allergen-like proteins in blood flukes (Platyhelminthes: Trematoda: Digenea)

 Breanna Sipley, Ash Bullard, Kenneth Halanych

Topic Sat-20 Hybridization

- 82 Genomic data distinguish phenotypically distinct birds despite low levels of divergence Stepfanie Aguillon, Richard Harrison, Irby Lovette
- A window into the introgression of male reproductive genes in a natural primate hybrid zone (Alouatta palliata x A. pigra)

 Molly Choi, Liliana Cortes Ortiz
- 84 Genetic and phenotypic evidence of a contact zone between populations of the red-eyed treefrog, Agalychnis callidryas

 Maria Akopyan, Erica Bree Rosenblum, Zachariah Gompert, Karina Klonoski, Andres Vega, Rachel Mackelprang, Jeanne Robertson
- 85 Fine scale admixture patterns across the genome of the Africanized honeybee (Apis mellifera)

 Daniela Zárate
- 86 Detecting introgressive hybridization in red-eyed treefrogs through genomics and color pattern distribution

 Meaghan Clark, Andres Vega, Jeanne Robertson
- 87 Inferring selection on hybrids in wild populations of Spadefoot toads Audrey Kelly, Karin Pfennig

Topic Sat-21 Invasive/invasion

- 89 Invasive rat colonization history and movement dynamics in Haida Gwaii Bryson Sjodin, Robyn Irivine, Gregg Howald, Michael Russello
- 90 A genetic reconstruction of the invasion of the calanoid copepod Pseudodiaptomus inopinus across the North American Pacific Coast

 Eric Dexter, Stephen Bollens, Jeffery Cordell, Ho So, Gretchen Rollwagen-Bollens, Susanne Pfeifer, Jérôme Goudet, Severine Vuilleumier

Topic Sat-22 Life history

- 91 The Genetic Basis for Inheritance of Life History Traits in Trinidadian Guppies Sarah Ruckman, Simon Zhu, Vincent Poon, Yuridia Reynoso, David Reznick
- 92 Sex ratio of mirid populations shifts in response to hostplant co-infestation or altered cytokinin signaling
 Nora Adam, Theresa Erler, Mario Kallenbach, Martin Kaltenpoth, Grit Kunert, Ian Baldwin, Meredith Schuman
- 93 Testing the effects of species, life stage and sex on Isoprene emission in mosses. *Timea Deakova*

Topic Sat-23 Macroevolution

- 94 Dating the terrestrial invasion of the Cyclophoroidea (Mollusca:Gastropoda) using the Fossilized Birth-Death model.

 Timothy Gervascio, Jesse Czekanski-Moir, Rebecca Rundell, Nicole Webster
- 95 Characterizing the performance of species diversification models Natya Hans, Jose Miguel Ponciano, J. Gordon Burleigh

Body shape modularity and morphological diversification rates in a hyperdiverse freshwater fish lineage Mike Burns

Topic Sat-24 Molecular evolution

- 97 Molecular underpinings of a novel defensive enzyme in tomato withdrawn 2017-06-23 18:09:11.0
 - Allan Castillo, Mark Rausher
- Comparative molecular analyses of the amniote p53 signaling network: Rapid evolution and positive selection on mammalian mediators of p53. Courtney Passow, Heath Blackmon, Suzanne McGaugh
- 99 Large-scale assessment of mitochondrial adaptation to high altitude in birds María Cecilia Estalles, Pablo D. Lavinia, Pablo Tubaro, Dario Lijtmaer

Topic Sat-25 Mutualism

- 100 Bacterial communities associated with butterflies and their effect on butterfly fitness Kruttika Phalnikar, Deepa Agashe, Krushnamegh Kunte
- 101 The density dependence of mutualistic interactions Christopher Moore, Karen Abbott
- 102 Microbes aid adaptation to a novel environment Aparna Agarwal, Deepa Agashe

Topic Sat-26 Paleobiology

- 103 Convergent body size evolution of crocodyliformes upon entering the aquatic realm William Gearty, Jonathan Payne
- 104 Total dental occlusal area as a constrained feature in modern walruses Win McLaughlin, Samantha Hopkins, Edward Byrd Davis, Caitlyn Boatman

Topic Sat-27 Pathogens

- 105 Acquisition of azithromycin resistance in Neisseria gonorrhoeae via intragenus recombination
 - Crista Wadsworth, Yonatan Grad
- 106 Horizontally acquired biosynthesis genes boost Coxiella burnetii's physiology Jess Millar, Abraham Moses, Matteo Bonazzi, Paul Beare, Rahul Raghavan
- 107 Evolutionary analysis of viral strains infecting a single anole species Sofia Prado-Irwin, Martijn van de Schoot, Anthony Geneva

Topic Sat-28 Phenotypic plasticity/GxE

- 108 Phenotypic Differences in a Single Clonal Type in Newly Colonized and Established **Populations**
 - Madison Armstrong, Mark Dybdahl
- 109 Physiological Plasticity and Divergence in Migrating vs Nonmigrating Monarch Butterflies Cole Julick, Delbert Green, Ayse Tenger-Trolander, Marcus Kronforst, Kristi Montooth

Topic Sat-29 Phylogenetic comparative methods

110 Morphological form is predicted by genetic evidence function of the mammalian vomeronasal system

Laurel Yohe, Simone Hoffmann, Liliana Davalos

- 111 Functional coupling and morphological integration in the evolution of suction feeding and gill ventilation of sculpins and relatives (Perciformes: Cottoidei)

 Stacy Farina, Katherine Corn, Matthew Knope, William Bemis, Adam Summers
- 112 Pair-living as a stepping stone to group-living in primate social evolution Luca Pozzi, Peter Kappeler

Topic Sat-30 Phylogenetic methods development

113 FastNet: Fast and accurate inference of phylogenetic networks using large-scale genomic sequence data

Hussein Hejase, Natalie VandePol, Gregory Bonito, Kevin Liu

- 114 A macroevolutionary simulator for phylogenomic data Wade Dismukes, Tracy Heath
- 115 Reducing the size of terraces in phylogenetic tree space through the removal of taxa from a supermatrix phylogeny

 Joseph Charboneau, MIchael Sanderson

Topic Sat-31 Phylogenomics

- 116 A Migratory Divide in the Painted Bunting (Passerina ciris)

 C.J. Battey, Ethan Linck, Kevin Epperly, Cooper French, David Slager, Paul Sykes, John Klicka
- 117 Reaffirming Hennig in the Genomics era: evolution of spider flies (Diptera, Acroceridae)

 Jessica Gillung, Shaun Winterton
- 118 Kryptolebias marmoratus phylogenomics and future laboratory applications Shawn Trojahn, Luana Lins, Joanna Kelley
- 119 Molecular evolution of the DNA Methyltransferase gene family in vertebrates withdrawn 2017-06-24 18:55:03.0

 Karen Bobier
- 120 A Phylogenetic Analysis of the Pannota Hayden Kallas, T. Heath Ogden
- 121 Phylogeny of the Amphisbaenia (Reptilia: Squamata) based on anchored phylogenomics and morphological dataset

 Roberta Graboski, Emily Lemmon, Alan Lemmon, Hussam Zaher, Tracy Heath
- 122 Combining genomic and morphological data in a model-base approach to ascertain species-delimitation in the South America marsh rat (Holochilus)

 Joyce Prado, L. Lacey Knowles, Alexandre Reis Percequillo

Topic Sat-32 Phylogeography

- 123 Phylogeography of Melanesian flycatchers (Aves: Monarchidae) based on ultraconserved elements

 Lukas Klicka, Brian Weeks, Chris Filardi, Robert Movle
- 124 Phylogegraphy of Micrurus surinamensis and Micrurus lemniscatus Renan Bosque, Hussam Zaher, Guarino Colli, Miguel Rodrigues, Omar Torres-Carvajal, Nelson Jorge da SIlva Junior, Ana Lucia da Costa Prudente, Felipe Grazziotin, Laurie Vitt, Brice Noonan
- 125 Trait-based population genomics across a biogeographic barrier Kaiya Provost, William Mauck III, Brian Smith

Topic Sat-33 Plant reproductive biology

- 127 Range-wide variation in selfing rate in American bellflower (Campanula americana) Yva Eline, Matthew Koski, Laura Galloway, Jeremiah Busch
- 128 Sterile marginal flowers increase insect visitation and fruit set in the hobblebush (Viburnum lantanoides)

 Brian Park, Michael Donoghue

<u>Topic Sat-34 Population genetics: inference of</u> selection

- 129 Investigating genetic adaptation to high altitude across Andean hummingbird species and populations
 - Marisa Lim, Christopher Witt, Catherine Graham, Liliana Davalos
- 130 The genomics of behavioral adaptation to photoperiodism in an Asian burying beetle. Yuki Haba, Hsiang-Yu Tsai, Dustin Rubenstein, Sheng-Feng Shen
- 131 Comparative species and population analyses identify human-specific patterns of selection at schizophrenia GWAS loci

 Mariam Sankoh, Brian Verrelli
- 132 Evidence of selection in extant and extinct monk seals
 Stephen Gaughran, Emily Lipstein, George Amato, Matthew Aardema, Alan Scott, Jeffrey Powell,
 Adalgisa Caccone

<u>Topic Sat-35 Population genetics: molecular ecology</u>

- 133 Population Genetics of Steller Sea Lions at a Single Haulout: Optimized Low Copy DNA Methods
 - Peter Han, Brad Swanson
- 134 Population genomic study of adaptation and selection in a dune sunflower April Goebl, Silas Tittes, Nolan Kane
- 135 Population genomics of the armored catfish Hypancistrus sp. from the Xingu River, in the Brazilian Amazonia
 - Marcella Santos, Leandro Sousa, Brice Noonan
- 136 Historical demography of Malagasy herpetofaunal assemblages Arianna Kuhn
- 137 Characterizing adaptive immunogenetic variation in the endangered salt marsh harvest mouse, Reithrodontomys raviventris

 Anastasia Ennis, C. Sarah Cohen
- 138 Genome-wide markers reveal the determinants of genetic divergence in a lowland Mesoamerican bird (Icterus gularis)

 Lucas Rocha Moreira, Blanca Hernández Baños, Brian Smith

Topic Sat-36 Quantitative genetics

139 Should I stay or should I go? Evaluating the genetic basis of migration-related traits in rainbow trout (Oncorhynchus mykiss).

Spencer Weinstein, Matt Hale, Frank Thrower, Krista Nichols

Topic Sat-37 Reproductive biology

- 141 The evolution of mating type loci in lichenized fungi. withdrawn 2017-06-10 10:25:18.0 Kristin White
- 142 Disrupted Genomic Imprinting Negatively Affects Maternal Behaviors in an Interspecific Mouse Cross

Sarah Gardner, Polly Campbell

143 Drosophila accessory gland secondary cells mediate between-male competition for fertilisations

Ben Hopkins, Irem Sepil, Tommaso Pizzari, Clive Wilson, Stuart Wigby

Topic Sat-38 Sensory systems

- 144 Heat of the moment? Effects of long term temperature exposure on cricket courtship song Hayden Waller, Kerry Shaw
- 145 Transcriptome sequencing reveals divergent selective pressures on snake visual transduction genes associated with rod-cone transmutation

 Ryan Schott, Alexander Van Nynatten, Daren Card, Todd Castoe, Belinda Chang
- 146 Pheromone and Vomeronasal Receptor Evolution in House Mice Caitlin Miller, Polly Campbell, Michael Sheehan

Topic Sat-39 Sex/recombination

- 147 Identifying a neo-sex chromosome via deep resequencing in a hybridizing primate species Marcella Nidiffer, Liliana Cortes Ortiz, Priscilla Tucker, Jacob Mueller
- 148 Estimating mutational fitness effects from growth data in budding yeast Linnea Sandell, Sally Otto, Anastasia Kuzmin

Topic Sat-40 Sexual selection

- 149 Predator-induced maternal and paternal effects independently alter female offspring mate choice
 - Whitley Lehto, Robin Tinghitella
- 150 Increased sperm transfer and fertilization success in the copulatory silk wrapping nursery web spider, Pisaurina mira

 Alissa Anderson*, Eileen Hebets
- 151 Determining the role of postcopulatory sexual selection in female sperm storage organ evolution in Drosophila.

 Tiffini Smith*, Mollie Manier
- 152 Female color variation and male harassment in the polymorphic damselfly Megalagrion calliphya

 Phoebe Cook, Rebecca Rasmussen, Anjali Batra, Jonathan Brown, Idelle Cooper
- 153 Quantifying Female Reproductive Cyclicity Across Four Peromyscus species William Weber, Heidi Fisher
- 154 Parental clutch traits as a source of transgenerational developmental stress: consequences for secondary sexual traits and reproductive success in the zebra finch *Kerianne Wilson*, *Nancy Burley*
- 155 Population variation in mating traits and preferences: character displacement or selective tradeoffs?

 Gina Calabrese, Karin Pfennig

Topic Sat-41 Social systems

156 Worker ovicide can drive the evolution of social polymorphism and split sex ratios Gil Henriques, Andrés Quiñones, Ido Pen

Topic Sat-42 Speciation

- 157 Subspeciation rates and diversity gradients Mauro Toshiro Sugawara, Matthew Pennell
- 158 Patterns of speciation under adaptation from standing genetic variation Ken Thompson, Matthew Osmond, Dolph Schluter
- 159 Genetic divergence and speciation of a generalist insect herbivore Mayra Vidal, Shannon Murphy
- 160 Hybrid speciation in Phlox
 Benjamin Goulet, Robin Hopkins
- 161 Studying genomic differentiation among Northern Goshawks in British Columbia Kenneth Askelson, Darren Irwin, Armando Geraldes
- 162 Phenotypic and phylogenetic homogeneity corroborate conservative macro-evolutionary dynamics in the Neotropical tree genus Myrcia

 Thais Vasconcelos, Eve Lucas, Gerhard Prenner, Marion Chartier

Topic Sat-43 Species delimitation

- 163 Hybridization and Species Delimitation in Oregon Buttercups

 Jessica Persinger, Pamela Thompson, Brendan F Kohrn, Mitch Cruzan
- 164 Differential fitness reveals a speciation process in karyotipic variants of the Brazilian rodent Akodon cursor (Sigmodontinae)

 Fernanda Zaidan, Cristina Nogueira-Massariol, L. Lacey Knowles, Valeria Fagundes

Topic Sat-44 Systematics

- 165 Molecular Analysis of Mayfly Family Baetidae (Ephemeroptera)

 Bryson Carrier, Daniel Ferguson, T. Heath Ogden
- 166 Population admixture in the cosmopolitan fern genus, Pteridium (Dennstaedtiaceae)
 Sylvia Kinosian, Paul Wolf, Carol Rowe, Joshua Der, Zachariah Gompert, Martin Schilling, Peter
 Lockhart, Lara Shepherd, Trish McLenachan, John Thomson

Topic Sat-45 Last minute additions

- 167 Population Genomics of Moss Campion Heather Gillette
- 168 Adaptation and demographic history of two tidal marsh species inferred from whole genome sequencing.
 - Gemma Clucas, Jennifer Walsh, Matthew MacManes, Adrienne Kovach
- 169 Sex and chromosome effects on transcriptome variation William Etges
- 170 Differentiation in seed endophytic communities among natural populations of C. bursa-pastoris

 Byungwook Choi, Seorin Jeong, Eunsuk Kim

171 Rapid adaptation in a contaminated environment: Evolutionary adaptive response of old field grass Andropogon virginicus to heavy metals in abandoned Lead and Zinc mines

Samantha Sharpe, Loretta Johnson, Nora Bello, Matthew Galliart, Olivia Parrish

Session Sun1-Ball202 SSB Symposium - A debate of conceptual issues surrounding genetic-based species delimitation in the genomic era 1

Sunday - 8:15 - 9:45

Room Oregon ballroom 202

- 8:15 Benefits of using the multi species coalescent for species delimitation Matthew Fujita
- 8:30 When sampling matters: biases in species delimitation. Kelly Zamudio, Nicholas Mason, Nicholas Fletcher
- 8:45 Moderated discussion: Does the multispecies coalescent delimit species?

 *David Hillis, L. Lacey Knowles**
- 9:00 The necessity of morphology for delimiting species in the molecular age Felipe Zapata
- 9:15 The superfluity of morphology for species delimitation in the molecular age SARA RUANE
- 9:30 Moderated discussion: Is species delimitation necessary for assessing the drivers of diversification?

 David Hillis, L. Lacey Knowles

Session Sun1-A105 Phylogenetic methods development 1 Sunday - 8:30 - 9:45 Room A105

- 8:30 Independent rate models are biologically unrealistic Mario dos Reis
- 8:45 Some additional new plotting methods for phylogenies & comparative data Liam Revell
- 9:00 Tools for assessing ancestral recombination graphs Mary Kuhner
- 9:15 Horseshoes and hand grenades: shrinking models with exploding numbers of parameters using horseshoe priors

 Andrew Magee, Adam Leache, Vladimir Minin
- 9:30 Flexible Modeling of Morphological Characters for Phylogenetic Inference April Wright, Walker Pett, Tracy Heath

Session Sun1-A106 Biogeography 1

Sunday - 8:30 - 9:45

Room A106

- 8:30 Filter-dispersal assembly of lowland Neotropical rainforests across the Andes

 Jordan Bemmels, Joe Wright, Nancy Garwood, Simon Queenborough, Renato Valencia, Christopher

 Dick
- 8:45 Global patterns and drivers of squamate body sizes Alex Slavenko, Feldman Anat, Shai Meiri
- 9:00 Unveiling geographical, ecological, and evolutionary relationships of a Neotropical species complex of unisexual lizards of genus Loxopholis (Squamata: Gymnophthalmidae) through GIS-based analysis

Tuliana Brunes, Katia Pellegrino, Levi Carina Terribile, Matheus Lima-Ribeiro, Sergio Marques de Souza, Miguel Rodrigues

- 9:15 Speciation and the Assembly of Global Vertebrate Diversity Daniel Rabosky
- 9:30 Biodiversity and Topographic Complexity: Modern and Geohistorical Perspectives *Tara Smiley*, Catherine Badgley, Rebecca Terry, Edward Byrd Davis, Brett Riddle, Brian Yanites, John
 Finarelli, Samantha Hopkins

Session Sun1-A107-109 Macroevolution 2 Sunday - 8:30 - 9:45 Room A107-109

8:30 The influence of wing morphology upon the dispersal, geographic distributions and diversification of a global radiation of passerine birds

Jon Kennedy, Michael Borregaard, Knud Jønsson, Petter Marki, Jon Fjeldså, Carsten Rahbek

8:45 Evolutionary and Ecologic Signal of Colors in Fungal Fruit Bodies
Franz Krah, Hanno Schaefer, Antje Heideroth, Lynne Boddy, Carrie Andrew, Einar Heegaard, Håvard
Kauserud, Jacob Heilmann-Clausen, Joerg Mueller, Robert Freckleton, Christoph Heibl, Claus Baessler

9:00 Major habitat transitions within Class Gastropoda Teresa Rose Osborne, David Bullis, Jesse Czekanski-Moir, Rebecca Rundell

- 9:15 Origins of diversity in caridean shrimps: insights from a time-calibrated supertree Katie Davis, Sammy De Grave, Cyrille Delmer, Matthew Wills
- 9:30 Evolution of extreme ontogenetic allometric diversity and heterochrony in pythons, a clade of giant and dwarf snakes

 Damien Esquerre*, Emma Sherratt, J. Scott Keogh

Session Sun1-B111-112 Quantitative genetics / adaptation Sunday - 8:30 - 9:45 Room B111-112

- 8:30 Resolving the Genetic Architecture Paradox in Complex Trait Evolution: A Simulation Study David Remington, Mario Augustinovic
- 8:45 Deeply conserved genetic constraints influence adaptive radiation of Anolis lizards Joel McGlothlin, Megan Kobiela, Helen Wright, Jason Kolbe, Jonathan Losos, Butch Brodie
- 9:00 Expression of additive genetic variance for fitness in a population of partridgepea grown in two field sites

 Ruth Shaw, Seema Sheth, Mason Kulbaba, Rachel Pain
- 9:15 Pollen dispersal slows geographical range shift and accelerates ecological niche shift under climate change
 Ophelie Ronce, Robin Aquilee, Gael Raoul, Francois Rousset
- 9:30 Can parental environment influence the potential for adaptation to global change? Evatt Chirgwin, Dustin Marshall, Carla Sgrò, Keyne Monro

Session Sun1-B113 Behavior 1

Sunday - 8:30 - 9:45

Room B113

- 8:30 Information Warfare: Mimicry & Signal Detection David Kikuchi
- 8:45 Behavioural adaptations in Darwin's finches Kiyoko Gotanda
- 9:00 A distance-performance trade-off in the phenotypic basis of dispersal in a stream salamander Brett Addis, Bret Tobalske, Winsor Lowe
- 9:15 Father-embryo coevolution in Neotropical glassfrogs Jesse Delia
- 9:30 The eco-evolutionary consequences of agonistic character displacement Kenichi Okamoto, Jonathan Drury, Greg Grether

Session Sun1-B114-115 Inbreeding / conservation biology Sunday - 8:30 - 9:45 Room B114-115

8:30 Genomic consequences of inbreeding in a wild wolf population

Toby Fountain, Marty Kardos, Mikael Åkesson, Øystein Flagstad, Olof Liberg, Pall Olason, Håkan Sand,
Petter Wabakken, Camilla Wikenros, Hans Ellegren

- 8:45 Genome-scale data reveals that endemic Poecilia populations from small sulfidic springs display no evidence of inbreeding
 - **Anthony Brown**, Ryan Greenway, Samuel Morgan, Corey Quackenbush, Luca Giordani, Lenin Arias-Rodriguez, Michael Tobler, Joanna Kelley
- 9:00 Reviving a Lost Species: The Case of the Floreana Galápagos Giant Tortoise Chelonoidis elephantopus

Joshua Miller, Maud Quinzin, Nikos Poulakakis, Jonathan Sandoval-Castillo, James Gibbs, Luciano Beheregaray, Ryan Garrick, Michael Russello, Claudio Ciofi, Danielle Edwards, Elizabeth Hunter, Washington Tapia, Dany Rueda, Jorge Carrión, Andrés Valdivieso, Adalgisa Caccone

9:15 Effect of ecological stress (host and temperature) on magnitude of inbreeding depression in Callosobruchus maculatus

Amv Springer, Zachariah Gompert, Frank Messina

9:30 Unequivalent lethal equivalents: Approaches for quantifying genetic load and inbreeding depression

Pirmin Nietlisbach, Lukas Keller

Session Sun1-B116 Hybridization 1

Sunday - 8:30 - 9:45

Room B116

- 8:30 Chemical compounds of pheromone signals and reproductive isolation in naturally-hybridizing swordtail fish.

 Chris Holland, Gil Rosenthal
- 8:45 Mechanical and tactile incompatibilities cause reproductive isolation in young damselfly species

 Alexandra Barnard, Ola Fincke, Mark McPeek, John Masly
- 9:00 The collapse of assortative mating uncovers a melanoma hybrid incompatibility Mateo Garcia, Daniel Powell, Gil Rosenthal, Manfred Schartl, Molly Schumer
- 9:15 Modeling local ancestry and Dobzhansky-Muller incompatibilities in a hybrid zone Joel Smith, Megan Frayer, John Novembre, Bret Payseur

Session Sun1-B117-119 Adaptation / genomics 1 Sunday - 8:30 - 9:45 Room B117-119

8:30 Elucidating the genetic control of long-term success in a near century long study of barley (Hordeum vulgare)

Jacob Landis, Angelica Guercio, Christopher Fiscus, Daniel Koenig

- 8:45 The Panicum hallii genome and the evolution of molecular drought-responses in C4 grasses.

 John Lovell, Thomas Juenger, Jeremy Schmutz
- 9:00 Polygenic adaptation in maize.

Emily Josephs, Jeremy Berg, Jeffrey Ross-Ibarra, Graham Coop

- 9:15 Population Genomics of Moss Campion withdrawn 2017-06-04 07:32:32.0

 Heather Gillette, Katrina Dlugosch, Brittany Barker, William Morris, Daniel Doak
- 9:30 Impact of Horizontal Gene Transfer on the Evolution of the Early Fungi Michał Ciach, Anna Muszewska, Paweł Górecki

Session Sun1-Ball203 Parallel / convergent evolution Sunday - 8:30 - 9:45 Room Oregon ballroom 203

- 8:30 Why convergence? Inferring constraint from comparative genomic studies of local adaptation Sam Yeaman, Michael Whitlock, Kathryn Hodgins, Aleeza Gerstein
- 8:45 What determines the distinct morphology of species with a particular ecology? The roles of many-to-one mapping and trade-offs in the evolution of frog ecomorphology and performance Daniel Moen
- 9:00 Ploidy drive in fungal microbes: an evolutionary mystery waiting to be solved Aleeza Gerstein, Meleah Hickman, Judith Berman, Kirsten Nielsen, Sally Otto
- 9:15 From simplicity to complexity: Evolution and development of flower color patterning in Mimulus

 Arielle Cooley

Session Sun1-Ball204 ASN Spotlight - Processes underlying pattern: considering the evolutionary mechanisms underlying population-level differentiation Sunday - 8:30 - 9:45 Room Oregon ballroom 204

- 8:30 Introduction: Connecting pattern to process in studies of speciation Rebecca Safran
- 8:45 Anthropogenic changes in nutrient availability alter sexual selection dynamics and population differentiation in a butterfly

 Emilie Snell-Rood, Anne Espeset, Kristin Sikkink, Megan Kobiela
- 8:50 Genetic basis and adaptive significance of population level and plastic variation in leaf shape Kathleen Ferris, Graham Coop, Jeffrey Ross-Ibarra, Johanna Schmitt
- 8:55 Differences in spatial autocorrelation between habitat types promote local adaptation in dispersal strategy

 Meredith Cenzer, Leithen M'Gonigle
- 9:00 Isolation by distance as a null hypothesis of population structure Gideon Bradburd
- 9:15 Surfing in pools of beetles: using replicated landscape experiments to disentangle signatures of selection and drift
 Silas Tittes, Ruth Hufbauer, Nolan Kane, Brett Melbourne, Christopher Weiss-Lehman
- 9:20 How hybridization structures the genomes of related birds David Toews
- 9:25 Coevolutionary diversification in the yucca-yucca moth interaction Kari Segraves
- 9:30 Correlated evolution of functional physiology and thermal niches in tropical freshwater fishes Zachary Culumber

Session Sun1-C120-122 Ecological genetics / adaptation Sunday - 8:30 - 9:45 Room C120-122

- 8:30 Evolution of daylength adaptation in temperate grasses Erica Leder
- 8:45 Natural selection on phenology across a seasonal gradient in elevation of an annual plant, Rhinanthus minor Dylan Sora, Christopher Eckert, Dave Ensing, Anna Hargreaves
- 9:00 Genomic evidence of rapid adaptation to a novel freshwater environment Janna Willoughby, Jacob Tennessen, Kim Scribner, Mark R. Christie
- 9:15 Circadian clock function correlates with diurnal changes in gas-exchange traits in Brassica rapa crop types and RILs in controlled environmental settings

 Yulia Yarkhunova, Matti Salmela, Brent Ewers, Ping Lou, C. Robertson McClung, Cynthia Weinig
- 9:30 Local adaptation and adaptive variation in distantly-related conifers Jeremy Yoder, Sally Aitken

Session Sun1-C123 Education

Sunday - 8:30 - 9:45

Room C123

- 8:30 GET CREATIVE TO MAXIMIZE AN EVOLUTION CLASS E Dale Broder
- 8:45 Feedback in Online Learning Activities: Should We K.I.S.S. or Try to Explain?

 David Ford, Anna Hiatt
- 9:00 Women in Academia: Trends in funding success rates and submissions across the sciences at the National Science Foundation

 Leslie Rissler

Session Sun1-C124 Genomics / expression studies Sunday - 8:30 - 9:45 Room C124

8:30 Differential gene expression across tissues, sex, and life stages in the sea urchin Tripneustes gratilla

Áki Jarl Láruson, Floyd Reed

- 8:45 Linking genomic transposable element evolution to germ-line expression in snakes.

 Giulia Irene Maria Pasquesi, Michael Vandewege, Audra Andrew, Nicole Hales, Andrew Corbin, Todd
 Castoe
- 9:00 Comparative functional genomics of osmotic physiology in 16 species of killifish Noah Reid
- 9:15 Seasonal variation in gene expression across a community of plants at Harvard Forest Hannah Marx, Eldridge Wisely, Stacy Jorgensen, Katrina Dlugosch, Michael Barker
- 9:30 The genomics of conflict: X-chromosome drive in the fly Drosophila neotestacea Kelly Dyer, Kathleen Pieper, Robert Unckless

Session American Naturalist editorial board meeting
Sunday - 8:30 - 10:15
Room C126

Session Taylor & Francis editorial board meeting
Sunday - 8:30 - 10:15 Room B110

Session Coffee break

Sunday - 9:45 - 10:15

Room Exhibit Hall A

Sponsored by



Session Sun2-A105 Phylogenetic methods development 2 Sunday - 10:15 - 11:30 Room A105

- 10:15 Sensitivity of quartet-based species tree methods to missing data Ian Gilman, David Tank
- 10:30 Utility for off-target DNA mining in non-model organisms and an assessment of the discriminatory power of DNA barcodes in plants and animals.

 Alisha Mechtley
- 10:45 The development of scientific consensus: analyzing conflict and concordance among Avian phylogenies
 - Joseph Brown, **Ning Wang**, Stephen Smith
- 11:00 Evaluating allopolyploid origins in strawberries (Fragaria) using haplotypes generated from target capture sequencing withdrawn 2017-06-20 12:55:44.0

 John Syring, Olga Kamneva, Aaron Liston, Noah Rosenberg

Session Sun2-A106 Biogeography 2

Sunday - 10:15 - 11:30

Room A106

- 10:15 An empirical test of the role of topographic complexity in the general dynamic model of oceanic island biogeography

 Yannik Roell
- 10:30 Islands in Idaho: Using a Naturally Fragmented Landscape to Investigate Communities Katie Peterson, Christine Parent
- 10:45 Convergent phenotypes obscure biogeographic history in Neotropical Seedeaters (Aves: Sporophila)

Nicholas Mason, Arturo Olvera-Vital, Irby Lovette, Adolfo Navarro-Sigüenza

11:00 Patterns of character evolution in a mega-diverse angiosperm clade (Asteridae) withdrawn 2017-06-25 11:14:10.0

Gregory Stull, Melanie Schori, Douglas Soltis, Pamela Soltis

11:15 The evolution of clonality in seasonal seas: the influence of temperature on geographic patterns of life cycle diversity in the sea anemone Diadumene lineata Will Ryan

Session Sun2-A107-109 Macroevolution 3 Sunday - 10:15 - 11:30 Room A107-109

- 10:15 Evolution of sexual dichromatism in the 'blue' cardinalids (Aves: Cardinalidae)
 Natalia Garcia, Ana Barreira, Pablo Tubaro
- 10:30 How ecology shapes adaptive phenotypic evolution in a eusocial lineage: the evolution of the specialized soldier caste in turtle ants

 Shauna Price. Scott Powell. Corrie Moreau
- 10:45 Extinction and time are drivers of the marine-terrestrial biodiversity gradient: Is the ocean a deathtrap?

 Elizabeth Miller, John Wiens
- 11:00 Disentangling the evolutionary history of nickel tolerance and hyperaccumulation in wild sunflowers (Helianthus): a function-valued phylogenetic comparative approach

 Eric Goolsby, Chase Mason, Lauren Brill, Gaven Meyers, Lisa Donovan
- 11:15 Testing sexual selection as a determinant of species extinction and origination in Late Cretaceous cytheroid ostracodes

 Maria Joao Fernandes Martins, Gene Hunt, T Markham Puckett, Rowan Lockwood, John P Swaddle

Session Sun2-B111-112 Quantitative genetics / genomics Sunday - 10:15 - 11:30 Room B111-112

10:15 Dissecting the genetic basis of learning, memory and thermal tolerance in a multi-parental population.

Patricka Williams-Simon, Elizabeth King, Troy Zars

10:30 Genetic constraints in the evolution of chemical defenses in multiple tissues of Boechera stricta

Rose Keith, Tom Mitchell-Olds

- 10:45 A genomic scan for flower shape variation in Penstemon virgatus: implications for transitions to hummingbird pollination

 Carolyn Wessinger, John Kelly, Lena Hileman
- 11:00 Heritability and Genomic Basis for Variation in Thermal Tolerance of the Coral Holobiont in Orbicella faveolata

 Katherine Dziedzic, Eli Meyer
- 11:15 Fine-mapping the genetic basis of extreme color diversity used for social recognition Michael Sheehan, Sara Miller, Andrew Legan, Sarah Bluher

Session Sun2-B113 Behavior 2

Sunday - 10:15 - 11:30

Room B113

- 10:15 Cooperative Breeding Influences Evolution of Communicative Complexity in Birds Gavin Leighton
- 10:30 It pays to wait: intragenerational social mobility key to long-term stability in Tibetan macaque society

Lixing Sun, Dongpo Xia, Shine Sun, Lori Sheeran, Jinhua Li

- 10:45 Fighting over food unites the birds of North America in a continental dominance hierarchy Eliot Miller, David Bonter, Charles Eldermire, Benjamin Freeman, Emma Greig, Luke Harmon, Curtis Lisle, Wesley Hochachka
- 11:00 Does boldness predict ecological outcomes in a clonal sea anemone **David Ayre**, Rick Grosberg
- 11:15 Geographical barriers as drivers for song and genetic divergence in the highland endemic Timberline Wren (Thryorchilus browni)

 Andrés Camacho

Session Sun2-B114-115 Invasive / invasion 1 Sunday - 10:15 - 11:30 Room B114-115

- 10:15 Genetic signatures of adaptation and gene surfing in replicated biological range expansions Christopher Weiss-Lehman, Silas Tittes, Nolan Kane, Ruth Hufbauer, Brett Melbourne
- 10:30 Effect of temperature and genetic structure on gypsy moth invasiveness at a range margin Trevor Faske, Lily Thompson, Chris Friedline, Salvatore Agosta, Derek Johnson, Kristine Grayson, Andrew Eckert
- 10:45 Impact of suburban landscape features on dispersal of Brachypodium sylvaticum, an invasive grass

Tina Arredondo, Mitch Cruzan

- 11:00 Distribution of co-invading specialist rhizobium mutualist limits the potential of Medicago polymorpha to invade via large and small scale dispersal events

 Zoie Lopez, Maren Friesen, Stephanie Porter
- 11:15 The Medicago polymorpha-rhizobia mutualism affects herbivores interactions Chandra Jack, Maren Friesen

Session Sun2-B116 Hybridization 2

Sunday - 10:15 - 11:30

Room B116

- 10:15 Genomic Patterns in a North American Pitcher Plant Hybrid Zone Katherine Bell, Kristen L. King, Allan Strand
- 10:30 No Easy D: Little Evidence for Introgression from Whole-Genome Data in Wild Solanum Jennafer Hamlin, Leonie Moyle
- 10:45 Functionality of genome wide introgression in old world mice Eslam Elshahat, Michael Kohn
- 11:00 Introns gained or borrowed in Daphnia: on the importance of considering evolutionary history France Dufresne
- 11:15 Population modeling reveals several layers of genetic structure in a hybridizing multispecies complex, California white oaks

 Prahlada Papper

Session Sun2-B117-119 Adaptation / genomics 2 Sunday - 10:15 - 11:30 Room B117-119

- 10:15 The timescales of selection in stickleback: Why rapid adaptation takes millions of years Thomas Nelson, William Cresko
- 10:30 Genomic signatures of local adaptation in Atlantic salmon Victoria Pritchard

- 10:45 The Gulf pipefish genome enriches our understanding of an evolutionary innovation Clayton Small, Susan Bassham, Julian Catchen, Angel Amores, Allison Fuiten, Adam Jones, William Cresko
- 11:00 Evolution after whole genome duplication insights from the salmonids Simen Sandve
- 11:15 Finding a Genomic Basis for Coral Thermal Tolerance and Assessing Adaptive Potential in Platygyra daedalea withdrawn 2017-06-21 21:38:32.0

 Holland Elder, Line Bay, Eli Meyer

Session Sun2-Ball203 SSE Spotlight - Speciation with and without sex chromosomes

Sunday - 10:15 - 11:30

Room Oregon ballroom 203

- 10:15 Speciation and sex chromosomes: (Wh)y X? Norman Johnson
- 10:30 Haldane's rule in Caenorhabditis nematodes Joanna Bundus, Asher Cutter
- 10:35 Does dominance theory explain Haldane's rule in marsupials? Eric Watson, Jeff Demuth
- 10:40 Exploring the relationship between ecological isolation and reproductive isolation in nudibranchs

 Allison Fritts-Penniman
- 10:45 Not playing by the rules: speciation genetics in the absence of sex chromosomes Suzanne Edmands
- 11:00 Are mito-nuclear incompatibilities especially important for the evolution of reproductive isolation in taxa without sex chromosomes?

 Thiago Lima, Ron Burton, Ricardo Pereira
- 11:05 The multifaceted nature of speciation: variation in reproductive isolation across a species range
 Karen Barnard-Kubow, Laura Galloway
- 11:10 The small majority: the microbiome as a unit of selection and speciation Robert Brucker
- 11:15 Patterns and drivers of reptilian speciation with and without sex chromosomes Nicole Valenzuela

Session Sun2-Ball204 Sex / recombination 1 Sunday - 10:15 - 11:30 Room Oregon ballroom 204

- 10:15 The evolution of recombination rate in finite populations during ecological speciation James Reeve, Daniel Ortiz-Barrientos, Jan Engelstädter
- 10:30 Multicellularity drives the evolution of anisogamy, sexual dimorphism Erik Hanschen
- 10:45 The measure of total recombination Carl Veller, Pavitra Muralidhar
- 11:00 Demography influences the detection of recombination hotspots Amy Dapper, Bret Payseur
- 11:15 Gametic selection, meiotic drive, sex ratio bias, and transitions between sex determination systems

Michael Scott, Matthew Osmond, Sally Otto

Session Sun2-C120-122 Ecological genetics 1 Sunday - 10:15 - 11:30 Room C120-122

10:15 Different landscape features influence seed and pollen movement in the small forb Lasthenia californica, as seen through whole chloroplast genome sequencing and nuclear genotyping-by-sequencing

Pamela Thompson, Mitch Cruzan

- 10:30 Genomic signals of selection predict climate-driven population declines in a migratory bird Rachael Bay
- 10:45 Targeted sequencing of venom genes from cone snail genomes reveals multiple genetic and ecological mechanisms shaping venom diversification

 Mark Phuong
- 11:00 Ecophysiological genomics and the desert adapted rodent, Peromyscus eremicus Matthew MacManes
- 11:15 Experimental evidence natural selection affects genome-wide geographic and host-related differentiation during sympatric divergence.

Jeffrey Feder, Meredith Doellman, Glen Hood, Gregory Ragland, Peter Meyers, Thomas Powell, Scott Egan, Mary Glover, Cheyenne Tait, Dan Hahn, Stewart Berlocher, Patrik Nosil

Session Sun2-C123 Education / outreach Sunday - 10:15 - 11:30 Room C123

- 10:15 Accepting Evolution: Factors that lead to change in students' acceptance of evolution.

 Daniel Ferguson, T. Heath Ogden*
- 10:30 Seeking Common Ground in World Views: Humility and Hubris in Science and Religion John Niedzwiecki
- 10:45 Virtual Biology Lab 2.0: improving and implementing an inquiry-based educational resource David Ford, Anna Hiatt, Thomas Jones
- 11:00 Evolutionary video games as vehicles for outreach and engagement.

 Barrie Robison, Terence Soule, Samantha Heck, Landon Wright
- 11:15 Using natural history collections to train students in informal science education Susan Masta, Annie Lindgren

Session Sun2-C124 Genomics

Sunday - 10:15 - 11:30

Room C124

- 10:15 DNA transposon dynamics among fungi with diverse lifestyles

 Anna Muszewska, Kamil Steczkiewicz, Marta Stepniewska-Dziubinska, Julia Pawlowska, Krzysztof

 Ginalski
- 10:30 Using Genomic Data to Inform Population Viability in a Long-Lived Endangered Vertebrate

 Jaime Ashander, Evan McCartney-Melstad, Peter Ralph, Bradley Shaffer
- 10:45 Evolutionary variation in growth and stress responses underlying kidney regenerative growth from snakes to human organoids

 Audra Andrew, Blair Perry, Daren Card, Drew Schield, Robert Ruggiero, Suzanne McGaugh, Amit Choudhary, Stephen Secor, Todd Castoe
- 11:00 Plastome evolution in Bignonieae (Bignoniaceae)
 Alexandre Zuntini, Lúcia Lohmann
- 11:15 Genome size evolution in neotropical salamanders. Paul Decena, Sean Rovito

Session Sun2-Ball202 SSB Symposium - A debate of conceptual issues surrounding genetic-based species delimitation in the genomic era 2

Sunday - 10:15 - 11:45

Room Oregon ballroom 202

- 10:15 When are standardized protocols for species delimitation problematic? David Hillis, E. Anne Chambers, Harry Greene
- 10:30 When are standardized protocols for species delimitation appropriate? Thomas Near
- 10:45 Moderated discussion: Should we seek a "standardized protocol" for species delimitation?

 David Hillis, L. Lacey Knowles

11:00 When are species delimitation methods misleading in the study of macroevolutionary patterns?

Catherine Wagner

- 11:15 When are species delimitation methods necessary for studying macroevolutionary patterns?

 Brian Smith
- 11:30 Moderated discussion: What is the role of morphology in species delimitation in the genomic era?

David Hillis, L. Lacey Knowles

Session Diversity in Science luncheon: Essentials of a successful diversity and inclusion program Sunday - 11:30 - 13:30 Room Oregon ballroom 201

Prior registration required (during conference registration).

Organizers: SSE Graduate Student Advisory Council with input from SSB and ASN

Topic: Join us for a discussion on identifying bias and increasing diversity across all dimensions.

Keynote speaker: <u>Lee Bitsoi</u> (Rush U.)

Session Lunch

Sunday - 11:30 - 13:30

Room On the town

Session SSE Education & Outreach committee meeting Sunday - 11:30 - 13:30 Room C126

Session Early career researchers workshop (2 of 2): Maximizing the impact of your research (British Ecological Society)

Sunday - 12:15 - 13:15

Room B113

Cost: free, no registration required. Bring your own lunch. Organizers: Chris Grieves (Methods in Ecology and Evolution)

In recent years there has been a significant increase in the number of academic articles published. At the same time, readers are changing how they find content – searching for individual articles rather than looking through journals. This interactive workshop will provide new authors with the necessary skills to maximize the impact of their research in the digital age. It will include guidance on optimising your articles for search engines, identifying the best forums to promote your work, pitching your paper to different audiences, and using social media.

Session Sun3-A105 Phylogenetic methods development 3 Sunday - 13:30 - 14:45 Room A105

- 13:30 Morphometrics on phylogenies: progress report on a likelihood approach Joe Felsenstein
- 13:45 Polymorphism-aware phylogenetic Models (PoMos) and their Applications withdrawn 2017-06-20 13:12:34.0

Carolin Kosiol, Dominik Schrempf

- 14:00 Budding or Anagenesis? Paraphyly and Ancestor-Descendant Relationships from Tip-Dating Phylogenies of Fossil Lineages

 David Bapst, Sandra Carlson
- 14:15 A FORMAL MODEL OF CLONAL EXPANSION IN BACTERIAL POPULATION GENETICS Alice Ledda, Xavier Didelot
- 14:30 A comprehensive view of the plant tree of life. Stephen Smith, Joseph Brown

Session Sun3-A106 Biogeography / evolutionary ecology Sunday - 13:30 - 14:45 Room A106

- 13:30 Niche conservatism or divergence: Insights into the evolutionary histories of Pinus taeda, P. rigida, and P. pungens

 Constance Bolte
- 13:45 Inferring diversification rates without molecular phylogenies or the fossil record: long-term diversity declines inferred for the Hawaiian archipelago

 Charles Marshall, Jun Ying Lim
- 14:00 Testing for latitudinal gradients in defence at the macroevolutionary scale Daniel Anstett, Jeff Ahern, Marc Johnson, Juha-Pekka Salminen
- 14:15 Ecological niche co-evolution through time between aye-ayes (Primates) and their major food plant, Canarium withdrawn 2017-06-13 18:44:55.0

 Jen Tinsman
- 14:30 Niche divergence in Mimulus: driven by range overlap and time Qin Li, Dena Grossenbacher, Amy Angert

Session Sun3-A107-109 Macroevolution 4 Sunday - 13:30 - 14:45 Room A107-109

- 13:30 Phylodiversity and evolution of tropical brittle-stars Guadalupe Bribiesca Contreras
- 13:45 Phylogenetic comparative methods and cultural diversity: what drives the evolution of human kinship systems?

 Catherine Sheard, Sam Passmore, Claire Bowern, Fiona Jordan
- 14:00 The Diversity of Avian Egg Shape through Time and Morphospace Mary Stoddard
- 14:15 Functional diversification of a skeletal novelty in bats Kathryn Stanchak, Sharlene Santana
- 14:30 Evaluating the evolution of trait seasonality with environment in dogwoods (Cornus): utility of a function-valued approach

 Chase Mason, Michael LaScaleia, Grey Monroe, Eric Goolsby

Session Sun3-B111-112 Quantitative genetics / life history Sunday - 13:30 - 14:45 Room B111-112

- 13:30 Genetic dissection of nutrition-induced plasticity in insulin/insulin-like growth factor signaling and median lifespan in a Drosophila multiparent population Elizabeth King, Patrick Stanley, Enoch Ng'oma, Siri O'Day
- 13:45 Evolutionary potential of nutrition-dependent effects on lifespan in a multi-parental population of Drosophila melanogaster

 Enoch Ng'oma, Anna Perinchery, Kevin Middleton, Elizabeth King
- 14:00 The Phenotypic Correlates and Quantitative Genetics of Masculinization in the Rodent,
 Octodon degus
 Derek Roff, Matthew Wolak, Loreto Correa, Mauricio Soto-Gamboa
- 14:15 Detection of candidate genes involved in life history tradeoffs in outcrossing Arabidopsis lyrata using diploid genome alignment method and allele-specific expression analyses.

 Bishwa Giri, David Remington
- 14:30 Effect of life history on the predictability of selection and adaptation to a fluctuating environment

 Olivier Cotto, Luis-Miguel Chevin

Session Sun3-B113 Behavior genetics

Sunday - 13:30 - 14:45

Room B113

- 13:30 Uncovering the genetic basis of social cognition in a 'smart' paper wasp Sara Miller, Michael Sheehan
- 13:45 The inheritance and neurogenomics of paternal care in sticklebacks

 Alison Bell
- 14:00 DNA methylation and the evolution of bee eusociality Natalia Araujo, Maria Cristina Arias
- 14:15 Genome-wide selection components analysis in a fish with male pregnancy Sarah Flanagan, Adam Jones

Session Sun3-B114-115 Invasive / invasion 2 Sunday - 13:30 - 14:45 Room B114-115

- 13:30 Evolution of an agricultural weed, Helianthus annuus Emily Drummond, Loren Rieseberg
- 13:45 Evidence for eco-evolutionary dynamics during aquatic invasions Michael Gillis, Matthew Walsh
- 14:00 Two types of cryptic invasion of the shrub Cytisus scoparius in Europe. *Ursula Brandes*, Siri Fjelheim, Beate Beatriz Furevik, Lene Rostgaard Nielsen, Erik Dahl Kjær
- 14:15 Tracking the evolution of a plant invasion through time and space Kathryn Turner
- 14:30 Hybridization with an invader and position along the watershed influence morphological variation in populations of Orconectes sanbornii

 Angela Roles, Elisa Henderson

Session Sun3-B116 Speciation / reproductive isolation Sunday - 13:30 - 14:45 Room B116

- 13:30 Choosy natives and industrious immigrants: Genome duplication and pollinator discrimination in an ecologically dominant southwestern desert shrub (Larrea tridentata; Zygophyllaceae)

 Robert Laport, Diana Pilson
- 13:45 Divergence before the host shift? Prezygotic reproductive isolation among three varieties of a specialist fly on a single host plant

 Alaine Hippee, Marty Condon, Andrew Forbes
- 14:00 Battle of the sexes: the relative importance of female versus male mate-choice in Drosophila speciation

 Roman Yukilevich, Elizabeth Peterson
- 14:15 Assortative mating by an obliquely transmitted local cultural trait reduces hybridization Justin Yeh
- 14:30 Male-driven reproductive and agonistic character displacement in darters Rachel Moran. Becky Fuller

Session Sun3-B117-119 Adaptation / genomics 3 Sunday - 13:30 - 14:45 Room B117-119

- 13:30 Variants of large-effect underlie sex-specific resistance to a transmissible cancer Mark Margres, Brendan Epstein, Menna Jones, Rodrigo Hamede, Paul Hohenlohe, Andrew Storfer
- 13:45 The genetics of rapid adaptation to extreme environments in house mice across the Americas Kathleen Ferris, Mallory Ballinger, Gabriela Heyer, Michael Nachman
- 14:00 The color code: a multi-trait perspective on color evolution in a diverse salamander lineage (Salamandra spp.).

 James Burgon, David Vieites, Sebastian Steinfartz, Miguel Vences, Barbara Mable, Kathryn Elmer
- 14:15 The predictability of genomic changes underlying a recent host shift in Melissa blue butterflies

 Samridhi Chaturvedi, Lauren Lucas, Matthew Forister, Zachariah Gompert

Session Sun3-Ball202 SSB Symposium - Phylogenetic approaches to connecting genotypes to phenotypes 1 Sunday - 13:30 - 14:45 Room Oregon ballroom 202

- 13:30 Introduction to phylogenetic approaches to associating genes and phenotypes Casey Dunn, Stacey Smith
- 14:00 Genome-wide association mapping in the context of complex evolutionary histories: new phylogenomic approaches

 Hussein Hejase, Natalie VandePol, Gregory Bonito, Patrick Edger, Kevin Liu
- 14:30 Promises and pitfalls of phylogenetic mapping in species radiations Fabio Kuriki Mendes, Matthew Hahn

Session Sun3-Ball203 SSE Spotlight -Sexual selection via male competition and the process of speciation Sunday - 13:30 - 14:45 Room Oregon ballroom 203

- 13:30 Male competition as the flipside of sexual selection in speciation Janette Boughman, Jason Keagy
- 13:45 Female competition in a hybrid zone between sex-role reversed species Sara Lipshutz, Elizabeth Derryberry
- 13:50 Sexual conflict can constrain the evolution of reinforcement Alison Wardlaw, Yaniv Brandvain
- 13:55 Divergent versus convergent sexual selection from male competition: ecology is key Alycia Lackey, Janette Boughman
- 14:00 Can male competition enhance reproductive isolation? Evaluating the association between success in intraspecific and heterospecific sperm competition

 Leonie Moyle, Jeremy Davis
- 14:15 Emergence and diversification of male grasping structures in the water strider genus Rheumatobates

 Augustin Le Bouquin, Abderrahman Khila, Locke Rowe
- 14:20 Phenotypic and genetic divergence in weaponry across a broad landscape in a cactus-feeding bug

 Pablo Allen, Christine Miller
- 14:30 Contrasting patterns of male competition and female choice in recently diverged threespine stickleback color types Robin Tinghitella

Session Sun3-Ball204 Sex / recombination 2 Sunday - 13:30 - 14:45 Room Oregon ballroom 204

13:30 Genome dynamics of the anciently asexual bdelloid rotifers

Reuben Nowell, Pedro Almeida, Christopher Wilson, Alan Tunnacliffe, Chiara Boschetti, Tim Barraclough

- 13:45 Phenotypic Versus Genotypic Sex: Characterization of Multiple Loci in a Complex Sex Determination System

 Ashley Elias
- 14:00 Sex chromosome evolution in voles
 Polly Campbell, Matthew B. Couger, Scott Roy
- 14:15 Sex reinforces DNA methylation in plants

 Adam Bewick, Alex Harkess, Zefu Lu, Paul Fourounjian, Brigitte Hofmeister, Robert Schmitz, Joachim Messing, Blake Meyers
- 14:30 Evolution of sexual dimorphism in the genome-wide recombination rate in house mice April Peterson, Bret Payseur

Session Sun3-C120-122 Ecological genetics 2 Sunday - 13:30 - 14:45 Room C120-122

- 13:30 Range-wide genetic diversity of the ringed salamander (Ambystoma annulatum) Jacob Burkhart, Chelsey Kroese, Emily Puckett, Raymond Semlitsch, Lori Eggert
- 13:45 Genetic diversity and hybridization in New Zealand's Selliera (Goodeniaceae) Vaughan Symonds, Kay Pilkington
- 14:00 Environmentally-driven morphological and genomic divergence in forest mice (Praomys sp.) across Central África Katy Morgan
- 14:15 Ecological Transcriptomics of Trophic Divergence in the Highly Polymoprhic Arctic Charr: Uncovering the Molecular Phenotype Madeleine Carruthers, Arne Jacobs, Kathryn Elmer, Colin E. Adams, Natalia V. Gordeeva, Sergey S. Aleksevev
- 14:30 Genetic and evolutionary bases for structural color shift in the Buckeye butterfly Rachel Thayer

Session Sun3-C123 Contemporary evolution 1 Sunday - 13:30 - 14:45 Room C123

- 13:30 Eco-evolutionary consequences of mosquitofish domestication Zachary Wood, David Fryxell, Eric Palkovacs, Michael Kinnison
- 13:45 Combining niche shift models and population genetics predicts rapid phenotypic evolution during invasion of a marine alga Erik Sotka
- 14:00 Rapid evolution following experimental introduction in Brassica rapa Steven Franks, Michael Sekor
- 14:15 Contemporary evolution and genetic variation in an invasive species affect biotic and abiotic resistance to invasion Casey terHorst
- 14:30 Adaptive potential of native and introduced ragweed populations Brechann McGoey, John Stinchcombe

Session Sun3-C124 Genomics / methods / theory Room C124 Sundav - 13:30 - 14:45

13:30 Genotyping in thousands by sequencing (GT-seq): A low cost, high-throughput, targeted SNP genotyping method

Nate Campbell, Stephanie Harmon, Shawn Narum

- 13:45 Expressed Exome Capture Sequencing (EecSeq): a method for cost-effective exome sequencing of non-model organisms **Jonathan Puritz**, Kathleen Lotterhos
- 14:00 Co-association networks reveal the genetic architecture underlying adaptation to a multivariate environment Kathleen Lotterhos, Sam Yeaman, Kathryn Hodgins, Jonathan Degner, Sally Aitken
- 14:15 The effects of low coverage sequence data on demographic reconstruction Angela Taravella, Melissa Wilson Sayres
- 14:30 Modeling next-generation sequencer sampling error dramatically reduces false positives in genetic structure tests Rebecca M. Hamner, Jason D. Selwyn, Evan Krell, Scott A. King, Chris Bird

Session Coffee break

Sunday - 14:45 - 15:15

Room Exhibit Hall A



Session Sun4-A105 Phylogeography

Sunday - 15:15 - 16:30

Room A105

- 15:15 Evaluating mechanisms of diversification in a Guineo-Congolian forest frog using demographic model selection

 Daniel Portik, Adam Leache, Danielle Rivera, David Blackburn, Matthew Fujita
- 15:30 Sequence analysis of California populations of redwood sorrel (Oxalis oregana) supports post-glacial recolonization of the Pacific Northwest from northern and southern refugia Chris Brinegar
- 15:45 Using ddRAD-seq phylogeography to test for genetic effects of headwater river capture in Hypostomus catfishes from central Brazil

 Justin Bagley, Pedro De Podestà Uchôa de Aquino, Tomas Hrbek, Sandra Hernandez, Francisco Langeani, Guarino Colli
- 16:00 Using hypothesis testing to unearth phylogeographic patterns within Mole Skinks Kathryn Mercier, Milagros Baylac, Christopher Parkinson
- 16:15 Mito-nuclear discordance in the evolutionary history of a widespread passerine (Troglodytes aedon)

 Dario Lijtmaer, Cecilia Kopuchian, Pablo Tubaro, Leonardo Campagna

Session Sun4-A106 Disease

Sunday - 15:15 - 16:30

Room A106

- 15:15 Host response to changes in the microbiome through symptom progression in Sea Star Wasting Disease

 Melanie Lloyd, Melissa Pespeni
- 15:30 Unlocking the story in the swab: A new genotyping assay for the amphibian chytrid fungus Allie Byrne, Erica Bree Rosenblum, Andrew Rothstein, Thomas Poorten, Matthew Settles, Jesse Erens
- 15:45 Measuring the within tumor clonal spatial mixing level with phylogenies of phased haplotypes Yuantong Ding
- 16:00 Population genetic analysis of the DARC locus (Duffy), involved in malaria resistance in humans
 - Omar Cornejo, Kimberly McManus, Martin Sikora, Carlos Bustamante, Brena Henn, Angela Taravella
- 16:15 Natural hybridization in poplars reveals new genomic architectures of disease resistance. *Karl Fetter, Stephen Keller*

Session Sun4-A107-109 Macroevolution 5 Sunday - 15:15 - 16:30 Room A107-109

- 15:15 Variation in population differentiation and speciation rate in a species-rich clade of lizards Sonal Singhal, Huanteng Huang, Maggie Grundler, Raquel Marchan, Pascal Title, Daniel Rabosky
- 15:30 Macroevolution of the dewlap and diversification of Anolis lizards Travis Ingram
- 15:45 Can we predict signal evolution in birds? Chris Cooney, Gavin Thomas

16:00 Hybridization accelerates speciation rates in salamanders

Austin Patton, Brendan Epstein, Jonathan Eastman, Luke Harmon, Andrew Storfer

16:15 Life in Trees: Macroevolution of Morphology and Diversity in Arboreal Plethodontid Salamanders

Erica Baken, Dean Adams

Session Sun4-B111-112 Quantitative genetics / sexual selection

Sunday - 15:15 - 16:30

Room B111-112

15:15 The quantitative genetics of sexual dimorphisms: how important are sex-linkage and dominance?

Daphne Fairbairn, Derek Roff

- 15:30 Using sexually antagonistic skewers to explore the genetic architecture of sexual dimorphism in Anolis lizards

 Robert Cox, Joel McGlothlin*
- 15:45 The genomic architecture of repeated sexual signal divergence Thomas Blankers, Kerry Shaw
- 16:00 Inter-population and genetic variation in the thermal sensitivity of reproductive behaviors of the vibrationally-singing insect Enchenopa binotata

 Kasey Fowler-Finn, Dowen Jocson, Morgan Smeester
- 16:15 The role of the Y-chormosome in the evolution of autosomally coded traits Ian Kutch, Kenneth Fedorka

Session Sun4-B113 Behavior / genomics Sunday - 15:15 - 16:30 Room B113

- 15:15 Adaptation and acclimation in a changing climate: integrating physiology, transcriptomics, and genomics of a thermal specialist

 Maria Tonione
- 15:30 Genetic architecture of social organisation in the ant Pheidole pallidula

 Emeline Favreau, Rodrigo Pracana, Claude Lebas, Eckart Stolle, Max Reuter, Christophe Eizaguirre,
 Yannick Wurm
- 15:45 Comparative population genomics in halictid bees provides insights into the evolution of social behavior

 Ben Rubin, Sarah Kocher
- 16:00 The genetic basis for song-preference coevolution in the Hawaiian cricket Laupala Mingzi Xu, Kerry Shaw
- 16:15 Genomic perspectives on the recurrent evolution of brood parasitic behavior Christopher Balakrishnan, Matthew Louder, Michael S. Brewer, Claire N. Spottiswoode, Michael Sorenson, Mark Hauber

Session Sun4-B114-115 Population genetics: inference of selection 1

Sunday - 15:15 - 16:30

Room B114-115

- 15:15 Can pooled sequencing provide evidence of viability selection in raptors?

 Jacqueline Doyle, J. Andrew DeWoody, Janna Willoughby, Nadia Fernandez, Doug Bell
- 15:30 Assessing differential selection on immunity-related genes across monarch butterfly populations

 Wen-Hao Tan, Nicole Gerardo, Jacobus de Roode
- 15:45 Adaptation and gene flow among tidal marsh sparrows withdrawn 2017-06-04 07:33:29.0 Gemma Clucas
- 16:00 Genome-wide evidence of evolution and adaptation in the invasive Florida python population Daren Card, Blair Perry, Richard Adams, Audra Andrew, Drew Schield, Acacia Young, Tereza Jezkova, Mike Rochford, Frank Mazzotti, Kristen Hart, Margaret Hunter, Todd Castoe

16:15 Genomic data provides new insights into the origins of adaptive variation in the bush monkeyflower (Mimulus)

Madeline Chase, Sean Stankowski, Matt Streisfeld

Session Sun4-B116 Speciation / hybridization Sunday - 15:15 - 16:30 Room B116

- 15:15 Genome-wide admixture, differential introgression, and signatures of reproductive isolation in a howler monkey hybrid zone

 Marcella Nidiffer, Liliana Cortes Ortiz
- 15:30 Tracing the footprints of a moving Setophaga warbler hybrid zone Silu Wang, Darren Irwin
- 15:45 Ancient and recent hybridisation and polyploidy in monkeyflowers (Mimulus) Mario Vallejo-Marin
- 16:00 Assortative mate choice and maintenance of reproductive barriers: lessons from a phenotypically diverse avian species

 Georgy Semenov, Elizabeth Scordato, David Khaydarov, Chris Smith, Nolan Kane, Alexander Badyaev, Evgeniy Koblik, Yaroslav Red'kin, Rebecca Safran
- 16:15 The genetic basis and evolutionary dynamics of a hybrid incompatibility in Mimulus Matthew Zuellig, Andrea Sweigart

Session Sun4-B117-119 Adaptation / genomics 4 Sunday - 15:15 - 16:30 Room B117-119

15:15 Whole genome scan reveals the multigenic basis of recent tidal marsh adaptation in a sparrow

Petra Deane-Coe, Bronwyn Butcher, Russell Greenberg, Richard Harrison, Irby Lovette

- 15:30 Herbarium accessions for temporal studies of adaptation to climate in Arabidopsis thaliana. Lua Lopez Perez, Stephanie Marciniak, Victoria DeLeo, George "PJ" Perry, Jesse Lasky
- 15:45 Characterizing potential adaptations of Tasmanian devil populations in the face of a transmissible cancer

 Alexandra Fraik, Brendan Epstein, Mark Margres, Rodrigo Hamede, Menna Jones, Paul Hohenlohe, Joanna Kelley, Andrew Storfer
- 16:00 The Genetic Mechanisms of Local Adaptation David Lowry, Billie Gould
- 16:15 LANDSCAPE GENOMICS OF STEELHEAD TROUT (ONCORHYNCHUS MYKISS) THROUGHOUT THE COLUMBIA RIVER BASIN.

 Steven Micheletti, Shawn Narum

Session Sun4-Ball203 SSE Spotlight - Evolution of hybrid incompatibilities

Sunday - 15:15 - 16:30

Room Oregon ballroom 203

- 15:15 Genetic and evolutionary causes of hybrid incompatibilities in Mimulus Andrea Sweigart
- 15:30 Causes of hybrid incompatibility polymorphism early in speciation Erica Larson, Jeff Good
- 15:35 Ecological divergence and the evolution of hybrid phenotypic incompatibilities Ken Thompson
- 15:40 Assortative mating and hybrid incompatibility in natural hybrid populations Molly Schumer
- 15:45 Gene regulation and speciation in house mice Michael Nachman, Katya Mack, Polly Campbell
- 16:00 Complex hybrid incompatibilities inferred from widespread pairwise interactions among sterility loci

 Rafael Guerrero, Christopher Muir, Sarah Josway, Leonie Moyle

- 16:05 Incompatibility between mutations from parallel adaptation to a common environment Jasmine Ono, Aleeza Gerstein, Sally Otto
- 16:10 Parental conflict, parent of origin effects, and the evolution of hybrid seed failure in Mimulus Jenn Coughlan, John Willis
- 16:15 Dissecting the role of sex chromosomes in Drosophila speciation: genetic mapping of hybrid male sterility genes

Colin Meiklejohn, Emily Landeen, Sarah Kingan, Anthony Geneva, Jeffrey Vedanayagam, Christina Muirhead, Kathleen Gordon, Daniel Garrigan, David Stern, Daven Presgraves

Session Sun4-Ball204 Sex / recombination 3 Sunday - 15:15 - 16:30 Room Oregon ballroom 204

- 15:15 Rapidly Evolving Recombination Hotspots in Threespine Stickleback Alice Shanfelter, Michael White, Madison Zwink, Lucas Nell
- 15:30 Hot chocolate: Recombination hotspots across the Theobroma cacao genome Enrique Jimenez Schwarzkopf, Omar Cornejo
- 15:45 An ancient UV sex chromosome system in mosses Sarah Carey, Adam Payton, Matthew Johnson, Stuart McDaniel
- 16:00 Sex determination in East African cichlid fishes: limited options or space for novelty?

 **Astrid Boehne*, Nicolas Lichilin, Athimed El Taher, Geoffrey Fucile, Walter Salzburger*
- 16:15 The effects of insulin signaling on sexually dimorphic gene expression in Drosophila Rita Graze, RueiYing Tzeng, Tiffany Howard, Michelle Arbeitman

Session Sun4-C120-122 Ecological genetics 3 Sunday - 15:15 - 16:30 Room C120-122

15:15 Population genomics of Boechera stricta reveals demographic history and signatures of selection

Baosheng Wang, Tom Mitchell-Olds

- 15:30 Conflict between symbionts: Can plants repel parasitic nematodes while attracting mutualistic bacteria?

 Corlett Wood, Bonnie Pilkington-Lindo, Priya Vaidya, John Stinchcombe
- 15:45 Coevolution of skeletal and behavioral traits in deer mice Emily Hager, Hopi Hoekstra
- 16:00 Biome and migratory behaviour significantly influence vertebrate genetic diversity

 Nadia Fernandez, Janna Willoughby, Mekala Sundaram, Bhagya Wijayawardena, Maureen Lamb,
 steven kimble, Yanzhu Ji, Jennifer Antonides, Nicholas Marra, J. Andrew DeWoody
- 16:15 Taking Isolation by Distance into the Temporal Dimension Arthur Weis, Madeline Peters

Session Sun4-C123 Contemporary evolution 2 Sunday - 15:15 - 16:30 Room C123

- 15:15 Eco-evolutionary feedbacks predict the time course of rapid life-history evolution Joseph Travis, Corey Handelsman, Cameron Ghalambor, Ron Bassar, David Reznick
- 15:30 Contemporary evolution of monarch butterflies in their introduced range Micah Freedman, Hugh Dingle
- 15:45 Predation induced life history evolution Kelsey Lyberger
- **16:00 Detecting short-term selection in a pedigreed natural population** *Nancy Chen, Ivan Juric, Reed Bowman, John Fitzpatrick, Andrew Clark, Graham Coop*
- 16:15 New methods for detecting selection on structural variants and introgressions Luca Ferretti

Session Sun4-C124 Genomics / gene flow

Room C124

- 15:15 Reconstructing the demographic history of Atlantic salmon across its distribution range using approximate Bayesian computations *Quentin Rougemont*, Louis Bernatchez
- 15:30 Genomics of adaptive lateral gene transfers in plants
 Luke Dunning, Jill Olofsson, Colin Osborne, Patrik Nosil, Pascal-Antoine Christin
- 15:45 Exploring the role of environmental bacteria in the spread of antibiotic resistance genes Lauren Brooks, Mark Sistrom
- 16:00 We're one, but we're not the same: Shallow evolutionary divergence between two andean hummingbirds

 Catalina Palacios, silvana Garcia, Juan Luis Parra, Andrés Cuervo, F. Gary Stiles, John McCormack,

Leonardo Campagna, Daniel Cadena

16:15 Genomics and gene flow in the Galapagos finches

Lucinda Lawson, John Niedzwiecki, Kenneth Petren

Session Sun4-Ball202 SSB Symposium - Phylogenetic approaches to connecting genotypes to phenotypes 2
Sunday - 15:15 - 16:45 Room Oregon ballroom 202

15:15 Applying stochastic models for expression evolution to identify genes with adaptive expression

Rori Rohlfs, Scott Roy, Ksenia Arzumanova, Manuel Irimia, Jamie Moon

- 15:45 Distinguishing among modes of convergent evolution using population genomic data Kristin Lee, Graham Coop
- 16:15 Evaluating the evidence for protein coding convergence in phenotypically convergent anoles Russell Corbett-Detig

Session SSB Social Mixer Sunday - 16:30 - 17:30 Room Oregon ballroom 201

Open to all SSB members - students, postdocs & faculty. Come chat with your SSB colleagues. Cash bar and SSB grad member free drink tickets accepted.

Session SSE Hamilton award judges meeting
Sunday - 16:30 - 17:30 Room B110

Session Sun5-Ball202 SSB Presidential address & award announcements

Sunday - 17:30 - 18:30

Room Oregon ballroom 202-204

One Tree to Rule Them All Anne Yoder

Session SSB Members business meeting
Sunday - 18:30 - 19:30 Room A105

Open to all SSB members.

Poster Session Sun-01 Undergrad Diversity Sunday - 18:30 - 20:30 Room EHA-01

- 1 Evolution of the functional determinants in epigenetics readers Cristobal Carrera, Jordon Rahaman, Jessica Siltberg-Liberles
- 2 No evidence for adaptation to the local mutualistic rhizobium in the legume Medicago lupulina Isabela Borges, Tia Harrison, Corlett Wood, John Stinchcombe
- 3 Testing whether coevolution occurred in a microbial mutualism
 Blanca Rodriguez, Norhan Algetany, Navriti Sharma, Stephen Wallen, Amina Becic, Nathan Kaneko,
 Claire Dang, Helen Triplett, Willie Christianto, Colin Feng, Kristina Hillesland
- 4 Quantifying the role lipids play in primate brain adaptation. Mariamagdalena Ortiz
- 5 Darwin's Demons: An Evolutionary Video Game Samantha Heck, Terence Soule, Landon Wright, Spencer Gomez, Barrie Robison
- 6 Return of the African Wild Dog Lycaon pictus to the Serengeti Anastasia Wlaschin-Wiest
- 7 Sex-Specific Dispersal of Urban Brown Rats (Rattus norvegicus)
 Destiny Mims, Emily Puckett, Matthew Combs, Jason Munshi-South
- 8 Variation in dominance traits and body condition in urban Anolis cristatellus Derek Briggs
- 9 Effects of Viral Mutations in the Evolution of Thermotolerance Erin McClure, Hannah Busch, Sonia Singhal, Benjamin Kerr
- 10 More than meets the eye? Protective functions of body color pigments in Hawaiian damselflies

 Edward Hsieh, Jongbeom Park, Idelle Cooper, Jonathan Brown
- 11 Assessment of prevalence and phylogenetic diversity of avian haemosporidians from Italy Taylor Verrett, Gary Voelker, Johanna Harvey
- 12 The distribution and diversity of symbiotic microbial communities within a widespread Neotropical ant host (Cephalotes atratus)

 Destiny Reeves
- 13 Unraveling the diversification of Equidae at different taxonomic levels and under different preservation models

 Rodolfo Graciotti, Mathias Pires, Tiago Quental
- 14 The Synonymous and nonsynonymous substitution ratios (dN/dS) associated with habitat transitions in the Gastropoda

 Alyssa Lau, Jesse Czekanski-Moir, Teresa Rose Osborne, David Bullis, Rebecca Rundell
- 15 Retention of the Symbiosis Island in a Mesorhizobium population Angeliqua Montoya, Stephanie Porter, Emily Helliwell, Amanda Khosravifar
- 16 Extracellular proteases: a potential Bacillus subtilis defense mechanism against bacteriophage infection

 Madison Strine, Jenna DeCurzio, Veronique Delesalle, Greg Krukonis
- 17 The genomic health of ancient hominins Taylor Cooper, Joseph Lachance
- 18 Morphological stasis despite deep genetic divergences in pseudoscorpions Emmanuelle Callaghan, Trov Olsen, Susan Masta
- 19 Understanding the origins and diversity of wild and inbred brown rats (Rattus norvegicus) using nuclear and mitochondrial genome analysis

 Olivia Micci-Smith, Emily Puckett, Jason Munshi-South
- 20 Isle Royale Populations of Primula mistassinica are Isolated by Geography, Genetics, and Pollinators
 Jonathan MacDougall, Luana Maroia, Joan Edwards
- 21 Approaches and troubleshooting using custom TaqMan in a non-model organism, Mimulus guttatus

 Toshiro Newsum*, Vanessa Koelling
- 22a Rapid, high-throughput quantification of phenotypes via crowdsourcing Emmanuel D'Agostino, Caitlin Lewarch, Hopi Hoekstra
- 22b Phenotypic consequences of sexual conflict in the context of reproductive mode polymorphism

 Emma Greimann, Maurine Neiman, Joel Sharbrough

Topic Sun-02 Adaptation

23 Adaptive response during historic drought in a Sierran monkeyflower: a resurrection experiment

Jason Sexton, Erin Dickman, Steven Franks

- 24 Acute and chronic artificial selection on heat tolerance result in different thermal performance curves in Drosophila subobscura.

 Andres Mesas, Angélica Jaramillo, Luis Castaneda
- 25 Convergence in Gecko Toe Pad Shape Travis Hagey
- 26 Heat tolerance responses to different warming scenarios in the invasive fly, Drosophila subobscura

 Luis Castaneda, Andres Mesas, Angélica Jaramillo

Topic Sun-03 Bioinformatics

- 27 Uncovering Constraints on the Evolution of Genome Architecture: A Computational Analysis of Models of Gene Rearrangement in Metazoan Mitochondrial DNA Abigail Bisesi, Angela Roles
- 28 Development of Software for Genome Sequence Analysis

 David Farr
- 29 An Analysis of Lossless Compression Algorithims for Genomic Data Abigail Howell
- 30 Degeneration of a nuclear rRNA group I intron in Teloschistes chrysophthalmus Dawn Simon, Audrey Codina, Dawson Johnson, Derek Kleier, Stefanie Hartmann, Jolanta Miadlikowska, Ester Gaya, François Lutzoni

Topic Sun-04 Coevolution

- 31 Fitness benefits of paternal mitochondrial transmission in hybrids Joseph Ross, Shadi Adineh
- 32 Are complementary male and female genital morphologies genetically coupled? Exploring the genetic architecture of sexually coevolved reproductive structures in a damselfly withdrawn 2017-06-19 17:01:01.0

 Alexandra Barnard, John Masly
- 33 Community-scale analysis of lizard gut microbiomes Iris Holmes
- 34 Novel compounds extracted from endophytes isolated from Juniperus osteosperma Schyler Ellsworth, Merrill Halling

Topic Sun-05 Computational biology

- 35 An integrated mathematical/computational approach to define the impact of TGF-Beta inhibition on the evolution of bone metastatic prostate cancer David Basanta, Leah Cook, Arturo Araujo, Conor Lynch
- 36 Stop and go, fast and slow: can the Covarion model of rate variation improve our understanding of phylogenies and molecular evolution?

 Ashley Schoonmaker*, Jeremy Brown, Lyndon Coghill, Lauren Rodriguez
- 37 Not all those who wander are lost: A simulation modelling framework for exploring the interplay between dispersal evolution and environmental adaptation in plants

 Courtney Van Den Elzen, Nancy Emery, Samuel Flaxman, Brett Melbourne

Topic Sun-06 Diversification

- 38 Reticulate evolution and species delimitation in Asian Spiranthes (Orchidaceae) Siddharthan Surveswaran, Mei Sun
- 39 Climate-driven diversification revealed by calibrating avian time-trees Santiago Claramunt, Emmanuel Paradis, Joel Cracraft
- 40 Patterns in global diversification of the hyperdiverse plant lineage Psychotrieae (Rubicaeae) withdrawn 2017-06-16 14:41:25.0

 John Paul, Elaine Zhang, Nila Le
- 41 Life history is associated with accelerated rates of climatic niche evolution in salamanders Samuel Weaver, Ken Kozak
- 42 Elaborating the evolution and phylogenetic distribution of fungal ecologies and nutritional modes

 Matthew Nelsen, C. Kevin Boyce

Topic Sun-07 Ecological genetics

- 43 Horizontal gene transfer as a mechanism for convergent evolution in divergent arthropods Jennifer Kovacs , Jack Werren
- 44 Sharp adaptive divergence of Hawaiian picture-wing Drosophila across an elevation gradient evidenced by whole transcriptome profiling and measures of physiology and behavior *M. Renee Bellinger*, Jonathan Eldon, Don Price
- 45 Evolutionary response of Carbonic Anhydrase (isoforms 1, 2, and 5) in Daphnia pulicaria populations across a pH gradient in three North American lakes

 Billy Culver, Francisco Acosta
- 46 Subsampling Loci From a Large RAD Dataset to Estimate Effective Population Size Joseph Braasch, Brittany Barker, Katrina Dlugosch

Topic Sun-08 Education

- 47 Changes in Acceptance of Evolution and Associated Factors During a Year of Introductory Biology Instruction
 Ryan Dunk, Jason Wiles
- 48 Teaching Evolution & Biology to a Diversity of Learners and Viewpoints: Accessibility, Inclusion, and Belief Cheryl Pinzone
- 49 Cretaceous Crusaders: Educational gaming as a tool to teach evolution Chandra Jack, Thomas Haynes, Mia Swanton, Maren Friesen, Barrie Robison, Terence Soule
- 50 Signs along the learner's way in evolutionary biology
 Víctor Rogelio Hernández Marroquín , Rodolfo Salas Lizana, Rosaura Ruiz Gutiérrez, Javier Fregoso
 Urrutia, Ana Escalante García, Gloria Montaño Aguilar, Eréndira Alvarez Pérez
- 51 A Course-based Undergraduate Research Experience (CURE) to Explore NextGen Sequences of Rhagoletis Mitogenomes, Personal Genotypes and Core Concepts in Evolution James Smith, Patrick Edger
- 52 QUBES: Building a community to promote undergraduate quantitative biology education Gabriela Hamerlinck
- **53** Evaluating the use of integrative Evo-Ed cases for evolution education *Alexa Warwick*, *Peter White, Louise Mead, James Smith*

Topic Sun-09 Evolutionary ecology

- Coexistence mechanisms as promoters of divergence Kelly Thomasson
- 55 Phenotypic, Genetic, and Epigenetic Population Structure of Olympia Oysters in Puget Sound, Katherine Silliman, Steven Roberts, Tynan Bower
- Variation of Leaf Hydraulic Conductance in Genus Rhododendron Sharon Danielson, Lawren Sack, Juliana Medeiros
- The Role of Urbanization & Community Assembly on the Microbial Defense Mechanisms of Plethodon cinereus Elle Barnes
- Rapid assessments of genetic viability predict long-term population trends in wood frogs Jonathan Richardson, David Skelly, Robin Waples
- Phenology of Drosophila species across a temperate growing season Jennifer Gleason, Paula Roy, Elizabeth Everman, Terry Gleason, Theodore Morgan

Topic Sun-10 Evolutionary theory

- Role of Sustainability in Evolution of Species 60 PANKAJ VALLABH
- **Environmental Robustness in Populations with Differential Adaptive Histories** 61 Ao Wang
- 62 **Aneuploidy and Sex Chromosome Evolution** Heath Blackmon
- Chaos and the predictability of evolution in a changing environment Artur Rego Costa, Florence Debarre, Luis-Miguel Chevin
- Insights from Fisher's geometric model on the likelihood of speciation under different environmental scenarios Ryo Yamaguchi, Sally Otto

Topic Sun-11 Experimental evolution

- 65 Experimental evolution of sexual dimorphism in the nematode C. remanei. Maria Modanu
- Effects of experimental drought on rapidly-flowering Brassica rapa Stephen Johnson, Steven Franks
- Adaptation to spatially structured environments results in longer generations and lower infectivity in T7 bacteriophage Richard Heineman
- 68 Autocatalytic Self-assembly of a Catalytic RNA from Many Small Fragments Niles Lehman, Tharuka Jayathilaka

Topic Sun-12 Expression studies/transcriptomics

- Comparative transcriptomics of Lithocarpus with and without leaf epicuticular wax 69 Chih-Wei Huang, Bing-Hong Huang, Pei-Chun Liao
- Evolution of gene expression following whole genome duplication Gareth Gillard, Rori Rohlfs, Torgeir R Hvidsten, Simen Sandve
- A global co-expression network approach for connecting genes to specialized metabolic pathways in plants Jennifer Wisecaver
- Comparing gene repertoires in annual and perennial soybean (Glycine) Michael Song, Jeremy Coate, Jeff Doyle
- Diet and life-stage remodeling of lipid metabolism regulation in the duplicated Atlantic salmon genome

Thomas Harvey, Gareth Gillard, Simen Sandve, Torgeir R Hvidsten

- 74 Transcriptional Activity of a Sex-specific B Chromosome in African Cichlids Frances "Betsy" Clark, Matthew Conte, Thomas Kocher
- 75 Evolutionary rates and expression patterns in allopolyploid brown algae Filipe Sousa, Joao Neiva, Linda Anderson, Peter Raimondi, Nelson Martins, Ester Serrao, Gareth Pearson

Topic Sun-13 Gene flow

- 76 Microgeographic analysis of gene flow across an elevational gradient in northeastern red spruce forests

 Brittany Verrico, Stephen Keller**
 - On a second to a select the tensor
- 77 On a puzzling relationship between a bacterium and its virus Olga Zhaxybayeva, Shannon Soucy, Migun Shakya, Xin Chen
- 78 Testing for adaptive introgression between Mus musculus domesticus and Mus spretus Sarah Banker, Michael Nachman
- 79 Landscape genetics of fine-scale seed dispersal in Lasthenia californica Avery Pheil, Mitch Cruzan, Pamela Thompson
- 80 Genomic regions with the genes related to species differences between two Sulawesi macaque species *Yohey Terai*
- 81 Inference of gene flow across spatial landscapes Erik Lundgren, Peter Ralph
- 82 The relative frequency of local adaptation to balanced polymorphism inferred from the Sunflower transcriptome.

 Mark Welch*, Michael Kartie
- 83 Effects of novel P transposable element invasion in Drosophila yakuba Antonio Serrato, Stephania Zhang, Daniel Matute
- 84 Inferring genetic connectivity in real populations, exemplified by coastal and oceanic Atlantic cod Ingrid Spies

Topic Sun-14 Genomics

- 85 Investigating gene loss in the reptilian linneage Erin Gilbertson, Heath Blackmon, Suzanne McGaugh
- 86 Fungal lifestyle reflected in serine protease repertoire
 Anna Muszewska, Marta Stepniewska-Dziubinska, Kamil Steczkiewicz, Julia Pawlowska, Agata
 Dziedzic, Krzysztof Ginalski
- 87 Exploring the oral microbiome of ancient humans Carlie Harding, Martin Sikora, Omar Cornejo
- 88 Genomic Architecture of Inbreeding Depression in C. remanei Paula Adams, Janna Fierst

Topic Sun-15 Geographic variation

- 89 Populations that vary in the ability to self-fertilize display similar levels of reproductive assurance under pollen limitation

 Lia Leibman, Anne Rowe, Matthew Koski, Laura Galloway
- 90 The Geography of Morphology in the Big Water Crayfish Species Complex Melyssa Minto, Bronwyn Williams
- 91 Phenotypic selection in Ponderosa Pine (Pinus ponderosa) under warm-dry climate.

 Marcus Warwell
- 92 Demographic History of the Drosophila subquinaria Complex Paul Ginsberg, Devon Humphreys, Brooke White, Kelly Dyer

Topic Sun-16 Host-parasite

- 93 Evolutionary History of Howler Monkeys and their Pinworms Xorla Ocloo, Liliana Cortes Ortiz
- 94 Evolution of antigenic diversity in the malaria parasite P. falciparum withdrawn 2017-06-07 15:13:10.0
 - **Mary Rorick**, Yael Artzy-Randrup, Shazia Ruybal-Pesántez, Kathryn Tiedje, Thomas Rask, Abraham Oduro, Anita Ghansah, Kwadwo Koram, Karen Day, Mercedes Pascual
- 95 Evolution of natural enemies of oak gall-forming wasps Anna Ward, Andrew Forbes
- 96 Helminth infections in male garter snakes (Thamnophis spp.) and their implications for host reproduction
 - Emily Uhrig, Christopher Friesen, Sean Spagnoli, Michael Kent, Robert Mason
- 97 Host population dynamics lead to diploid male production and decline of a natural hyperparasitoid population
 Abhilash Nair, Etsuko Nonaka, Saskya van Nouhuys

Topic Sun-17 Hybridization

- 98 Do mitochondrial-nuclear interactions influence nuclear allele frequencies as a result of hybrid breakdown? A test across candidate pathways in the copepod Tigriopus californicus. Daniel Schneck, Felipe Barreto
- 99 A comparative approach to studying the effects of climate change and landcover change on hybrid zones in a suture zone
 Shawn Billerman, Irby Lovette
- 100 Heterogeneous patterns of genetic differentiation across the genomes of hybridizing woodrats (Neotoma)

 Josh Jahner, Marjorie Matocq, Thomas Parchman
- 101 deNovo assembly and annotation of Ambystoma laterale and Ambystoma jeffersonianum transcriptomes: the first steps toward investigating polyploid salamander expression Sheri Sanders, Michael Pfrender

Topic Sun-18 Invasive/invasion

- 102 Rapid Adaptation to Cold in the Invasive Cane Toad Rhinella marina Cinnamon Mittan
- 103 Multiple introductions of the dengue vector, Aedes aegypti, into California Evlyn Pless, Andrea Gloria-Soria, Benjamin Evans, Vicki Kramer, Jeffrey Powell
- 104 Does genome size variation have ecological consequences in an invasive plant? Alice Cang, Katrina Dlugosch
- 105 Birds of Borrowed Paradise: comparing independently feral Gallus gallus from Bermuda and Kauai

 Eben Gering, Dominic Wright, Thomas Getty
- 106 Precise adaptation without loss of adaptive potential: An example from the avian beak evolution

 Georgy Semenov, Robert Hollingsworth, Amin Abtahi, Alexander Badyaev

Topic Sun-19 Life history

107 Minding the gap: evolutionary transitions between ecto- and endoparasitism in the Chalcid wasps and ramifications for host specificity

Rebecca A Boulton, George Heimpel

- 108 Does the Cost of Reproduction Increase with Age? A Test Using Burying Beetles Ethan Damron, Dane Jo, Mark Belk, Ashlee Smith
- 109 Inefficient energy metabolism disrupts nutrient sensing in Drosophila Justin Buchanan, Kristi Montooth
- 110 The equilibrium between aging and rejuvenation in bacterial lineages Audrey Menegaz Proenca
- 111 Life in the mud: Amphipod growth and reproduction under chronic arsenate stress *Ioanna Visviki*, Michael Judge
- 112 The evolution of diapause, developmental arrest during embryology, in the Killifish genus Rivulus
 - Andrew Furness, David Reznick, Andrey Tatarenkov, John Avise
- 113 Estimating performance curves in fluctuating environments Frank Shaw, joel kingsolver
- 114 Kinship interacts with density and nutrient availability to influence fitness components in the American toad (Bufo americanus)

 Sara Garnett, Thomast Getty

Topic Sun-20 Macroevolution

- 115 Doing it with a spermatostyle: preliminary insights into the diveristy and evolution of sperm in ground beetles (family Carabidae)

 Antonio Gomez, David Maddison
- 116 Massive increase in visual range preceded the origin of terrestrial vertebrates Malcolm MacIver, Lars Schmitz, Ugurcan Mugan, Todd Murphey, Curtis Mobley
- 117 Biogeography modeling and plumage color pattern evolution in birds Chad Eliason, Michael Andersen, Shannon Hackett
- 118 Evolution of Body Size and Cranial Modularity in Caviomorph Rodents Genevieve Perdue, Samantha Hopkins, Samantha Price

Topic Sun-21 Molecular evolution

- 119 Ancient and recent duplications support functional diversity of Daphnia opsins Christopher Brandon, Matthew Greenwold, Jeff Dudycha
- 120 Molecular evolution of the PEVK region of titin, a giant sarcomeric protein Kathleen Muenzen, Findley Finseth, Jenna Monroy
- 121 Tempo and mode of Peto's Paradox: a molecular evolutionary approach to understanding cancer prevention

 Marc Tollis, Aika Schneider-Utaka, Carlo Maley
- 122 Evolution of secretin-containing secretion systems

 Davi Ortega, Grant Jensen

Topic Sun-22 Phylogenomics

- 123 Understanding the effects of recombination on phylogenetic analysis Michael Conry, Alan Lemmon
- 124 Phylogenomic appraisal of species limits and hybridization in the endemic Lake Baikal sculpins (genus Limnocottus) withdrawn 2017-06-10 09:29:57.0

 Andres Aguilar, Michael Sandel, Frank Gutierrez, David Neely, Sergei Kirilchik
- 125 Phylogenomics and Macroevolutionary Dynamics of Diplodactyloid Geckos Phillip Skipwith, Paul Oliver
- 126 Mitochondrial transfer RNA genes became drastically reduced early in spider evolution Deborah Catanese, Susan Masta
- 127 Evolution of vernalization response in the PACMAD clade of the grass family (Poaceae)

 Martin Paliocha, Marian Schubert, Jill Preston, Siri Fjelheim
- 128 Phylogenomics And Proteomics Of Longamoebia: Flat Amoebae Without Center Yonas Tekle

- 129 Phylogenomics and the complexity of millipede chemical warfare through time

 Juanita Rodriguez, Tappey Jones, Paul Marek, Petra Sierwald, William Shear, Kevin Kocot, Jason Bond
- 130 Biologically inspired models of rate variation for ultraconserved elements Jeremy Brown, David Morris
- 131 Widespread Site-to-Site Synonymous Substitution Rate Variation in the Metazoans Sarah Wisotsky, Spencer Muse, Rachel Marceau West
- 132 Dating the Diversification of the Paper Wasp (Genus Polistes) Under the Fossilized Birth-Death Process

 Kevin Quinteros, Amy Toth, Tracy Heath
- 133 Genomic insights into the evolution of mud turtles (Kinosternidae)

 Brendan Reid

Topic Sun-23 Phylogeography

- 134 Landscape genomics of Pinus tabuliformis reveals deep divergence beyond the LGM Hanhan Xia, Baosheng Wang, Xiao-Ru Wang, Jian-Feng Mao
- 135 Foundations for Defining Species Boundaries in a Four Species Contact Zone (Hylidae: Pseudacris)

 Mysia Dye, Emily Lemmon
- 136 Genetic and genomic evidence suggest incipient speciation with phenotypic divergence across an environmental barrier in a forest passerine (Pipraeidea melanonota)
 Pablo D. Lavinia, Ana Barreira, Leonardo Campagna, Pablo Tubaro, Dario Lijtmaer
- 137 Phylogeography of the circumtropical shrimp, Stenopus hispidus (Crustacea: Decapoda), reveals divergent Atlantic and Pacific lineages

 Alealani Dudoit, Richard Coleman, Matthew Iacchei, Michelle Gaither, William Browne, Brian Bowen, Rob J. Toonen
- 138 On the demography of colilargo rodents in the Central Andes Natali Hurtado
- 139 Climate events and the lineages of the Neotropical spider Nephila clavipes Vera Solferini, Luiz Bartoleti
- 140 Genetic assessment reveals intraspecific differentiation in the Southern lapwing (Vanellus chilensis) in South America

 Belen Bukowski, Pablo Tubaro, Dario Lijtmaer

Topic Sun-24 Plant reproductive biology

- 141 Soil traits impacts on female frequency of a gynodioecious prairie plant, Lobelia spicata. Diane Byers, Tyler Rippel, Robert Philips
- 142 Evolution of energy supply from haploid gametophytes to diploid sporophytes and genomic conflict in plants

 Kazuhiro Bessho, Hisashi Ohtsuki, Akira Sasaki
- 143 Analyzing Potential for Gametophytic Selection & Selective Ovule Abortion in Mimulus guttatus

 Elizabeth Pérez, Mitch Cruzan, Jaime Schwoch
- 144 When to Move on: Generalist Pollinators Exhibit a Temporal Decrease in Floral Fidelity Towards a Common Prairie Perennial, Echinacea angustifolia Leah Prescott, Jennifer Ison
- 145 The fitness effects of unusually early flowering time in Leavenworthia stylosa following a mild winter.
 Caitlin Banaszak, Chris Herlihy

Topic Sun-25 Population genetics: inference of selection

- 146 The genetics of environmental adapation: house mice in the Americas Megan Phifer-Rixey, Felipe Martins, Ke Bi, Michael Nachman
- 147 Seasonally fluctuating predation pressures and the maintenance of genetic diversity in Daphnia

Karen Barnard-Kubow, Dörthe Becker, Alan Bergland

- 148 Detecting adaptive genetic variation among cryptic species of the microinvertebrate Euchlanis dilatata, using RAD-Seq
 Azar Kordbacheh, Elizabeth J. Walsh, Jonathon Mohl
- 149 Population genomic signatures of selection associated with freshwater invasion by the saline copepod Eurytemora affinis

 Tiago da Silva Ribeiro, Martin Bontrager, Carol Eunmi Lee
- **150** Modeling the Evolution of Alternative Splicing *Ksenia Arzumanova*, Jamie Moon, Scott Roy, Manuel Irimia, Rori Rohlfs

<u>Topic Sun-26 Population genetics: molecular ecology</u>

- 151 On the Impact of Adhesion in Geckos: Species Limits and Phylogeography of the South American Dwarf Gecko, Gonatodes humeralis

 Brendan Pinto, Guarino Colli, Timothy Higham, Anthony Russell, Laurie Vitt, Tony Gamble
- 152 Population genomics of stream salamanders in an urban versus suburban watershed Nicole Fusco, Jason Munshi-South
- 153 Assessing aquatic diversity through space and time using next-gen sequencing Sergio Samaniego, Elizabeth J. Walsh
- 154 Genetic monitoring of a sockeye salmon repatriation to improve interactive fisheries management

 Lucas Elliott, Michael Russello
- 155 sppIDer: Species Identification and Hybrid Detection with Short-Read Data Quinn Langdon, David Peris, Chris Hittinger

<u>Topic Sun-27 Population genetics: theory and methods</u>

- 156 Forward-time simulation in Python using fwdpp Kevin Thornton, Jaleal Sanjak
- 157 Using genealogical properties to identify selective sweeps in the human thousand genomes Martina Rauscher, Johannes Wirtz, Thomas Wiehe
- 158 Investigating Western rattlesnake population genetics and demographics to inform management and conservation

 Danielle Schmidt*, Purnima Govindarajulu, Karl Larsen, Michael Russello
- 159 Can we predict geographic range expansions? Carly Prior, Jeremiah Busch
- 160 Chromosome Drive and the Survival Probability of New Inversions Spencer Koury

Topic Sun-28 Sexual selection

- 161 The genetic basis for song-preference coevolution in the Hawaiian cricket Laupala Mingzi Xu, Kerry Shaw
- 162 Ornamental Plumage in the Non-breeding Season is Associated with Behavioral Change in a Tropical Passerine

Trey Hendrix, Sarah Toner, Samantha Lantz, Joseph Welklin, Facundo Fernandez-Duque, John P Swaddle, Michael Webster , Jordan Karubian

Topic Sun-29 Speciation

163 The geographic context of speciation by sexual selection or Let's talk about speciation in crickets

Thomas Blankers, Kerry Shaw

- 164 Parallel ecotypic divergence on its way to new species? A case study of autotetraploid Cochlearia officinalis from Northern Norway

 Marie Kristine Brandrud, Ovidiu Paun, Maria Lorenzo, Inger Nordal, Anne Brysting
- 165 Identifying genome-scale structural differences between ecologically divergent populations of cactophilic flies

 Coline Jaworski, Carson Allan, Luciano Matzkin
- 166 Has climatic oscillation influenced the P. axillaris evolutionary history?

 Giovanna Giudicelli, Caroline Turchetto, Gustavo Adolfo Silva-Arias, Loreta Brandão de Freitas
- 167 Defining a species: Genomic differentiation between Rhagoletis mendax and the undescribed sparkleberry fly

 Katherine Inskeep, Meredith Doellman, Scott Egan, Peter Meyers, Thomas Powell, Stewart Berlocher, James Smith, Jeffrey Feder

Topic Sun-30 Last minute additions

- 168 Weed Adaptation: Genetic Differentiation vs. Phenotypic Plasticity Ava Garrison, Jeffrey Conner
- 169 Predator cues decrease development time and reduce sexual size dimorphism in the mosquito, Aedes aegypti

 Brandon Pope, Robert Cox, Aaron Reedy
- 170 Playing it safe? Behavioral responses of a mosquito larvae encountering a fish predator Avehi Singh, Karthikeyan Chandrasegaran, Moumita Laha, Suhel Quader

Session 7th annual Evolution Film Festival/Video contest Sunday - 19:30 - 21:30 Room A107-109

Cost: free, no registration required.

One of the highlight of the conference that should not be missed! Join us for an evening of terrifically informative and entertaining videos, while enjoying popcorn and beer/wine/soda (courtesy of HHMI). During the first hour, event co-sponsor HHMI Biointeractive will screen some of the wonderful evolution videos they offer, and you will have a chance to ask questions of scientists and filmmakers involved in producing their videos. During the second hour, we will screen the top entrants to this year's contest. We asked the science and science education communities to tell a story about evolution in 3 min. or less, using animation, art, music, dance, etc. We have some terrific entries and you will be able to see them all and vote on your favorite. (To learn more and see past years' entries, go to evolutionfilmfestival.org and follow #evofilmfest on Twitter.)

Sponsored by the Duke Initiative for Science & Society, the Howard Hughes Medical Institute (HHMI), the Society for the Study of Evolution, and the BEACON Center for the Study of Evolution in Action.

Session SSE/BEACON Undergraduate Diversity Program: Undergraduate breakfast social Monday - 8:00 - 9:30 Room Oregon ballroom 201

Open to undergraduates and their advisers and mentors.Cost: free, but attendees had to pre-register. Organizer: Alexa Warwick

Come to meet and mingle with other undergrads, their advisors/mentors, and alumni of the Undergrad Diversity Program. This social is a great opportunity to discuss your research, career plans, etc. and expand your network in an informal setting while enjoying free continental breakfast!

Session Mon1-A105 Phylogenomics 1

Monday - 8:30 - 9:45

Room A105

8:30 Hidden diversity in the Vangas of Madagascar

Jane Younger, Lynika Strozier, Matt Bonfitto, Chris Kyriazis, Arpad Nyari, J Dylan Maddox , Neha Siddiqui, Steve Goodman, Marie-Jeanne Raherilalao, Sushma Reddy

8:45 In situ diversification is an important driver of parachuting frog diversification on the Sunda Shelf

Kyle O'Connell, Eric Smith, Utpal Smart, Nia Kurniawan, Matthew Fujita

- 9:00 Diversification of suboscine birds based on a genome-wide species-level phylogeny Michael Harvey, Gustavo Bravo, Graham Derryberry, Santiago Claramunt, Terry Chesser, Andrés Cuervo, Jessica Shearer, Robert Moyle, Frederick Sheldon, Joel Cracraft, Brian Smith, Luís Fábio Silveira, Robb Brumfield, Elizabeth Derryberry, Daniel Rabosky
- 9:15 The challenge of testing introgression when introgression obscures phylogeny: The case of the South American siskins.

Elizabeth Beckman, Christopher Witt

9:30 Evolution of the genes and genomes of 76 arthropod species Gregg Thomas, Matthew Hahn, Stephen Richards

Session Mon1-A106 Development

Monday - 8:30 - 9:45

Room A106

- 8:30 Relationship between ontogenetic trajectory and evolutionary variability in avian limb skeleton: implications for evolutionary diversification in birds

 Junya Watanabe
- 8:45 The role of a complex life cycle in facilitating morphological divergence Emma Sherratt, Marta Vidal-García, Marion Anstis, J. Scott Keogh
- 9:00 Egg teeth of squamate reptiles embryonic adaptation to hatching
 Mateusz Hermyt, Paweł Kaczmarek, Katarzyna Janiszewska, Hubert Okła, Weronika Rupik
- 9:15 Embryology of the vomeronasal organ and associated structures in the context of squamate evolution: a 3D perspective

 Paweł Kaczmarek, Katarzyna Janiszewska, Mateusz Hermyt, Weronika Rupik
- 9:30 When are maternal effects maternal strategies? Environmental mismatch in a passerine bird **Ahva Potticary**, Renee Duckworth

Session Mon1-A107-109 Macroevolution / diversification Monday - 8:30 - 9:45 Room A107-109

- 8:30 Diversification in time and space: disentangling the radiation of New World pitvipers Laura Alencar, Irina Birskis, Marcio Martins, Tiago Quental
- 8:45 Do macrophylogenies yield stable macroevolutionary inferences? An example from squamate reptiles

Pascal Title, Daniel Rabosky

- 9:00 The Nearshore Cradle and Museum of Early Vertebrate Diversification Lauren Sallan, Matt Friedman, Robert Sansom, Ivan Sansom
- 9:15 Rates of speciation and extinction are highly correlated in mammals Tiago Quental, Mathias Pires
- 9:30 Dimensions of morphological diversification: an ecomorphological perspective from carnivorans.

Graham Slater, Anthony Friscia

Session Mon1-B111-112 Life history / plasticity Monday - 8:30 - 9:45 Room B111-112

8:30 Genetic Basis of Variation in Life Cycle Gabriel Golczer, Erastus Thuo, Genevieve Kozak, Steve Bogdanowicz, Erik Dopman

- 8:45 Plasticity is a locally adapted trait and has consequences for responses to novel environments Tom Cameron, Matthew Bond, Stuart Piertney, Tim Benton
- 9:00 Multiple paths to success: fitness consequences of tiger salamander morphs with different life history strategies

 Alycia Lackey, Howard Whiteman, Michael Moore
- 9:15 Generalized adaptive responses to novel mutagens: costs of up-regulated antioxidant pathways?

Emilie Snell-Rood, Reilly Hostager, Nathanial Fremling, Megan Kobiela

9:30 Temperature tipping points: plasticity, adaptation, and trade-offs shape range-wide climate tolerances of a long-lived alpine plant Megan Peterson, Daniel Doak, William Morris

Session Mon1-B113 Genetics: behavior & disease Monday - 8:30 - 9:45 Room B113

- 8:30 Examining the genetic structure and physiological mechanisms underlying an evolutionarily novel, complex biochemical trait, mycotoxin tolerance, in a Drosophila tripunctata population Clare Scott Chialvo, Olivia Sorrell, Logan Griffin, Laura Reed
- 8:45 Genes underlying parasitoid-induced behavior in the fruit fly Michael Martin, Todd Schlenke
- 9:00 Neural basis of female receptivity in Drosophila melanogaster and D. simulans Pria Mahabir, Amanda Moehring
- 9:15 The genetic basis for cuticular hydrocarbon biosynthesis in Drosophila Heather Ward, Amanda Moehring
- 9:30 Mapping resistance to naturally coevolving viruses in the genetic model organism Drosophila melanogaster
 Rodrigo Cogni, Francis Jiggins

Session Mon1-B114-115 Population genetics: inference of selection 2

Monday - 8:30 - 9:45

Room B114-115

- 8:30 Footprints of Adaptive Introgression

 Joachim Hermisson, Derek Setter, Michael DeGiorgio, Rasmus Nielsen
- 8:45 The Selective Maintenance of Allelic Variation Under Generalized Dominance Hamish Spencer, Cuilodair Mitchell
- 9:00 Determining the factors driving selective effects of new nonsynonymous mutations across multiple species

 Kirk Lohmueller, Bernard Kim, Clare Marsden, Ying Zhen, Christian Huber
- 9:15 The Temporal Signature of Linked Selection Vince Buffalo
- 9:30 Is Background Selection a Confounding Effect to Local Selection? Remi Matthey-Doret, Michael Whitlock

Session Mon1-B116 Speciation / gene flow Monday - 8:30 - 9:45 Room B116

- 8:30 Characterizing the evolutionary dynamics of an hybrid sterility allele in Mimulus guttatus Rachel Kerwin, Andrea Sweigart
- 8:45 Reproductive isolation and gene flow vary among contact zones between incipient species Catherine Debban, Laura Galloway
- 9:00 Contrasting genetic profiles of a contact zone in California voles.

 Dana Lin, Ke Bi, Christopher Conroy, Rauri Bowie*
- 9:15 Population transcriptomics of ecologically differentiated, partially reproductively isolated Anolis lizards

 Anthony Geneva, Fatih Sarigol

9:30 Integration of the genomic and geographic landscapes reveals the impact of an ecological barrier to gene flow in the bush monkeyflower (Mimulus)

Sean Stankowski, Madeline Chase, Matt Streisfeld

Session Mon1-B117-119 Evolutionary ecology 1 Monday - 8:30 - 9:45 Room B117-119

- 8:30 Computational Food Web Models Gavin Abernethy
- 8:45 Testing character displacement as a potential driver in the evolution of root traits Sara Colom, Regina Baucom
- 9:00 Evolution of chemical signals in Neotropical Piper within the context of ecological interactions with bats

 Zofia Kaliszewska, Sharlene Santana, Jeff Riffell
- 9:15 Ecosystem consequences of rapid evolution in a coastal sedge Jason McLachlan, Michael Blum, Rachel Gentie
- 9:30 The effects of consumer satiation and interference on the strength of disruptive selection. Kyle Coblentz, Mark Novak

Session Mon1-Ball202 SSB Spotlight - Uncertainty in the face of strong support: new approaches and examples in the identification of signal, noise, and conflict in phylogenetic datasets

Monday - 8:30 - 9:45

Room Oregon ballroom 202

- 8:30 Detecting causes of conflict in phylogenomic datasets Robert Thomson, Lyndon Coghill, Jeremy Brown
- 8:45 Mito-nuclear discordance: distinguishing between contemporary hybridization, ILS and introgression

 Alana Alexander, Yong-Chao Su, Carl Oliveros, Karen Olson, Scott Travers, Rafe Brown
- 8:50 Gene tree conflict in neobatrachian frogs.
- 8:55 Understanding ancestry when canned programs give strange results Suzanne McGaugh
- 9:00 Beyond supermatrix resolution: Deep metazoan phylogeny from gene content Walker Pett
- 9:15 The effect of heterotachy on phylogenomics.

 Lyndon Coghill, Ashley Schoonmaker, Lauren Rodriguez, Robert Thomson, Jeremy Brown
- 9:20 Missing data in RADseq phylogenetics reduces branch length accuracy and increases computational cost

 Devon Humphreys, E. Anne Chambers, Katie Lyons
- 9:25 Phylogenomics of Neotropical knifefishes (Gymnotiformes): reconciling phylogenetic signal and electric signals

 Fernando Alda
- 9:30 Testing the limits of phylogenomics for resolving the bird tree of life Sushma Reddy, Rebecca Kimball, Ed Braun

Session Mon1-Ball203 SSE Symposium - Evolution in urban ecosystems 1

Monday - 8:30 - 9:45

Room Oregon ballroom 203

8:30 Urban Evolutionary Biology: novel selection pressures and a new subdiscipline for the Anthropocene

Jason Munshi-South

- 8:45 Pigeon Sex, Stress, and Reproduction: Developing Genomic Tools to Study Urban Impact Suzanne Austin, Matthew MacManes, Andrew Lang, Rebecca Calisi
- 9:00 The anthropogenically driven evolution of a globally reemerging ecotoparasite Warren Booth, Christopher Lawrence, Ondrej Balvin, Edward Vargo, Coby Schal
- 9:15 Phylogeography and demography of the global spread of the brown rat (Rattus norvegicus)

 Emily Puckett, Jason Munshi-South
- 9:30 Urbanization facilitates gene flow in the Western black widow spider, Latrodectus hesperus Lindsay Miles, Brian Verrelli

Session Mon1-Ball204 Predation Monday - 8:30 - 9:45 Room Oregon ballroom 204

8:30 Interactions between color and behavior for aposematic and thermoregulatory functions in a caterpillar

Matthew Nielsen, Johanna Mappes

- 8:45 Theoretical Shell Models: Mathematical and Biological Trade-offs Erynn Johnson, Lauren Sallan
- 9:00 More than Skin Deep: variation in behavior as a result of eumelanin-based coloration in the eastern mosquitofish

Eve Humphrey, Joseph Travis, Kimberly Hughes

- 9:15 Natural selection in behavior? A field experiment with Anolis lizards from the Caribbean Oriol Lapiedra, Jonathan Losos, Jason Kolbe
- 9:30 The evolution of vertebrate eye size across an environmental gradient: phenotype does not predict genotype in a Trinidadian killifish

 Shannon Beston, Elijah Wostl, Matthew Walsh

Session Mon1-C120-122 Sexual selection / conflict 1 Monday - 8:30 - 9:45 Room C120-122

- 8:30 The physical environment mediates male harm and its effect on selection in females Li Yun, Patrick Chen, Amardeep Singh, Aneil Agrawal, Howard Rundle
- 8:45 Sexual Selection Accelerates the Rate of Adaptation in a Complex Mating Environment Li Yun, Patrick Chen, Michelle Yang, Malak Bayoumi, Howard Rundle, Aneil Agrawal
- 9:00 Testing for sexually antagonistic selection across ontogeny in a sexually dimorphic lizard Aaron Reedy, Ariel Kahrl, Cara Giordano, Heidi Seears, Daniel Warner, Robert Cox
- 9:15 Females lose potential fitness benefits through male anti-cannibalism strategy in the nursery web spider, Pisaurina mira.

 Alissa Anderson, Eileen Hebets**
- 9:30 Remodeling and deweaponization as an alternative to the arms race of sexually antagonistic coevolution

 Samuel Snow, Richard Prum

Session Mon1-C123 Coevolution 1 Monday - 8:30 - 9:45 Room C123

- 8:30 Plant defenses and herbivores independently alter selection on reproductive traits in white clover (Trifolium repens)

 James Santangelo, Marc Johnson
- 8:45 The evolution of coexistence: reciprocal adaptation promotes the assembly of a simple community

 Ron Bassar, Joseph Travis, David Reznick*
- 9:00 Evolutionary transitions in fruit colors with respect to climate Miranda Sinnott-Armstrong, Michael Donoghue
- 9:15 Highly dynamic genomic architecture of cooperation in wild populations of Mesorhizobium Stephanie Porter, Joshua Faber-Hammond, Emily Helliwell, Cynthia Sackos, Maren Friesen
- 9:30 The Measurement of Coevolution in Mutualisms Bob Week, Scott Nuismer

Session Mon1-C124 Genomics / reproductive systems Monday - 8:30 - 9:45 Room C124

- 8:30 Genomic insights into the evolutionary transition to selfing Allan Castillo, Joanna Rifkin, Irene Liao, Mark Rausher
- 8:45 Patterns of adaptive molecular evolution in a New Zealand freshwater snail Laura Bankers, Jeffery Boore, John Logsdon, Maurine Neiman
- 9:00 Genomic signatures of asexual and sexual reproduction within the colonial hydrozoan Ectopleura larynx.

 E. Sally Chang, Maria E. Orive, Paulyn Cartwright
- 9:15 Self-fertility triggers rapid genome shrinkage and loss of sperm competition proteins withdrawn 2017-06-22 00:44:51.0

 Da Yin
- 9:30 Genomic consequences of asexuality *Maurine Neiman*, John Logsdon, Jeffrey L Boore, Laura Bankers, Peter Fields, Joseph Jalinsky, Kyle
 McElroy, Joel Sharbrough, Peter Wilton

Session Executive joint council (ASN/SSB/SSE) exit meeting Monday - 8:30 - 10:15 Room C126

Session Coffee break

Monday - 9:45 - 10:15

Room Exhibit Hall A

Session Mon2-A105 Phylogenomics 2

Monday - 10:15 - 11:30

Room A105

- 10:15 Deep Sea Lights: Evolution of bioluminescence and light detection in the twilight zone Heather Bracken-Grissom
- 10:30 The tuatara genome sheds light on phylogenetics and rates of evolution during the amniote radiation

 Marc Tollis, Chris Organ, Stefan Prost, David Winter, Neil Gemmell
- 10:45 Conflicting phylogenomic signals reveal a pattern of reticulate evolution in a recent high-Andean diversification (Asteraceae: Astereae: Diplostephium)

 Oscar Vargas, Edgardo M Ortiz, Beryl Simpson
- 11:00 Wading into beetle phylogenomics: application of the Coleoptera UCE bait set to Hydrophiloidea

 Alana Alexander, Andrew Short
- 11:15 Phylogeny and patterns of gene flow among fruit fly species of the fraterculus group inferred from Transcriptomic data
 Carlos Congrains, Roberto A Zucchi, Reinaldo A de Brito

Session Mon2-A106 Development / expression studies Monday - 10:15 - 11:30 Room A106

- 10:15 Genomic and transcriptomic changes underlying the transition to an experimentally evolved multicellular lifestyle

 Noah Gettle, Michael Travisano
- 10:30 The analysis of early embryonic transcriptomes in rapidly-diversifying Drosophila species Madeline Chenevert, Joel Atallah
- 10:45 Floral symmetry in Asparagales and the evolution of candidate TCP genes Yesenia Madrigal Bedoya, Natalia Lucia Pabón-Mora, Juan Fernando Alzate

- 11:00 The developmental and genetic trajectory of coloration in the guppy (Poecilia reticulata) Cynthia Dick, Jeff Arendt, David Reznick, Cheryl Hayashi
- 11:15 Gene expression evolution during sunflower domestication Ray Watson, Benjamin Blackman

Session Mon2-A107-109 Macroevolution / paleobiology Monday - 10:15 - 11:30 Room A107-109

- 10:15 The changing functional composition of the North American species pool: modeling species origination-extinction as a function of functional group and environmental context Peter Smits
- 10:30 Pervasive diversity imbalance among sister clades of ray-fin fishes John Sime, Jonathan Chang, Michael Alfaro, Lauren Sallan
- 10:45 The fossil record of marine planktonic protists: resources, applications and potential for macroevolutionary research

 David Lazarus
- 11:00 Diversity dynamics of Phanerozoic terrestrial tetrapods at the local-community scale Roger Close, Roger B. J. Benson, Juan Benito, Emma Dunne, Richard Butler
- 11:15 Fossils flip phylogenetically-estimated extinction rates in salamander clades Jonathan Mitchell, Daniel Rabosky, Joanna Larson

Session Mon2-B111-112 Adaptation / plasticity Monday - 10:15 - 11:30 Room B111-112

- 10:15 Epigenetic differences associated with phenotypic divergence in a widespread invasive species

 Jennifer Madrid Thorson
- 10:30 Plasticity and Specialization in Variable Environments: Adaptation in Vernal Pool Goldfields Nancy Emery, Raffica La Rosa
- 10:45 Phenotypic plasticity can help or hinder adaptation to climate change: a simulation model Sam Scheiner, Michael Barfield, Robert Holt
- 11:00 Ancestral plasticity and salt marsh colonization in the Savannah Sparrow. *Phred Benham*, Zachary Cheviron
- 11:15 Effects of road salt on butterfly life history and potential for adaptation to high sodium levels Megan Kobiela, Emilie Snell-Rood

Session Mon2-B113 Behavior / ecology Monday - 10:15 - 11:30 Room B113

- 10:15 Temporal variation of anthropogenic perch use by populations of forest and urban lizards Kevin Aviles Rodriguez, Liam Revell
- 10:30 The influence of relatedness and size on spatial structure in an urban population of Anolis carolinensis lizards

 William Weber, Simon Lailvaux, Nicola Anthony
- 10:45 Assessing epigenetic and genetic diversity across a range expansion using bisulfite ddRADseq Tricia Rubi
- 11:00 When is bigger better? A test of the relationship between body size and starvation resistance, competitive ability, and fitness in a burying beetle

 Ashlee Smith, Mark Belk
- 11:15 Pharaoh ant colony productivity is linked to colony personality Justin Walsh, Tim Linksvayer

Session Mon2-B114-115 Population genetics: inference of selection 3

Monday - 10:15 - 11:30

Room B114-115

- 10:15 Pooled capture sequencing of sea urchin larvae reveals consistent changes in allele frequency in response to experimental acidification

 April Garrett, Lauren Ashlock, Heidi Hargarten, Kaitlin Huber, Melissa Pespeni
- 10:30 Evolutionary genomics of glucosinolate defensive chemistry in Boechera stricta Julius Mojica, Tom Mitchell-Olds
- 10:45 Gene-specific introgression drives local climatic adaptation in a conifer hybrid complex Jonathan Degner, Sam Yeaman, Kathryn Hodgins, Kristin Nurkowski, Loren Rieseberg, Sally Aitken
- 11:00 Local adaptation in a sensory complex: insight from the Cape horseshoe bat system Lizelle Odendaal

Session Mon2-B116 Speciation / genomics 1 Monday - 10:15 - 11:30 Room B116

- 10:15 The genomic landscape of prezygotic reproductive isolation in a sympatric species complex of intertidal isopods (Jaera albifrons)

 Marius Wenzel, Alex Douglas, Stuart Piertney
- 10:30 Genome-wide association mapping of seasonal mating time Genevieve Kozak, Crista Wadsworth, Brad Coates, Erik Dopman
- 10:45 Mito-nuclear incompatibilities contribute to hybrid inviability in early stages of divergence *Thiago Lima*, Ron Burton, Christopher Willett
- 11:00 From candidate variants to fitness landscapes: investigating the rare origins of novel trophic specialists during adaptive radiation Christopher Martin
- 11:15 Recombination, gene flow, and selection drive a heterogeneous pattern of divergence between ecotypes of Mimulus aurantiacus

 Matt Streisfeld, Sean Stankowski, Madeline Chase

Session Mon2-B117-119 Evolutionary ecology 2 Monday - 10:15 - 11:30 Room B117-119

- 10:15 Diversification of crustacean visual opsins revealed using phylotranscriptomics Danielle DeLeo, Jorge Pérez-Moreno, Hernan Vazquez Miranda, Heather Bracken-Grissom
- 10:30 An R2R3MYB involving an anther-color polymorphism in a southeastern trout lily Rong-Chien Lin, Mark Rausher
- 10:45 Wild Danio rerio a model for immunology, population genetics and ecology *Thomas Wiehe*, Jaanus Suurvaeli, Christopher Kraus, Johannes Schoeneich
- 11:00 Differential altitudinal migration and elevational range size are associated with differences in thermal tolerance in birds

 Shane DuBay, Yongjie Wu, Zachary Cheviron, Graham Scott, Yanhua Qu, Fumin Lei, John Bates
- 11:15 Time to tree-think the genetic effects of forest management: Assessing evolutionary patterns and processes beyond heterozygosity indices

 Brandon Lind, Malcolm North, Andrew Eckert

Session Mon2-Ball202 Systematics 1 Monday - 10:15 - 11:30 Room Oregon ballroom 202

- 10:15 Ancient DNA analysis of the extinct diving ducks Chendytes lawi and the Labrador duck reveal convergent evolution of diving in boreal habitats

 Janet Buckner, Terry Jones, David Gold, Ryan Ellingson, David Jacobs
- 10:30 A global phylogeny of turtles and how it informs conservation planning Bradley Shaffer, Robert Thomson, Janet Vu
- 10:45 A new and freely available exome sequence capture probe set of 10,000 loci for frogs: testing its effectiveness across multiple phylogenetic scales

 Carl Hutter, Kerry Cobb, Scott Travers, Rafe Brown
- 11:00 Tusk evolution in the burrowing mayflies: traditional genes vs. phylogenomics. Dustin Miller, T. Heath Ogden

11:15 Leveraging the multispecies coalescent model for inferences of selection and its impact on species tree estimation and species delimitation

Richard Adams, Daren Card, Drew Schield, Heath Blackmon, Todd Castoe

Session Mon2-Ball203 SSE Symposium - Evolution in urban ecosystems 2

Monday - 10:15 - 11:30

Room Oregon ballroom 203

- 10:15 Adaptive and non-adaptive genomic changes during urbanization in burrowing owls **Jakob C. Mueller**, Martina Carrete, Heiner Kuhl, Stefan Boerno, Jose Tella, Bart Kempenaers
- 10:30 Rapid evolution of ant thermal tolerance within urban heat islands Ryan Martin, Lacy Chick, Abe Perez, Stephanie Strickler, Sarah Diamond
- 10:45 Urban adaptation in Lizards: Connecting phenotypic shifts with performance and survival Kristin Winchell, Liam Revell
- 11:00 Is Convergence at the Level of the Phenotype Mirrored Across the Genome? The Repeated Evolution of Glyphosate Resistance in the Common Morning Glory, Ipomoea purpurea Regina Baucom
- 11:15 Plant evolution in the urban jungle Marc Johnson

Session Mon2-Ball204 ASN Spotlight - Eco-evolutionary dynamics

Monday - 10:15 - 11:30

Room Oregon ballroom 204

- 10:15 Eco-evolutionary Dynamics

 Andrew Hendry
- 10:30 The role of behaviour in eco-ecolutionary dynamics Kiyoko Gotanda, Pierre-Olivier Montiglio
- 10:35 Eco-evolutionary dynamics in model systems: It's very competitive Tom Cameron
- 10:40 Eco-evolutionary trophic cascades

 Zachary Wood, Eric Palkovacs, Michael Kinnison
- 10:45 Ecosystem evolutionary dynamics: Integrating indirect genetic effects and niche construction Joe Bailey, Jen Schweitzer, Michael Van Nuland, Ian Ware
- 11:00 Experimental test of how eco-evolutionary dynamics impact the maintenance of species diversity using duckweed.

 Martin Turcotte, Simon P. Hart, Jonathan M. Levine
- 11:05 What rescues the voles. Demographic consequences of contemporary adaptive evolution in a wild rodent population

 Timothée Bonnet
- 11:10 Ecological consequences of sexually selected traits: from populations to ecosystems Sean Giery
- 11:15 Eco-evolutionary dynamics even in humans Fanie Pelletier

Session Mon2-C120-122 Sexual selection / conflict 2 Monday - 10:15 - 11:30 Room C120-122

- 10:15 Molecular Evolution of the Caenorhabditis Nematode Sperm Proteome Katja Kasimatis, Patrick Phillips
- 10:30 Sexually conflicting selection on floral design of the hermaphroditic plant Silene stellata Juannan Zhou, Michele Dudash, Charles Fenster, Elizabeth Zimmer
- 10:45 Sexually antagonistic polymorphism and the evolution of dimorphic sexual systems Colin Olito, Tim Connallon
- 11:00 The sexual preferences of selfish sex chromosomes Pavitra Muralidhar, Robert Trivers

Session Mon2-C123 Coevolution 2

Monday - 10:15 - 11:30

Room C123

- 10:15 High-throughput investigation of species-specific binding of abalone reproductive proteins Jolie Carlisle, Damien Wilburn, Willie Swanson
- 10:30 Associating host genomic variation with gut microbiota in wild house mice

 Taichi Suzuki, Megan Phifer-Rixey, Kathleen Ferris, Andreas Chavez, Michael Sheehan, Dana Lin,
 Michael Nachman
- 10:45 A time-calibrated natural history of Proteobacteria

 Joanna Wolfe
- 11:00 Searching for Microbe Coevolution in the Uncharted Venomous Territories of Marine Cone Snails

Sabah Ul-Hasan, Robert Bowers, Tanja Woyke, Mark Sistrom

11:15 Coevolution drives influenza antiviral resistance evolution in the absence of antivirals Samuel Scarpino

Session Mon2-C124 Molecular ecology / genomics Monday - 10:15 - 11:30 Room C124

- 10:15 Exploring the use of random forest for SNP selection in genetic population assignment Emma Sylvester, Paul Bentzen, Ian Bradbury, Marie Clément, Jon Pearce, John Horne, Robert Beiko
- 10:30 Molecular diversity of mangrove rivulus (Kryptolebias marmoratus) using genomic data Luana Lins, Shawn Trojahn, Muh-Ching Yee, Alexandra Sockell, Carlos Bustamante, Ryan Earley, Joanna Kelley
- 10:45 A Functional Genomic Study of Alba: The Locus Affecting Life History and Color in Colias Butterflies

 Alyssa Woronik
- 11:00 Genome-wide signatures of divergent selection in highly differentiated populations of Theobroma cacao

 Joel Nelson, Omar Cornejo
- 11:15 Direct estimation of germline mutation rate in mouse lemurs, genus Microcebus

 C. Ryan Campbell, Kelsie Hunnicutt, Rachel Williams, Peter Larsen, Mario dos Reis, Anne Yoder

Session Ecology and Evolution editorial board meeting Monday - 11:30 - 13:30 Room B114-115

Session Lunch

Monday - 11:30 - 13:30

Room On the town

Session SSE Executive council exit meeting
Monday - 11:30 - 13:30 Room C126

Session Systematic Biology editorial board meeting
Monday - 11:30 - 13:30 Room B110

Session NSF Information session

Cost: free, no registration required.

A information session on NSF funding opportunities and program changes. Lunch is not provided (you can bring your own).

Host: Samuel Scheiner & George Gilchrist (NSF Program Directors)

Session Mon3-A105 Phylogenomics / plants Monday - 13:30 - 14:45 Room A105

13:30 The Origin of land plants: Genomes and rocks

Mark Puttick, Jennifer Morris, Harald Schneider, Philip Donoghue

- 13:45 A fruitful phylogeny: using RADseq to reveal the evolutionary relationships of the mango genus (Mangifera)

 Emily Warschefsky, Eric von Wettberg
- 14:00 Constructing phylogenies with hundreds of transcriptomes: a case study in Caryophyllales Joseph Walker, Ya Yang, Michael J. Moore, Samuel F. Brockington, Ning Wang, Alfonso Timoneda, Tao Feng, Sonia Ahluwalia , Stephen Smith
- 14:15 The origins of Coca: identifying progenitors of a South American domestic shrub (Erythroxylum spp.)
 - **Dawson White**, Roberta Mason-Gamer
- 14:30 Resolving the relationships of Jatropha using RADseq Nathan LeClear, Beryl Simpson, Craig Linder

Session Mon3-A106 Development / genetics Monday - 13:30 - 14:45 Room A106

- 13:30 The development, diversity, and ecology of solar tracking in sunflowers Benjamin Blackman
- 13:45 Multiple Mechanisms Allow Rapid and Repeated Floral Diversification Jamie Kostyun, Jill Preston, Leonie Moyle
- 14:00 Gene-By-Environment Interactions Drive Developmental Responses to Antibiotic Exposure in Threespine Stickleback Fish
 Emily Lescak, Ryan Lucas, Kenneth Sparks, Koral Campbell, Kelly Ireland, Kat Milligan-Myhre
- 14:15 Genetic basis of reduced nectar production in a selfing morning glory Irene Liao, Joanna Rifkin, Mark Rausher
- 14:30 Straight from the fish's mouth: Genetic variants influencing adaptive craniofacial divergence and speciation in Caribbean pupfish

 Joseph McGirr, Christopher Martin

Session Mon3-A107-109 Paleobiology / paleontology Monday - 13:30 - 14:45 Room A107-109

- 13:30 Paleodiversity of a newly excavated cave site in Puerto Rico: Evidence of extinction and continuity

 Melissa Kemp
- 13:45 Employing models of stratigraphic preservation in phylogenetic inference Caroline Parins-Fukuchi
- 14:00 Dynamics of diversity and distribution patterns across geologically short time scales:
 Evidence from West Indian fossils
 Jessica Oswald, Nancy Albury, Robert Guralnick, David Steadman
- 14:15 Sexual dimorphism is generally stable over the lifetime of fossil ostracode species
 Gene Hunt, Maria Joao Fernandes Martins, T Markham Puckett, Rowan Lockwood, John P Swaddle,
 Jack Shaw
- 14:30 Dynamic trait covariances despite stationary univariate patterns on the approach to speciation

 Thomas Ezard

Session Mon3-B111-112 Life history Monday - 13:30 - 14:45 Room B111-112

13:30 From hostile host to mercurial mosquito: parasite strategies for maximizing transmission to a dynamic vector population

Megan Greischar, Lindsay M. Beck-Johnson, Nicole Mideo

- 13:45 Sex-specific effects of compromised energy metabolism on immunity and life-history Justin Buchanan, Kristi Montooth
- 14:00 Does an introduced parasite cause life history trade-offs in Galapagos mockingbirds? Sabrina McNew, Dale Clayton
- 14:15 The Unicellular Origin of Biological Aging
 Lin Chao, Ulla Camilla Rang
- 14:30 Reevaluating the cost of reproduction Wendy Hood

Session Mon3-B113 Experimental evolution 1 Monday - 13:30 - 14:45 Room B113

- 13:30 The emergence, maintenance and demise of diversity in a spatially variable antibiotic regime Alanna Leale, Rees Kassen
- 13:45 Experimental Evolution of host-attachment in a facultative parasite: an evolutionary stepping-stone towards parasitism

 Emily Durkin, Lien Luong
- 14:00 Brrrr and beer: Cold adaptation in hybrid yeast Caiti Smukowski Heil, Chris Large, Maitreya Dunham
- 14:15 Evidence of intralocus sexual conflict in experimentally evolved S. cerevisiae populations. Katie Fisher, Sean Buskirk, Greg Lang

Session Mon3-B114-115 Pop gen: theory and methods Monday - 13:30 - 14:45 Room B114-115

- 13:45 Rampant heterosis on the fitness landscape of regulatory genetic interactions Adam Porter, Norman Johnson, Alexander Tulchinsky
- 14:00 Using local PCA to summarize how relatedness varies along the genome Peter Ralph, Han Li
- 14:15 Detecting selection for ancestry in admixed populations with arbitrary population structure Erin Calfee, Graham Coop
- 14:30 Testing how migratory fidelity influences connectivity: A novel Bayesian population assignment framework that incorporates proxies for migratory traditions Emma Carroll, Oscar Gaggiotti

Session Mon3-B116 Speciation / genomics 2 Monday - 13:30 - 14:45 Room B116

- 13:30 Integrating micro- and macroevolutionary approaches to understand the role of ecology in nudibranch diversification

 Allison Fritts-Penniman, Ngurah Mahardika, Terrence Gosliner, Paul Barber
- 13:45 Repeated divergent selection on pigmentation genes in a rapid finch radiation Leonardo Campagna, Márcio Repenning, Luís Fábio Silveira, Carla Suertegaray Fontana, Pablo Tubaro, Irby Lovette
- 14:00 Parallel transcriptomic landscape across continents within the Coregonus species complex. Clément Rougeux, Pierre-Alexandre Gagnaire, Kim Praebel, Louis Bernatchez
- 14:15 Phylogenomics of the Andean Plant genus Burmeistera (Campanulaceae) Simon Uribe-Convers, Nathan Muchhala

Session Mon3-B117-119 Evolutionary ecology 3 Monday - 13:30 - 14:45 Room B117-119

13:30 Estimating the benefits of plasticity in ectotherm heat tolerance under natural thermal variability

Alex Gunderson, Michael Dillon, Jonathon Stillman

- 13:45 Working with what you have: Exploring the role of Chlamydomonas defensive colonies in the development of permanent volvocine multicellularity

 Nicholas Lorusso
- 14:00 Historical distribution of diploid and autotetraploid Callisia graminea (Commelinaceae) and implications for climate change
 Kate Hertweck, Andrea Ortiz
- 14:15 Living within your means: low variability in Rhododendron wood traits constrains hydraulic supply but some clades mitigate through increased leaf variability or face increased hydraulic risk

 Juliana Medeiros. Charlotte Hewins
- 14:30 Functional variation in a key defense gene structures herbivore communities and alters plant performance

Nora Adam, Mario Kallenbach, Stefan Meldau, Daniel Veit, Nicole van Dam, Ian Baldwin, Meredith Schuman

Session Mon3-Ball202 SSB Spotlight - Impact of gene flow and reticulation in phylogenetics Monday - 13:30 - 14:45 Room Oregon ballroom 202

- 13:30 The effect of gene flow on coalescent-based species tree inference Laura Kubatko
- 13:45 HyDe: a Python package for genome-scale hybrid detection Paul Blischak
- 13:50 Evidence of intracontinental hybridization in the baobab trees
 Nisa Karimi, noah Stenz, Corrinne Grover, Joseph Gallagher, Jonathan Wendel, David Baum
- 13:55 Unraveling the effects of gene flow on species limits and phylogenetic signal in Myotis bats Ariadna Morales, Brian O'Meara, Nathan Jackson, Bryan Carstens
- 14:00 Modeling hybridization with incomplete lineage sorting james degnan
- 14:15 Trait evolution on phylogenetic networks Brian O'Meara, Dwueng-Chwuan Jhwueng
- 14:23 Phylogenetic comparative methods on species networks
 Cécile Ané. Paul Bastide
- 14:30 The reticulating phylogenetic history of the kingsnakes Frank Burbrink , Marcelo Gehara

Session Mon3-Ball203 SSE Symposium - The impact of stress on genetic variation 1

Monday - 13:30 - 14:45

Room Oregon ballroom 203

- 13:30 Lessons learned from a century of meiotic recombination rate plasticity studies in Drosophila Laurie Stevison, Rita Graze, Stephen Sefick
- 14:00 Stress induced mutation and recombination population genetic models Lilach Hadany

14:30 Evolution in the dark: the role of large effect variants in shaping phenotypes in cavefish Suzanne McGaugh, Adam Herman, James Weagley, Richard Borowsky, Yaniv Brandvain, William Jeffery, Erin Gilbertson, Courtney Passow, Wesley Warren, Masato Yoshizawa, Kelly O'Quin, Luis Espinasa

Session Mon3-Ball204 Sexual selection / speciation Monday - 13:30 - 14:45 Room Oregon ballroom 204

13:30 Reinforcement of conspecific sperm precedence weakens sexual selection in sympatric populations

Dean Castillo, Leonie Moyle

- 13:45 Male-male competition and its role in phenotypic divergence in a polymorphic poison frog Yusan Yang, Corinne Richards-Zawacki, Matthew Dugas
- 14:00 Patterns of divergence over time: the effects of mate choice mechanisms Maria Servedio, Reinhard Bürger
- 14:15 Building female mate preferences: gene to neuron to behavior.

 Amanda Moehring, Ryan Calhoun, Tabashir Chowdhury
- 14:30 Male mate choice and the evolution of behavioral isolation among sexually dimorphic species

 Tamra Mendelson, Jennifer Gumm, Michael Martin, Patrick Ciccotto

Session Mon3-C120-122 Sexual selection / behavior Monday - 13:30 - 14:45 Room C120-122

13:30 The Evolution of Decadence by Mate Choice in the Club-winged Manakin (Machaeropterus deliciosus)

Richard Prum, Samuel Snow, Jacob Berv, Matthew Hack

13:45 Obligate, but not facultative, satellite males prefer same male sexual signal characteristic as females

Rachel Olzer. Marlene Zuk

14:00 Acoustic experience alters exploratory behavior and reproductive investment in Pacific field crickets

Justa Heinen-Kay, Daina Strub, Marlene Zuk

14:15 Mate choice in the dark: The role of visual and acoustic signals in female mate choice in red-eyed tree

Jeanne Robertson. Edauri Navarro Pérez. Chloe Boehkle. Andres Vega. Kristine Kaiser

14:30 Do guppies have a 'bro-code'? Testing for altruism and spite in male-male competition Mitchel Daniel, Helen Rodd

Session Mon3-C123 Coevolution 3

Monday - 13:30 - 14:45

Room C123

- 13:30 Phylogenomics of Sodalis symbionts: models for repeated replacement

 Bret Boyd, Kevin Johnson, Ian James, Adam Clayton, Andrew Sweet, Julie Allen, Robert Weiss, Colin
 Dale
- 13:45 UCEs and tangled trees: elucidating co-phylogenomic patterns in Panamanian figs and fig wasps

Jordan Satler, Tracy Heath, Deren Eaton, Allen Herre, Charlotte Jander, Carlos Machado, John Nason

- 14:00 Demographic differences predict patterns of local adaptation in a venomous snake and its resistant prey

 Matthew Holding, Michael Sovic, Lisle Gibbs
- 14:15 Adaptive variation in the Sierra garter snake (Thamnophis couchii): The influence of biogeography and genetic structure on patterns of TTX-resistance

 Joshua M. Hallas, Chris Feldman, Butch Brodie, Michael Pfrender, Edmund D. Brodie, Jr., Thomas Parchman
- 14:30 Finding disease genes in the face of the Red Queen Ailene MacPheson, Sally Otto, Scott Nuismer

Session Mon3-C124 Mitochondria

Monday - 13:30 - 14:45

Room C124

13:30 If you can't stand the heat, get your X to the tropics Joseph Ross, Rania Haddad, Enrique Cazares-Navarro

13:45 Mitonuclear coevolution in the absence of sex

Joel Sharbrough, Daniel Sloan, Kristi Montooth, Laura Bankers, Peter Fields, Kyle McElroy, Jeffrey L Boore, John Logsdon, Maurine Neiman

14:00 Mitochondrial genome evolution within the Paramecium genus

Parul Johri, Georgi K. Marinov, Michael Lynch

14:15 Differential effects of thermal stress in populations of the intertidal copepod Tigriopus californicus

Alice Harada, Ron Burton

14:30 The rhetorical and empirical extremes of mito-nuclear discordance **David Toews**

Session Coffee break

Monday - 14:45 - 15:15

Room Exhibit Hall A

Session Meet the Editors: A question and answer session Monday - 15:15 - 16:30 **Room Oregon ballroom 201**

Cost: free, no registration required. Open to all conference attendees. An informal question and answer session with a panel of editors from the leading journals in the field including:

- Judith Bronstein, American Naturalist
- Allen Moore, Ecology and Evolution
- Mohamed Noor, Evolution
- Zach Gompert, Evolution Letters (Assoc. editor)
 Barbara Mable, Heredity
 Thomas Near, Systematic Biology

Session Mon4-A105 Phylogenomics / systematics Monday - 15:15 - 16:30 **Room A105**

- 15:15 Evolutionary origins and lifestyle transitions of Vespidae (Hymenoptera) Elizabeth Murray
- 15:30 Rapidly Evolving Long Exon Capture: A New Sequence Capture Dataset for Phylogenomics in **Amniotes**

Benjamin Karin, Tony Gamble, Todd Jackman

- 15:45 The Evolution of Mayflies: Ephemeroptera Phylogenomics T. Heath Ogden, Dustin Miller, Jesse Breinholt
- 16:00 Interrogating difficult nodes in the placental mammal tree **Jacob Esselstyn**, Carl Oliveros, Mark Swanson, Brant Faircloth
- 16:15 Genomic evidence for gene flow between species of magic flowers (Achimenes, Gesneriaceae) Wade Roberts, Eric Roalson

Session Mon4-A106 Development / genomics Monday - 15:15 - 16:30 **Room A106**

15:15 Evo-devo of mimetic color diversity in bumble bees Heather Hines, Li Tian, Sarthok Rahman

15:30 Developing Humanized Mouse Models to Study Human Evolution Emily Dutrow, James Noonan

Session Mon4-A107-109 Comparative biology / macroevolution

Monday - 15:15 - 16:30

Room A107-109

- 15:15 Developmental bias primes anurans for ear loss Molly Womack, Jennifer Stynoski, Kim Hoke
- 15:30 Ecomorphological implications of presacral morphology in Temnospondyli Aja Carter, Lauren Sallan, Peter Dodson, S-Tonia Hsieh
- 15:45 Comparing comparative methods: An evaluation of high-dimensional multivariate phylogenetic comparative approaches

 Dean Adams, MIchael Collyer
- 16:00 Functional controls on the morphological diversification of turtle shells C. Tristan Stayton
- 16:15 Stepwise evolution of floral pigmentation predicted by biochemical pathway structure Julienne Ng, Stacey Smith

Session Mon4-B111-112 Life history / evolutionary ecology Monday - 15:15 - 16:30 Room B111-112

- 15:15 Selection on correlated life history traits under the risk of mortality Veronica Chong, John Stinchcombe
- 15:30 Measuring the cost of whole-organism performance traits: an example with lizard life histories

 Jerry Husak, Simon Lailvaux
- 15:45 Competition mediates selection on metabolic rates in the field Amanda Pettersen, Craig White, Dustin Marshall
- 16:00 Should mothers provision their offspring equally?

 Hayley Cameron, Keyne Monro, Dustin Marshall
- 16:15 Are we underestimating the strength and variability of selection?

 Dustin Marshall

Session Mon4-B113 Experimental evolution 2 Monday - 15:15 - 16:30 Room B113

- 15:15 Population size determines the sign of indirect selection acting on mutation rate modifiers Yevgeniy Raynes, C. Scott Wylie, Paul Sniegowski, Daniel Weinreich
- 15:30 Sounds of silence: the distribution of fitness effects of synonymous mutations Eleonore Lebeuf-Taylor, Rees Kassen
- 15:45 Venom and Drosophila: Response Variation in an Experimental Prey Type Micaiah Ward, Darin Rokyta
- 16:00 "Exploring the relationship between Epistasis and Selection" Kedar Karkare
- 16:15 An experimental test of the effects of rapid evolution on species coexistence in the field.

 Martin Turcotte, Simon P. Hart, Jonathan M. Levine

Session Mon4-B114-115 Population genetics / gene flow Monday - 15:15 - 16:30 Room B114-115

15:15 Selective introgression of exons across a mule deer hybrid swarm Margaret Haines, Gordon Luikart, Stephen Amish, Seth Smith, Emily Latch

- 15:30 Let the seeds fall where they may: Investigating the influence of landscape features on fine-scale seed dispersal
 - Monica Grasty, Mitch Cruzan, Pamela Thompson, Brendan F Kohrn, Elizabeth Hendrickson
- 15:45 Population genetics and gene flow of the arctic moist tundra sedge, tussock cottongrass (Eriophorum vaginatum), in the context of local adaptation and climate change **Elizabeth Stunz**, Jonathon Mohl, Ned Fetcher, Jim Tang, Michael Moody
- 16:00 Chromosome-wide divergence underlies local adaptation along an environmental cline in **Atlantic Silversides**
 - Aryn Wilder, Stephen Palumbi, David Conover, Nina Overgaard Therkildsen
- 16:15 Population genomics and genomic architecture of central Texas cave and spring salamanders
 - Andrew Corbin, Matthew Moseley, Giulia Irene Maria Pasquesi, Drew Schield, Richard Adams, Daren Card, Todd Castoe, Paul Chippindale

Session Mon4-B116 Speciation / genomics / hybridization Monday - 15:15 - 16:30 **Room B116**

- 15:15 RAD-seg as a genome-wide approach to understand allopolyploid evolution in Dactylorhiza (Orchidaceae)
 - Marie Kristine Brandrud, Juliane Baar, Maria Lorenzo, Emiliano Trucchi, Mikael Hedren, Ovidiu Paun
- 15:30 Adaptive introgression from distant Caribbean islands contributed to diversification of novel trophic specialist pupfishes Emilie Richards, Christopher Martin
- 15:45 Prononced epigenetic differentiation and asymmetric hybrid methylome disturbance in Lake Whitefish (Coregonus clupeaformis) species complex Martin Laporte, Jérémy LeLuyer, Madoka Krick, Anne-Marie Dion-Côté, Louis Bernatchez
- 16:00 Exploring the surprisingly finely dissected genomes of the Helianthus hybrid species. Gregory Owens, Loren Rieseberg
- 16:15 Genome-wide patterns of introgression and differentiation in two tidal marsh sparrows Jennifer Walsh, Adrienne Kovach, Brian Olsen, Greg Shriver, Irby Lovette

Session Mon4-B117-119 Evolutionary ecology / diversification Room B117-119 Monday - 15:15 - 16:30

- 15:15 A comparative analysis of the filling of ecological 'space' by marine animals Matthew Knope, Andrew Bush, Luke Frishkoff, Noel Heim, Jonathan Payne
- 15:30 Phylogenetic relationships and the evolution of wing shape tradeoffs in the subfamily Arsenurinae (Lepidoptera: Saturniidae) Chris Hamilton, Nathalie Keller, Jesse Breinholt, Jesse Barber, Akito Kawahara
- 15:45 Rewarding mimicry in a sunbird-pollination system in the Cape Floristic Region Anina Coetzee, Claire N. Spottiswoode, Colleen L. Seymour
- 16:00 The evolution of sexual dichromatism in large carpenter bees (genus Xylocopa) Bonnie Blaimer, Jonathan Mawdsley, Seán Brady
- 16:15 Modeling community phylogenetic structure in adaptive and nonadaptive radiations Jesse Czekanski-Moir

Session Mon4-Ball202 Systematics 2 Monday - 15:15 - 16:30 Room Oregon ballroom 202

- 15:15 Whose ancestor is it anyway? Unraveling the history of polyploid speciation in the North American gray treefrog complex (Hyla chrysoscelis/versicolor) William Booker, Emily Lemmon, Alan Lemmon, Carl Gerhardt, Alyssa Hassinger, Johannes Schul
- 15:30 Phylogenetic position and evolutionary origin of capped-golden langur lineage Kunal Arekar, Praveen Karanth
- 15:45 Laboulbeniomycetes fungal ectoparasites of cockroaches: three-gene phylogeny and implications

Danny Haelewaters, Valter Pfliegler, Michał Gorczak, Donald Pfister

16:00 3-D morphology of pharyngeal dentition of the genus Capoeta (Teleostei: Cyprinidae), its taxonomic and possible phylogenetic significance

Anna Ayvazyan, Davit Vasilyan, Madelaine Böhme

Session Mon4-Ball203 SSE Symposium - The impact of stress on genetic variation 2

Monday - 15:15 - 16:30

Room Oregon ballroom 203

- 15:15 Gene regulation and the repeated evolution of hydrogen sulfide loving fish Joanna Kelley, Courtney Passow, Michael Tobler
- 15:30 The expression of deleterious alleles under environmental stress Crispin Jordan, Kathryn Hodgins, Dilara Ally, Laramy Enders, Charles Fox
- 16:00 Adaptive remodeling of the transcriptional response to drought stress in Arabidopsis lyrata

 Juliette de Meaux, Fei He, Kim Steige, Maroua Bouzid Elkhessairi, Andreas Beyer

Session Mon4-Ball204 ASN Spotlight - '25 Years of Sensory Drive'

Monday - 15:15 - 16:30

Room Oregon ballroom 204

15:15 Past, present & future of Sensory Drive—predicting signal variation from the seen to the unseen

Molly Cummings

15:30 Uniting comparative and experimental approaches to test for a foraging-based sensory bias in live-bearing fishes

Michael Foisy, Jeremy Beaulieu, Luke Frishkoff, David Reznick, Mark Springer, Arcadio Gonzalez,

Michael Folsy, Jeremy Beaulieu, Luke Frishkolf, David Reznick, Mark Springer, Arcadio Gonzalez Helen Rodd, D. Luke Mahler

- 15:35 The Efficient Coding Hypothesis and Signal Design Samuel Hulse, Tamra Mendelson
- 15:40 The pervasive effects of lighting environments on the evolution of color patterns and color vision

Becky Fuller, Lisa Mitchem

- 15:45 The diversity of visual perception in animals: more than just differences in color vision Sönke Johnsen, Eleanor Caves
- 16:00 Rapid and reversible visual plasticity across cichlid habitats Miranda Yourick, Sri Pratima Nandamuri, Karen Carleton
- 16:05 Olfactory receptor evolution shows association with dietary radiation in neotropical Leaf-nosed bats

 Laurel Yohe, Liliana Davalos
- 16:10 Sensory Space of the Threespine Stickleback Robert Mobley, Janette Boughman
- 16:15 Does sensory drive tell us anything about visual-signal diversity in terrestrial habitats?

 Leo Fleishman

Session Mon4-C120-122 Reproductive systems Monday - 15:15 - 16:30 Room C120-122

- 15:15 Evolutionary lag and rescue in stage-structured clonal populations Maria E. Orive
- 15:30 Overview of breeding systems in Fabaceae and Cactaceae withdrawn 2017-06-23 17:12:47.0 Lucy Delaney
- 15:45 The effects of historic range expansion on drift load, inbreeding depression, and autonomous selfing

 Matthew Koski, Jeremiah Busch, Laura Galloway
- 16:00 Differences in Early Life History Traits Between Sexual and Asexual Easter Daisies Evan Hersh, Alberto Ruiz-Larrea Medina, Jeannette Whitton

Session Mon4-C123 Coevolution / host-parasite Monday - 15:15 - 16:30 Room C123

- 15:15 The strange behaviour of genotype frequencies in a hierarchically structured host-parasite metapopulation

 Robert Dünner
- 15:30 Infection dynamics of an invasive trematode in the Upper Mississippi River Region Jennifer Koop, Rebecca Bachtel, Madison Bailey-Schofield
- 15:45 The effect of parasite virulence and host-parasite coevolution on the maintenance of obligate outcrossing in host populations

 Jae Hoon Cho, McKenna Penley, Sathvik Namburar, Levi Morran
- 16:00 Tree shape-based approaches for the comparative study of cophylogeny Mariano Avino, Art Poon
- 16:15 Sexual Conflict and STIs

 Alison Wardlaw, Aneil Agrawal

Session Mon4-C124 Molecular evolution / mammals Monday - 15:15 - 16:30 Room C124

- 15:15 The Molecular Evolution of Human Breasts
 Reed Cartwright, Joanna Malukiewicz, David Winter
- **15:30** Transposable Elements and the Evolution of Mammalian Pregnancy *Katelyn Mika*, *Vincent J. Lynch*
- 15:45 Hox clusters molecular evolution provides hints for mammalian evolution

 Mariana Nery, Lucas Canesin, Esther Camilo, Juan Pablo Torres-Florez, Tiana Kohlsdorf
- 16:00 Baby genomics: tracing the evolutionary changes that gave rise to placentation Yue Hao, Hyuk Jin Lee, Michael Baraboo, Katie Burch, Taylor Maurer, Gavin Conant
- 16:15 Functional changes in bat dim-light visual pigment evolution are associated with differing echolocation abilities

 Eduardo de Almeida Gutierrez, Gianni Castiglione, James Morrow, Belinda Chang

Session SSE Student-Faculty Mixer Monday - 16:30 - 17:30 Room Oregon ballroom 201

Open to students & faculty SSE members. Come chat with your SSE colleagues. Cash bar and SSE grad member free drink tickets accepted.

Session Mon5-Ball202 SSE Presidential address & award announcements

Monday - 17:30 - 18:30 Room Oregon ballroom 202-204

Linking levels of selection with genetic modifiers Sally Otto

Session SSE Members business meeting Monday - 18:30 - 19:30 Room A105

Poster Session Mon-01 Adaptation Monday - 18:30 - 20:30 Room EHA-01

- 1 Drought Resistance Not Lost in Cultivated Sunflower (Helianthus annuus L.) Victoria Burns, Lisa Donovan
- 2 The Evolution of Saccharomycotina Yeasts
 Dana Opulente, Emily Rollinson, Amanda Hulfachor, Cleome Bernick-Roehr, Antonis Rokas, Cletus
 Kurtzman, Chris Hittinger
- Widespread duplication and gonad-specific expression of nuclear import genes in stalk-eyed flies (Diopside)

 Philip Johns, Nawat Bunnag, Rodzay Wahab, Richard Baker

Topic Mon-02 Bioinformatics

- The effect of common inversion polymorphisms In(2L)t and In(3R)Mo on patterns of transcriptional variation in Drosophila melanogaster

 Erik Lavington, Andrew Kern
- 5 The Supertree Toolkit Jon Hill, Katie Davis
- An efficient pipeline for assaying whole-genome plastid variation for population genetics and phylogeography

 Brendan F Kohrn, Jessica Persinger, Mitch Cruzan

Topic Mon-03 Comparative biology

- 7 The evolutionary origin of variation in song length and frequency in the avian family Cettiidae Chentao WEI, Trevor Price, Jiayu Liu, Per Alstrom, Yanyun Zhang
- 8 Phylogenetic trends and environmental correlates of genome size variation in Helianthus sunflowers
 - Fan Qiu, Eric Baack, Dan Bock, Loren Rieseberg, Kenneth Whitney, Mark Ungerer
- 9 Life history correlates of adult attendance in New World temperate and tropical songbirds Suzanne Austin, W. Douglas Robinson, Tara Rodden Robinson, Robert Ricklefs
- 10 Evolution of diet and visual acuity in damselfishes (Pomacentridae)
 Sarah Williams, Madison Wiltse, Jessica Valenzuela, Lars Schmitz
- 11 Morphological divergence and integration among subcastes in the hyper-diverse ant genus ants
 Nicholas Friedman, Beatrice Lecrog, Evan Economo

Topic Mon-04 Ecological genetics

- 13 Resurrected "ancient" Daphnia genotypes show reduced thermal stress tolerance compared to modern descendants
 - **Lawrence Weider**, Aime'e Yousey, Priyanka Roy Chowdhury, Nicole Biddinger, Jennifer Shaw, Punidan Jevasingh
- 14 Exploring Leaf Associated Fungal Communities of Grassland Plants Ryan Estes, Mark Ungerer, Ari Jumpponen
- 15 Genetic structure and diversity through time for northern California Peromyscus maniculatus Robert Boria, Sarah Brown, Jessica Blois

Topic Mon-05 Education

- 16 ETC (Explorer of Taxon Concepts) Pipeline for Text Mining, Ontology Building, Matrix Construction, and Taxon Concept Comparison

 Hong Cui, James Macklin, Nico Franz, Bertram Ludäscher, Lorena Endara, Lisa Moore, Carrine Blank, Robert Morris, Thomas Rodenhasen, Joel Sachs, J. Gordon Burleigh
- 17 Undergraduate Student Conceptions About Randomness and Mutation.

 Michael Wiser, Robert Pennock, Louise Mead, Cory Kohn, James Smith
- 18 Students Discover: ANTS Connecting Science & Education
 Daniela Magdalena Sorger, Paige Derouin, Michelle Hafey, Maggie McKinley, Robert Dunn
- 19 Efficacy of a Travel Award Program to Increase Diversity in Evolutionary Science Hollie Heape
- 20 Procedural generation of video game creatures using evolution of digital genomes. Landon Wright, Barrie Robison, Terence Soule, Samantha Heck
- 21 Biology Majors Acceptance of Evolution Chad Talbot, T. Heath Ogden
- 22 Bringing Food Trucks Into the Classroom: A Lesson on Ecology, Evolution, and Community Assembly Theory Elizabeth Carlen, Elle Barnes

Topic Mon-06 Experimental evolution

- 23 Dynamics of Adaptive Allele Frequency Change in Experimental Populations of Saccharomyces cerevisiae

 Benjamin Weeder, Eugene Housman, Molly Burke
- 24 Motile Bacteria Evolve in Response to Bacteriophage Spatial Structure Michael Blazanin, Michael Travisano
- 25 Fitness Consequences of Cryopreservation on Diverse Natural Isolates of Saccharomyces Cerevisiae

 Kieslana Wing, Molly Burke
- 26 Effects of anthropogenic noise on mating behavior and fitness of an acoustically signaling invertebrate

 Gabby Gurule-Small
- 27 Adaptation to Discovered and Self-constructed Niches During Incipient Speciation in an Experimental Population of E. coli

 Zachary Blount, Caroline Turner, Richard Lenski, Kiyana Weatherspoon, Brooke Sommerfeld, Maia Rowles

Topic Mon-07 Expression studies/transcriptomics

- 28 Physiological and genetic intersections between drought and freezing tolerance in Pooideae Aayudh Das, Natalie Dedon, Jill Preston
- 29 Dissecting the causes of natural variation in protein expression dynamics Ciara Asamoto, Daniel Pollard
- 30 Differential gene expression and hybrid male sterility in Drosophila Alberto Civetta, Doaa Alhazmi
- 31 Comparative transcriptomics of the cellular response to hypoxia in a hibernating primate Kelsie Hunnicutt, Adam Brown, Peter Larsen, Sheena Faherty, C. Ryan Campbell, Rachel Williams, Anne Yoder
- 32 Evolved differences in gene expression underlie geographic variation in desiccation resistance in Rhagoletis flies

 Christa Kohnert, Dietmar Schwarz
- 33 Vision loss and transcriptome evolution in cave-adapted crayfish David Stern, Keith Crandall

Topic Mon-08 Genomics

- 34 Analysis of Host Range Mutants of Bacillus subtilis Bacteriophages Alexandra Agesen, Veronique Delesalle, Greg Krukonis
- 35 MycoMeth: the comparison and evolution of DNA methylation across fungi Adam Bewick, Brigitte Hofmeister, Rob Powers, Zachary Lewis, Timothy James, Robert Schmitz, Jason Stajich
- 36 Identifying the location of phototransduction genes in the rainbow trout genome Christy Smith
- 37 Natural genomic variation underlying stress tolerance and metabolic traits in rhizobia Brendan Epstein, Liana Burghardt, Joseph Guhlin, Matt Nelson, Reda Abou-Shanab, Abdelaal Shamseldin, Michael Sadowsky, Peter Tiffin
- 38 Roots To The Evolution Of Photosynthesis? Biofilm Communities In A Manganese Depositing Hot Spring Wendy Smythe

Topic Mon-09 Host-parasite

- 39 Whole genome and transcriptome sequencing of a major crop pest in Sub-Saharan Africa, Busseola fusca (Lepidoptera) withdrawn 2017-06-23 20:03:01.0

 Kayla Hardwick, Sarah Schaack
- 40 Macroevolutionary patterns of susceptibility to avian malaria Lisa Barrow, Sabrina McNew, Spencer Galen, Holly Lutz, Thomas Valqui, Heather Skeen, Jason Weckstein, Christopher Witt
- 41 Bioinformatic analysis for viral sequences detection in the transcriptome of five species of bats from Yucatan, Mexico.

 Diana Moreno , Jorge Ortega, Georgina Hernández-Montes

Topic Mon-10 Hybridization

- **42 Genomic analysis of ancestry in hybrid mice** *Megan Frayer*, Leslie Turner, Bettina Harr, Diethard Tautz, Bret Payseur
- 43 Investigating F1 hybrid seed inviability in subspecies of Mimulus tilingii Gabrielle Sandstedt
- 44 MHC genetic variation in hybridizing species, the case of the Alouatta hybrid zone in Mexico Adrian Melo Carrillo, Liliana Cortes Ortiz
- 45 Inter-specific population dynamics of the Common goldeneye and Barrow's goldeneye Joshua Brown, Sarah Sonsthagen, Robert Wilson, Sean Boyd, Sandra Talbot, Philip Lavretsky
- **46** Evolution of novel mimicry rings facilitated by adaptive introgression in tropical butterflies *Juan Enciso*, Carolina Pardo-Diaz, Camilo Salazar, Simon Martin, Chris Jiggins, Owen McMillan, Carlos Arias, Mauricio Linares

Topic Mon-11 Invasive/invasion

- 47 Fitness cost of glyphosate resistance in hairy fleabane (Erigeron bonariensis) under drought conditions
 - Ankit Pathak, **Adam Avila**, Katherine Waselkov
- 48 Testing Darwin's Naturalization Hypothesis: Phylogenetic Community Structure Among Native and Non-Native Species
 William Weaver, Robert Laport, Julienne Ng
- 49 Invasion dynamics of wavyleaf basketgrass (Oplismenus undulatifolius), an emerging threat to US mid-Atlantic forests

 Carrie Wu, Anna Hakkenberg, Emma Dressel
- 50 Exploring the epigenetic reasons behind invasive species success

 Andrea Sequeira, Flavia Mendonca de Sousa, Mary Dornon, Viviana Confalonieri, Marcela Rodriguero,
 Noelia Guzman, Analia Lanteri
- 51 Reciprocal transplantation of Japanese knotweed (Reynoutria japonica) reveals phenotypic differentiation, but no local adaptation across a wide latitudinal range.

 Acer VanWallendael, Elena Hamann, Steven Franks**

Topic Mon-12 Macroevolution

- 52 What Do We Know About Cockroach Diversification? Andres Velez, Juan M. Daza
- 53 Miocene adaptive radiation in grasshoppers revealed by a species-level supertree of Orthoptera

Katie Davis, Adam Bakewell, Ed Baker, Jon Hill, Peter Mayhew, David Chesmore

- 54 Insights into feather diversity from the King of Saxony Bird of Paradise and other bizarre feathers

 Teresa Feo, Vinodkumar Saranathan, Richard Prum
- 55 Pigeon domestication, what can artificial selection tell us about evolutionary constrains Cesar Augusto Espinoza Campuzano
- 56 Accelerated diversification in lichen forming fungi after the K-Pg boundary Jen-Pan Huang

Topic Mon-13 Molecular evolution

- 57 Mutation-selection models for the detection of adaptation in protein-coding genes Nicolas Rodrigue
- The evolution of metabolic networks in high-altitude deer mice withdrawn 2017-06-19 10:56:45.0

Rena Schweizer, Zachary Cheviron

- 59 Linking genotype to phenotype: Using CRISPR/Cas9 to alter the wing pattern coloration of Heliconius butterflies

 David Tian, Owen McMillan*, Carolina Concha*
- 60 The risk of using amplicon size of chloroplast SSR marker for genetic relationship studies Sang Min Chung
- 61 Numts and heteroplasmy disrupting mitochondrial markers on evolutionary studies Elaine Françoso, Paulo Cseri, Maria Cristina Arias

Topic Mon-14 Mutation

- 62 Do somatic mutations speed up the evolution of Sitka spruce? Vincent Hanlon, Sally Aitken
- 63 Investigating the genetic basis of glyphosate resistance in the San Joaquin Valley agricultural weed hairy fleabane (Erigeron bonariensis)
 Rigoberto Molina, Ankit Pathak, Katherine Waselkov
- 64 Evolvability of emerging host range mutants of RNA virus phi6 Lele Zhao, Siobain Duffy
- Accumulation and transmission dynamics of a naturally-occurring mtDNA deletion in Caenorhabditis briggsae

 Jennifer Sullins, Anna Coleman-Hulbert, Suzanne Estes
- 66 The Effects of Spontaneous Mutation on Neural Function
 Sarah Eberle, Djeneba Dezoumbe, Rhegan Humphrey, Shane Kinzer, Whitney Raver, Sarah Schaack,
 Leigh Latta IV

Topic Mon-15 Mutualism

67 Distribution of a Host range restriction peptidase (hrrP) in populations of symbiotic Ensifer medicae across the native and invaded regions of its host legume Emily Helliwell, Joel Griffitts, Maren Friesen, Stephanie Porter

68 Effects of antibiotic exposure on growth of bacterial strains isolated from threespine stickleback gut

Kenneth Sparks, Emily Lescak, Ruth Isenberg, Ryan Lucas, Kelly Ireland, Koral Campbell, Kat Milligan-Myhre

Topic Mon-16 Paleobiology

- 70 Phenotypic innovation and adaptive constraints in the evolutionary radiation of Paleozoic crinoids (Echinodermata)
 David Wright
- 71 The NSB (Neptune) marine microfossil occurrences database: design, content and use in paleobiologic research

 David Lazarus, Johan Renaudie
- 72 Body size evolution and homology of parafrontal bones in sphaerodactylid geckos (Reptilia: Squamata)

 Aaron Griffing, Tony Gamble, Aaron Bauer
- 73 Inferring Behavioral and Social Organization in Fossil Antilocapridae
 Holley Flora, Edward Byrd Davis
- 74 Revised Evolutionary History of the Sabertooth Carnivoran Barbourofelis (Nimravidae)
 Paul Barrett
- 75 Machine reasoning about phenotypes: enhancing expert knowledge about the genetics of a fossil transition

 Todd Vision, James Balhoff, Wasila Dahdul, Alex Dececchi, Pasan Fernando, Hilmar Lapp, Paula Mabee, Prashanti Manda, Kellen Mastick, Monte Westerfield, Erliang Zeng
- 76 Diversification rates vary more through time than between sub-clades for fossil Carnivorans Daniel VARAJAO DE LATORRE

Topic Mon-17 Pathogens

- 77 Comparative Genomics and Evolution of Pseudomonas syringae Bacteriophages

 Jenna DeCurzio, Rebecca King, Britt Koskella, Norma Morella, Veronique Delesalle, Greg Krukonis
- 78 Genomic signatures of pathogen-mediated selection across the bird tree of life Allison Shultz, Julia Yu, Tim Sackton
- 79 Population Genetics of Complex Human Disease Jeremy Berg, Guy Sella
- 80 Metabarcoding for streamlining vector surveillance: Culicoides spp. (Diptera: Ceratopogonidae) as a test case

 Matthew Hopken, Zaid Abdo, Antoinette Piaggio

Topic Mon-18 Phenotypic plasticity/GxE

- 81 Genome-wide methylation patterns are associated with parasite load in guppies (Poecilia reticulata)

 JUNTAO HU, FELIPE PÉREZ-JVOSTOV, Léa Blondel, Rowan Barrett
- 82 Untangling Morphotype and Latitudinal Variation in Spartina Alterniflora August Guang, Lillian Hancock, David Rand, Emily Hollenbeck
- 83 Temperature effects on Oncorhynchus mykiss muscle physiology Erin Wiese, Patrick Carter
- 84 Rapid plasticity of opsin expression in adult African cichlids Miranda Yourick, Sri Pratima Nandamuri, Karen Carleton
- 85 Resource quality and predation risk influence prey eye size and allometry Jared Goos, Matthew Walsh
- 86 Plastic induction of cellular differentiation in an undifferentiated species Dinah Davison, Richard Michod
- 87 Evolution of plasticity: mechanistic link between development and reversible acclimation Julian Beaman, Craig White, Frank Seebacher

Topic Mon-19 Phylogenetic comparative methods

- 88 The phylogenetic distribution of climate change vulnerability in California's birds Shelby Moshier, Joshua Reece, Gurjap Dhaliwal
- 89 Phylogenetic relationships of Crocodyliformes based on morphology and RAG1 sequence data Alexandria DiGiacomo, Robert Meredith, Tracy Heath, Daniel Ksepka
- 90 A novel approach for deciphering cryptic species problems using large scale RNA-seq data Fiona Wood, Yonas Tekle
- 91 Morphological rate shifts and disparity in the postcrania of Rodentia Carl Saltzberg, Scott Steppan
- 92 Phylogeny of Eucnide (Loasaceae): Taxonomic implications Josh Brindley, Larry Hufford

Topic Mon-20 Phylogenetic methods development

- 93 Graph Splitting Method: A New Phylogenetic Approach to Resolve the Early Evolution of Protein Superfamilies

 Motomu Matsui, Wataru Iwasaki
- 94 Efficiency and accuracy of phylogenetic community detection using Treescaper Genevieve Mount, Diego Elias, David Morris, Jeremy Ash, Wen Huang, Melissa Marchand, Kyle Gallivan, James Wilgenbusch, Jeremy Brown
- 95 Probabilistic Path Hamiltonian Monte Carlo for Bayesian Phylogenetic Inference Vu Dinh, Arman Bilge, Chenq Zhang, Frederick Matsen
- 96 Faster likelihood computation for sparse alignment matrices Benjamin Redelings
- 97 A complete ray-finned fish phylogeny using taxonomy and birth-death models Jonathan Chang, Michael Alfaro

Topic Mon-21 Phylogenomics

- 98 Phylogenomics of Formica wood-ants reveals single origins of dependent colony foundation and frequent dispersal between the Old and New World

 Marek Borowiec, Christian Rabeling
- 99 Phylogenomics using target capture of ultraconserved elements and exons reveals deep relationships within the Neotropical manakins (Aves: Pipridae)

 Rafael Leite, Rebecca Kimball, Ed Braun, Elizabeth Derryberry, Peter Hosner, Graham Derryberry, Marina Anciães, Diogo Meyer, Alexandre Aleixo, Camila Ribas, Robb Brumfield, Joel Cracraft

Topic Mon-22 Phylogeography

- 100 Covicariance Revisited: Using whole plastome sequence and RAD seq data to estimate the timing of speciation in Joshua Trees.
 Christopher Smith
- 101 Phylogeography of seadragons inferred from genome-wide data Josefin Stiller, Nerida G. Wilson, Greg Rouse
- 102 Testing diversification processes of Neotropical open biomes: phylogeography of the Narrow-Billed Woodcreeper (Lepidocolaptes angustirostris).

 Amanda Rocha, Gustavo Sebastian Cabanne, Alexandre Aleixo, Luís Fábio Silveira, Pablo Tubaro, Renato Caparroz
- 103 Phylogeny and biogeography of deep-sea wood boring Xylophagainae (Mollusca, Bivalvia) Chiara Romano, Greg Rouse, Christian Borowski, Verena Tunnicliffe, Daniel Martin

- 104 Population genetic analysis of Amusium pleuronectes in the Coral Triangle to identify distinct populations or a species complex

 Jessica Matadamas, Alvin Alejandrino
- 105 Phylogeography, population genomics and adaptive variation on species complex Pitcairnia lanuginosa (Bromeliaceae)

 Clarisse Palma, Barbara Leal, Vanessa Graciano Araujo, Cleber Chaves, Luis Huacre, Myriam Heuertz
- 106 Effects of Andean geographic history on the population history of the plant-ant Azteca.

 Maria Fernanda Torres Jimenez, James Richardson, Graham Stone
- 107 An Investigation of the Impact of Phylogeny on Continental Scale Phenotypic Variation in Garter Snakes (Thamnophis)

 Leonard Jones, Frank Burbrink, Adam Leache

Topic Mon-23 Plant reproductive biology

- 108 Weed versus wildflower: Does habitat encroachment by an exotic dandelion species disrupt the breeding system of its native alpine congener?

 Austin Lynn, Candace Galen
- 109 Selection on floral traits in two species of Alpine orchids (Gymnadenia) Kelsey Byers, Roman Kellenberger, Philipp Schlüter
- 110 Pollen viability varies with elevation in high temperature conditions Caitlyn Byron, David Lowry
- 111 Does widespread purging of inbreeding depression occur in populations of the common yellow monkeyflower, Mimulus guttatus?: a field-based approach.

 Vanessa Koelling, Toshiro Newsum, John Kelly

<u>Topic Mon-24 Population genetics: inference of</u> selection

- 112 Evolutionary rescue of bat populations devastated by white-nose syndrome? Giorgia Auteri, L. Lacey Knowles
- 113 Investigating Expression Adaptation and Conservation in Alternative Splicing Jamie Moon, Ksenia Arzumanova, Scott Roy, Manuel Irimia, Rori Rohlfs
- 114 Examining models of recurrent hitchhiking in Drosophila melanogaster and their effects on demographic inference

 Jeremy Lange, John Pool
- 115 Looking for signals of selection in the ochre sea star genome following a mass mortality event Dannise Ruiz-Ramos, Lauren Schiebelhut, John Wares, Michael Dawson

Topic Mon-25 Population genetics: molecular ecology

- 116 Population genomics supports speciation with gene flow, not genomic islands of differentiation, in sky-island populations of southwestern white pine

 Justin Bagley, Mitra Menon, Chris Friedline, Amy Whipple, Anna Schoettle, Alejandro Sáenz, Christian Wehenkel, Daniel McGarvey, Lluvia Flores-Renteria, Richard Sniezko, Sam Cushman, Kristen Waring, Andrew Eckert
- 117 Comparative spatial genomics of urban brown rats: From studying populations to understanding species

 Matthew Combs, Jason Munshi-South, Michael Blum, Chelsea Himsworth
- 118 Genetic consequences of range expansions in North American amphibians and reptiles Jeffrey Streicher
- 119 Speciation despite gene flow in two owls (Aegolius ssp.): evidence from 2,210 loci with ultraconserved elements (UCEs)

 Kevin Winker, Travis Glenn, Jack WIthrow, Spencer Sealy, Brant Faircloth

<u>Topic Mon-26 Population genetics: theory and</u> methods

- 120 On the joint distribution of coalescent tree height and tree length *Ilana Arbisser*
- 121 On the joint distribution of coalescent tree height and tree length withdrawn 2017-06-23 11:43:14.0

Ilana Arbisser

- 122 MIG-seq: efficient PCR-based method for genome-wide sequencing Yoshihisa Suyama, Chika Mitsuyuki, Yoshihiro Tsunamoto
- 123 Inversion polymorphism in populations with sexual antagonism and reproductive skew Christopher McAllester, John Pool
- 124 Characterizing adaptive Neanderthal introgression in modern humans Sivan Yair, Graham Coop
- 125 How do inversions affect recombination? A study in the gibbon Debora Brandt, Lucia Carbone, Jeffrey Wall

Topic Mon-27 Quantitative genetics

- 126 Morphological, genetic and evolutionary bases for structural color shift in the Buckeye butterfly Rachel Thayer
- 127 A quantitative genetics approach to studying metabolite contributions to an ancient plant insect interaction

 Leslie Kollar, Todd Rosenstiel, Sarah Eppley, Stuart McDaniel
- 128 The Footprints of Hybrid Zone Movement Remi Matthey-Doret, Silu Wang

Topic Mon-28 Reproductive biology

- 129 Norms of reaction for pheromone response in Saccharomyces cerevisiae isolates Shelby Duffy, Daniel Pollard
- 130 Influence of weight and age on seminal parameters of captive golden-headed lion tamarin Patricia Hergert Bacher1, Paloma Rocha Arakaki, Bruno Sauce, Rodrigo del Rio do Valle, Andrea C Peripato

Topic Mon-29 Reproductive isolation

- 131 Variation in drought-response traits in sympatric Mimulus Samuel Mantel
- 132 Speciation in the fast lane: Rapid genetic and reproductive isolation in a young Drosophila species complex Roman Yukilevich, Luana Maroja
- 133 Mate -choice mechanisms and the dynamics of reproductive isolation: a simulation study withdrawn 2017-06-22 08:59:50.0

 Daniel Powell
- 134 Quantifying behavioral isolation between closely related species of mice *Tyler Ryan*, *Polly Campbell*
- 135 Testing for postzygotic isolation in a Joshua tree hybrid zone Anne Royer, Christopher Smith

Topic Mon-30 Sensory systems

- 137 Dim-light vision in ancient whales: Reconstructing the ancestral cetacean rhodopsin Sarah Dungan, Belinda Chang
- 138 Molecular phylogeny and behavioral trait evolution of singing flies Kaila Colyott, Jennifer Gleason
- 139 Increasing structural complexity is associated with decreasing molecular complexity in the evolution of the shell-eyes of chitons

 Alexandra Kingston, Daniel Speiser**

Topic Mon-31 Sex/recombination

- 140 An ancient UV sex chromosome system in mosses
 Sarah Carey, Adam Payton, Matthew Johnson, Roth Conrad, J. Gordon Burleigh, Jeremy Schmutz,
 Jerry Jenkins, Kerrie Barry, Jane Grimwood, Stuart McDaniel
- 141 The effects of reproductive mode and ploidy level on transposable element evolution Kyle McElroy, Emily Opfer, Jorge Moreno, Jeffrey L Boore, John Logsdon, Maurine Neiman
- 142 Polygenic Sex Determination in Astatotilapia burtoni Erin Peterson, Ashley Elias, Reade Roberts

Topic Mon-32 Sexual conflict

144 Parent Offspring Recognition in Helobdella Leeches

James Boothroyd, Angela Rincon, Hannah Combs, Rebecca Beresic-Perrins, Stephen Shuster

Topic Mon-33 Sexual selection

- 145 Increased fitness with brighter juvenile color in collard lizards (Crotaphytus collaris)

 Jodie Wiggins
- 148 Sexual Dimorphism and the Evolution of Sex-Limited Genes in the Dioecious Tree Populus balsamifera

 Matt Olson, Brian Sanderson, Peter Tiffin, Li Wang, Zhi-qiang Wu
- 149 Sex-specific effects of serotonin on aggressive behaviors in stalk-eyed flies Jaime Grace, Nadia Lovko
- 150 Relative Importance of Natural and Sexual Selection in Speciation Megha Suswaram, Danielle Edwards, Justin Yeakel

Topic Mon-34 Social systems

- 151 Foundations of chemosensation in paper wasps Andrew Legan, Michael Sheehan
- 152 Society at the edge of collapse: Social mobility, opportunity, and adaptive strategy during group fission in a nonhuman primate

 Lixing Sun, Dongpo Xia, Shine Sun, Jinhua Li
- 153 Shifting cichlid color vision: Tbx2a acts in trans on LWS opsin expression

 Ben Sandkam, Sri Pratima Nandamuri, William Gammerdinger, Connor O'Brien, Matthew Conte, Karen
 Carleton

- 154 Understanding the evolution of social behavior: The regulation of estrogen receptor alpha expression in a species with alternative behavioral phenotypes

 Kathleen Grogan, Jennifer Merritt, Wendy Zinzow-Kramer, Suzanne Mays, Eric Ortlund, Donna Maney
- 155 The influence of sociality on the evolution of defensive morphology in butterflyfishes Jennifer Hodge, Peter Wainwright

Topic Mon-35 Speciation

- 156 Hybrid deficiency in learning and memory: A novel reproductive isolating barrier?

 Michael McQuillan, Amber Rice
- 157 Genome-wide congealing and the spatial scale of differentiation Martin Schilling, Samuel Flaxman, Jeffrey Feder, Patrik Nosil
- 158 Exploring reinforcement and genetic mechanisms of incipient speciation in prairie sunflower (Helianthus petiolaris)

 Kaichi Huang, Kate Ostevik, Rose Andrew, Loren Rieseberg
- 159 Widespread species shed light on the process of speciation in Neotropical fishes Jack Craig, James Albert, William Crampton

Topic Mon-36 Species delimitation

- 160 Identifying the loci that underlie a multifaceted crossing barrier Kate Ostevik, Joanna Rifkin, Mark Rausher
- 161 Low-cost genomic architecture as a species delimitation tool using rDNA fingerprints John Sproul, David Maddison
- 162 A day-flashing Photinus firefly (Coleoptera: Lampyridae) from central Panamá: an emergent shift to predator-free space?

 Luana Maroja, Fredric Vencl, Xin Luan, Xinhua Fu
- 163 Genetic, morphological, and ecological species delimitation of the greenthroat darter (Etheostoma lepidum)

 Daniel MacGuigan, Thomas Near

Topic Mon-37 Systematics

- 164 Molecular phylogeny of a morphologically cryptic frog genus Fejervarya from South Asia reveals unrecognized diversity prudhvi raj
- 165 Tyrophagus curvipenis (Acari: Acaridae): investigating species boundaries using DNA and morphology

 PAMELA MURILLO, Pavel Klimov, Barry OConnor
- 166 Effects of orthology assessment and gene occupancy on the phylogenomic analysis of the orb-weaving spider family Araneidae (Arthropoda, Arachnida, Araneae).

 Robert Kallal, Rosa Fernandez, Gonzalo Giribet, Gustavo Hormiga
- 167 Phylogenetics in the speication "grey zone": disentangling relationships among the paintbrushes (Castilleja; Orobanchaceae) in the midst of rampant hybridization and introgression

 Maribeth Latvis, Paul Blischak, David Tank
- 168 Phylogeny of Mayfly Subfamily Ecdyonurinae (Heptageniidae: Ephemeroptera)

 Ishaah Talker, T. Heath Ogden
- 169 Evaluating Conflicting Topologies of the Genus Juniperus using Plastome Sequences Therese Balkenbush, Linda Raubeson, Rhiannon Peery, Julie Morris, Robert Adams, Andrea Schwarzbach
- 170 Revisiting the architecture of sciurid phylogeny John Mercer, V. Louise Roth

- 8:30 Evolutionary correlates of seasonal migration in a family of avian aerial insectivores withdrawn 2017-06-24 10:33:10.0 Clare Brown, Frederick Sheldon
- 8:45 Pattern and paradox in the macroecology and evolutionary history of Passerine nest types Jay McEntee, J. Gordon Burleigh
- 9:00 A slimy problem: body size and shape evolution in hagfish Joshua Reece, Theodore Uyeno, Toby Daly-Engel
- 9:15 Macroevolution and morphological diversification of Neotropical headstanding fishes (Characiformes: Anostomoidea) as illuminated by a new ultraconserved element phylogeny Brian Sidlauskas, Brant Faircloth, Michael Alfaro, Kendra Hoekzema, Bruno Melo, Mike Burns, Casey Dillman, Ben Frable, Mark Sabaj, Claudio Oliveira, Richard Vari
- 9:30 An evolutionary shift in photosynthetic pigment composition facilitated functional diversification in cryptophytes Matthew Greenwold, Brady Cunningham, Tammi Richardson, Jeff Dudycha

Session Tue1-A106 Bioinformatics

Tuesday - 8:30 - 9:45

Room A106

- 8:30 Authors in the driver's seat: fast, consistent, computable phenotype data and ontology production
 - **Hong Cui**, James Macklin, Julian Starr, Anton Reznicek, Joel Sachs, Bruce Ford, Lyubo Penev, Hsinliang
- 8:45 Blue tail, red tail: a solution to inapplicable discrete characters in phylogenetics Thomas Guillerme, Martin Smith, Martin Brazeau
- 9:00 A computer vision for animal ecology Ben Weinstein
- 9:15 Simpler is better: Machine learning using amino acid composition to classify CRESS viruses to aenus. Erik Lavington, Lele Zhao, Siobain Duffy
- 9:30 aTRAM 2.0: Targeted assembly of loci from both reduced and whole genome NGS data Julie Allen, Raphael LaFrance, Ryan Folk, Kayce Bell, Kevin Johnson, Robert Guralnick

Session Tue1-A107-109 Comparative biology 1 Tuesday - 8:30 - 9:45 Room A107-109

- 8:30 Of cavefish and cavemen. What can we learn from blind, fat fish about ourselves? Nicolas Rohner
- 8:45 The evolution of size and shape of insect eggs across eight orders of magnitude Samuel Church, Seth Donoughe, Bruno de Medeiros, Cassandra Extavour
- 9:00 Comparative Analysis of Parental Effects on Developmental Period Length in Anseriform Birds Daniel Barton
- 9:15 The evolution of craniofacial changes in the blind Mexican cavefish Amanda Powers, Erin Davis, Shane Kaplan, Tyler Boggs, Joshua Gross
- 9:30 Big mammals with small voices comparison of vocalization frequency limits shows how aquatic species diverged from expected trends. Kobe Martin, Marlee Tucker, Tracey Rogers

Session Tue1-B111-112 Microbes 1

Tuesday - 8:30 - 9:45

Room B111-112

- 8:30 Salamanders of the Trans-Mexican Volcanic Belt: the microbiome in a phylogenetic context Julio Cesar Garcia Sanchez, Sean Rovito
- 8:45 Genome evolution and gene loss in the luminous symbionts of deep sea anglerfish Tory Hendry

- 9:00 The microbe advantage: Adaptation to novel diet through a change in the microbiome Aparna Agarwal, Deepa Agashe
- 9:15 Inferring microbial gene family evolution despite phylogenetic uncertainty Mukul Bansal
- 9:30 The holobiont of domesticated and wild Brassicas Elizabeth Mittell

Session Tue1-B113 Adaptation / experimental evolution Tuesday - 8:30 - 9:45 Room B113

- 8:30 Experimental adaptation of a non-host-associated bacterium to the zebrafish digestive tract Jarrett Lebov, Brendan Bohannan
- 8:45 Strategies for investigating resistance evolution in complex organisms

 Barbara Mable, Stephen Doyle, James Cotton, Alan Reynolds, Eileen Devaney
- 9:00 Mitochondrial mutation compensates for nuclear-encoded ETC dysfunction during C. elegans experimental evolution
 Stephen Christy, Suzanne Estes, Jennifer Sullins
- 9:15 Genomic Basis of Rapid Adaptation to a Low Quality Host Plant in Callosobruchus maculatus Alexandre Rego, Samridhi Chaturvedi, Frank Messina, Zachariah Gompert
- 9:30 Reversal of response to artificial selection on body size in a wild passerine bird Thomas Kvalnes, Thor Harald Ringsby, Henrik Jensen, Ingerid Julie Hagen, Bernt Rønning, Henrik Pärn, Håkon Holand, Steinar Engen, Bernt-Erik Sæther

Session Tue1-B114-115 Molecular ecology / gene flow Tuesday - 8:30 - 9:45 Room B114-115

- 8:30 Population structure and genetic diversity of two amphibian species in an urbanizing landscape
 - Jared Homola, Michael Kinnison, Cynthia Loftin
- 8:45 Inferring Demographic and Evolutionary History in a Coral Reef Goby (Coryphopterus glaucofraenum)

 Daniel Volk, Carlos E. L. Ferreira, Eric Hoffman
- 9:00 Investigating Population Structure and Barriers to Gene Flow in the American Badger of British Columbia (Taxidea taxus jeffersonii) to Inform Conservation Action

 Brett Ford, Rich Weir, Karl Larsen, Michael Russello
- 9:15 Deconstructing dispersal potential Lauren Schiebelhut, Laura Jurgens, Brian Gaylord, Rick Grosberg, Michael Dawson
- 9:30 Phylogeography and population genomics of the Olympia oyster (Ostrea lurida)

 Katherine Silliman, Doug Eernisse

Session Tue1-B116 Hybridization / reproductive isolation Tuesday - 8:30 - 9:45 Room B116

- 8:30 Cultural, genetic, and behavioral divergence coincide in an avian hybrid zone
 Sara Lipshutz, Isaac Overcast, Michael Hickerson, Robb Brumfield, Elizabeth Derryberry
- 8:45 Mate-choice mechanisms and reproductive isolation in swordtail fish Gil Rosenthal
- 9:00 The chemical ecology of a hybridizing chickadee system: olfaction and reproductive isolation Alex Huynh
- 9:15 Repeated hybridization between Yellowstone cutthroat trout and rainbow trout Elizabeth Mandeville
- 9:30 Mate choice in early generation swordtail fish hybrids Daniel Powell, Gil Rosenthal

Session Tue1-B117-119 Adaptation / evolutionary ecology 1

8:30 Evolutionary constraint on range limits in Boechera stricta: Tradeoff linked to signaling network

Jason Olsen, David Siemens

- 8:45 Little plant, big city: testing for urban adaptation in ragweed Amanda Gorton, David Moeller, Peter Tiffin
- 9:00 Climate change generates maladaptation along an elevational gradient Susana Wadgymar, Jill Anderson
- 9:15 Evolutionary responses to repeated drought episodes in Brassica rapa Elena Hamann, Steven Franks
- 9:30 A dynamic eco-evolutionary model predicts slow response of alpine plants to climate warming Olivier Cotto, Max Schmid, Stefan Dullinger, Wilfried Thuiller, Frédéric Guillaume

Session Tue1-Ball202 Mutation 1 Tuesday - 8:30 - 9:45 Room Oregon ballroom 202

- 8:30 Mutation load across mating systems: how does load change and how do we best estimate it? Kimberly Gilbert, Aneil Agrawal, Stephen Wright
- 8:45 The effect of ploidy level on the rate and spectrum of new mutations Nathaniel Sharp, Linnea Sandell, Chris James, Sally Otto
- 9:00 Direct estimate suggests remarkable mitochondrial mutation rate variation within the genus Daphnia

Fenner Macrae, Maia Benner, Anna Ritz, Sarah Schaack

- 9:15 The effects of cell-lineage selection on clonal evolution and adaptation in Mimulus guttatus Jaime Schwoch, Mitch Cruzan
- 9:30 Patterns of mutation match physical branching topology in Eucalyptus Adam Orr, Robert Lanfear, Reed Cartwright

Session Tue1-Ball203 SSE Spotlight - Genetic transmission at the population level

Tuesday - 8:30 - 9:45

Room Oregon ballroom 203

- 8:30 From haploid to diploid, from diploid to haploid, complex life cycles are everywhere Stacy Krueger-Hadfield
- 8:45 Population-genetic expectations for trait filtering of self-incompatibility on islands Jeremiah Busch, Nathan Layman
- 8:50 Inbreeding depression and polyploidy as a genetic island Nathan Layman, Jeremiah Busch
- 8:55 How to make an apomict: Asymmetrical crossing success between Boechera species Catherine Rushworth, Tom Mitchell-Olds
- 9:00 Chromosome gone wild: centromeric drive in Mimulus Lila Fishman, Findley Finseth
- 9:15 The evolution of biparental organelle inheritance: a rescue mechanism for cytonuclear incompatibility?

 Karen Barnard-Kubow, Laura Galloway
- 9:20 Attack of the Clones Asymmetrical reproductive interference in the Crepis agamic complex Evan Hersh, Jaime Grimm, Jeannette Whitton
- 9:25 The effects of haploid selection on plant Y chromosome evolution Georgy Sandler
- 9:30 How the mode of genetic transmission impacts introgression Yaniv Brandvain

Session Tue1-Ball204 Mutualism Tuesday - 8:30 - 9:45 Room Oregon ballroom 204

- 8:30 Ecological context influences mutualism abandonment in the mycorrhizal symbiosis Hafiz Maherali, Brad Oberle, Gijsbert Werner, Daniel McGlinn, Kabir Peay, Dylan Schwilk
- 8:45 Population genomics of cryptic species in Caribbean Panama supports habitat specialization by symbiotic sea anemones

 Emily Bellis, Dee Denver
- 9:00 The enemy of my enemy is my friend: nematode infection of non-pollinating fig wasp antagonists and the consequences for the fig-fig wasp pollination mutualism Justin Van Goor, John Nason
- 9:15 The evolution of obligate endosymbiosis across the insects LEE HENRY, Anouk van't Padje, Toby Kiers, Ellers Jacintha
- 9:30 Host-derived resources may support long-term saprophytic survival of dormant rhizobia Katherine Muller, Ford Denison

Session Tue1-C120-122 Plant reproductive biology Tuesday - 8:30 - 9:45 Room C120-122

- 8:30 Genomic journeys of a strawberry sex locus

 Jacob Tennessen, Na Wei, Shannon Straub, Rajanikanth Govindarajulu, Aaron Liston, Tia-Lynn
 Ashman
- 8:45 The ecological consequences of pollen color variation in trout lily (Erythronium americanum) withdrawn 2017-06-25 10:09:41.0

 Emily Austen, Shang-Yao Lin, Jessica Forrest
- 9:00 Environmental influences on the mating system of the common morning glory Diego Alvarado-Serrano, Shu-Mei Chang, Regina Baucom
- 9:15 The effective pollinators of Silene vulgaris do not respond to variation in sex ratio Brian Sanderson, Malcolm Augat, Anita Enriquez, Butch Brodie
- 9:30 Phenotypic selection on floral traits by three bee species Johanne Brunet, Austin Bauer

Session Tue1-C123 Species interactions Tuesday - 8:30 - 9:45 Room C123

- 8:30 Character displacement in evolutionary-novel Anolis lizards James T. Stroud, Kenneth Feeley
- 8:45 Microbes, lagomorphs, & the geographic range limit of a California endemic wildflower John Benning, David Moeller
- 9:00 Does competition between the Dominican native Anolis oculatus and the invasive Anolis cristatellus drive changes in ecological, agonistic and reproductive traits?

 Claire Dufour, Anthony Herrel, Jonathan Losos
- 9:15 Phylogenetic approaches to studying behavioral interference shed light on interspecific territoriality in North American passerines

 Jonathan Drury, Madeline Cowen, Greg Grether
- 9:30 Domestication & Evolution of Eudicot defenses against a generalist fungal pathogen Celine Caseys, Nicole Soltis, Gongjun Shi, Daniel Kliebenstein

Session Tue1-C124 Molecular evolution 1 Tuesday - 8:30 - 9:45 Room C124

- 8:30 Conserved rates of molecular evolution among duplicate gene copies after whole genome duplication

 George Tiley, J. Gordon Burleigh
- 8:45 Nitrogen conservation pervades the transcriptional response to nitrogen starvation in yeast Claudia Acquisti
- 9:00 Population genomics of Anolis carolinensis transposable elements: insertion polymorphisms are abundant but rarely approach fixation

 Robert Ruggiero, Yann Bourgeois*, Stephane Boissinot*

- 9:15 Selective pressures on the fidelity of transcription.

 Jean-Francois Gout, Weiyi Li, Michael Lynch, Marc Vermulst
- 9:30 Strong and widespread action of site-specific positive selection in the snake venom Kunitz/BPTI protein family

 Dusan Kordis

Session ASN Executive council exit meeting Tuesday - 8:30 - 10:15 Room C126

Session SSB Executive council exit meeting Tuesday - 8:30 - 10:15 Room B110

Session Coffee break

Tuesday - 9:45 - 10:15

Room Exhibit Hall A

Session Tue2-A105 Phylogenetic comparative methods 3 Tuesday - 10:15 - 11:30 Room A105

- 10:15 Deriving the appropriate protein substitution matrix to describe the single-stranded DNA virus tree of "life"

 Lele Zhao, Erik Lavington, Siobain Duffy
- 10:30 Macroevolutionary analysis of discrete character evolution using BAMM Michael Grundler, Daniel Rabosky
- 10:45 Some problems with model-fitting and model-selection in the analysis of discrete character evolution using likelihood Klaus Schliep, Liam Revell
- 11:00 The impact of conflicting signal in protein structural classes on phylogenetic inference Akanksha Pandey, Ed Braun
- 11:15 Scalable approximate inference for the phylogenetic threshold model Nathaniel Pope, Eric Goolsby

Session Tue2-A106 Bioinformatics / genomics Tuesday - 10:15 - 11:30 Room A106

- 10:15 Genome assembly strategies for outbred non-model organisms James Lewis, Charles Danko, Robert Reed
- 10:30 Influence of reproductive mode on genome architecture in a freshwater snail Jorge Moreno, Kyle McElroy, Maurine Neiman
- 10:45 Gene duplications associated with phytochemistry in Cannabis sativa Daniela Vergara, Ezra Huscher, Kyle Keepers, Reggie Gaudino, Nolan Kane
- 11:00 The evolution of LTR retrotransposons in the genomes of two heterosporous ferns *Mathew Simenc*, Koppers Nils, Alex Harkess, Aleksandra Berick, Shifeng Cheng, Gane Wong, Brautigam Andrea, Henriette Schluepmann, Kathleen Pryer, Paul Wolf, Fay-Wei Li, Joshua Der
- 11:15 Leveraging transcriptomic data to study regulatory motif evolution in Drosophila species Joel Atallah, Ahmad Karkoutli, Colin Shew, Susan Lott

Session Tue2-A107-109 Comparative biology 2 Tuesday - 10:15 - 11:30 Room A107-109

- 10:15 Physiological and transcriptional responses to drought stress and recovery in grasses Fan Qiu, Seton Bachle, Ryan Estes, Jesse Nippert, Mark Ungerer
- 10:30 Reinventing the limpet: Shell patterning and experimental morphology of a slipper-shell snail Darcy Ross
- 10:45 Placental invasiveness and Eutherian mammal diversification
 Andrew Furness, Mark Springer, William Freyman, Sebastian Hoehna
- 11:00 An evolutionary framework for dissecting and understanding regenerative intestine growth in snakes

Blair Perry, Audra Andrew, Daren Card, Drew Schield, Giulia Irene Maria Pasquesi, Abu Hena Mostafa Kamal, Robert Ruggiero, Saiful Chowdhury, Stephen Secor, Todd Castoe

11:15 The genetic factors of bilaterian evolution.

Peter Heger, Wen Zheng, Maria Thieser, Anna Schlizio, Thomas Wiehe

Session Tue2-B111-112 Microbes 2 Tuesday - 10:15 - 11:30 Room B111-112

- 10:15 Avian influenza virus and the cloacal microbiome of wild avian hosts Sarah Hird
- 10:30 The bite of the finch: Gut microbiome sequencing of Darwin's Finches reflects divergence of vampire finches

Jaime Chaves, Lewis Ward, Alice Michel, Kiyoko Gotanda, John McCormack, Shana Goffredi, J. Albert C. Uy, Daniel Baldassarre

10:45 Determining the impact of metabolic role in symbiont genome decay in Adelgidae (Insecta: Hemiptera)

Kathryn Weglarz, Gaelen Burke, Carol von Dohlen

- 11:00 Examining the role of maternal microbes in the insect immune response: Lycaeides melissa as a model for wild immunology

 Su'ad Yoon, Josh Harrison, Matthew Forister, Angela Smilanich
- 11:15 Bacterial communities associated with butterflies and their effect on butterfly fitness Kruttika Phalnikar, Deepa Agashe, Krushnamegh Kunte

Session Tue2-B113 Adaptation / life history Tuesday - 10:15 - 11:30 Room B113

- 10:15 Effects of diadromous life histories on genetic diversity and population structure in fishes. Joel Corush, Evin Carter, Ben Fitzpatrick
- 10:30 Predation risk, body condition and immunity: Life history-related resource allocation in a dragonfly
 Rosalind Murray, Locke Rowe, Shannon McCauley
- 10:45 Updated Phylogeny and Ancestral State Reconstruction of Brooding in Crinoids Kathleen Torrence, Valeska Stamatti, Scott Steppan
- 11:00 Evolution of emergence strategies in a dynamic climate Collin Edwards, Louie Yang
- 11:15 Collective dispersal leads to variance in fitness and maintains offspring size variation Scott Burgess, Robin Snyder

Session Tue2-B114-115 Molecular ecology Tuesday - 10:15 - 11:30 Room B114-115

10:15 The evolutionary history of Madagascar's dwarf lemurs

Rachel Williams, Marina Blanco, Sheena Faherty, C. Ryan Campbell, Kelsie Hunnicutt, Peter Larsen, Anne Yoder

- 10:30 Population genomics of lodgepole and jack pines assessed with >20k SNPs Rhiannon Peery, Catherine Cullingham, Kate St. Onge, Janice Cooke, David Coltman
- 10:45 Genomic insights into the evolutionary history of the taxonomically controversial Northwestern Crow (Corvus caurinus)

 David Slager, Kevin Epperly, Renee Ha, Sievert Rohwer, Caroline Van Hemert, John Klicka

- 11:00 It's about time: Insights into temporal genetic patterns from biodiversity indices Matthew Iacchei, Elizabeth Butcher, Elan Portner, Erica Goetze
- 11:15 Phycodnavirus phylodynamic diversity across Pacific Northwest Cascadian lakes Michael Zanis, Carolyn Stenbak, Lindsay Whitlow

Session Tue2-B116 Gene flow / hybridization Tuesday - 10:15 - 11:30 Room B116

- 10:15 Genomic insights to diversification and speciation using hybrid zones of widespread Australian geckos (Heteronotia binoei)

 James Titus-McQuillan
- 10:30 The effects of admixture on deleterious genetic variation Bernard Kim, Jacqueline Robinson, Kirk Lohmueller
- 10:45 Detecting hybridization using phylogenetic invariants Paul Blischak, Julia Chifman, Andrea Wolfe, Laura Kubatko
- 11:00 Revisiting a wild perennial sunflower hybrid swarm

 Adam Herman, Jill Ekar, Mikey Kantar, Robert Stupar, Yaniv Brandvain
- 11:15 Divergence and introgression between closely-related spiny lizard (Sceloporus) species

 Shea Lambert

Session Tue2-B117-119 Adaptation / evolutionary ecology 2 Tuesday - 10:15 - 11:30 Room B117-119

- 10:15 Peeing Poison: The biochemistry of bamboo lemur cyanide survival M. Elise Lauterbur, Lydia Tongasoa, Jeremy Peralta, Marta Concheiro-Guisan
- 10:30 From sex pheromones to nestmate recognition: Deciphering evolutionarily conserved signaling properties of cuticular hydrocarbons

 Jan Buellesbach, Thomas Schmitt, Jürgen Gadau, Neil Tsutsui
- 10:45 A widespread salamander does not show predicted morph frequency shifts despite 40 years of climate change

 Annette Evans, Brenna Forester, Elizabeth Jockusch, Mark Urban
- 11:00 Response to drought in an assemblage of stream fishes: general predictors or species-specific responses?

 Mark Belk, Richard Simkins, R. Cary Tuckfield
- 11:15 Acclimation, body size, and thermal tolerance in a freshwater invertebrate Brooks Miner, Gabriella Ampem-Darko, Kathryn Dubyk

Session Tue2-Ball202 Mutation 2 Tuesday - 10:15 - 11:30 Room Oregon ballroom 202

- 10:15 Copy-Number Changes Under Varying Intensities of Selection During Mutation Accumulation and Fitness Recovery in C. elegans Experimental Evolution Lines

 Vaishali Katju, Anke Konrad, Ulfar Bergthorsson
- 10:30 rDNA Copy Number Loss in Mutation Accumulation Lines of Daphnia pulex Eleanor Harvey, Melania Cristescu, Teresa Crease
- 10:45 The Convergent Cryptosporidium parvum Lactate Dehydrogenase: a blacksheep in the Apicomplexa family Jacob Wirth
- 11:00 Evolution of the substitution gradient in ratsnake mt genomes Carolyn Sy, Frank Burbrink , Marcelo Gehara
- 11:15 Evaluation of epistatic interactions reveals possible functional constraints in the evolution of TTX-resistant sodium channels in garter snakes.

 Gabriela Toledo, Butch Brodie, Charles Hanifin, Shana Geffeney

Session Tue2-Ball203 SSE Spotlight - Sex in the wild

Tuesday - 10:15 - 11:30

Room Oregon ballroom 203

10:15 Keeping up with pathogens for fifty million years without sex: ecological and evolutionary insights from bdelloid rotifers

Christopher Wilson, Antonios Kriezis, Joshua Potter, Timothy Penny, Cuong Tang, Sophie Gresham, Nicholas Tew, John Paterson, Tabassum Mujtaba, Isobel Eyres, Reuben Nowell, Constantinos Xenophontos, Tim Barraclough

10:30 Secondary contact and rare sex between distant lineages of the amphibian-killing chytrid fungus

Thomas Jenkinson, Kelly Zamudio, Felipe Toledo, Timothy James

- 10:35 Sexual conflict and the maintenance of sex in facultative parthenogens Mercedes Burns
- 10:40 Sperm dependence limits niche expansion in unisexual salamanders of varying ploidy Robert Denton, Lisle Gibbs, James Bogart
- 10:45 Ecological differences and reproductive interactions between sexuals and their related apomicts in the sunflower family Jeannette Whitton
- 11:00 Macroevolutionary dynamics of flowering plant sexual systems

 Emma Goldberg, Tia-Lynn Ashman, Itay Mayrose, Ray Ming, Sally Otto, Niv Sabath, Jana Vamosi
- 11:05 Mutation accumulation in an asexual plant John Lovell, Robert Williamson, Stephen Wright, Tim Sharbel
- 11:10 Balancing selection maintains sex/asex polymorphism in natural plant populations Catherine Rushworth, Tom Mitchell-Olds
- 11:15 Doing research in the wild: Why does it matter? Stephanie Meirmans

Session Tue2-Ball204 Mutualism / evolutionary ecology Tuesday - 10:15 - 11:30 Room Oregon ballroom 204

- 10:15 Context dependence in a beneficial nematode-bacterium interaction Kim Hoang, Nicole Gerardo, Levi Morran
- 10:30 Evolution of a Microbial Symbiont While Outside its Native Animal Host Diminishes the Negative Correlation Between Two Traits, Promotes Host Range Expansion, and Improves Host Colonization

 WILLIAM SOTO, Michael Travisano, Michael Nishiguchi
- 10:45 Select and re-sequence reveals fitness tradeoffs and loci underlying symbiotic and free-living fitness in a legume-rhizobium mutualism

 Liana Burghardt, Brendan Epstein, Joseph Guhlin, Matt Nelson, Margaret Taylor, Nevin Young, Michael Sadowsky, Peter Tiffin
- 11:00 Natural selection in a declining mutualism: How nitrogen addition causes the evolution of less cooperative rhizobia.

 Jennifer Lau, Dylan Weese, Katy Heath
- 11:15 Molecular dynamics of a nascent microbial mutualism Yee Mey Seah, Kristina Hillesland

Session Tue2-C120-122 Pollination Tuesday - 10:15 - 11:30 Room C120-122

10:15 The early flower attracts the bee: consequences and mechanisms of phenological isolation in plants

Jennifer Ison, Leah Prescott, Amy Waananen, Scott Nordstrom, Stuart Wagenius

- 10:30 The influence of distinct pollinators on male and female reproductive success in alfalfa. Emmanuel Santa-Martinez, Johanne Brunet
- 10:45 Drought effects on floral traits and their heritabilities in Phlox drummondii Sevan Suni, Ben Ainsworth, Robin Hopkins
- 11:00 The adaptive function of heterostyly investigated with quantum dots Marinus de Jager, Corneile Minnaar, Bruce Anderson
- 11:15 A meta-analysis of the causes of natural selection on floral traits Christina Caruso, Katherine Eisen, Ryan Martin, Nina Sletvold

Session Tue2-C123 Geographic variation Tuesday - 10:15 - 11:30 Room C123

- 10:15 Selection across a light level spectrum explains the maintenance and spatial structure of colour morphs in a polymorphic bird

 Jacqueline Bishop
- 10:30 Evolution in continuous space using SLiM 2 models Benjamin Haller, Philipp Messer
- 10:45 Patterns of clinal variation in populations of Drosophila simulans from North and Central America

Alisa Sedghifar, Julie Cridland, Perot Saelao, David Begun

- 11:00 Macroecology of benthic hydroids (Cnidaria) in the deep Atlantic Ocean Marina Fernandez, Antonio Marques
- 11:15 Species' range shifts drive the evolution of spatial difference in phenotypic plasticity and environmental tolerance

 Max Schmid, Ramon Dallo, Frédéric Guillaume

Session Tue2-C124 Molecular evolution 2 Tuesday - 10:15 - 11:30 Room C124

- 10:15 Using recursive dynamic Markov clustering (RD-MCL) for fine-grained classification of orthogroups

 Stephen Bond, Andy Baxevanis
- 10:30 Co-option of an ancient metabolic sensor in the evolution of polyphenism regulation in nematodes

 Joana Projecto Garcia, Erik Ragsdale
- 10:45 New molecular clocks for Arctic and tropical marine invertebrates

 Tzitziki Loeza-Quintana, Yash Bhatt, Christina Carr, Tooba Khan, Samantha Lyon, Sarah J.

 Adamowicz
- 11:00 Molecular evolution of voltage-gated sodium channels in herons and relatives (Ardeidae)

 Angela Hornsby, John Abramyan, Joel McGlothlin
- 11:15 Asian Ranid species possessing gastromyzophorous tadpoles (Amphibia, Anura): their phylogeny and biodiversity

 Umilaela Arifin, Stefan Hertwig, Eric Smith, Djoko Iskandar, Alexander Haas

Session Lunch

Tuesday - 11:30 - 13:30

Room On the town

Session Tue3-Ball202 Young Investigator / Dobzhansky Awards symposium 1 Tuesday - 13:30 - 14:45 Room Oregon ballroom 202-204

ASN Jasper Loftus-Hills Young Investigators Award

The Jasper Loftus-Hills Young Investigators Award was established in 1984 to recognize outstanding and promising work by investigators who received their doctorates in the three years preceding the application deadline or who are in their final year of graduate school. Jasper Loftus-Hills (1946-1974) was an Australian biologist of exceptional promise, whose career was cut short just three years after receiving his degree when he was killed by a hit-and-run driver while tape-recording frog calls along a Texas highway. The award includes presentation of a research paper at the annual meeting of the American Society of Naturalists, an award of \$500, a travel allowance of \$700, the cost of registration at the meetings, and a supplement of \$500 in case of intercontinental travel.

The 2017 winners are: Sarah Fitzpatrick, Anna Hargreaves, Martha Muñoz, and Alison Wright.

13:30 Young Investigator Prize announcement Kathleen Donohue

- 13:45 Linking evolution & demography through genetic rescue of small populations Sarah Fitzpatrick
- 14:15 Dynamics at the edge: species interactions vs. species distributions

 Anna Hargreaves

Session Coffee break

Tuesday - 14:45 - 15:15

Room Exhibit Hall A

Session Tue4-Ball202 Young Investigator / Dobzhansky Awards symposium 2 Tuesday - 15:15 - 16:45 Room Oregon ballroom 202-204

ASN Jasper Loftus-Hills Young Investigators Award

The Jasper Loftus-Hills Young Investigators Award was established in 1984 to recognize outstanding and promising work by investigators who received their doctorates in the three years preceding the application deadline or who are in their final year of graduate school. Jasper Loftus-Hills (1946-1974) was an Australian biologist of exceptional promise, whose career was cut short just three years after receiving his degree when he was killed by a hit-and-run driver while tape-recording frog calls along a Texas highway. The award includes presentation of a research paper at the annual meeting of the American Society of Naturalists, an award of \$500, a travel allowance of \$700, the cost of registration at the meetings, and a supplement of \$500 in case of intercontinental travel.

The 2017 winners are: Sarah Fitzpatrick, Anna Hargreaves, Martha Muñoz, and Alison Wright.

SSE Theodosius Dobzhansky Prize

The Theodosius Dobzhansky Prize is awarded annually by the Society for the Study of Evolution to recognize the accomplishments and future promise of an outstanding young evolutionary biologist. The prize was established in memory of Professor Dobzhansky by his friends and colleagues, and reflects his lifelong commitment to fostering the research careers of young scientists.

The 2017 winner is Molly Schumer, Harvard University.

- 15:15 Janzen's Hypothesis Meets the Bogert Effect: A Synthesis Nearly 100 Years in the Making Martha Munoz
- 15:45 Routes to the resolution of sexual conflict Alison Wright
- 16:15 Dobzhansky prize announcement Cynthia Weinig
- 16:25 Hybridization shapes the evolution of genomes and species: insights from swordtail fish *Molly Schumer*

Session Super Social

Tuesday - 18:30 - 23:59

Room Oregon Zoo

The conference-ending Super Social will be held at an indoor/outdoor venue at the <u>Oregon Zoo</u> starting at 6:30 pm. Dinner is provided and a DJ for dancing will be available later in the evening. The zoo is located at 4001 SW Canyon Rd and is accessible via the MAX light rail using your free conference pass. From the Convention Center, take the Blue or Red line westbound and get off at Washington Park Station (the only one that is underground). Take the elevator at the east end of the platform. The train ride takes approximately 25 min.

Advance online ticket purchase is required for the Super Social. Tickets will NOT be available for purchase during the conference or at the zoo. Lost tickets will not be replaced. Your ticket will also get you entrance to the zoo any time during the day on Tuesday, should you want to visit earlier.

Session American Institute of Biological Sciences workshop:
Informing and Engaging Decision-makers
Wednesday - 9:00 - 17:00
Room B110

The is an optional post-conference event. Prior application was required (during conference registration). Lunch is 'own your own' and is not included in registration cost. Sponsor: SSE

Organizer: Janette Boughman (Michigan State U.)

Topic: Rarely has the need for effective and influential communication about science been more important than it is today. Politicians and political interests are redefining and reinterpreting science—with great persistence and impact. The majority of the public still respect scientists and value science, but they often find it challenging to discern who and what is legitimate.

Simply adding to the noise will not provide individuals with the understanding they need to inform deliberations and decisions. Scientists must become more skilled at communicating with policymakers in impactful ways.

The American Institute of Biological Sciences (AIBS) will deliver this program. AIBS has a long and successful track record of engaging, informing, and influencing the public and science policy decision-makers. Audiences AIBS routinely work with include members of Congress, federal agency heads and program managers, state officials, and university administrators. AIBS has played important roles, for example, in helping to defeat state-level initiatives to introduce creationism into state curricula. AIBS has also been a leader in national efforts that have led to new federal investments in biology.

Participants will learn:

- How to tell a resonant story that informs decision-makers
- How to prepare for and engage in a meeting with a decision-maker
- How to protect your scientific reputation
- How to identify and define the audience you need to reach
- What policymakers want and need to know from a scientist