



### **University of Groningen**

### Winter moth adaptation to climate change

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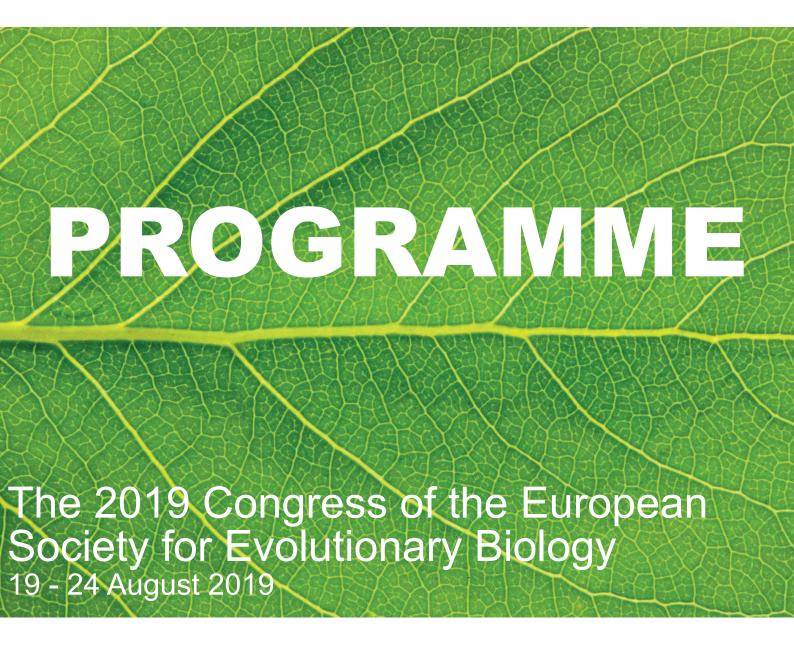
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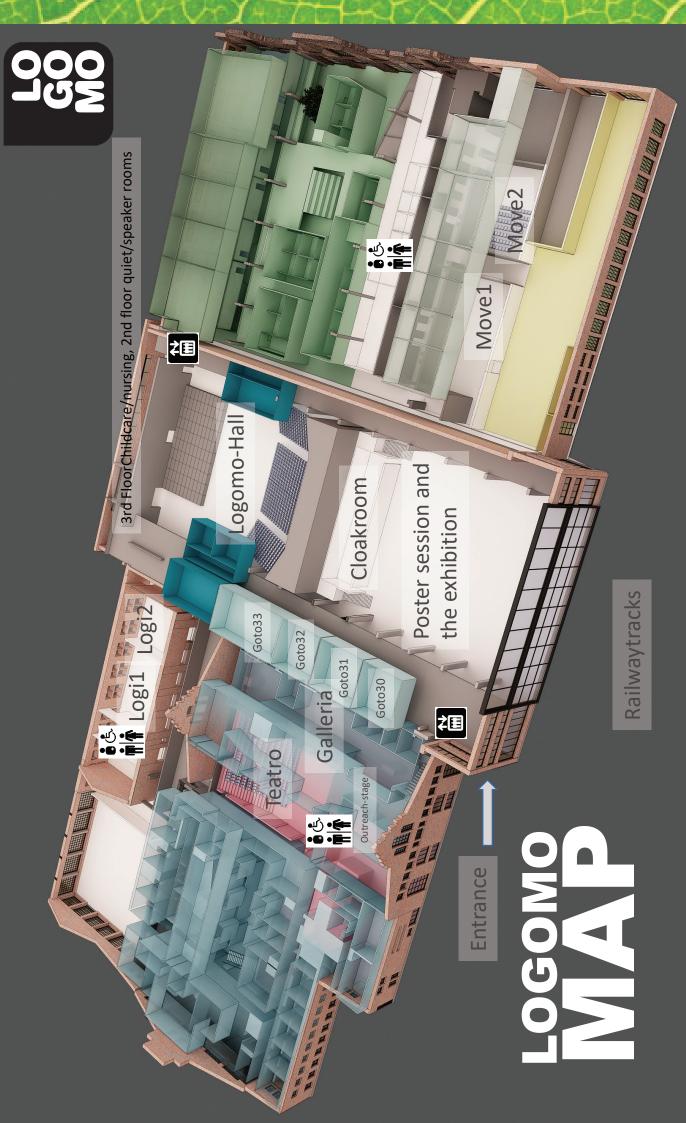












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### LCOME

Dear colleagues,

It is a pleasure to welcome you to the Logomo entertainment centre in Turku, Finland, for the 17th Congress of the European Society for Evolutionary Biology! More than 1300 people have registered for the conference, of which 543 researchers are presenting their research in one of the oral sessions, and an additional 570 will present posters.

The five-day programme follows the traditional ESEB format with 35 themed symposia proposed by members of the ESEB community. As a new innovation, we have classified abstracts submitted to the Open Symposium into five broad sub-themes (36a. Sexual selection and reproductive strategies; 36b. Phylogeography, biogeography, Speciation, systematics; 36c. Species interactions; 36d. Genome Evolution; 36e. Phenotypic Evolution) that we hope will make it easier for delegates to identify sessions and talks matching their interests.

In line with ESEB values, we have also placed emphasis on achieving gender balance amongst speakers and ensuring the environmental impact of the meeting is minimized. 53.7% of all speakers, and 57.7% of invited speakers who reported their gender at registration are female (compared to 52.9% of all abstract submitters). To reduce environmental impact, we have minimized the use of plastic throughout the conference (meals and coffee are served with reusable crockery and cutlery, no plastic cups are on offer for water, no plastic cover for badges etc.) and have focused on sourcing food with a low carbon footprint (e.g. 2 days with vegetarian only, 2 days with locally sourced fish). The conference t-shirt and bag are also made from recycled cotton.

We have also arranged active outreach and social programs. Outreach includes a school children mini-conference that will bring over 40 high school students to Logomo on Thursday morning, while on the social side, the conference pub (Koulu a.k.a "The Old School") will be open until 2am each day (kitchen open until 23:30). There is a large area upstairs reserved for congress participants where you can taste two of their craft beers that have been re-branded for the congress (Beer Reviewed and Drinkage Disequilibrium).

Finally, we would like to thank the 39 volunteers and the exhibitors and sponsors for helping to make this event possible, and we wish you all a productive, supportive and enjoyable meeting!

On behalf of the organizing and scientific committees

**Craig Primmer** (main organizer)

## PRERENC

### **CONGRESS VENUE**

The congress venue, Logomo, is a centre for culture, arts and creative economy operating from the defunct locomotive workshop near the main railway station, the bus station and close to Turku city centre. A pedestrian bridge provides access from the main railway station to Logomo. Walking from the Market Square (which unfortunately is currently under construction) to Logomo takes about 20 minutes (1.5 km). All session rooms are located on the ground floor, except for GOTO rooms, which are located on the third floor, accessible by elevator and Logomo Hall stairs.

### **REGISTRATION AND INFORMATION**

The registration & information desk will be open on Monday, August 19, from 15:00 until 20:00 on the entrance hall, and during the scientific programme on the following congress days, from 7:30 (Tue) and from 8:00 (Wed-Sat). Participant registration fees include attendance at the scientific sessions, coffee and lunch on the congress days as marked on the programme, congress bag, congress app, access to live stream of all sessions, as well as the Welcome reception on 19 August. Last minute registration is possible at the information desk.

### **LUNCHES AND COFFEE**

A cooked lunch on each day is included as a part of the congress registration fee. Coffee and tea are available all day (starting 30 mins before the first session commences), while additional snacks are served during the morning and afternoon breaks (except Saturday afternoon). To reduce the environmental impact of the meals we will serve vegetarian only options on two days, and two days will have locally sourced abundant fish as the meat source. All meals and drinks are served with reusable crockery and cutlery. For the refill water stations, we recommend bringing your own bottle. There is also a restaurant and a bar located in Logomo. Restaurant Kitchen is open daily 9:00-16:00 and Teatro bar is open on Monday 16:00-21:00, and on following days from 12:30 onwards.

### **CONGRESS STAFF & VOLUNTEERS**

Besides registration and info desk staff, there are congress volunteers who can be identified by their white congress T-shirts at the congress venue. Feel free to ask them for any assistance you may need.

### QUIET/REFLECTION ROOM

There is a quiet/reflection room available on the second floor (indicated with signs).

### SPEAKERS' PRACTICE ROOM

There is a speakers' room on the second floor (indicated with signs) where speakers can practice their talks. If you need to borrow a computer to practice in the room, please ask the info desk. There is a time reservation sheet on the room's door.

### **CHILDCARE & NURSING ROOM**

Childcare will be organized at the 3rd floor "backstage" rooms (take an elevator at the far end of the long hallway on the right side of the Logomo hall). Opening hours are Tue 8:30-17:45, Wed 8:45-17:45, Thu 8:45-13:00, Fri 8:45-17:45 and Sat. 8:45-16:15.

There is also a room for nursing or bottle feeding your child available on the second floor (indicated with signs). The room includes a shower and washbasin, a kitchen sink, refrigerator and microwave. A sign for indicating that the room is in use will be available if you wish to feed in private.

### **PRESENTATIONS**

Regular oral presentations will be 10 min long 3 additional minutes for discussions and then 2 minutes for changing rooms. Invited symposiums presentations will be 23 min long 5 minutes for discussion and 2 minutes for changing rooms. Presenters must follow the assigned times to ensure the eight concurrent sessions remain synchronised. Loud music will play during the 2 min transition to ensure speakers cannot speak overtime. The next speaker's timeslot starts as soon as the music stops. To enable the staff to handle the technical aspects in an efficient way, all presentations must be prepared according to the guidelines listed on the congress website.

### **POSTER SESSIONS**

There will be two poster sessions during ESEB2019 on two evenings (17:20-19:20), after the parallel symposia: Poster Session 1 on Tuesday, 20 August and Poster Session 2 on Friday, 23 August. Presenters can check their poster session in the programme on the congress website. Poster boards will be marked with poster codes. Poster presenters are required to be by their poster for at least one hour during the poster session designated to their poster. Poster presenters also have the opportunity to invite up to 3 attendees of their choice to visit their poster through the Postvites system. Poster presenters will serve wine to their poster visitors.

### **POSTER PRIZES**

There will be prizes awarded for the best student poster in both poster session one and poster session two (by public vote), as well as a "jury's choice" poster prize chosen by members of the scientific committee that recognizes a student poster (across both poster sessions) that honestly and clearly presents complicated/non-significant/counter-intuitive results. The prizes will be announced at the closing ceremony.

### **EVOKE OUTREACH STAGE**

The EvoKE team is arranging a series of events to get evolutionary biology researchers more involved in outreach. These events will mostly be held during lunch breaks on the EvoKE outreach stage in the Teatro café and bar.

The events are listed under "Satellite events".

The EvoKE (Evolutionary Knowledge for Everyone) network is funded by ESEB. EvoKE seeks to contribute to a world where people understand evolution and can use scientific knowledge and skills to make informed decisions that address societal problems thereby contributing to an inclusive, sustainable and resilient future. See https://evokeproject.org/ for more details

### **STREAMING**

In order to encourage open science, and to allow people not able to attend ESEB2019 to have access to presentations, we are providing presenters the opportunity for their presentations to be live streamed and also available for viewing following the conference. All presenters are asked if they do not wish to give permission to allow the streaming and recording of their talk on the presentation upload form. Only those who have given the permission will be streamed/recorded.

### WIFI

In Logomo, the wifi network is LogomoPublic and password loGOmo2012.

### **CONGRESS APP**

Aboa Events Congress app is available for the ESEB2019 congress participants and it is free to download from Google Play and App Store. The Aboa Events app contains features such as abstracts, programme, information about the transportation service Föli, notifications about the possible updates in the programme, maps, venue information and other useful features.

### **BUS (FÖLI) PASS**

Delegates will receive complimentary weekly bus passes (QR-codes in the name badges) with FÖLI public transportation from Saturday 17th until Sunday 25th of August. FÖLI Turku region traffic allows you to use local bus services in the city of Turku, without limitation (www.foli.fi).

### **NAME BADGE**

Entrance to the congress venue and upstairs at the conference pub requires wearing your name badge. The conference dinner ticket, if you have purchased one, is also on your name badge, as well as FÖLI pass (QR code).

### **DELEGATE BAGS**

Delegate bags are made in the Turku Work Centre as a part of rehabilitative services for the special needs unemployed. Bags are made of waste fabric and other recycled materials, they all come in different colours and patterns. This new project is inspired by ESEB 2019, which it is the first congress to receive these bags. Take one if you like from the City of Turku stand in the lobby. Also city maps are available.

The Turku Work Centre will also have a pop-up shop in the lobby on Friday, 23 August at 12:00-16:30. They sell lovely handmade, local, Scandinavian style products, toys, wool socks, small purses and bags.

### **PRINTING**

There is no printing possibility at the venue of the congress. Your hotel may have a business centre where you can print. Closest printing place Niini, address Laivurinkatu 1, 20810 Turku, open 8:00-17:00 on weekdays.

### **FIRST AID**

If you need first aid, please contact any staff member or volunteer and you will be directed to first aid room.

### **ELECTRICITY**

There are sockets in the "street area" (furnished with tables and chairs) by the entrance hall. The Voltage: 220-240 Volts. Electrical sockets (outlets) in Finland are one of the two European standard electrical socket types: "Type C" Europlug and "Type E/F" Schuko.

### **CITY OF TURKU**

You can find useful information and get inspired about the city of Turku by visiting the congress website, www.visitturku.fi/en or the congress app Aboa Events.

### **ENVIRONMENTAL IMPACT**

In line with ESEB values, we have placed emphasis on ensuring the environmental impact of the meeting is as low as possible. For example, we provided details of strategies for reaching Turku without flying, as well as options for compensating flight carbon footprints. The City of Turku has also provided all delegates with a weekly bus pass free of charge for moving around the city.

Further, we have minimized the use of plastic throughout the conference (e.g. meals and coffee are served with reusable crockery and cutlery, no plastic cups are on offer for water, no plastic cover for badges etc.) and have focused on sourcing food with a low carbon footprint. There will be two vegetarian lunches and two lunches with locally sourced fish. Berries and herbs in salads come direct from the Finnish nature. The conference t-shirt and optional conference bag are also made from recycled cotton, and are sourced from local companies.

Finland offers the best quality water straight from a tap, so there is no need for bottled water. You can fill your own mug or bottle making use of the water filling stations in Logomo. Finland uses a deposit-based efficient return system for beverage bottles and cans, so do not throw them into trash, but return them to a store and get money back. Logomo also provides recycling centers for other waste.

### **DIVERSITY OF PRESENTATIONS**

In line with ESEB values, we have also placed emphasis on promoting diversity in gender, career stage and nationalities amongst speakers. 54% of all speakers, and 58% of invited speakers who reported their gender at registration are female (compared to 53% of all abstract submitters). ECR and mid-career scientists make up 84% of symposium organisers and 60% of invited speakers. 28 nationalities are represented amongst the symposium organisers and 19 amongst the 75 invited speakers.

### **SOCIAL MEDIA POLICY**

ESEB supports open communication of science. Therefore, in addition to offering a live streaming opportunity for all oral presentations, the default assumption is that information presented at the congress (in oral or poster format) may be reported and discussed, and images of slides posted, by attendees in social media and blogs unless presenters specifically state otherwise. If a presenter does not want information from their presentation to be broadcast and/or photographed they should make this clear in their talk/poster, for example by including one or both of the following images.





We expect delegates to respect the rights of presenters. Any clear breaches of this policy should be reported to the congress desk.

### CODE OF CONDUCT

The ESEB Congress is intended to foster the exchange of scientific ideas, providing participants with an opportunity to network with an international community of evolutionary biologists. ESEB is 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 will not be tolerated. Unacceptable behaviour includes (but is not limited to) unwanted verbal attention, unwanted touching, intimidation, stalking, shaming, or bullying. Blatant 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 constitutes unacceptable behaviour. Retaliation for reporting harassment is also unacceptable, as is reporting an incident in bad faith.

Please note that the use of certain language or images in oral or poster presentations may contravene the Code of Conduct if they represent disrespectful criticism of individuals or teams rather than valid criticism of their science, if they are seen to objectify or demean individuals or groups. It is important to recognise that sensitivity to such aspects of communication varies, and what might be acceptable or humorous to some people might not be to others.

The meeting organizers 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, and/or (c) prohibited from attending future meetings. Establishing this Code of Conduct is intended to maintain the high quality of scientific discourse that members have come to expect from our meetings.

If you experience any form of inappropriate behaviour, you may wish to contact and speak with an experienced external Human Relations counsellor that ESEB has contracted to help in such situations. You may also speak with the ESEB Office Manager, Dr Ute Moniatte, who can liaise with the external counsellor on your behalf. Either way, all communication will be held in strict confidence.

If you contact our counsellor, you will be asked the following:

- to give your name
- to describe the events or behaviour that took place, and any other relevant circumstances surrounding the incident
- if relevant or appropriate, to identify the perpetra-
- if relevant or appropriate, to identify any witnesses.

### **Important**

Nothing will be undertaken without your consent, nor will your name be communicated to anyone without your consent.

Our external HR Adviser is Joanne Harding, at Workforce Window Ltd, a Human Resources company based in the UK with many years' experience in dealing with individual complaints and breaches of codes of conduct. Joanne will handle your issues both sensitively and confidentially.

The Workforce Window website is: www.workforcewindowltd.co.uk

To contact Joanne Harding, either send her an email (joanne@workforcewindow.co.uk) or phone/text her (+44 792 009 46 63).

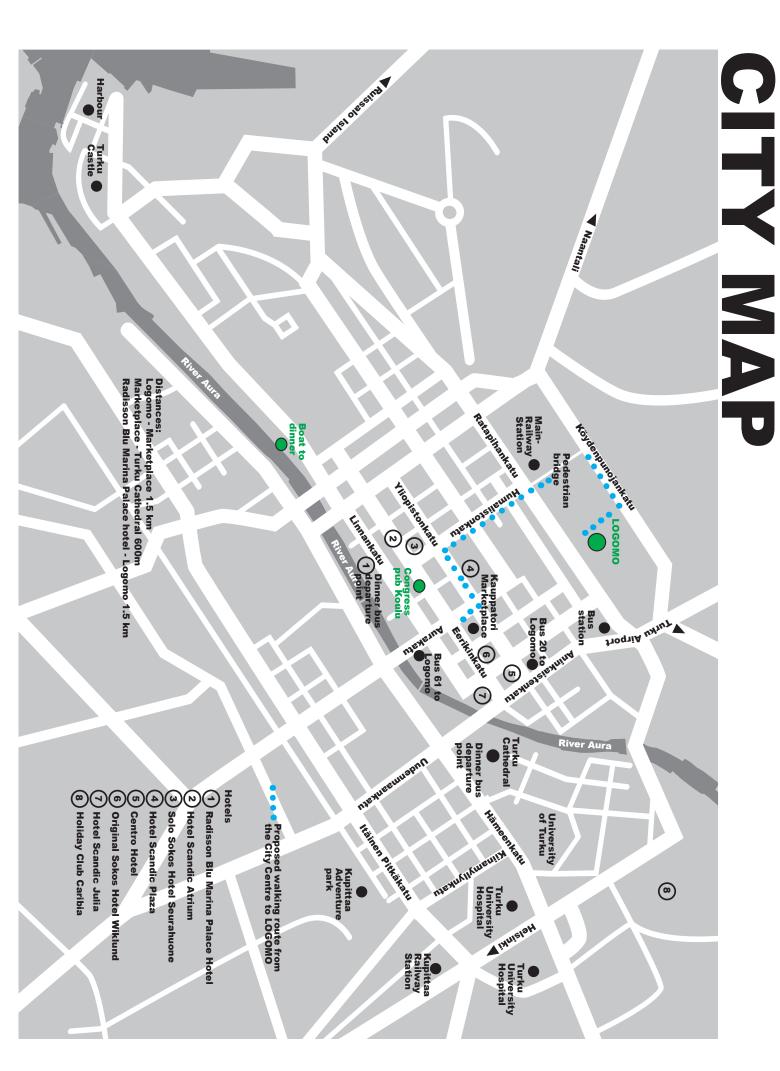
To contact Ute Moniatte, either send her an email (office@eseb.org) or phone/text her (+49 160 524 3050).

Workforce Window Ltd follow the General Data Protection Regulations and are registered with the Information Commissioners Office. The company has no other links with ESEB.

### **EXHIBITION**

Exhbition is open throughout the congress in the entrance hall. List of exhibitors:

- 1. Peer Community In
- 2. Guarant International
- 3. Frontiers
- 4. Wiley
- 5. Oxford University Press
- 6. Royal Society Publishing
- 7. Cambridge University Press
- 8. Current Zoology
- 9. The New Phytologist Trust
- 10. Nordic Society Oikos
- 11. University of Helsinki / HiLIFE
- 12. EvoKE Outreach activities
- 13. ESEB organisation





PAT MONAGHAN

(Glasgow)

Title: Bad beginnings and untimely ends: environments, telomeres and life history variation

Tuesday, August 20

09.00-09.45 Room: LOGOMO HALL

Pat Monaghan is an evolutionary ecologist, based at the University of Glasgow, where she holds the Regius Chair in Zoology. She did her PhD at Durham University on seabird ecology, followed by work on the interactions between seabirds and fisheries management. She then began research on the effect of early life conditions in shaping individual life histories, involving studies at many different biological levels from physiology and molecular biology to ecology and behavioural biology. Her work is mainly on birds, with related work in other taxa. A current major focus is on telomere dynamics, and the extent to which this system of genome protection influences life history evolution and ageing patterns.



SINEAD COLLINS

(Edinburgh)

Title: Understanding evolution in life-giving slime

Wednesday, August 21

09.05-09.50 Room: LOGOMO HALL

Sinead Collins is a Reader at the Institute of Evolutionary Biology at the University of Edinburgh. Her research focuses on building the theory needed to predict trait evolution in large populations of photosynthetic microbes, particularly those in the ocean. To do this, she and her group use microbial experimental evolution, make models, and collaborate closely with marine microbiologists and oceanographers. Experimental evolution is a field that rarely considers marine systems, and Sinead has spent much of the past decade working with others to create a field of "marine microbial experimental evolution" that pulls together the strengths of marine and evolutionary biologists.



**DAVID QUELLER** 

(Washington)

Title: Evolutionary conflict and molecular arms races in cooperative systems

Thursday, August 22

09.05-09.50 Room: LOGOMO HALL

David Queller is a professor at Washington University in St. Louis. His dissertation investigated sexual selection and kin selection in plants. He subsequently worked for many years, together with Joan Strassmann, on social insects, showing the importance of relatedness in both cooperation and conflict. They later switched to studying social amoebas, especially the evolution of cheating in and its control by high relatedness, kin discrimination, pleiotropy, and resistance. His theoretical work includes methods for estimating relatedness, models of kin selection and other social forces, the evolution of eusociality via demographic advantages, evolutionary conflict, and fundamental theorems of natural selection.



ANNA-LIISA LAINE

(University of Zurich, Helsinki University)

Title: What keeps pathogens in check in the wild?

Friday, August 23

09.05-09.50 Room: LOGOMO HALL

Anna-Liisa Laine is an evolutionary ecologist who is broadly interested in the eco-evolutionary feedback loops that drive species interactions. She is a professor of ecology at the University of Zurich, and a visiting professor at the University of Helsinki. She received her PhD at the University of Helsinki in 2005 and continued to do post doctoral research at the University of California, Santa Cruz, and CSIRO Canberra. Much of her work is focused on uncovering the mechanism that enable coexistence of host and their parasites in natural populations, and the mechanism which maintain diversity in host-parasite interaction traits. Currently, her work is addressing these questions within a community ecology framework.

### RASMUS NIELSEN

(UC Berkley/Copenhagen)

Title: Human adaptation in time and space Saturday, August 24 09.05-09.50 Room: LOGOMO HALL



Dr. Nielsen's work is on statistical and population genetic analyses of genomic data, in particular methods for detecting natural selection, describing population genetic variation, inferring demography, and methods for association mapping. Much of his current research concerns statistical analysis of next-generation sequencing data, both in the context of medical genetics and population genetics. Many of the methods he has developed are heavily used by other researchers, including the phylogeny based methods for detecting positive selection implemented in PAML, the methods for inferring demographic histories implemented in the IM and IMa programs, the method for detecting selective sweeps implemented in the SweepFinder programs, and the methods for analysing Next Generation Sequencing (NGS) data implemented in ANGSD.

# SOCIAL

### **WELCOME RECEPTION**

Monday, 19 August, 18:00-21:00

Join your colleagues for a welcome reception at the congress venue Logomo on Monday evening, at 18:00.

Finger is food served until 19:30. The event includes 2 complimentary drinks, one offered by the City of Turku. Additional drinks available for purchase until 21 after which we will move to the conference pub (Koulu).

### **CONGRESS DINNER AT MOOMIN WORLD**

Saturday, 24 August, 18:30-02:00

The congress will be concluded with the conference dinner held in Moomin World. This children's culture classic operates in Naantali, where the Archipelago Sea and Naantali's Old Town with its wooden houses meet.

Tove Jansson was a Finnish writer and a visual artist, whose Moomin characters are known all over the world. Moomin World is an ode to fairy tales and a tribute to Tove Jansson's imagination, located in south-west coast of Finland, in the island of Kailo, in Naantali. Moomin World can be reached within 20 minutes from the downtown of Turku.

Who knows, maybe we will get to meet the Moomins during the dinner, come along to spend a memorable evening!

### Bus transportation to dinner

Buses to dinner leave from two locations (see below) in short intervals between 17:15 and 17:40. Families with children are recommended to take the first bus in order to give them time to explore the island before larger crowds arrive:

- 1. Turku Cathedral, address Tuomiokirkonkatu 1
- 2. Hotel Radisson Blu Marina Palace, address Linnankatu 32

In Naantali, there is about 10 minute walk from the bus to Moominworld island. Congress volunteers will show the way.

Return buses will leave starting from 20.00 in about 30 minute intervals. All return buses will stop at Hotel Radisson Blu Marina Palace and near Turku Cathedral.

### Boat transportation (one way, return by bus)

If you have booked a ticket for the boat:

The boat leaves at 17:15 sharp from the River Aura, address: Läntinen Rantakatu 37

### **RUNNING SOCIAL**

Thursday, 22 August, at 07:00

A running social will be organized early Thursday morning, meeting in the front of the Cathedral (Tuomiokirkonkatu 1, 20500 Turku). The routes will follow the river Aura. We aim to have two groups (different paces and distances).

### **FAMILY SOCIAL**

Thursday, 22 August, at 15:00

A family social will be organized on Thursday afternoon at the Seikkailupuisto adventure park (Kupittaankatu 2, 20520 Turku; meeting point at the big moose statue ), and in case of rain, in a child-friendly museum of history Aboa vetus (Itäinen Rantakatu 4-6, 20700, Turku) at 15:00.

### **LGBTQ SOCIAL**

Friday, 23 August, at 19:30

The ESEB 2019 LGBTQ social will feature a collaboration with Turku Pride! Following the poster session on Friday August 23, we will meet near the main door in Logomo at 19:30 and head over to Saaristobaari (Aurakatu 14, Turku) together to grab a bite to eat and get to know each other. At 10pm, there will be a drag show (5€ cover) featuring Finland's fiercest underground drag and burlesque artists and DJ Slaya Bit. The show is the official Turku Pride pre-party, so let's get there early! For more information, please check @ESEB2019LGBTQ on Twitter or email evolcongen1@gmail.com - the first 100 ESEB members attending the congress who RSVP by 22 August will get in free to the show.

### **CONGRESS PUB**

### Panimo ravintola Koulu

Address: Eerikinkatu 18 (the second floor of the restaurant is reserved for ESEB participants) Open every day 11:00-02:00, kitchen open until 21:30 (Mon-Thu), until 23:30 (Fri-Sat).

Look out for two evolutionary themed congress beers on tap in the upstairs bar of Panimoravintola Koulu; 'Beer Reviewed' and 'Drinkage Disequilibrium'. Our beer names were chosen following a twitter poll on a shortlist of names, whittled down from an extensive list of submissions of varying creativity (it turns out evolutionary biologists love both procrastinating and puns). Credit goes to Martin Seltmann who came up with 'Beer Reviewed' and Will Buswell for 'Drinkage Disequilibrium', as well as to Océane Liehrmann for the great logo adaptation and designs!

Panimoravintola Koulu (Brewery restaurant School in English) is a former elementary school, now Finland's largest brewery restaurant serving large selection of beers, wine and delicious food too!





# SATE

### PEER COMMUNITY IN (PCI) - PEER COMMUNITY IN...THE BEGINNING OF A **REVOLUTION IN OPEN ACCESS?**

Where: MOVE 1

When: Tuesday 20 August, 13:00-13:50 (lunch time)

Who: Researchers

Interested by discovering/joining/using the next generation publishing experiment with the "Peer Community In" (PCI, https://peercommunityin.org) project? In a few words: PCI is a non-profit scientific organization that aims to create specific communities of researchers reviewing and recommending, for free, unpublished preprints in their field (i.e. unpublished articles deposited on open online archives like arXiv.org and bioRxiv. org). Evaluations and recommendations by a PCI are free of charge for authors and readers. The first PCI, Peer Community in Evolutionary Biology (PCI Evol Biol), has been launched in 2017 and now counts >400 Editors. Other PCIs (eg PCI Ecology, PCI Paleontology, PCI Entomology...) have been created and several PCIs will probably open soon. Come along and meet Thomas Guillemaud & Denis Bourguet - co-funders of PCI - and many reasearchers already involved as editors @PCI Evol Biol. See also https://youtu.be/4PZhpnc8wwo, @ PCIEvolBiol & @PeerCommunityIn.

### THE EUROPEAN RESEARCH COUNCIL - FUNDING OPPORTUNITIES FOR BRIGHT **MINDS**

Where: MOVE 1

When: Wednesday 21 August, 13:15-14:05 (lunch time)

Who: Researchers

Is an ERC grant for you? You will be explained what the European Research Council is, who can benefit from its funding opportunities and what to expect in the application and selection process. The ERC supports researchers performing interesting and ambitious fundamental research. This could be you!

The mission of the European Research Council is to encourage the highest quality research in Europe. The concept is simple: competitive individual funding for researchers with a great idea, across all fields. ERC grants are awarded through open competition to projects headed by starting and established researchers of any nationality and age, who are working or moving to work in Europe or an associated country. The sole criterion of choice is scientific excellence.

### **NETHERLANDS EVOLUTIONARY BIOLOGY GET-TOGETHER**

Where: Congress pub Panimoravintola Koulu, address: Eerikinkatu 18

When: Wednesday 21 August, 18:00-19:30

Who: All scientists working in the Netherlands or of Dutch origin

The Netherlands society for evolutionary biology (NLSEB) aims to build a community of all evolutionary biologists in the Netherlands. NLSEB therefore welcomes all scientists working in the Netherlands or from Dutch origin for drinks. Come and (re-)connect to Dutch evolutionary biology!

### **MEET THE EDITORS - A ROYAL SOCIETY PUBLISHING WORKSHOP**

Where: MOVE 1

When: Friday 23 August, 13:00-13:50 (lunch time)

Who: Everybody interested

Presenters: Editors from the Royal Society journals Proceedings B, Philosophical Transactions B and Biology

Letters, including ESEB President Professor Nina Wedell.

Have you ever wondered what happens to a paper submitted to a Royal Society journal? This is your chance to find out. This one-off event offers an excellent opportunity to gain valuable insight into the peer review and processes behind the scenes at Royal Society Publishing. Come along and meet some of the highly experienced and reputable editors working for the Royal Society journals, and hear more about their expectations and top tips for compiling high quality articles. There will also be plenty of time to discuss topical publishing issues, and questions and feedback from the audience will be encouraged.

### **OUTREACH EVENTS**

### - OPEN TO EVERYBODY INTERESTED IN OUTREACH

### **HOW TO FOSTER PUBLIC ENGAGEMENT AT CONFERENCES**

Where: Outreach Stage

When: Tuesday 20 August, 12:45-13:55 (lunch time)

In this session, you will be presented with a few examples of how outreach and public engagement were fostered at scientific conferences and participate to a brainstorming session on how to do in the future, to be able to go back home with plenty of concrete ideas! (Héloïse Dufour)

### A CITIZEN-SCIENCE WORKSHOP

Where: GOTO 31 (3rd floor)

When: Tuesday 20 August, 12:45-13:55 (lunch time)

A citizen-science workshop showcasing an amazing and successful Droseu citizen-science initiative (Roberto Torres)

### **HOW TO PITCH YOUR SCIENCE TO NON-SPECIALIST AUDIENCES**

Where: GOTO 31 (3rd floor)

When: Wednesday 21 August, 12:55-13:55 (lunch time)

In this workshop, you will get tips on how to discuss your science with non-specialist audiences and actually build and practice on YOUR pitch(es). (Héloïse Dufour)

### SCISPARKS, HOW TO ORGANISE SPEED MEETINGS IN HIGH-SCHOOLS

Where: Outreach Stage

When: Friday 23 August, 12:45-13:55 (lunch time)

In this session, you will learn how to easily organise effective encounters between researchers and highschool students using speed-meetings, and how to get support to start your own. They are fun ways to create engaging links between students, teachers, and researchers! This session is also for you if you want to become part of a European coordinated activity dedicated to evolution! (Héloïse Dufour)

### **ART-UP YOUR EVOLUTION**

Where: Outreach Stage

When: lunchtime + coffee-breaks + continued moderation throughout the conference on a flexible basis, lasts until the end of the last coffee break

Unleash your artistic side! Take the paint, crayons, paper, brushes and whatever else you need - and show us your artistic vision of your research, results - or yourself as a scientist! All materials will be provided - just come and express yourself. (Szymek Drobniak)

### **COME AND MEET EVOKE!**

Where: Exhibition area

When: Throughout whole conference

Come share with us what outreach activities you are involved in and why! You will also learn about examples of activities you can get involved in or use. Last but not least, you will hear about EvoKE, the network aiming at Evolutionary Knowledge for Everyone, to get in touch with a diversity of people with the same goal!

### LIST OF SYMPOSIA

### S1. Trans generational plasticity in animals (Trans gen plast)

Organisers: Dalial Freitak, Olivia Roth Invited: Marjo Saastamoinen, Seth Barribeau

### **S2.** Evolution in real time: experimental evolution approaches

(Exp evol)

Organisers: Biljana Stojković, Uroš Savković, Mirko

Đorđević

Invited: Göran Arngvist, Tadeusz Kawecki

### **S3.** Exploring the role of nongenetic inheritance in evolution (Non-gen inherit)

Organisers: Pim Edelaar, Russell Bonduriansky, Troy Day

Invited: Itamar Lev, Sonia Sultan

### **S4.** Cognitive evolution and environment

(Cognition)

Organisers: Antonin Crumiere, Manuel Nagel Invited: Reuven Dukas, Gabrielle Davidson

### **S5. Aging & Cancer through the lens of** evolution (Aging & cancer)

Organisers: E. Yagmur Erten, Matthias Galipaud, Robert

Noble

Invited: Vera Gorbunova, Joao Pedro de

Magalhaes

### S6. Eco-evolutionary approach to the antimicrobial resistance problem

(Anti-micro resist)

**Organisers:** Teppo Hiltunen, Lutz Becks Invited: Danna R Gifford, Dan Andersson

### **S7. Human-induced evolution** (Human-induced)

Organisers: Miguel Baltazar-Soares, Kristien Brans,

Christophe Eizaguirre

Invited: Fanie Pelletier, Mikko Heino

### **S8. Genetics of small populations**

(Small pop gen)

Organisers: Alina Niskanen, Lumi Viljakainen, Henrik

Invited: Richard Frankham, Nancy Chen & Jane Reid (Ex-

ternally sponsored)

### S9. Microbial genome and community evo**lution in food environments** (Microbes & food)

Organisers: Jeanne Ropars, Ricardo Rodriguez de la Vega Invited: Delphine Sicard, John Gibbons

### \$10. Rapid evolutionary adaption: potential and constraints

(Rapid adapt)

Organisers: Carolin Wendling, Jürgen Gadau

Invited: Alison Feder, Lutz Becks

The symposium is sponsored by the DFG priority program

SPP1819

### S11. Quantitative trait effect size distributions and their impact on evolutionary processes (Quant traits)

Organisers: Arild Husby, Anna Santure Invited: John Kelly, Mirte Bosse

### S12. Quantifying selection and evolvability in wild plant populations: methods and measurements (Wild plant sel)

Organisers: Øystein H. Opedal, Rocío Pérez-Barrales Invited: Benoit Pujol, Maria Clara Castellanos

### S13. Genetics and genomics of adaption

(Adapt gen)

Organisers: Carmelo Fruciano, Paolo Franchini, Julia C.

Jones

Invited: Kathryn Elmer, Henrique Teotónio

### **S14. The mechanisms of evolutionary** change: moving from genomic signatures to functional validation (Genome funct)

Organisers: Darren J. Parker, Nicola Cook Invited: Alistair P. McGregor, Megan Neville

### **\$15. Tracing evolution through time using** ancient DNA (Ancient DNA)

Organisers: Päivi Onkamo, Verena Schünemann, Elina

Invited: David Wegmann, Johannes Krause

### \$16. Mito-nuclear interactions across levels of biological organisation (Mito-nuclear)

**Organisers:** Florencia Camus, Hernan Morales Invited: Ronald S. Burton, Kristi Montooth

### **S17. Selfish genetic elements** (Selfish GEs)

Organisers: Robert Kofler, Kirsten A. Senti Invited: Catherine Montchamp-Moreau, Arturo

Mari-Ordonez

### S18. The genetic architecture of polygenic adaption: sweeps, small shifts and everything in between (Polygen arch)

Organisers: Christian Schlötterer, Neda Barghi Invited: Catherine Peichel, Joachim Hermisson The symposium is supported by Molecular Ecology

### **MOLECULAR ECOLOGY**

### S19. Gene-phenotype associations across evolutionary scales (Geno-pheno)

Organisers: Jo Baker, Stephen Montgomery, Francesco

Cicconardi

Rapid

Evolutionary

L\_Adaptation

Invited: Nicola Nadeau, Itay Mayrose

### **S20.** The evolutionary consequences of social transmission and animal culture

(Social trans)

Organisers: Rose Thorogood, Neeltje Boogert

Invited: Lucy Aplin, Sasha Dall

### **S21.** Colour across the evolutionary spectrum: from production to perception (Colour)

Organisers: Hugo Gruson, Amélie Fargevieille, Nicola

Nadeau

Invited: Edwige Moyroud, Martine Maan

### **\$22.** Evolution of host-plant use in arthropods (Host-plant)

Organisers: Ernesto Villacis-Perez, Nicky Wybouw

Invited: Silke Allmann, Noah Whiteman

### **S23. Parasite community dynamics and** their role in the evolution of host immunity

(Parasite comm dyn)

Organisers: Tobias Lenz, Sébastien Calvignac-

Spencer

Invited: Anssi Karvonen, Elin Videvall

### S24. Microbial evolution under biotic stress (Microbial stress)

**Organisers:** Marie Vasse, Antoine Frenoy Invited: Olaya Rendueles, Ville-Petri Friman

### **S25.** Assortative mating for quantitative traits: mechanisms, estimation and evolutionary consequences (Assort mating)

Organisers: Niels Dingemanse, Barbara Class Invited: Wolfgang Forstmeier, Roger Butlin

### S26. Sexual conflict: linking behavior, genetics and ecology (Sex conflicts)

Organisers: Kenyon Mobley, Jessica Abbott, Stephen

De Lisle

Invited: Jen Perry, Howard Rundle

### S27. Design of social traits: genes, individuals and social groups (Social traits)

Organisers: Gonçalo S. Faria, Thomas Hitchcock,

Jasmeen Kanwal

Invited: Susanne Shultz, Alan Grafen

### **S28. Evolutionary game theory: modern** development and interdisciplinary applications (Game theory)

Organisers: Xiang-Yi Li, Vlastimil Křivan, Christian Hilbe

Invited: Katerina Stankova, Redouan Bshary

### **\$29.** Moving beyond a quantification of eco-evolutionary dynamics (Eco-evo)

Organisers: Lynn Govaert, Marjolein Bruijning

Invited: Jelena Pantel, Tim Coulson

### S30. Eco-evolutionary feedback between pollinator behaviour and floral evolution

(Pollinator)

Organisers: Mario Vallejo-Marin, Avery Russell

Invited: Aimee Dunlap, Allan Ellis

### S31. Life history evolution: bridging theory and data (Life history)

Organisers: Piret Avila, Mauricio González-Forero Invited: Alexei Maklakov, Irja Ida Ratikainen

### \$32. Niche width evolution and its (mal)adaptive significance (Niche width)

Organisers: Maud Charlery de la Masselière, Virginie

Ravigné, Vincent Calcagno

Invited: Claus Rueffler, Michael Singer

### S33. Evolutionary ecology of ageing: from mechanisms to life-history consequences

Organisers: Sophie Reichert, Hannah Froy, Antoine Stier

Invited: Sandra Bouwhuis, Tonia Schwartz

### S34. Mathematical models in evolutionary biology (Math models)

Organisers: Guy Cooper, Matishalin Patel, Tom Scott,

Asher Leeks

Invited: Hanna Kokko, Florence Débarre

### S35. Evolution outreach projects: keep SCREAMing (Science Communication Research Empowers AMazing outreach)

(Evol outreach)

Organisers: Dragana Cvetković, Szymon M. Drobniak

Invited: Pedro Russo, Héloïse Dufour

### 36a. Sexual selection and reproductive strategies (Sex select & mating)

Organizers: Natalie Pilakouta, Murielle Ålund, Colin Olito

### 36b. Phylogeography, biogeography, speciation, systematics (Phylogeo & syst)

Organizers: Bjarki Eldon, Niklas Wahlberg

### S36c. Species interactions (Spp interact)

Organizers: Alexandre Figueiredo, Jos Kramer, Elisa

Granato

### 36d. Genome evolution (Genome evol)

Organizers: Alexander Nater, Wen-Juan Ma

### 36e. Phenotypic evolution (Phenotypic evol)

Organizer: Dany Garant

### AT A GLANCE

# PROGRAMME

| MON            | DAY AUGUST 19     |
|----------------|-------------------|
| 15.00<br>18.00 | Registration      |
| 18.00<br>21.00 | Welcome reception |

| TUE            | SDAY AUGUST 20   |
|----------------|--|
| 7.30           | Registration   |
| 8.30           | Opening of conference and practical information  |
| 9.00<br>9.45   | Keynote I Pat Monaghan, Bad beginnings and untimely ends: environments, telomeres and life history variation   |
| 10.00          | SYMPOSIA S10: Rapid Adapt, S31: Life History, S34: Math Models, S8: Small pop gen, S7: Human-induced, S23: Parasite com dyn, S18: Polygen arch, S32: Niche width       |
| 11.00          | Coffee & Exhibition & Outreach   |
| 11.30          | SYMPOSIA S10: Rapid Adapt, S31: Life History, S34: Math Models, S8: Small pop gen, S7: Human-induced, S23: Parasite com dyn, S18: Polygen arch, S32: Niche width       |
| 12.30          | Lunch & Exhibition & Satellite events/outreach   |
| 14.00          | SYMPOSIA S10: Rapid Adapt, S31: Life History, S34: Math Models, S8: Small pop gen, S7: Human-induced, S23: Parasite com dyn, S18: Polygen arch, S20: Social trans      |
| 15.30          | Coffee & Exhibition & Outreach   |
| 16.00<br>17.15 | SYMPOSIA S10: Rapid Adapt, S31: Life History, S36d: Genome evol, S8: Small pop gen, S7: Human-induced, S36e: Phenotypic evol, S36b: Phylogeo & syst, S20: Social trans |
| 17.20<br>19.20 | POSTER SESSION I   |

| WEDI           | WEDNESDAY AUGUST 21  |  |  |  |  |
|----------------|--|--|--|--|--|
| 08.55          | ESEB initiatives and practical information   |  |  |  |  |
| 9.05<br>9.50   | Keynote II Sinead Collins, Understanding evolution in life-giving slime  |  |  |  |  |
| 10.00          | SYMPOSIA S10: Rapid Adapt, S31: Life History, S34: Math Models, S14: Genome funct, S33: Aging, S4: Cognition, S35: Evol outreach, S11: Quant traits  |  |  |  |  |
| 11.00          | Coffee & Exhibition & Outreach   |  |  |  |  |
| 11.30          | SYMPOSIA S10: Rapid Adapt, S31: Life History, S34: Math Models, S14: Genome funct, S33: Aging, S4: Cognition, S35: Evol outreach, S11: Quant traits  |  |  |  |  |
| 12.45          | Lunch & Exhibition & Satellite events/outreach   |  |  |  |  |
| 14.15          | SYMPOSIA S13: Adapt gen, S21: Colour, S6: Anti-micro resist, S14: Genome funct, S17: Selfish GEs, S4: Cognition, S25: Assort mating, S22: Host-plant |  |  |  |  |
| 15.45          | Coffee & Exhibition & Outreach   |  |  |  |  |
| 16.15<br>17.30 | SYMPOSIA S13: Adapt gen, S21: Colour, S6: Anti-micro resist, S14: Genome funct, S17: Selfish GEs, S33: Aging, S25: Assort mating, S22: Host-plant    |  |  |  |  |

| THURSDAY AUGUST 22 |   |  |  |  |  |
|--------------------|---|--|--|--|--|
| 08.55              | ESEB initiatives and practical information  |  |  |  |  |
| 9.05<br>9.50       | Keynote III David Queller, Evolutionary conflict and molecular arms races in cooperative systems  |  |  |  |  |
| 10.00              | SYMPOSIA \$13: Adapt gen, \$21: Colour, \$26: Sex conflict, \$3: Non-gen inherit, \$29: Eco-evo, \$6: Anti-micro resist, \$12: Wild plant sel, \$16: Mito-nuclear |  |  |  |  |
| 11.00              | Coffee & Exhibition & Outreach  |  |  |  |  |
| 11.30              | SYMPOSIA S13: Adapt gen, S21: Colour, S26: Sex conflict, S3: Non-gen inherit, S29: Eco-evo, S7: Human induced, S12: Wild plant sel, S16: Mito-nuclear             |  |  |  |  |
| 12.45              | Lunch & Exhibition & Satellite events/outreach  |  |  |  |  |
| 13.45<br>18.00     | Excursions  |  |  |  |  |

| FRIDAY AUGUST 23 |   |  |  |  |  |
|------------------|---|--|--|--|--|
| 8.55             | ESEB initiatives and practical information  |  |  |  |  |
| 9.05<br>9.50     | Keynote IV Anna-Liisa Laine, What keeps pathogens in check in the wild?   |  |  |  |  |
| 10.00            | SYMPOSIA S13: Adapt gen, S2: Exp evol, S26: Sex conflict, S15: Ancient DNA, S27: Social traits, S1: Trans gen plast, S19: Geno-pheno, S24: Microbial stress     |  |  |  |  |
| 11.00            | Coffee & Exhibition & Outreach  |  |  |  |  |
| 11.30            | SYMPOSIA S13: Adapt gen, S2: Exp evol, S26: Sex conflict, S15: Ancient DNA, S27: Social traits, S1: Trans gen plast, S19: Geno-pheno, S24: Microbial stress     |  |  |  |  |
| 12.30            | Lunch & Exhibition & Satellite events/outreach  |  |  |  |  |
| 14.00            | SYMPOSIA S13: Adapt gen, S2: Exp evol, S26: Sex conflict, S15: Ancient DNA, S27: Social traits, S1: Trans gen plast, S19: Geno-pheno, S24: Microbial stress     |  |  |  |  |
| 15.30            | Coffee & Exhibition & Outreach  |  |  |  |  |
| 16.00<br>17.15   | SYMPOSIA S13: Adapt gen, S2: Exp evol, S26: Sex conflict, S3: Non-gen inherit, S27: Social traits, S36b: Phylogeo & syst, S36d: Genome evol, S36c: Spp interact |  |  |  |  |
| 17.20<br>19.20   | POSTER SESSION II   |  |  |  |  |

| SATU           | RDAY AUGUST 24   |
|----------------|--|
| 8.55           | ESEB initiatives and practical information   |
| 9.05<br>9.50   | Keynote V Rasmus Nielsen, Human adaptation in time and space   |
| 10.00          | SYMPOSIA S13: Adapt gen, S2: Exp evol, S36a: Sex select & mating, S28: Game theory, S27: Social traits, S30: Pollinator, S9: Microbes & food, S5: Aging & cancer   |
| 11.00          | Coffee & Exhibition & Outreach   |
| 11.30          | SYMPOSIA S13: Adapt gen, S2: Exp evol, S36a: Sex select & mating, S28: Game theory, S27: Social traits, S30: Pollinator, S9: Microbes & food, S5: Aging & cancer   |
| 12.45          | Lunch & Exhibition   |
| 13.30          | ESEB members meeting   |
| 14.30          | Incoming president's address, Ophelie Ronce, Integrating niche evolution with life history theory can help us better understand the consequences of climate change |
| 15.10          | Leg stretching break   |
| 15.20          | JMS award winner 2019, Karl Grieshop, Sexual conflict and the maintenance of genetic variance in fitness   |
| 15.50<br>16.20 | Closing ceremony   |
| 18.30<br>02.00 | Congress dinner at Muuminworld   |

|        |       | LOGOMO HALL   | TEATRO  | GALLERIA  | LOGI1  |  |
|--------|-------|---|---|---|--|--|
|        | 7.45  |   | REGIST  | RATION  |  |  |
|        | 8.30  | OPENING OF CONFERENCE AND PRACTICAL INFORMATION   |   |   |  |  |
|        | 9.00  | KEYNOTE I Pat Mona  | ghan, Bad beginnings and unti   | mely ends: environments, telor  | meres and life history variation   |  |
| 0      |       | S10: RAPID ADAPT  | S31: LIFE HISTORY   | S34: MATH MODELS  | S8: SMALL POP GEN  |  |
|        | 10.00 | \$10.01 Slowing the rapid evolution of HIV drug resistance A. Feder   | S31.O1 Why do organisms age: Beyond energy trade-offs A. Maklakov   | S34.O1 Why you might want to care about population regulation, no matter what your question is H. Kokko                                     | S8.01 What sizes are required for populations to be genetically viable? Re-evaluation of the 50/500 rules R. Frankham        |  |
| AUGUST | 10.30 | S10.O2  Host virus coevolution – demography versus selection in the face of multiple stressors  L. Becks  | S31.O2 Co-evolution of life history traits in variable environments I. I. Ratikainen  | S34.O2  Reconciling different modelling approaches in evolutionary theory  F. Débarre   | S8.O2 Genetic and fitness consequences of dispersal in a small pedigreed population N. Chen                                  |  |
| 4      | 11.00 | CC  | FFEE & EXHIBIT<br>(Art up your evolution, Out   | TION & OUTREAC<br>treach stage, Teatro lobby)   | СН   |  |
| SDAY,  | 11.30 | S10.O3  Tracking viral life history during experimental coevolution with their hosts  E. J. P. Lievens  | \$31.03  Ageing and the fecundity/longevity trade-off in social insects: a comparative approach  J. Korb                        | S34.O3  Does ecology matter in evolutionary models?  B. Ashby   | S8.O3  Complexities of inbreeding, outbreeding and inbreeding depression in a song sparrow meta-population  J. Reid          |  |
|        | 11.45 | S10.04 Rapid resource use specialisation leads to increased virulence in plant pathogenic Ralstonia solanacearumbacterium L. Mikonranta   | S31.04 The effect of environmental stress on ageing in social insects V. Rau  | S34.O4 Individual-based models improve understanding of evolutionary dynamics: examples from female multiple mating and dispersal G. Bocedi | S8.04 Genetic load accumulation from the perspective of post-bottleneck populations of Galapagos Mockingbirds. J. Vlček      |  |
| 英      | 12.00 | S10.05 Changes in allelic frequencies of Brassica rapa under experimental evolution with selection by bumblebees L. Frachon   | S31.05 The cost of longevity: Transgenerational effects of parental lifespan extension under dietary restriction E. Ivimey-Cook | S34.05  Dynamic invariance of evolutionary models  J. Otsuka  | S8.05  Patterns of genetic variation across the genome in bottlenecked populations of Eurasian and Iberian lynx  J. A. Godoy |  |
|        | 12.15 | S10.06  The genomics of rapid adaptation to climate change: host preference evolution increases short-term ecological resilience J. Bridle  | S31.06 Social context does not modulate age fitness effects in Drosophila melanogaster Z. Sultanova                             | S34.06 Predicting evolution: combining developmental biology and quantitative genetics L. Milocco   | \$8.06 Founder-specific inbreeding depression in an island bird population P. Nietlisbach                                    |  |
|        | 12.30 | LUNCH & EX  | HIBITION & SAT  | ELLITE EVENTS   | / OUTREACH   |  |
| X      |       | Satellite events  Peer Community in (PCI) – Peer Community In the beginning of a revolution in Open Access?, MOVE 1 at 13:00-13:50  How to foster public engagement at conferences, Outreach Stage in Teatro lobby at 12:45-13:55  A citizen-science workshop, GOTO 31 at12:45-13:55  Art-up your evolution, Outreach Stage, Teatro lobby |   |   |  |  |

|       | MOVE1   | MOVE2   | LOGI2   | <i>GOТО33</i>   |  |
|-------|---|---|---|---|--|
| 7.45  |   | REGIST  | RATION  |   |  |
| 8.30  | OPENING OF  | CONFERENCE AI   | ND PRACTICAL I  | NFORMATION  |  |
| 9.00  | KEYNOTE I Pat Mona  | ghan, Bad beginnings and unti   | mely ends: environments, telor  | meres and life history variation  |  |
|       | S7: HUMAN-INDUCED   | S23: PARASITE COM DYN   | S18: POLYGE ARCH  | S32: NICHE WIDTH  |  |
| 10.00 | S7.01 Hunting regulation and the dynamic of selection in large mammals F. Pelletier   | S23.O1  Dynamics of parasite co-infections – why do they matter?  A. Karvonen   | \$18.01 Genetic and genomic architecture of polygenic adaptation in lake-stream sticklebacks C. Peichel           | S32.01 Evolutionary diversification driven by competition for resources - does organismal complexity matter? C. Rueffler                          |  |
| 10.30 | S7.02 Fisheries-induced evolution in the wild and in the lab M. Heino   | S23.O2  Dual transcriptomics of avian malaria  E. Videvall  | \$18.02  Polygenic adaptation: The adaptive architecture of a quantitative trait  J. Hermisson                    | S32.O2 Colonizations and host shifts cause diversification of preference and expansion of diet breadth M. Singer                                  |  |
| 11.00 | CC  | FFEE & EXHIBIT<br>(Art up your evolution, Out   | TION & OUTREAC  | СН  |  |
| 11.30 | S7.03  Anthropogenic hybridization between red deer and sika in Kintyre results in many backcrossed individuals  S. E. McFarlane  | S23.O3  Virulence-transmission relationships under competition in the spider mite Tetranychus urticae  A. Duncan                        | S18.O3  Proper Treatment of Haplotype Structure and LD Reduces Error in Sequence Data Analysis S. Belohlavy       | S32.O3  Habitat choice meets thermal niche specialization: competition with specialists may drive suboptimal preferences in generalists  S. Jacob |  |
| 11.45 | S7.04 Evolutionary rescue through hybridization triggered by predator introduction in a Daphnia population K. Enberg  | S23.04 Wolbachia incidence and host shift in scale insects E. Sanaei  | \$18.04 Genetic redundancy fuels polygenic adaptation in Drosophila R. Kofler                                     | S32.O4 Expression of phenotypic plasticity in multi-dimensional environments N. Schtickzelle  |  |
| 12.00 | S7.05 The Global Urban Evolution Project: Parallel Adaptation To The World's Urban Jungles M. Johnson   | S23.05 Characterization of the human pathogen peptidome and specialization in peptide binding among MHC class-I alleles O. Özer         | S18.05  Response from standing variation at linked loci in the highly polygenic/infinitesimal limit.  H. Sachdeva | S32.05  Not a generalist after all?  Life history genomic regions explain differences in Atlantic salmon diet.  T. Aykanat                        |  |
| 12.15 | S7.06 Contrasting body-size shifts in urban communities T. Merckx   | S23.06  How decreased parasite diversity affects host immunity: Approaching "Old Friends" with the cavefish, Astyanax mexicanus R. Peuß | S18.06  Detecting the signature of epistatic selection in subdivided populations  K. Csilléry                     | S32.06 Are differences in incubation behavior and niche use linked in two sympatric flycatcher species? P.M. Sirkiä                               |  |
| 12.30 | LUNCH & EX  | HIBITION & SAT  | ELLITE EVENTS   | / OUTREACH  |  |
|       | Satellite events  Peer Community in (PCI) – Peer Community In the beginning of a revolution in Open Access?, MOVE 1 at 13:00-13:50  How to foster public engagement at conferences, Outreach Stage in Teatro lobby at 12:45-13:55  A citizen-science workshop, GOTO 31 at12:45-13:55  Art-up your evolution, Outreach Stage, Teatro lobby |   |   |   |  |

|       | LOGOMO HALL   | TEATRO  | <b>GALLERIA</b>   | LOGI1   |
|-------|---|---|---|---|
|       | S10: RAPID ADAPT  | S31: LIFE HISTORY   | S34: MATH MODELS  | S8: SMALL POP GEN   |
| 14.00 | S10.07 Regulatory networks link phenotypic plasticity to evolvability F. Weissing   | \$31.07 Limits to post-reproductive fitness benefits in humans S. Chapman   | S34.07 Free-riding, exclusion, and congestion in a sequential teamwork dilemma J. Peña  | S8.07 Genetic diversity and connectivity in wetland plant meta-populations depend on the degree of clonality S. Donna Lozada-Gobilard |
| 14.15 | S10.08 Assessing genetic constraints on the evolution of plasticity in multiple stressor environments A. Hudak                                      | S31.08 Child volunteers in World War II have accelerated reproduction and higher lifetime reproductive success R. Lynch | S34.O8 Selection and Polymorphism at Two Loci H. Spencer  | S8.08 Effects of non-random mating and Haldane's Sieve on floral polymorphisms in plant metapopulations J. Pannell                    |
| 14.30 | S10.09  Evolution of physiological plasticity and selection from balanced polymorphisms during rapid habitat invasions  C.E. Lee                    | S31.09  The antagonistic pleiotropy riddle for populations along the slow-fast continuum  C. Coste                      | S34.09  Modeling antimicrobial cycling, mixing, and combination therapy: Why is it so difficult to draw conclusions?  H. Uecker | S8.09 Fitness, life-histories, and ageing in small populations of Daphnia C. Haag   |
| 14.45 | S10.010  Plasticity in evolutionary potential under environmental variation in a population of pied flycatchers, Ficedula hypoleuca  J. Le Vaillant | S31.010  Live fast, die old: Oxidative stress as a potential mediator of an unexpected life-history evolution  N. Tüzün | \$34.010  The evolution of self-incompatible mating types  J. Christie  | S8.010  Understanding contemporary levels of genetic diversity in populations of silver fir(Abies Alba Mill.)  B. Trubenová           |
| 15.00 | S10.011 Impact of maternal genetic effects on the evolutionary potential of a red deer population J. Gauzere  | S31.011  Does the life history response to dietary restriction persist with infection or injury?  E. Savola             | S34.011  Kin selection of function-valued traits  P. Avila  | S8.011 On the generality of the diploid male vortex in parasitoids with single-locus complementary sex determination E. Nonaka        |
| 15.15 | S10.012  Somatic mutation and cell lineage selection during vegetative growth promotes rapid adaptation in plants  J. Schwoch                       | S31.012  Diet-based developmental plasticity and fitness in a detritivorous isopod (Asellus aquaticus)  M. Lürig        | S34.012  Emergence of diverse life cycles and life histories at the origin of multicellularity  M. Staps                        | S8.012  Genomic signatures of critically-endangered bird Chinese Crested Tern (Thalasseus bernsteini)  G. Chen                        |
| 15.30 |   | FFEE & EXHIBIT  |   |   |

|       | MOVE1  | MOVE2  | LOGI2  | <i>GOTO33</i>   |
|-------|--|--|--|---|
|       | S7: HUMAN-INDUCED  | S23: PARASITE COM DYN  | S18: POLYGE ARCH   | S20: SOCIAL TRANS   |
| 14.00 | S7.07  Anthropogenic Pb driving selection in urban adapted population of Drosophila subobscura  A. Patenković              | S23.07  The determinants of pathogen communities in wild plant populations  H. Susi                                    | \$18.07 Selective sweep at at QTL in a randomly fluctuating environment LM. Chevin   | S20.O1 Animal Culture in Changing Environments L. Aplin                                       |
| 14.15 |  |  |  |   |
| 14.13 | S7.08 Going to the dogs? — Human-induced evolution in the grey wolf M. Pilot   | S23.08  Population genomics of Gyrodactylus bullatarudis reveals molecular basis of adaptation to the host  M. Konczal | S18.08 Wild wild test: Release-recapture genomic experiment reveals within-generation polygenic adaptation in stickleback fish T. Laurentino |   |
| 14.30 | \$7.09   | \$23.09  | S18.09   | \$20.02   |
|       | Genomics of adaptation of<br>Penicilliumfungi used for<br>blue cheese and dry-cured<br>meat production<br>A. Branca        | Within-host pathogen diversity: how it forms and what are the fitness consequences for the host S. Sallinen            | The genomic basis of parallel adaptation  A. M. Westram  | Does cultural transmission<br>evolve because it is<br>Lamarckian?<br>S. Dall                  |
| 14.45 | \$7.010  House sparrows evolved human commensalism with  | S23.O10  Disease-induced diversity of a crustacean iridescent  | \$18.010  Contemporary Atlantic salmon domestications  |   |
|       | the development of<br>agriculture<br><b>M. Ravinet</b>   | virus<br><b>V. G. Faria</b>  | reveal the architecture of<br>polygenic adaptation<br>N. J. Barson   |   |
| 15.00 |  |  |  |   |
| 13.00 | S7.011 Can angling-induced evolution be counteracted by releasing hatchery-reared fish? A. Vainikka                        | S23.011  Manipulated geographic mosaics: disentangling prevalence of infection and strength of selection  F. Feijen    | S18.011 Efficiency of outlier methods for detecting loci involved in a polygenic trait under divergent selection L. Bouteille                | S20.03  How do predators use social information about defended prey in the wild L. Hämäläinen |
| 15.15 | S10.012  | S23.O12  | S18.012  | \$20.04   |
|       | Rapid niche expansion in<br>European whitefish<br>following a eutrophica-<br>tion-induced species<br>collapse<br>A. Jacobs | Fitness effects of wild<br>Drosophila viruses<br><b>M. Wallace</b>   | Genomic prediction from<br>pool-seq to understand<br>ash dieback susceptibility<br>in fraxinus excelsior<br>C. Metheringham                  | Payoff- and sex-biased social learning interact in wild primate population  E. van de Waal    |
| 15.30 | CC   | DEFEE & EYHIRIT  | TION & OUTREAC   | CH CH   |

|                | LOGOMO HALL   | TEATRO  | GALLERIA   | LOGI1   |
|----------------|---|---|--|---|
|                | S10: RAPID ADAPT  | S31: LIFE HISTORY   | S36d: GENOME EVOL  | S8: SMALL POP GEN   |
| 16.00          | S10.013 Environmental integrons, drivers of microbial adaptation in an acidic extreme environment E. Sandoval-Quintana                            | S31.013 The evolution of variance control M. Bruijning  | S36d.O1  Extreme differences in recombination rate between the genomes of a solitary and a social bee J. Jones   | S8.013  Eco-Evolutionary feedbacks between genetic diversity and varying population sizes can lead to an extinction vortex P. Nabutanyi |
| 16.15          | S10.014 Predicting adaptive evolution in heterogeneous environments from standing genetic variation J. Engelstädter                               | S31.014 Life history evolution under cancer risk: linking cell-level strategies to organismal traits E. Y. Erten                    | S36d.O2 Shared ancient sex chromosomes in varanids, beaded lizards, and alligator lizards M. Rovatsos  | S8.014  Demography affects the likelihood of genetic convergence and our ability to detect it in nature  J. Whiting                     |
| 16.30          | S10.015  Beneficial mutations have greater fitness effects at higher temperatures, supporting the temperature-selection speed hypothesis  XL. Chu | \$31.015 Insulin-like growth factor 1 and the evolution of vertebrate life histories J. Lodjak                                      | S36d.O3 Germline-restricted chromosomes are widespread in songbirds and contain dozens of developmental genes A. Suh   | S8.015 High population divergence at small spatial scales – the joint role of population size and migration A. Nair                     |
| 16.45          | S10.016  Mechanisms of rapid adaptive responses to arid environments in long-lived organisms  O. Razgour  | S31.016  Maturation probability and condition correlate genetically for a major-effect locus (vgll3) in Atlantic salmon P. V. Debes | S36d.O4  Rearranged and relocated:     chromosome-level     assemblies and comparative genomics of two     pelagic freshwater herring     species     L. Milec | S8.016  Mechanisms and consequences of balancing selection in a model cyclic parthenogen living in ephemeral habitats  A. Bergland      |
| 17.00          | S10.017 Predicting adaptive dynamics in different habitats using ancestral trait values and demographic events V. Ravi Kumar                      | \$31.017  Adult male size in a sexually dimorphic spider depends on genetic factors and food availability  S. Quiñones-Lebrón       | S36d.O5 Epigenetic modification associated with ZEB2 provides a key evidence for the human evolution JE. Lee   | \$8.017  Bypassing summary statistics: a deep learning approach to infer population size history  T. Sanchez                            |
| 17.20<br>19.20 |   | POSTER  | SESSION I  |   |

|                | MOVE1   | MOVE2  | LOGI2  | <i>GOTO33</i>   |
|----------------|---|--|--|---|
|                | S7: HUMAN-INDUCED   | S36e: PHENOTYPIC EVOL  | S36b: PHYLOGEO & SYST  | S20: SOCIAL TRANS   |
| 16.00          | S7.O13 Evolution in salmon life-history induced by direct and indirect effects of fishing Y. Czorlich   | S36e.O1 Exploring patterns of additive genetic, mutational and environmental (co)variance across traits J. G. King           | \$36b.01 Global diversification patterns of crangonid shrimps (Decapoda, Caridea, Crangonidae) K. H. Chu   | S20.05 Cultural Transmission of Mating Preferences in Fruit Flies S. Nöbel                                    |
| 16.15          | S7.014  Harvest-associated selection and population density effects in fisheries-induced evolution  A. Crespel                                    | S36e.O2  Dissecting phenotypic integration and connecting micro- and macro-evolutionary time scales  C. Fruciano             | S36b.O2  Evo-devo approach to study asexual development and whole body regeneration: insights from tunicates  S. Tiozzo  | S20.06 A new perspective of social population networks in a reproductive context M. Plaza                     |
| 16.30          | S7.O15 Understanding the effect of multiple anthropogenic stressors on freshwater organisms from an evolutionary perspective M. Cuenca Cambronero | S36e.O3  An integrated approach to understanding the evolution of flight and wing shape in heliconius butterflies  L. Queste | S36b.O3  Phylogeography of a widespread spider: admixture across geographical barriers shapes the diversification of Gasteracantha cancriformis  F. C. Salgado-Roa | S20.07  On social transmission, individual agency, and a generalised theory of adaptive evolution  P. Edelaar |
| 16.45          | S7.016  Selection for small size affects the pace-of-life syndrome in medaka impacting the invertebrate community  B. Diaz Pauli                  | S36e.O4 Intraspecific variation in floral scent in the perennial herb Arabis alpina H. Petrén                                | S36b.O4 Incipient hybrid speciation in young and rapidly speciating neotropical cichlid fish?  M. Olave  | S20.08  Social network structure and infectious disease transmission in group-living animals  M. Silk         |
| 17.00          | S7.017  Applying the Anna Karenina principle to the bank vole gut microbiota in a disturbed environment  A. Lavrinienko                           | S36e.O5 Evolution of fork tails in aerial insectivorous birds M. Hasegawa  | S36b.O5  Patterns consistent with Darwin's corollary in a Ficedulaflycatcher hybrid zone C. Segami Marzal  | S20.09 Social transmission in avian brood parasitism systems D. Campobello                                    |
| 17.20<br>19.20 |   | POSTER   | SESSION I  |   |

|       | LOGOMO HALL   | TEATRO   | GALLERIA  | LOGI1  |  |  |
|-------|---|--|---|--|--|--|
| 8.55  | ESEI  | B initiatives and  | practical informa   | ation  |  |  |
| 9.05  | KEYNOTE II Sinead Collins, Understanding evolution in life-giving slime   |  |   |  |  |  |
|       | S10: RAPID ADAPT  | S31: LIFE HISTORY  | S34: MATH MODELS  | S14: GENOME FUNCT  |  |  |
| 10.00 | S10.018  Disparate signatures of rapid adaptation and genomic divergence in Nicaraguan Midas cichlid fishes  A. Nater                       | S31.018  Evolutionary constraints persist through a major life history event: metamorphosis  J. Collet                   | S34.O13 Extended haplodiploidy hypothesis P. Rautiala   | S14.O1  Differences in tartan underlie the evolution of male genital morphology between Drosophila species  A. P. McGregor |  |  |
| 10.15 | S10.019 Using whole genome sequences of newly introduced populations reveals rapid genetic adaptation in Trinidadian guppies M. van der Zee | S31.019 Disparity in diapause and its effects on insect movement V. Bhaumik  | S34.014 Evolution of the irreversible somatic differentiation Y. Gao  |  |  |  |
| 10.30 | \$10.020  Genetics and genomics of parallel evolution without gene flow Y. Yamasaki   | S31.O20 Locally adapted plasticity maintains geographic variation in life history strategies in a butterfly O. Lindestad | \$34.015 Flows of information in evolution A. Pocheville  | S14.02 Understanding the neural circuits that encode sex-specific behaviours in Drosophila melanogaster M. Neville         |  |  |
| 10.45 | S10.021  A tale of many flounders: the genomics of rapid adaptation in Platichthys spp. P. Momigliano                                       | S31.O21 Constrained evolution of instar-level characteristics of larval growth in Lepidoptera S. Kivelä                  | S34.016  Abstraction for dealing with the multiple realizability of evolution: the ultimate constraint of computation  A. Kaznatcheev |  |  |  |
| 11.00 |   |  | TION & OUTREAC  |  |  |  |

|       | MOVE1   | MOVE2  | LOGI2  | <i>GOTO33</i>  |  |  |  |  |
|-------|---|--|--|--|--|--|--|--|
| 8.55  | ESEE  | B initiatives and  | practical informa  | ation  |  |  |  |  |
| 9.05  | KEYNOTE II Sinead Collins, Understanding evolution in life-giving slime   |  |  |  |  |  |  |  |
|       | S33: AGING S4: COGNITION S35: EVOL OUTREACH S11: QUANT TRAITS   |  |  |  |  |  |  |  |
| 10.00 | S33.01 Understanding senescence and trans-generational parental age effects in a long-lived seabird S. Bouwhuis | S4.01 Evolutionary biology of expertise R. Dukas   | S35.01 The public and researchers: It's complicated P. Russo                               | S11.01 Inversions as large effect loci in quantitative genetics J. Kelly     |  |  |  |  |
| 10.30 | Sas.02 Senescence in reptiles: from mechanisms to life-history consequences T. Schwartz                         | S4.O2  The interplay between environment, gut microbiome and host cognition  G. Davidson | S35.O2 Willing to promote evolutionary knowledge for everyone? Join communities! H. Dufour | S11.O2 The clawprint of selection in wildlife and livestock genomes M. Bosse |  |  |  |  |
| 11.00 | CO  | FFEE & EXHIBIT<br>(Art up your evolution, Out  | FION & OUTREAC<br>treach stage, Teatro lobby)  | CH   |  |  |  |  |

|                                    | LOGOMO HALL  | TEATRO   | GALLERIA  | LOGI1   |
|------------------------------------|--|--|---|---|
|                                    | S10: RAPID ADAPT   | S31: LIFE HISTORY  | S34: MATH MODELS  | S14: GENOME FUNCT   |
| 11.30 <b>L S S S S S S S S S S</b> | S10.O22  Genomic divergence of rapidly evolving populations of Italian wall lizards  A. Štambuk                              | S31.022 Sex-specific effects of maturation timing on reproductive fitness in wild Atlantic salmon K. Mobley                | S34.017 Transcriptional cross-talk varies between regulatory networks designs T. Friedlander                      | S14.O3  Functional significance and evolutionary mechanisms of VMAT1genetic variants underlie psychological diversity in humans  D. Sato          |
| 11.45 AUGUAY, AUGUANA 12.00        | S10.023 Characterising genetic diversity and differentiation in multiple phenotypes of a marine invasive species M. Prentice | S31.023 Predation risk drives the evolution of placentas in live-bearing fish populations (family Poeciliidae) A. Hagmayer | S34.018  Evolutionary dynamics of plasticity in a mechanistic gene-network model  A. Odorico                      | S14.04 The genomic and transcriptomic basis of carotenoid-based sexual dichromatism in Finches M. Gazda   |
|                                    | S10.024 Why does male-biased gene expression evolve so rapidly? R. Griffin   | S31.024 Individual differences in carry-over effects on fitness: the role of personality S. M. Harris                      | S34.019 Flexible, realistic, fast evolutionary simulations with SLiM B. Haller                                    | S14.O5 The evolution of lifespan: from whole genomes to SNPs K. Hoedjes   |
| 12.15<br><b>W</b>                  | S10.025 Testing the factors promoting recurrent, convergent, and rapid adaptation in a wild insect  J. Rayner                | S31.O25 Environmental drivers of phenotypic selection in a small passerine species M. Gamelon                              | \$34.020 What can machine learning teach us about evolutionary ecology data?  J. Morimoto                         | S14.06  Molecular diversity and developmental expression of the master regulator doublesex in the sexually dimorphic Papilio polytes  R. Deshmukh |
| 12.30                              | S10.026 Identifying the evolutionary dynamics and genetics of rapid evolutionary rescue in Callosobruchus maculatus  A. Rêgo | S31.026 Evolutionary consequences of cryptobiosis on male reproduction M. Vecchi   | S34.021  Speciation, extinction and environmental change: from fossil data to mathematical modelling  J. Toivonen | S14.07 Key physiological genes important for freshwater adaptation and life history evolution in sticklebacks A. Ishikawa                         |

### **LUNCH & EXHIBITION & SATELLITE EVENTS / OUTREACH** 12.45

### Satellite events

The European Research Council – funding opportunities for bright minds, MOVE 1 at 13:15-14:05 How to pitch your science to non-specialist audiences, GOTO31 at 12:55-13:55 Art-up your evolution, Outreach Stage in Teatro lobby

|       | MOVE1  | MOVE2  | LOGI2   | <i>GOTO33</i>  |
|-------|--|--|---|--|
|       | S33: AGING   | S4: COGNITION  | S35: EVOL OUTREACH  | S11: QUANT TRAITS  |
| 11.30 | \$33.03 Sex-biased ageing in the invertebrate Tigriopus californicus and the role of mito-nuclear interactions S. Edmands                  | S4.03 A social perspective on the microbiota-gut-brain axis: ants as a model organism S. Teseo                           | S35.O3  Melanogaster Catch The Fly: a citizen science project on adaptation genomics J. Gonzalez                  | S11.O3 The response of a population to a change in optimum N. Barton   |
| 11.45 | S33.O4  Sex differences in functional and reproductive senescence in African annual killifish  M. Reichard                                 | S4.04  Benefits of working memory depend upon forage availability for bumblebees (Bombus terrestris)  E. Leadbeater      | s35.04 evALLution: can we make fundamental evolution concepts accessible to the blind community? T. G. Laurentino | S11.04 Disentangling the roles of mutation, selection, and genetic drift, on cis- and trans- regulatory evolution M. Hill                                      |
| 12.00 | S33.05  How the queen manages to stay young: orchid bee queens maintain young transcriptomes throughout life  A. Séguret                   | S4.05 Selective social information use in the nest choice of solitary bees O. Loukola                                    | S35.O5 Science and Community: evolutionary facts for an inclusive society J. R. Torres Miranda                    | S11.05 Dissecting evolution of adaptive traits in Arabidopsis after island colonization C. Neto  |
| 12.15 | S33.06  Long live the queen: eusociality and the evolutionary theory of ageing B. H. Kramer  | S4.06 Environmental complexity and the correlated evolution of (social) behaviour and (social) cognition S. A. M. Varela | S35.06  Evolution in action – project: How to impact society through science and art education?  C. Lindstedt     | S11.06 Genomic Prediction in a wild mammal population J. Slate   |
| 12.30 | S33.07  Extreme lifespan extension in tapeworm-infected ants facilitated by increased care and upregulation of longevity genes  S. Foitzik | S4.07  Heritability and co-variation among cognitive abilities in pheasants; an animal model approach  E. Langley        | \$35.07 The "WOW effect" of Evolution T. Adnađević  | S11.07  Beyond large-effect loci: large-scale GWAS reveals mixed large-effect and polygenic architecture of Atlantic salmon age-at-maturity M. Sinclair-Waters |
| 12.45 | LUNCH & EX   | HIBITION & SAT   | ELLITE EVENTS   | / OUTREACH   |
|       | The European De  | Satellit<br>search Council – funding oppo  | e events  | /E 1 at 12:15 1/:05  |
|       |  | pitch your science to non-spec   | =   |  |

How to pitch your science to non-specialist audiences, GOTO31 at 12:55-13:55 Art-up your evolution, Outreach Stage in Teatro lobby

|                |       | LOGOMO HALL   | <b>TEATRO</b>   | <b>GALLERIA</b>   | LOGI1  |
|----------------|-------|---|---|---|--|
| <b>T</b> _     |       | S13: ADAPT GEN  | S21: COLOUR   | S6: ANTI-MICRO RESIST   | S14: GENOME FUNCT  |
|                | 14.15 | S13.01 Adaptation and evolution of alternative reproductive modes K. Elmer  | S21.01 Painting by numbers: understanding the eco-evo-devo mechanisms of petal patterning E. Moyroud        | S6.01  Mutators drive evolution of resistance to multiple antibiotics during single-drug and combination treatment  D. R. Gifford | S14.08 Effects of random coding sequences on Escherichia coli D. Bhave   |
| Y, AUG         | 14.30 |   |   |   | S14.09 Colour encoded in innate immune gene? Accumulating evidence for Hamilton-Zuk 'Good genes' in great tits M. Vinkler              |
| NESDAY, AUGUST | 14.45 | S13.O2  Natural selection explains parallel evolution of locomotion bias, genetic drift variable interdependence of component traits  H. Teotónio | S21.02 Colour vision and the origin of species: what you see is who you are? M. Maan                        | S6.O2 Unstable antibiotic resistance D. Andersson   | S14.O10  Applying gene manipulation approaches for characterizing the evolution, dynamics and complexity of venom production  Y. Moran |
|                | 15.00 |   |   |   | S14.O11  Molecular mechanisms and evolution of a novel floral volatile biosynthesis in wild tobacco  S. Xu                             |
|                | 15.15 | S13.03  Parallel clines in iridescence in butterfly co-mimics despite different levels of genomic divergence and selection  E. Curran             | S21.03 Inter-chromosomal coupling between vision and pigmentation genes during genomic divergence O. Puebla | S6.O3  To establish, or not to establish – testing the probability of antibiotic resistance emergence  M. Saebelfeld              | S14.012  Molecular and phenotypic characterization of roo elements inserted in a unique insertional cluster  M. Merenciano             |
|                | 15.30 | S13.O4  Population genomics in a case of rapid, parallel adaptation: Cape Verde Islands Arabidopsis thaliana  A. Fulgione                         | S21.O4  Fine-mapping of color variation in a butterfly shed light on the evolution of supergenes  P. Jay    | S6.04 Repeatable ecological dynamics govern antibiotic response of experimental microbial community  J. Cairns                    | \$14.013 The contribution of novel genes to the development of novel traits R. Arbore  |
| M              | 15.45 | CO  | FFEE & EXHIBIT<br>(Art up your evolution, Out   | FION & OUTREAC<br>treach stage, Teatro lobby)   | H  |

|       | MOVE1   | MOVE2   | LOGI2   | <i>GOTO33</i>  |
|-------|---|---|---|--|
|       | S17: SELFISH GEs  | S4: COGNITION   | S25: ASSORT MATING  | S22: HOST PLANT  |
| 14.15 | S17.O1  X chromosome drive and intragenomic conflict: a textbook case in Drosophila simulans  C. Montchamp-Moreau | S4.08  Community diversity affects categorization by foragers: implications for signal evolution  D. Kikuchi                                      | S25.O1 Assortative mating from humans to birds – the role of mate choice W. Forstmeier                            | S22.O1  Getting tuned: Understanding specificity in plant volatile signaling  S. Allmann                                 |
| 14.30 |   | S4.09 Widespread learned predator recognition and amphibian resilience to alien predators N. Polo-Cavia   |   |  |
| 14.45 | \$17.02   | \$4.010   | \$25.02   | S22.O2   |
|       | Recognition and silencing<br>of active retrotransposons<br>in arabidopsis<br>A. Mari-Ordonez                      | Anti-predatory behaviour, sensory systems and brain transcriptomics in Icelandic threespine stickleback adapting to turbid environments  M. Ålund | Assortative mating, sexual selection and their consequences for gene flow in Littorina R. Butlin                  | Genome engineering as a<br>tool for studying host plant<br>specialisation<br>N. Whiteman                                 |
| 15.00 |   | S4.O11 Cognitive ontogeny: environmental effects on brain size divergence in developing sunfish ecotypes C. Axelrod                               |   |  |
| 15.15 | S17.03  Evolutionary dynamics of transposable elements in asexual bdelloid rotifers  R. Nowell                    | S4.012 Experimental support for the mosaic brain evolution hypothesis S. Fong   | S25.O3  Decomposing social genetic effects on phenology and assortative mating in a long-lived seabird  M. Moiron | S22.O3  Effect of plant inhibitory proteins on pectinases in herbivorous beetles  W. Häger                               |
| 15.30 | S17.04  Molecular dissection of a natural transposable element invasion  C. Schlötterer                           | S4.013  Annual predation risk relates to the direction of selection for brain size in the wild  M. Öst  | S25.O4  Reproductive isolation driven by ecological adaptation in Gambusia hubbsi  V. Pärssinen                   | S22.O4  Beyond target-site insensitivity - the role of ABCB transporters in adaptations to cardiac glycosides  S. Dobler |
| 15.45 | CO  | FFEE & EXHIBIT<br>(Art up your evolution, Ou  | TION & OUTREAC<br>treach stage, Teatro lobby)   | СН   |

|                                   | LOGOMO HALL  | TEATRO   | GALLERIA   | LOGI1  |
|-----------------------------------|--|--|--|--|
|                                   | S13: ADAPT GEN   | S21: COLOUR  | S6: ANTI-MICRO RESIST  | S14: GENOME FUNCT  |
| 16.15<br><b>L</b>                 | S13.05  Parallel evolution of reproductive timing in Atlantic and Pacific herring E. Petrou                                    | S21.O5  To change or not to change: evolution of seasonal colour polymorphism in the least weasel  I. Miranda                      | S6.05 On the evolutionary ecology of multidrug resistance in bacteria S. Lehtinen  | S14.O14  Evolution of male pregnancy reveals remodelling of vertebrate adaptive immunity  O. Roth                  |
| NESDAY, AUGUSTA 16.30 16.30 16.45 | S13.06 Geographic heterogeneity in parallel evolution – three spined sticklebacks revisited B. Fang                            | S21.06 Paint it red: co-option of MYB transcription factors shift color hue in a hummingbird-pollinated species A. Berardi         | S6.06 Associations between sensitivity to antibiotics and non-antibiotic antibacterials in natural and clinical escherichia coli isolates A. Bischofberger | S14.O15 Wide pleiotropic effects of melanin pathway genes on mating behaviour and life-history traits V. Tyukmaeva |
|                                   | \$13.07 Searching for signatures of genetic adaptation to climate in bank voles R.Folkertsma                                   | S21.07  More than meets the eye?  Protective functions of red pigments in endemic Hawaiian damselflies  I. Cooper                  | S6.07  Microbiome suppresses growth and resistance evolution of Escherichia coli in a human gut microcosm  M. Baumgartner                                  | S14.O16  Evolution and function of the key digestive enzymes sucrase and maltase in vertebrates  D. Mendez-Aranda  |
| 17.00<br><b>1</b> 7.00            | S13.08 Assessing genomic vulnerability to climate change in Canada's northernmost freshwater fish, Arctic charr K. K.S. Layton | S21.08  Mitochondria-targeted molecules determine the redness of thezebra finch bill  A. Cantarero                                 | S6.08  Quantifying the impact of treatment history on plasmid-mediated resistance evolution in human gut microbiota  B. Tepekule                           | S14.O17  Developmental mechanisms of Arctic charr (Salvelinus Alpinus) adaptive divergence  K. H. Kapralova        |
| 17.15                             | S13.09 Adapting to a warming world; the molecular basis of seasonal timing in a song bird M. Visser                            | S21.09 Breaking the back of the parasite: reducing early-life burden affects nestling and adult feather colouration E. Perez-Badas | S6.09  Variation in collateral sensitivity phenotypes of Escherichia coli across genotypes and growth environments  R. Allen                               | S14.018 ENHANCing the limb: from micro to macroevolution J. P. L. Castro   |

|       | MOVE1  | MOVE2   | LOGI2  | <i>GOTO33</i>   |
|-------|--|---|--|---|
|       | S17: SELFISH GEs   | S33: AGING  | S25: ASSORT MATING   | S22: HOST PLANT   |
| 16.15 | S17.O5 Kirc, a new superfamily of massive DNA transposons A. A. Vogan  | S33.08  Ageing in free-living great tits: multimarker evidence for age-related increase in oxidative and physiological stress  M. Těšický   | S25.O5 A theoretical study of the effects of assortative mating on adaptive potential under climate change C. Godineau | S22.O5 Interactions between metal-based and organic defences: Alternative weapons against spider mites attacking tomato plants D. Prino Godinho |
| 16.30 | S17.06 T-lex3: an accurate tool to genotype and estimate population frequencies of transposable elements M. Bogaerts Márquez | S33.09 Linking early-life environment to ageing rate: the role of prenatal thyroid hormones? S. Ruuskanen                                   | S25.06  The timing of attraction as a driver of species diversification in the fall armyworm  S. Hänniger              | S22.06 Urban environments select for higher growth potential but lower herbivore resistance in Arabidopsis thaliana J. Qu                       |
| 16.45 | S17.07 The selfish endosymbiont Wolbachia exploits the sex determination of its host to achieve maximal transmission F. Chen | \$33.010  Early-life environmental quality and variability reflected in telomere lengths and lifespan in a wild mammal  S.H.J. van Lieshout | S25.07 The genetics of visual preferences in a hybrid species A. E. Hausmann   | S22.O7 Opposite Responses to Drought Induced Changes in Host Plant Quality within a Butterfly Metapopulation A. Kahilainen                      |
| 17.00 | S17.08  Dynamics of prokaryotic cell differentiation during horizontal gene transfer R. Miyazaki                             | S33.011  Fitness consequences of germline mutation accumulation: the hidden cost of lifespan extension?  E. Duxbury                         | S25.08  Factors mediating reproductive isolation between related species at contact zones  A. Kirschel                 | S22.08 Chemical defences in a Heliconius butterfly and its Passiflora host A. Mattila   |
| 17.15 | S17.09  Molecular evolution of the Greenbeard Social b supergene in the fire ant Solenopsis invicta Q. Helleu                | S33.012 Using Wild Crickets to test key predictions of life-history theories of senescence T. Tregenza                                      | S25.09  Reinforcement and assortative mating between incipient outcrossing and selfing Clarkia species  D. Moeller     | S22.09 From monophagy to oligophagy, ecological and genetic variation affect host-associated diversification of butterfly species R. Mattos     |

| 8.55  | ESE  | 3 initiatives and   | practical informa  | ation  |  |  |
|-------|--|---|--|--|--|--|
| 9.05  | ESEB initiatives and practical information  KEYNOTE III David Queller, Evolutionary conflict and molecular arms races in cooperative systems |   |  |  |  |  |
|       | S13: ADAPT GEN   | S21: COLOUR   | S26: SEX CONFLICTS   | S3: NON-GEN INHERIT  |  |  |
| 10.00 | \$13.010 Altitude shapes local adaptation in Heliconius butterflies G. Montejo-Kovacevich  | S21.010  Evolutionary decoupling of larval and adult colour in shield bugs: comparative and experimental evidence  I. Medina  | S26.O1 Sexual conflict in ecological context in a semiaquatic bug J. Perry                               | S3.O1 Transgenerational inheritance of small RNAs in C.elegans I. Lev  |  |  |
| 10.15 | \$13.011  An integrative perspective of adaptation to different altitudes in an alpine plant  A. Szukala                                     | S21.011  Red or dead: imperfect   Müllerian mimicry between burnet and red,   not yellow, wood tiger   moths   B. Rojas       |  |  |  |  |
| 10.30 | S13.O12  Bacterial adaptations –  NOT what you thought  O. Avram   | S21.O12  Beyond the 'red edge': does visual sensitivity to long wavelengths facilitate resource location in beetles? LY. Wang | S26.O2  The ecology of sexual conflict and the population genetic consequences of mate choice  H. Rundle | S3.O2 Inherited effects of parental environment: Multi-generation GxE and the unscripted phenotype S. Sultan |  |  |
| 10.45 | S13.O13  Back to the future of bacterial population genomics  J. Cury  | S21.013 Iridescence as camouflage K. Kjernsmo   |  |  |  |  |
| 11.00 | CC   | <br> <br>  DFFEE & EXHIBIT  | ION & OUTREA   | <br>   |  |  |

|       | MOVE1  | MOVE2  | LOGI2   | <i>GOTO33</i>   |  |  |  |
|-------|--|--|---|---|--|--|--|
| 8.55  | ESEI   | B initiatives and  | practical informa   | ation   |  |  |  |
| 9.05  | KEYNOTE III David Queller, Evolutionary conflict and molecular arms races in cooperative systems                         |  |   |   |  |  |  |
|       | S29: ECO-EVO   | S6: ANTI-MICRO RESIST  | S12: WILD PLANT SEL   | S16: MITO-NUCLEAR   |  |  |  |
| 10.00 | S29.01  Quantitative eco-evolutionary dynamics: Numerical signatures of varying sources of phenotypic novelty  J. Pantel | S6.O10  Microbiota inoculum composition affects holobiont assembly and host growth in Daphnia  E. Decaestecker   | S12.01 Evolvability, selection, and disrupting mechanisms in the wild: A roadmap for evaluating adaptive evolution B. Pujol | S16.O1  Effects of mitonuclear genomic interactions on ATP synthesis and developmental time  R. S. Burton |  |  |  |
| 10.15 |  | S6.O11 Prophages increase bacterial fitness in the presence of high antibiotic concentrations C. Wendling  |   |   |  |  |  |
| 10.30 | S29.02 When do eco-evolutionary feedbacks aid adaptation, and when do they hinder them? T. Coulson                       | S6.O12 Evolutionary rescue in the face of an arbitrarily moving optimum in asexuals G. Martin  S6.O13 Do antibiotic treatments accelerate evolution? Population dynamics matter! A. Frenoy | Flower evolution in the wild under stable and changing pollination environments  M. C. Castellanos                          | S16.O2 The impact of mito-nuclear interactions from OXPHOS to genome evolution K. Montooth                |  |  |  |
| 11.00 | CC   | OFFEE & EXHIBIT<br>(Art up your evolution, Out   | TION & OUTREAC<br>treach stage, Teatro lobby)   | СН  |  |  |  |

|       | LOGOMO HALL   | TEATRO  | <b>GALLERIA</b>  | LOGI1   |
|-------|---|---|--|---|
|       | S13: ADAPT GEN  | S21: COLOUR   | S26: SEX CONFLICTS   | S3: NON-GEN INHERIT   |
| 11.30 | S13.014 Linking allele-specific expression and natural selection in wild populations R. Laso-Jadart | S21.014  Variation in thin film structure produces diverse visual appearances in Christmas beetles (Scarabeidae - Rutelinae)  L. Ospina | S26.O3  Ecology and sexual conflict drive the macroevolutionary dynamics of female-limited colour polymorphisms  B. Willink                | S3.O3  The ecological consequences and evolutionary potential of transgenerational temperature plasticity in Mimulus  J. Colicchio          |
| 11.45 | \$13.015 Linking a mutation to survival in wild mice S. Laurent                                     | S21.015  The hidden side of wing transparency in Lepidoptera C. Pinna   | S26.O4  The role of alternative splicing in the evolution of sexual dimorphism  T. Rogers  | S3.O4  The role of epigenetic mechanisms in within and between generation phenotypic plasticity in Neurospora crassa  I. Kronholm           |
| 12.00 | S13.016 The speciation supergene in wild Petunia: structure and evolution T. Tenreira               | S21.016  Different ways to make red flowers: Colour evolution in the New World Gesneriaceae  E. Ogutcen                                 | S26.05 Inter-population variation in morphology reflects different trajectories of sexually antagonistic coevolution in a beetle C. Koshio | S3.O5 Genotype-specific integration of genetic, nongenetic and environmental cues shapes water flea development and life history  E. Harney |
| 12.15 | S13.017  Adaptation in the wild - a systems genetics approach using Daphnia  D. Becker              | S21.017 Evolving rainbows: deriving a spectrum of phylogenetic signals in avian colour evolution S. M. Drobniak                         | S26.O6  Dynamics of sex biased gene expression during development in a hemimetabolous insect  J. Djordjevic                                | \$3.06  DNA methylation facilitates adaptation to ocean salinity change  M. J. Heckwolf   |
| 12.30 | S13.018  Contemporary natural selection on transcript abundance in wild brown trout  F. Ahmad       | S21.018 Climate shapes near-infra- red reflectance properties in birds and butterflies D. Stuart-Fox                                    | S26.O7  Toxic males to gentle courters: evolutionary reduction in sexual antagonism due to shift in life-history  B. Nandy                 | \$3.07 The role of DNA methylation in adaptation – social spiders as a case study T. Bilde  |
|       |   |   | i .  | İ   |
| 12.45 | LUNCH & EX  | HIBITION & SAT  | ELLITE EVENTS  | / OUTREACH  |
| 12.45 | LUNCH & EX  | Satellit  | ELLITE EVENTS te events treach Stage in Teatro lobby   | / OUTREACH  |

|       | MOVE1  | MOVE2  | LOGI2  | <i>GOTO33</i>   |
|-------|--|--|--|---|
|       | S29: ECO-EVO   | S7: HUMAN-INDUCED  | S12: WILD PLANT SEL  | S16: MITO-NUCLEAR   |
| 11.30 | S29.O3 Rapid Change in Mammalian Eye Shape Is Explained by Activity Pattern J. Baker   | S7.018  Fluctuating selection and enhancing diversity to overcome insecticide resistance evolution  R. Mangan                  | S12.O3  Benefits of using non-linear path analysis for estimating natural selection  G. H. Bolstad                               | S16.O3  Do mitolineages and sex-linked mitonuclear genotypes impact respiration, metabolic performance and hybrid fitness?  A. Pavlova    |
| 11.45 |  |  |  |   |
|       | S29.04 Extinction and the temporal distribution of macroevolutionary bursts S. De Lisle  | S7.019  The consequences of domestication to the wheat microbiome biodiversity  E. Özkurt                                      | S12.04  Does selection on plants defense strategies vary along a successional gradient?  A. Kalske                               | S16.04 Divergent mitochondrial and nuclear OXPHOS gene are candidates for genetic incompatibilities in Ficedula Flycatchers A. Qvarnström |
| 12.00 | S29.05 Intraspecific variation alters ecological properties and fosters transgenera- tional carry-over effects as much as temperature variation A. Raffard | S7.O20 Climate change and Green Sea Turtle sex ratio: preventing possible extinction J. Blechschmidt                           | S12.O5  Measuring viability selection from prospective cohort mortality studies in wild plant populations J. J. Robledo-Arnuncio | S16.05 The genetics of sex-biased hybrid incompatibility in Tigriopus californicus E. Watson  |
| 12.15 | S29.06  Density-dependent selection on exploration behaviour across multiple great tit populations  A. Mouchet   | S7.021 Expanding thermal breadth facilitates adaptation of Daphnia to raising temperature M. Dziuba                            | S12.06 Fitness consequences of hybridization between fully inbred lines from natural predominantly selfing populations J. Clo    | \$16.06  SmithRNAs, a new arena for mito-nuclear interaction and coevolution  M. Passamonti   |
| 12.30 | S29.07  Dynamic of introgression during density-dependent range expansion: European wildcats as a case study  C. S. Quilodrán                              | S7.O22 Invasion success of the Asian tiger mosquito in Europe: pre-adaptation, post-introduction evolution, or both? S. Sherpa | S12.07 Herbivory drives evolution of genetic architecture for plant defense and competition traits  A. Uesugi                    | S16.07 Mito-nuclear interactions in an emerging hybrid species - Insights form a time series transcriptomic study E. Iwaszkiewicz         |
| 12.45 | LUNCH & EX   | HIBITION & SAT   | ELLITE EVENTS  | / OUTREACH  |
|       |  | Satellit   | e events   |   |
| 13.45 |  |  | treach Stage in Teatro lobby   |   |
| 18.00 |  | EXCU   | RSIONS   |   |

|              |       | LOGOMO HALL  | <b>TEATRO</b>  | <b>GALLERIA</b>   | LOGI1  |  |  |
|--------------|-------|--|--|---|--|--|--|
| _            | 8.55  | ESEE   | 3 initiatives and  | practical informa   | ation  |  |  |
| _            | 9.05  | KEYNOTE IV Anna-Liisa Laine, What keeps pathogens in check in the wild?  |  |   |  |  |  |
|              |       | S13: ADAPT GEN   | S2: EXP EVOL   | S26: SEX CONFLICT   | S15: ANCIENT DNA   |  |  |
| T 23         | 10.00 | S13.019 The genetic and physiological basis of local adaptation across latitudinal range in 360 Arabidopsis accessions Y. Yarkhunova | S2.O1 Replaying the tape of life: the experimental study of adaptive evolution in seed beetles G. Arnqvist | S26.08 The genetic architecture of sexually dimorphic traits: gene knock-outs and sex-specific genetic variance W. van der Bijl | S15.O1  Modern tools for ancient Data: Quantifying evolution from paleogenomes D. Wegmann  |  |  |
| IDAY, AUGUST | 10.15 | \$13.020 Repeated Genomic Signatures of Local Selection in Atlantic Salmon V. Pritchard  |  | S26.09 Sex-specific transcriptomic responses to changes in the nutritional environment F. Camus                                 |  |  |  |
| IIDAY, /     | 10.30 | S13.021 Dissecting the transcriptomic basis of phenotypic evolution in an aquatic keystone grazer D. Frisch                          | S2.O2  Experimental adaptation to juvenile malnutrition: insights from and challenges of omics  T. Kawecki | S26.O10  Male sexual trait interacts with environment in determining female fitness A. Plesnar-Bielak                           | S15.02 The genetic history of the Plague: From the Stone Age to the 18th century J. Krause |  |  |
|              | 10.45 | S13.022 Genetic and morphological bases of a complex innovation – pelvic brooding in Sulawesi ricefishes  J. Schwarzer               |  | S26.011 Substantial sex differences in recombination in a threatened passerine with high levels of sexual conflict A. Santure   |  |  |  |

COFFEE & EXHIBITION & OUTREACH
(Art up your evolution, Outreach stage, Teatro lobby)

11.00

|       | MOVE1   | MOVE2   | LOGI2  | <i>GOTO33</i>   |  |  |  |
|-------|---|---|--|---|--|--|--|
| 8.55  | ESEB initiatives and practical information  |   |  |   |  |  |  |
| 9.05  | KEYNOTE IV Anna-Liisa Laine, What keeps pathogens in check in the wild?   |   |  |   |  |  |  |
|       | S27: SOCIAL TRAITS  | S1: TRANS GEN PLAST   | S19: GENO-PHENO  | S24: MICROBIAL STRESS   |  |  |  |
| 10.00 | S27.O1 The evolutionary implications of sociality: Population structuring associated with shifts in life history and behavior S. Shultz | S1.01 Stress responses within and across generations: From epigenetic regulation to selection in the wild M. Saastamoinen | S19.O1 Colour evolution in birds and butterflies: From macro to micro and back again N. Nadeau                             | S24.O1  Bacterial capsules as key referees in adaptation  O. Rendueles  |  |  |  |
|       |   |   |  |   |  |  |  |
| 10.30 | S27.02 Epistemology and non-discrimination: Inclusive fitness still on top A. Grafen  | S1.02 Insect immune memory, how does it work and why should we care? S. Barribeau   | S19.O2  A phylogenetic framework for the detection of trait-dependent shifts in patterns of sequence evolution  I. Mayrose | S24.02  Phage-bacteria coevolution in the rhizosphere:  Consequences for microbiome functioning and plant disease outbreaks  VP. Friman |  |  |  |
| 10.45 |   |   |  |   |  |  |  |
| 11.00 | CC  | OFFEE & EXHIBIT<br>(Art up your evolution, Out  | TION & OUTREA<br>treach stage, Teatro lobby)   | СН  |  |  |  |

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|       | LOGOMO HALL  | <b>TEATRO</b>   | <b>GALLERIA</b>  | LOGI1  |
|-------|--|---|--|--|
|       | S13: ADAPT GEN   | S2: EXP EVOL  | S26: SEX CONFLICT  | S15: ANCIENT DNA   |
| 11.30 | S13.023 Extreme morphological and genomic divergence underlies deep-water adaptation in Arctic charr (Salvelinus alpinus) morphs T. Kess | S2.03  Larval resource competition alters capability of adult reproductive interference  W. Mukaimine                                 | S26.012 Sexual conflict in the light of Caenorhabditis nematodes J. Palka  | \$15.03 6,500-year-old Salmonella enterica genomes link human-host adaptation to animal domestication A. Herbig                              |
| 11.45 | S13.024 Exploring the joint effects of global and local selection on the emergence of reproductive barriers G. Bisschop                  | S2.04 Microevolutionary genomic signatures of sexual selection R. R. Snook  | S26.013  Sex-specific adaptation to a high temperature in Drosophila SK. Hsu   | S15.04  2,000-year-old pathogen genomes reconstructed from mummies provide insights into the health status of ancient Egyptians  J. Neukamm  |
| 12.00 | S13.025 Genetic effects on phenotypic 'predictability' of guppy stress-response behaviour P. M Prentice                                  | S2.05 Experimental evolution study on Drosophila melanogaster: manifold consequences of adaptation to unfavourable diets E. lakovleva | S26.014  Sexual conflict over genes related to immunity: Evidence from a species with strong sexual selection J. Roved               | S15.O5 Studying the evolution of host-associated microbiome through time using ancient dental calculus K. Guschanski                         |
| 12.15 | \$13.026  On (small) step at a time: Measuring adaptive potential of yeast populations under different stresses I. Fragata               | S2.06 Consequences of adaptation to juvenile malnutrition on adult metabolism C. Dupuis   | S26.015  Uncovering the role of sexually antagonistic selection on sex differences in immunity in Drosophila melanogaster  S. Sharda | S15.06  Von Linné to today: -omics-based investigations of fungal adaptations to extreme environments with herbarium specimens  B. H. Conlon |
| 12.30 | LUNCH & E  |   | TELLITE EVENTS   | / OUTREACH   |
|       | Meet   | <b>Sate</b> i<br>he editors – a Royal Society Pu:   | llite events   | 12.00 12.50  |

SciSparks, how to organise speed meetings in high-schools, Outreach stage in Teatro lobby at 12:45-13:55

|       | MOVE1  | MOVE2  | LOGI2   | <i>GOTO33</i>  |
|-------|--|--|---|--|
|       | S27: SOCIAL TRAITS   | S1: TRANS GEN PLAST  | S19: GENO-PHENO   | S24: MICROBIAL STRESS  |
| 11.30 | S27.03 The evolution of mechanisms to divide labour G. Cooper  | S1.O3  Does maternal behavioural plasticity facilitate the evolution of viviparity?  A. Pettersen                                    | S19.03 Gene expression evolution in Lake Tanganyika cichlid fishes: Novel insights through data integration A. El Taher                       | S24.O3 Antibiotic stresses modify the evolution of Pseudomonas aeruginosa phage resistance T. Dimitriu                                       |
| 11.45 | S27.04  Molecular signatures of kin selection: Are caste-associated genes nearly neutral?  G. Thompson                   | S1.04  Paternal contribution to transgenerational plasticity of the freshwater snail Physa acuta in response to predation  J. Tariel | S19.04 Differential gene expression underlying caste- and sex-specific gonad development in the honey bee (Apis mellifera) D. Cavalcante Lago | S24.O4  Bacterial biodiversity drives the evolution of CRISPR-based resistance against phage E. Alseth                                       |
| 12.00 | S27.05  Benefits of cooperation and its life-history costs in complex environments in a social pine sawfly  C. Lindstedt | S1.05 Adaptive significance of Anticipatory Maternal Effects in Drosophila melanogaster P. Kohlmeier                                 | \$19.05 A genetic and evolutionary perspective on foot feathering in a domestic avian species C. Bortoluzzi                                   | S24.05 Eco-evolutionary dynamics in a simple Cystic Fibro- sis-like bacterial communi- ty treated with a low antibiotic concentration J. Law |
| 12.15 | S27.06 Dispersal strategies of sessile superorganisms: the evolution of dispersal in ants S. Hakala                      | S1.06  The role of genetic adaptation and phenotypic plasticity in response to changing salinity conditions  H. Goehlich             | S19.06 The molecular basis of phenotypic evolution across a genus: cold acclimation in Drosophila N. Cook                                     | S24.06 The Evolutionary Design of the Type-6 Secretion System W. Smith   |
| 12.30 | LUNCH & EX   | (HIBITION & SAT  | ELLITE EVENTS   | / OUTREACH   |
|       |  | Satel  | lite events   |  |
|       | Meet   | the editors – a Royal Society Pu   | blishing workshop, MOVE 1 at  | 13:00-13:50  |

SciSparks, how to organise speed meetings in high-schools, Outreach stage in Teatro lobby at 12:45-13:55

|       | LOGOMO HALL  | TEATRO  | GALLERIA   | LOGI1   |
|-------|--|---|--|---|
|       | S13: ADAPT GEN   | S2: EXP EVOL  | S26: SEX CONFLICT  | S15: ANCIENT DNA  |
| 14.00 | S13.O27 Interplay of microbiome and transcriptome shapes fitness in response to environmental change  J. Beninde | S2.07 Experimentally altered sex ratios and the evolution of sex-specific life histories J. Stångberg                   | S26.016 Sex-limited experimental evolution on a simultaneous hermaphroditic flatworm leads to differential responses of sex allocation Q. Li | S15.07 The demographic history of woolly rhinoceros E. Lord   |
| 14.15 | S13.028 Identification of chromosome subpopulations by recombination differences C. Ruiz-Arenas                  | S2.08  Evolution of reproductive efficiency in Caenorhabditis elegans under introduced obligatory outcrossing  W. Antoł | S26.017 Intersexual conflict over seed size is stronger in more outcrossed populations of a mixed-mating plant A. Raunsgard                  | S15.08  Discovering the Legacy of Atlantic cod exploitation using ancient DNA  G. Ferrari   |
| 14.30 | \$13.029  Understanding sex differences in crossing-over patterns  M. Kivikoski                                  | S2.O9 Parental care relaxes selection and increases genetic variation S. Pascoal  | S26.O18 Coevolution of female fidelity and male help under interactions between intra- and inter-locus sexual conflict Xiang-Yi Li           | S15.09  The aboriginal heritage project and the modern human colonization of Australia  J. Teixeira                                   |
| 14.45 | \$13.030 Structural variants in a haplotype-resolved hybrid rabbit genome E. Enbody                              | S2.O10  Dynamic phenotypic plasticity evolves in response to experimental environmental predictability  C. Leung        | S26.O19 Sexual conflict and the diversity of warning patterns in Heliconius butterflies M. Freire  | S15.O10  The population dynamics of eastern Siberia revealed by Lake Baikal region H. Yu  |
| 15.00 | S13.O31  Positive selection on sociobiological traits in invasive fire ants  E. Privman                          | S2.011 The Genomics of Selfing in Maize (Zea mays ssp. mays): Catching Purging in the Act A. Muyle                      | S26.O20 Sexually antagonistic coevolution between the sex chromosomes of Drosophila melanogaster C. Olito                                    | S15.O11 Genome-wide ancient-DNA investigation characterizes a genetic contact point in the Eneolithic southwestern Russia K. Majander |
| 15.15 | S13.032 Genomic architecture underlying the evolution of a novel form of social organisation R. Pracana          | S2.012 Optimizing the power to identify the genetic basis of complex traits with E&R studies C. Vlachos                 | S26.O21 Sexually-antagonistic selection on dispersal in a cooperatively-breeding bird J. Green   | \$15.012  Genes and language in the prehistory of Uralic-speaking peoples  O. Vesakoski   |
| 15.30 | CO   | FFEE & EXHIBIT<br>(Art up your evolution, Out   | TION & OUTREAC<br>creach stage, Teatro lobby)  | СН  |

|       | MOVE1   | MOVE2   | LOGI2   | <i>GOTO33</i>  |  |
|-------|---|---|---|--|--|
|       | S27: SOCIAL TRAITS  | S1: TRANS GEN PLAST   | S19: GENO-PHENO   | S24: MICROBIAL STRESS  |  |
| 14.00 | S27.07 Helping Results in Indirect Fitness Gains in Cooperative Birds P. Downing  | S1.07 Evolutionary insights into transgenerational effects of pesticides V. Castaño-Sanz  | S19.07 The evolutionary history of Alba, a trans-specific Alternative life history strategy K. Tunström                 | S24.07 Long lasting infections select for poorly transmitted bacterial variants M. Cambon                                      |  |
| 14.15 | S27.O8  | \$1.08  | S19.08  | \$24.08  |  |
|       | The design of the social hierarchy in spotted hyenas A. Courtiol  | Longer life span is associated with elevated immune activity in a seasonally polyphenic butterfly  T. Esperk                    | Evolution of photoperiodic<br>flowering and the VRN2/-<br>CO9 genes in temperate<br>Pooideae grasses<br>S. Fjellheim    | Lysed bacterial cells inhibit population growth in multiple bacterial species  F. Smakman                                      |  |
| 14.30 | S27.09 Social organization in ungulates: revisiting Jarman's hypotheses K. Szemán   | S1.09 Trans-generational effects of prenatal thyroid hormones in a wild bird species T. Sarraude                                | S19.09 The genetic underpinnings of bird beak shape morphological evolution on a macroevolutionary scale T. Gossmann    | S24.09 Artificial selection for cooperative degradation of toxins in small bacterial communities B. Vessman                    |  |
| 14.45 | S27.O10 The fitness benefits of   | \$1.010  The effect of early-life   | \$19.010<br>Many options, few   | S24.O10 The evolution of mass  |  |
|       | living with kin in a long-lived, social mammal  E. Lynch  | stress on DNA methylation<br>and exploratory behaviour<br>in wild great tits<br><b>B. Sepers</b>                                | solutions: over 60 million<br>years snakes converged on<br>few optimal venom<br>formulations<br>A. Barua                | suicide in bacterial warfare  E. Granato   |  |
| 15.00 | S27.011   | \$1.011   | \$19.011  | \$24.011   |  |
|       | Towards richer game-theo-<br>retical models: How does<br>uncertainty about the<br>social environment<br>influence reproductive<br>skew?<br>L. Olivier | Symbiont-mediated<br>maternal effects on<br>pathogen resistance in the<br>pea aphid, Acyrthosiphon<br>pisum<br><b>M. Hasoon</b> | A codon model for associating phenotypic traits with altered selective patterns of sequence evolution  K. Halabi        | Positive linkage between public goods suggests that generalist producers prevail in natural Pseudomonas communities  J. Kramer |  |
| 15.15 | S27.012  Human behaviour in economic games/social-dilemmas: designed to benefit the group, or the actor?  M. Burton-Chellew                           | S1.012  Role of epigenetic mechanisms during evolutionary adaptation to chronic malnutrition  B. Erkosar                        | S19.012  Phylogenetic comparative approaches to uncover the genomic basis of species' phenotypic differences  M. Hiller | S24.O12 Eco-evolutionary approach to species coexistence T. Hiltunen   |  |
| 15.30 | CC  | FFEE & EXHIBIT  | TION & OUTREAC  | СН   |  |
|       | (Art up your evolution, Outreach stage, Teatro lobby)   |   |   |  |  |

|  |              | LOGOMO HALL  | TEATRO   | GALLERIA   | LOGI1   |
|--|--------------|--|--|--|---|
|  |              | S13: ADAPT GEN   | S2: EXP EVOL   | S26: SEX CONFLICT  | S3: NON-GEN INHERIT   |
| 7 N  | 6.00         | S13.O33  Predation effects on fitness: genotype-phenotype mapping in Daphnia  M. Cordellier  | S2.013 Identifying the mechanisms that underlie adaptation against oral bacterial infection in D. melanogaster T. Paulo  | S26.O22 The genetic architecture of intra-locus sexual conflict in a pedigreed wild population L. Peters   | S3.08 What is 'non-genetic' inheritance? Insights from Molecular-Evolutionary Crosstalk I. Adrian-Kalchhauser                             |
|  | 6.15         | \$13.034  The contribution of pleiotropy to repeatable patterns of genomic divergence in threespine stickleback  D. Rennison                   | S2.014 Environmental heterogeneity disrupts the symmetry of host-parasite reciprocal selection, driving predictable variation in coevolutionary outcomes S. Auld | \$26.023 Sex differences in genetic underlying of personality traits S. Kralj-Fišer  | S3.09 Horizontal transmission and evolution of microbe-induced cooperation O. Lewin-Epstein   |
| TODAY, AUGUST  | 6.30         | S13.O35 Independent evolutionary trajectories underlie winter coat colour polymorphism in mountain hares I. Giska                              | S2.015  The feedback between selection and demography shapes coevolutionary genetic change  C. Retel   | S26.O24 Temperature as a modulator of sexual selection and sexual conflict P. Carazo   | \$3.010 (In)exhaustible suppliers for evolution? Epistatic selection tunes the adaptive potential of non-genetic inheritance  \$. Charlat |
|  | 6.45         | S13.O36 Characterizing the genetic basis of adaptation to arid environments in Drosophila melanogaster European populations V. Horvath         | S2.016  The home advantage: Ancestral microbes aid host adaptation to novel environments A. Agarwal  | S26.O25 Sex-biased gene expression is repeatedly masculinized in asexual females D. Parker   | S3.011  The impacts of epigenetic variation on the rate of speciation with gene flow P. Greenspoon  |
| 1  | 7.00         | S13.O37  Combining drug metabolism phenotypes and genomic diversity to understand evolution in metabolism of exogenous substances  M. Mouterde | S2.017  Can parasite evolution reinforce the effects of climate warming?  J. Wolinska  | S26.O26  Contrasting rates of molecular evolution in reproduction-related genes in Macrostomum flatworms with different reproductive strategies  R. A. W. Wiberg | S3.012  Early-exposure to new sex pheromone blend alters mate preference in butterflies and in their offspring  E. Dion                   |
| THE RESERVE OF THE PARTY OF THE | 7.20<br>9.20 |  | POSTER S   | SESSION II   |   |

|                | MOVE1   | MOVE2   | LOGI2  | <i>GOTO33</i>  |
|----------------|---|---|--|--|
|                | S27: SOCIAL TRAITS  | S36b: PHYLOGEO & SYST   | S36d: GENOME EVOL  | S36c: SPP INTERACT   |
| 16.00          | S27.013 A trait-based approach to map behaviour across species M. E. Herberstein                          | S36b.O6 Admixture among North American Canids: coyotes, wolves and the beasts between A. Carmagnini                     | S36d.O6  Polyploidy and floral evolution in a highly variable, coevolving plant species  K. Gross  | S36c.O1  Herbivores and plant defences affect selection on plant reproductive traits more strongly than pollinators  J. Santangelo               |
| 16.15          | S27.014  Bellicose bias: how sex differences in dispersal influence intrasexual aggression  E. Bath       | S36b.O7 Enriching conserved genomic elements to resolve relationships among sawflies W. Saskia                          | \$36d.07  Mutation-rate plasticity and the germline of unicellular organisms  D. Aanen   | S36c.O2 Evidence for a chemical arms race: Lections from a chemical mimicry system of cuckoo wasps T. Schmitt                                    |
| 16.30          | S27.015 The Strategic Reference Gene: an organismal theory of inclusive fitness L. Fromhage               | S36b.08  ddRAD sequencing reveals the evolutionary history of the snail Charpentieria itala in the Southern Alps  J. Xu | S36d.O8 Genomic introgression through newt hybrid zones – evidence from replicated transects P. Zieliński                                | S36c.O3  Mutualism mediates infection risk by an antagonist in experimental populations  J. Eck  |
| 16.45          | S27.016  Genotype-by-environment interactions on sociability in threespine sticklebacks  N. Pilakouta     | S36b.O9 Environmental variables shaping the distribution and hybridization in Heliconius butterflies N. Rueda           | S36d.O9  The role of chromosomal inversions in the speciation history of two Drosophila virilis group species  N. Poikela                | S36c.O4  Fitness trade-offs associated with host resistance in a natural insect host-ectoparasite symbiosis  M. Polak                            |
| 17.00          | S27.017  Ecological and social constraints promote social evolution in the clown anemonefish  R. Branconi | S36b.O10  Phylogenomics of the Hyalella (Amphipoda: Crustacea) species-flock in Lake Titicaca, High Andes F. Zapelloni  | S36d.O10  Exposure to environmental radionuclides associates with altered metabolic and immunity pathways in a wild rodent  J. Kesäniemi | S36c.O5  High conspecific density reduces hoarding success and affects sex-specific spatial distribution among wintering pygmy owls  E. Koivisto |
| 17.20<br>19.20 |   | POSTER S  | SESSION II   |  |

|       | LOGOMO HALL  | TEATRO   | GALLERIA  | LOGI1   |  |
|-------|--|--|---|---|--|
| 8.55  | ESEI   | 3 initiatives and  | practical informa   | ation   |  |
| 9.05  | KEYNOTE V Rasmus Nielsen, Human adaptation in time and space   |  |   |   |  |
|       | S13: ADAPT GEN   | S2: EXP EVOL   | 336a: SEX & SELECT MATING   | S28: GAME THEORY  |  |
| 10.00 | S13.038  The impact of protein architecture on adaptive evolution  A. F. Moutinho                        | S2.018 The genetic and molecular bases of real-time bacterial tRNA evolution J. Gallie             | S36a.O1  How diversity in parental care evolves: a phylogenetic comparative study in amphibians  A. Furness             | S28.01 Improving treatment of metastatic cancers through evolutionary game theory K. Stankova               |  |
| 10.15 | \$13.039  Adaptation to high soil trace metal element concentrations in Arabidopsis arenosa  C. Sailer   | S2.019  Bacterial predator-prey coevolution selects on virulence-associated prey defences  R. Nair | S36a.O2 Cobreeding females adjust their reproductive decisions by investing more in eggs and less in care J. Richardson |   |  |
| 10.30 | \$13.040  Regulatory evolution of metabolic adaptations in cavefish  N. Rohner                           | S2.O20 Forecasting experimental evolution in Pseudomonas P. Lind                                   | S36a.O3 Cannibalism rescues fitness impacts of skewed sex-ratios in red flour beetle Tribolium castaneum I. Khan        | S28.02  N-player collaborative hunting in yellow yellow saddle goatfish (Parupeneus cyclostomus)  R. Bshary |  |
| 10.45 | S13.041  True survivors: response to bat fungal pathogen varies according to exposure history  T. Lilley | S2.O21 Evolution of Multicellularity: Cheating Done Right W. Veit                                  | S36a.O4  Parental investment and sexual dimorphism in immunity  V. Revathi Venkateswaran                                |   |  |
|       |  | i  |   | I   |  |

|       | MOVE1  | MOVE2  | LOGI2   | <i>GOTO33</i>   |  |  |  |
|-------|--|--|---|---|--|--|--|
| 8.55  | ESEB initiatives and practical information   |  |   |   |  |  |  |
| 9.05  | KEYNOTE V Rasmus Nielsen, Human adaptation in time and space                                     |  |   |   |  |  |  |
|       | S27: SOCIAL TRAITS   | S30: POLLINATOR  | S9: MICROBES & FOOD   | S5: AGING & CANCER  |  |  |  |
| 10.00 | S27.018 Cooperative adaptations and exploitation resistance in social amoebae J. Strassmann      | \$30.01 Preparedness and contra- preparedness in pollinator learning A. Dunlap   | S9.01  Domestication of microbial communities for bread making: insights from a participatory research project  D. Sicard | S5.O1  Cancer resistance mechanisms in long-lived mammals  V. Gorbunova |  |  |  |
| 10.15 | S27.019 Siderophore investment strategies in Pseudomonas aeruginosa S. Mridha                    |  |   |   |  |  |  |
| 10.30 | S27.O20 The social control of virulence and the mystery of defective viruses A. Leeks            | S30.02  Eco-evolutionary feedbacks between floral traits and pollinator behaviour in deceptive pollination interactions A. Ellis | S9.O2 The fungal genus Aspergillus as a model to study microbial domestication J. Gibbons                                 | S5.O2 Evolutionary genomics, aging and cancer J. P. de Magalhaes        |  |  |  |
| 10.45 | S27.O21 Evolutionary Forces Behind the Diversification of Public Goods in Bacteria A. Figueiredo |  |   |   |  |  |  |
| 11.00 | COFFEE & EXHIBITION & OUTREACH (Art up your evolution, Outreach stage, Teatro lobby)             |  |   |   |  |  |  |

|                | LOGOMO HALL   | <b>TEATRO</b>   | <b>GALLERIA</b>  | LOGI1  |  |
|----------------|---|---|--|--|--|
|                | S13: ADAPT GEN  | S2: EXP EVOL S  | 36a: SEX & SELECT MATING   | S28: GAME THEORY   |  |
| 11.30          | S13.O42 Genomic introgression facilitated adaptation of European aspen to short growing seasons in northern Scandinavia M. Rendón-Anaya                           | S2.022 Spatial selection and experimental evolution of parasite dispersal strategies G. Zilio                         | S36a.O5  How does the environment influence the expression of animal mate choice and sexual signalling?  L. Dougherty    | S28.03  Microbial public goods games in a toxic environment: to degrade or to resist?  S. Shibasaki            |  |
| 11.45          | S13.043  Hitch-hiking laterally-acquired genes contribute to delayed adaptation  J. Olofsson  | S2.O23  The effects of predation on body and fin morphology in replicated mesocosms  N. Alioravainen                  | S36a.O6  Genetic architecture of reproductive performance in response to thermal stress  M. Zwoinska                     | S28.O4  Adaptive dynamics in spatially structured populations  T. Priklopil                                    |  |
| 12.00          | S13.044 Evidence that viruses, particularly SIV, drove genetic adaptation in natural populations of eastern chimpanzees A. Andrés                                 | S2.024  Reproductive interference as a driver of species exclusion and evolution in spider mites  M. Cruz             | S36a.O7  Experimental evidence for genetic and phenotypic effects of sexual selection on germline mutation rate  J. Baur | S28.O5 Evolutionary Dynamics of Coordinated Cooperation H. Ohtsuki   |  |
| 12.15          | S13.O45  The role of sRNA dominance modifiers in transitions to selfing in Capsella  J. Bachmann  | S2.O25  Multi dimensional niche evolution of a crop pest (Callosobruchus maculatus) under climate change  A. Leonard  | S36a.O8 Evolution of sexual signals in closely related frog species occurring in sympatry S. Goutte                      | S28.06 Effects of uncertainty and learning on the behaviour predicted by evolutionary game theory A. Higginson |  |
| 12.30          | S13.046 Environmentally dependent rewiring of epistatic networks and their contributions to quantitative trait plasticity Y. Zan                                  | S2.026  Eco-Evolutionary feedbacks in range expanding food webs: experimental evidence from small worlds E. Fronhofer | S36a.O9 Evolution of female promiscuity in songbirds J. T. Lifjeld   | S28.07 Reinforcement learning leads to bounded rationality in a public goods game O. Leimar                    |  |
| 12.45          | LUNCH & EXHIBITION  |   |  |  |  |
| 13.30          | ESEB members meeting  |   |  |  |  |
| 14.30          | Incoming president's address ophelie Ronce, Integrating niche evolution with life history theory can help us better understand the consequences of climate change |   |  |  |  |
| 15.10          | Leg stretching break  |   |  |  |  |
| 15.20          | JMS award winner 2019  Karl Grieshop, Sexual conflict and the maintenance of genetic variance in fitness  |   |  |  |  |
| 15.50<br>16.20 | Closing ceremony  |   |  |  |  |
| 18.30          | Congress dinner at Muuminworld  |   |  |  |  |
| 02.00          |   |   |  |  |  |

|                | MOVE1  | MOVE2   | LOGI2  | <i>GOTO33</i>  |  |  |  |
|----------------|--|---|--|--|--|--|--|
|                | S27: SOCIAL TRAITS   | S30: POLLINATOR   | S9: MICROBES & FOOD  | S5: AGING & CANCER   |  |  |  |
| 11.30          | S27.O22 Farming plant cooperation for more sustainable agriculture G. Montazeaud                         | \$30.03  Nectar chemistry changes pollinator behavior with implications for plant fitness  P. Jones   | S9.O3 Rapid pathogen resistance evolution can shape the biocontrol efficiency of plant growth promoting Pseudomonas bacteria S. Clough | S5.O3 Competition and cancer invasiveness in ageing landscapes S. P. Castillo                                  |  |  |  |
| 11.45          | S27.O23  | \$30.04   | \$9.04   | \$5.04   |  |  |  |
|                | Cooperation and cheating among germinating spores S. Pande   | Pollinator preference and flowering phenology: how to solve reproductive conflicts between species that share pollinators  R. Pérez-Barrales                      | Study of the domestication<br>in the blue cheese fungus<br>Penicillium roqueforti<br><b>T. Caron</b>                                   | Cancer evolution in<br>hierarchal organised<br>tissues<br>P. Ashcroft  |  |  |  |
| 12.00          | S27.024 Social plasticity in the wild K. Strickland  | S30.05  Mimicry and competition drive flower colour polymorphisms in sunbird-pollinated Erica  A. Coetzee   | S9.05  New model to assess genomic and functional effects of microbial domestication in food environments  K. Chacon-Vargas            | S5.O5 Choose your death: adaptive cell senescence predicts a late-life decrease of cancer prevalence T. Tissot |  |  |  |
| 12.15          | S27.O25  | \$30.06   | \$9.06   | \$5.06   |  |  |  |
|                | The evolution of social bet-hedging strategies <b>T. Aubier</b>  | Foraging preferences of bees and birds – assessing the adaptive value of heteranthery in Merianieae flowers  A. Dellinger   | Water kefir: metagenomic<br>analysis of a drinkable<br>symbiotic communities of<br>bacteria and yeast<br>JB. Boulé                     | Lifelong telomere dynam-<br>ics in wild Soay sheep<br><b>H. Froy</b>   |  |  |  |
| 12.30          | \$27.026 Greenbeard genes: theory and reality P. Madgwick  | S30.07 Should I stay or should I go? Diascia plants frequently shift their Rediviva pollinators  B. Kahnt   | S9.07 Characterisation of microbial communities on different apple varieties and orchard management practices E. Britt                 | S5.07 Limited longevity in a finite world J. Lehtonen  |  |  |  |
| 12.45          | LUNCH & EXHIBITION   |   |  |  |  |  |  |
| 13.30          | ESEB members meeting   |   |  |  |  |  |  |
| 14.30          | Incoming preside theory can help us better und   | Incoming president's address Ophelie Ronce, Integrating niche evolution with life history theory can help us better understand the consequences of climate change |  |  |  |  |  |
| 15.10          | Leg stretching break   |   |  |  |  |  |  |
| 15.20          | JMS award winner 2019  Karl Grieshop, Sexual conflict and the maintenance of genetic variance in fitness |   |  |  |  |  |  |
| 15.50<br>16.20 | Closing ceremony   |   |  |  |  |  |  |
| 18.30<br>02.00 | Congress dinner at Muuminworld   |   |  |  |  |  |  |
| 02.00          |  | -   | 2040.6   |  |  |  |  |

# **POSTER LIST**

### **POSTER SESSION TUESDAY 17.20-19.20**

### 4. Cognitive evolution and environment

#### **S4.P1**

Do developmental changes in fitness trade-offs predict mechanosensory cues for escape-hatching decisions? Chloe Fouilloux

#### S4.P3

Predator identification from salivary DNA left on artificial prey Daniela Rößler

Ecology of cognitive evolution in Heliconiini butterflies Fletcher Young

### **S4.P5**

The sensory basis of distance estimation in a coral reef

Cecilia Karlsson

#### **S4.P6**

Visual specialisation and expansion of Heliconius mushroom bodies

Stephen Montgomery

#### **S4.P7**

Brain size affects responsiveness in mating behavior to variation in predation pressure and sex-ratio Alberto Corral-Lopez

#### **S4.P8**

Proteomic profiling of cerebrospinal fluid in cognitively advanced birds: comparative approach Eleni Voukali

#### **S4.P9**

Evolution of emotions and learning - a neural network model

Magdalena Kozielska

Artificial selection for schooling behaviour decreases individual learning ability in fish

Regina Vega-Trejo

#### S4.P11

The evolution of foraging innovation following colonisation of a less variable environment

Gábor Herczeg

#### S4.P12

Non-nestmate templates improve nestmate recognition Volker Nehring

#### **S4.P14**

Effects of mating on female immune defence in a fruit fly Keiko Oku

Head measures as promising indices of sensory capacity: a study on geometrid moths

Juhan Javoiš

#### **S4.P16**

Causes and consequences of individual variation in cognitive ability

Krista van den Heuvel

#### S4.P17

Decision-making in wild great tits, with real world consequences

Shana Caro

#### S4.P18

Norm followers, cheaters and costly signallers in a sport charity campaign

**Judit Mokos** 

### 6. Eco-evolutionary approach to the antimicrobial resistance problem

#### S6.P1

Exploring the role of bacteria and phage genetic diversity for CRISPR-phage coevolution Jack Common

#### S6.P2

Evolution of antibiotic resistance investigated by single cell genomics

Manu Tamminen

#### S6.P3

Ecology and evolution of plasmid-mediated antimicrobial resistance (pAMR) transfer in the chicken microbiome Sarah Duxbury

#### S6.P4

Biotic stress response in Fagaceae: Focus on antimicrobial peptides

Tetyana Nosenko

#### S6.P5

Fight AMR evolution: predictive phage cocktails, plasmid-dependent phages and plasmids that re-sensitize bacteria to antibiotics

Matti Jalasvuori

### **S6.P6**

Antibiotic resistance plasmids spread at diverse rates through recipient populations, in the absence of selection

Fabienne Benz

#### S6.P7

Evolutionary instability of collateral susceptibility networks in clinical Escherichia coli strains Vidar Sørum

#### **S6.P8**

Fungal antimicrobial resistance towards termite mound defences

Nils Peereboom

Resistance management in a hospital setting: limited impact of a single drug intervention Clare Kinnear

### 7. Human-induced evolution

Breeding in an agricultural land: effects on evolutionary potential of a wild bird population Dany Garant

#### **S7.P2**

Toads respond to anthropogenic change by adjusting their chemical defence Bálint Üveges

Cat behavior: an evolutionary perspective Milla Salonen

#### **S7.P4**

Behavior and personality differences between cat breeds

Salla Mikkola

Is temperature-induced sterility important for predicting species' responses to climate change? Steven Parratt

#### S7.P6

Intra-species diversification using "elite plants" reduces herbivory and increases resilience

Tuuli-Marjaana Koski

#### **S7.P7**

Genomic consequences of eutrophication induced speciation reversal in Alpine whitefish

Philine Feulner

#### **S7.P8**

Mining herbaria and roaming the forests: Land-use and climate change affect plant phenology Franziska Merle Willems

Roles of acetylcholinesterase genes in organophosphate and carbamate resistance in Leptinotarsa decemlineata Aigi Margus

#### **S7.P10**

Population genomic differentiation of the Asian longhorned beetle (Anoplophora glabripennis) Iris Häußermann

#### S7.P11

Population genomics of the invasive Argentine ant Ida Holmberg

### **S7.P12**

Evolution of phenotypic variation of an invasive lizard following experimental introduction on small islands Amelie Fargevieille

### **S7.P13**

The ecological and genetic impact of interactions between domesticated and wild Atlantic salmon Joshka Kaufmann

### **S7.P14**

For better or for worse: How city life can alter the gut microbiome of species

Bethan Littleford-Colquhoun

#### S7.P15

Coping with change – how can chemical communication systems overcome disruptions?

Emily Burdfield-Steel

Winners and losers of the Anthropocene: evolutionary and ecological factors predict species invasion and extinction

Jacintha Ellers

#### **S7.P17**

Size-dependent harvesting modified the trophic niche of medaka Oryzias latipes in a mesocosm experiment Charlotte Evangelista

Impact of host plant range on the evolution of insecticide resistance, and vice-versa, in insects Kristina Karlsson Green

### 8. Genetics of small populations

#### S8.P1

Whole-genome sequencing helps unravel the evolutionary and demographic history of an endangered primate Catalina Gonda

#### S8.P2

Clonal architecture in the endangered populations of the shrub birch Betula humilis Schrk Agnieszka Bona

### **S8.P3**

Assessing the genomic diversity and signatures of selection in Arabian Peninsula and African dromedary camels Hussain Bahbahani

#### **S8.P4**

Is there a goose on the loose? investigating introgression into the Swedish Lesser White-fronted Goose David Diez-del-Molino

**Genetic Structure of Aedes albopictus from Asia** Jiyeong Shin

Capturing genetic variation in natural and planted stands of Picea abies using probes and WGS Helena Eklöf

Identification of the homogametic sex chromosome Charles Christian Riis Hansen

Non-parallel evolution of pelvic reduction in nine-spine sticklebacks

Petri Kemppainen

Genetic analyses of archaeological and historical barley grains

Mia Lempiäinen-Avci

#### **S8.P10**

Genetic consequences of geographical isolation: a case study of Betula nana in Poland

Katarzyna Jadwiszczak

### S8.P11

Habitat change leads to shifting allopatric boundaries in coastal and pelagic island seabirds Robin Cristofari

#### **S8.P12**

Selection efficiency in social and solitary Hymenoptera Arthur Weyna

#### S8.P13

Genetic drift during a biological invasion Eric J. Petit

#### **S8.P14**

Causes and consequences of TLR variation in a bottlenecked population

Charli Davies

Revisiting the role of inversions in maintaining genomic differentiation after secondary contact

Marina Rafajlović

#### S8.P16

Fitness consequences of dispersal in a house sparrow metapopulation

Dilan Saatoglu

#### **S8.P17**

Investigating adaptation in Swedish sand lizards Mette Lillie

#### 10. Rapid Evolutionary Adaptation: **Potential and Constraints**

Recipe for a rapid radiation: population divergence and repeated behavioral isolation through parallel genetic mechanisms

Thomas Blankers

#### S10.P2

Rapid adaptation of stress related traits in Drosophila melanogaster to seasonal changing environment Banu Sebnem Onder

#### S10.P3

Understanding rapid evolution of insecticide resistance using genomic data from 100-year old pest moths Angela McGaughran

#### S10.P4

Interspecific competition as a driver of ecological divergence in a songbirds secondary contact zone Camille Sottas

Interaction between sex and gene flow modulates speed of adaptation during range expansions Felix Moerman

### **S10.P6**

Ecological opportunity promotes diversifying selection and facilitates rapid phenotypic divergence in Icelandic Arctic charr

Matthew Brachmann

Digging up rapidly evolved traits in Itallian Wall lizard (Podarcis siculus)

Óscar Mira

Real-time evolution under climate warming: an experimental approach in populations of contrasting biogeographical history

Pedro Simões

#### S10.P9

Investigating the adaptive role of noise in gene expression

Pierre Lave

Is timing of spring arrival genetically controlled in a long-distant migrant?

Miloš Krist

#### S10.P11

Rapid phenotypic diversification among new stickleback populations created by a huge earthquake and subsequent tsunamis

Takuya Hosoki

#### S10.P12

Raccoon MHC diversity in native and introduced ranges: reduced allele diversity but not allele divergence Aleksandra Biedrzycka

#### S10.P13

The evolution of the genetic architecture of traits under artificial selection

Yvonne Wientjes

#### S10.P14

Perceived risk affects the genetic integration of behaviour and morphology in two stickleback populations Niels J. Dingemanse

#### S10.P15

How to deal with environmental changes: Molecular characterisation of (non) genetic mechanisms on invasive species

Pierre Marin

#### S10.P16

Visual habitat choice in East African cichlids Elodie Wilwert

#### S10.P17

Adaptive evolution and functional differentiation of testis expression genes in Theria

Yukako Katsura

### S10.P18

Rapid evolution in highly fecund populations Bjarki Eldon

#### S10.P20

The ghosts of evolutionary past: Phytoplankton's adaptive potential in a crowded changing world Maria Elisabetta Santelia

#### S10.P21

Sympatric stickleback in space: the role of selection and gene flow

Thijs M.P. Bal

The role of inversions in driving local and sex-specific adaptation Filip Ruzicka

### S10.P23

Effects of fungicide on a nontarget species, Colorado Potato Beetle (Leptinotarsa decemlineata) Shahed Saifullah

### S10.P24

Investigation of immune response in a seasonal Drosophila melanogaster population Ekin Demir

#### S10.P25

Genetic perspective of seasonal adaptation in Drosophila phalerata Chedly Kastally

Adaptation to seasonal fluctuations in desiccation tolerance in a natural population of Drosophila melanogaster Nur Seda Coşkun

#### S10.P27

Plasticity and epigenetic inheritance in the aphid parasitoid Aphidius ervi

Mark Lammers

#### S10.P28

Unexpected evolutionary dynamics of phenotypically important tandem repeats: a model of flocculin evolution in brewing

Colette St. Mary

### 11. Quantitative trait effect size distributions and their impact on evolutionary processes

#### **S11.P1**

Investigating genetic bases of hybrid sterility using wild-derived inbred strains from mouse hybrid zone populations

Paigan Aspinall

#### S11.P2

Genomic region including major seed dormancy QTL strongly contributes to local adaptation in Arabidopsis thaliana

Giulia Zacchello

#### S11.P3

Estimation of proportions of additive and non-additive genetic variance components

Anna-Margarete Staehler

Unravelling the mechanisms of population phenotypic changes: does individual differences matter for colonisation?

Marion Nicolaus

#### S11.P5

A polygenic network for pupation site choice behavior in Drosophila melanogaster

Wenyu Zhang

Genetic architecture of sexual dimorphism in the seed beetle Callosobruchus maculatus

Philipp Kaufmann

### **S11.P7**

Reconstructing the evolutionary dynamics of a functional variant in a spatially distributed natural population Ahmed Elfarargi

### 12. Quantifying selection and evolvability in wild plant populations: methods and measurements

#### S12.P1

Genomic quantification of selection on regions of open chromatin in Capsella grandiflora

Robert Horvath

#### **S12.P2**

Evolution of plant phenotypic plasticity to grassland management

Anna Kirschbaum

#### S12.P3

Linking dendroecology and association genetics in forest trees

Katrin Heer

#### S12.P4

Quantum speciation and genetic swamping in a range-limited Species

Mitch Cruzan

Assessing differences on evolutionary rates between haploid and diploid tissues in Pinus sylvestris Sandra Cervantes

#### S12.P6

The effects of natural and artificial selection on seedling traits in Scots pine

Outi Savolainen

#### S12.P7

Using seed banks to investigate shifts in drought resistance in four Mediterranean herbs Robert Rauschkolb

#### S12.P8

**Evolutionary-based resilience of wild populations** Ivan Scotti

### 14. The mechanisms of evolutionary change: moving from genomic signatures to functional validation

#### S14.P1

Seasonal and allele-specific expression dynamics of a major age-at-maturity gene in Atlantic salmon Jukka-Pekka Verta

#### S14.P2

Six6 expression in Atlantic salmon suggests a role in the development of body shape

Jacqueline Moustakas-Verho

#### S14.P3

Dissecting the genetic architecture underling mouth dimorphism in Pristionchus pacificus identifies a supergene locus

Mohannad Dardiry

Association study reveals genes underlying flower size plasticity in Arabidopsis thaliana

Roosa Laitinen

#### S14.P5

From GWAS to function: expression patterns of age-at-maturity-associated genes in Atlantic salmon Johanna Kurko

#### S14.P6

Genetic and nutrition effects on lipid amount and composition in juvenile Atlantic salmon

Andrew House

### S14.P7

miRNA expression during the development of different **Arctic charr morphologies** 

Dagny A. Runarsdottir

### S14.P8

Understanding functional changes of cold acclimation using ribosomal footprint profiling Jenni Prokkola

Functional validation of genes involved in fat storage in Atlantic salmon adipocytes using CRISPR Erica H Leder

### S14.P10

Bacteria-derived tail anchors target to discrete organelles when expressed in eukaryotic cells Cory Dunn

#### S14.P11

Investigating germ cell specific gene expression across panarthropods

Matthias Janeschik

#### S14.P12

Functional investigation of plasticity in wing morph determination in a water strider Erik Gudmunds

#### S14.P13

Unexpected fructan motifs in Nardus stricta: cloning, purification, and functional analysis of an invertase enzym Camilla Lorange Lindberg

#### S14.P14

Narrowing in on the Red Queen: QTL and fine mapping parasite resistance in Daphnia magna Maridel Fredericksen

#### S14.P15

Genotype-phenotype mapping of skull development and adaptation in squamate reptiles Joni Ollonen

#### S14.P16

The genetic underpinnings of bill length variation in great tits (Parus major) Judith Risse

### 17. Selfish genetic elements

### S17.P1

Sex chromosomes suppress vertical transmission of feminizing Wolbachia symbionts in an isopod Richard Cordaux

#### S17.P2

The genetic basis of meiotic drive in Podospora S. Lorena Ament Velasquez

#### S17.P3

Burst of transposable elements is associated with vertebrate differentiation Feng Shao

### S17.P4

Population genetics of a segregation distorter in fungal systems

Ivain Martinossi-Allibert

Do spider mites evolve antagonistic traits against cytoplasmic incompatibility induced by Wolbachia? Flore Zélé

Contribution of TEs mobilization to tomato improvement

Marisol Dominguez

#### **S17.P7**

The extent of DNA transfer between plasmids and chromosomes in prokaryotes

Ahmad Samer Kadib Alban

#### S17.P8

Tracing the onset of the 'green beard' signal in fire ant Solenopsis invicta Qiaowei Pan

Allorecognition genes in Basidiomycetes - a genetic stalemate? Benjamin Auxier

### 18. The genetic architecture of polygenic adaptation: sweeps, small shifts and everything in between

Tracking allele trajectories over 20 generations of selection for long limbs in mice Layla Hiramatsu

#### S18.P2

Genomic footprints of polygenic adaptation in Theobroma cacao

Tuomas Hämälä

#### S18.P3

Natural selection and the lengths of fixed chromosomal inversions

Tim Connallon

#### S18.P4

Origin and dynamics of adaptive alleles in physiological adaptation in sticklebacks Jun Kitano

#### S18.P5

A chromosomal rearrangement explains variation in age of seaward migration in Atlantic salmon Sarah Lehnert

#### S18.P6

The resolution of genomic conflicts following admixture in a polygenic hybrid incompatibility model Flávia Schlichta

### S18.P7

Polygenic adaptation of a quantitative trait Benjamin Wölfl

### 20. The evolutionary consequences of social transmission and animal culture

#### **S20.P1**

To be or not to be...when individual performance interacts with group composition Frederic Mery

#### S20.P2

Role of different types of social information in learning to avoid aposematic prey Alice Exnerova

#### S20.P3

Host group relatedness and social transmission of the gut microbiome: an experimental study on ostriches Hanna Bensch

### S20.P4

Informed movement: how information shapes the co-evolutionary outcomes of range-expansion Katja Rönkä

#### S20.P5

Social information transfer during dispersal in Zootoca vivipara: mechanisms and implications for population dynamics

Mathieu Brevet

#### S20.P6

The Neolithic transition to large-scale societies is favoured by the co-evolution of cooperation and institu-

Claire Guérin

#### **S20.P7**

Evolution of personal and social immunity in the context of family life

Michelle Ziadie

### 21. Colour across the evolutionary spectrum: from production to perception

#### S21.P1

Patterns of sex-specific selection and inheritance of a colour pattern polymorphism in an Australian lizard Genevieve Matthews

Examining the link between relaxed predation and bird colouration on islands

Louis Bliard

#### S21.P3

Sparring stomatopods: Do coloured patches signal weapon performance?

Amanda Franklin

#### S21.P4

The molecular basis of continuous flower colour in Oncocyclus irises

Esther Senden

#### S21.P5

The repeated evolution of wasp colour-pattern mimicry in hoverflies

Tom Reader

#### S21.P7

Uncovering the genomic basis of an aposematic colour polymorphism in the wood tiger moth Eugenie Charley Yen

Potential and realized costs associated with ultraviolet signals in a lizard

Arnaud Badiane

#### **S21.P9**

Changing colour in a polluted environment Asma Althomali

Sexual selection, predation, and the maintenance of polymorphic Y-linked colour genes in the Trinidadian guppy Josephine Paris

#### S21.P11

Color evolution in European butterflies evolved via Darwin's, not Wallace's, model of evolution Christopher W. Wheat

#### S21.P12

The genetics and condition-dependence of structural colour in mimetic Heliconius butterflies Melanie Brien

#### S21.P13

Sexual selection predicts the rate and direction of colour evolution in a large avian radiation Christopher Cooney

#### S21.P14

Micro-scale architecture of the blue tit feathers Katarzyna Janas

#### S21.P15

Phenotypic variation in poison frogs: From predator perception to the molecular basis of color variation Heike Pröhl

#### S21.P16

Carotenoid coloration signals a males' tendency to invest in parental care in passerines

Alejandro Gonzalez-Voyer

#### S21.P17

Genetics and selection of ventral colouration in oviparous and viviparous common lizards Hans Recknagel

#### S21.P18

Evaluation of phenotypic resemblance across multiple mimicry rings in Heliconius

Maria González-Rojas

### S21.P19

Revisiting male colour and opsin polymorphism along the predation regime continuum in the Trinidadian guppy

Lengxob 'Lenny' Yong

#### S21.P20

Stabilizing selection on individual, conspicuous colour pattern elements of an aposematic nudibranch Anne Winters

#### S21.P21

Red and green plumage colouration in a wild x domestic intercross

Jesper Fogelholm

#### S21.P22

Continuous variation in scale ultrastructure in two Heliconius mimics

Juan Enciso-Romero

The effects of genetic and environmental factors on Drosophila body color components Lafuente Elvira

### S21.P24

"Fretted with golden fire": structural colouration based on micron and sub-micron structures in selected insects Dragana Cvetković

### S21.P25

Gene expression profiling of aposematism in Anisomorpha stick insects

Victor Soria-Carrasco

#### S21.P26

The evolutionary maintenance of ontogenetic colour polymorphism in wood frogs (Rana sylvatica) Debora Goedert

Effect of natural and sexual selection on wing colour variation of butterflies Bhavya Dharmaraj

### 23. Parasite community dynamics and their role in the evolution of host immunity

Parasite-mediated sexual selection in moths Ke Gao

#### S23.P3

Social antimicrobial wound care in a predatory ant Erik Thomas Frank

#### S23.P4

Pathogens, species distribution and migration: molecular evolution of genes of immunity in cetaceans Maria Luiza Andreani

#### S23.P5

Apparent manipulation: How parasites may modify their host's behaviour without using any tricks Camilla Håkonsrud Jensen

#### S23.P6

Towards the understanding of zoonotic events in infectious diseases: Tracing animal reservoirs in Switzerland Christian Urban

#### S23.P7

Humic-acid-driven escape from eye parasites Kristina Noreikiene

#### S23.P9

Seasonal variation in endoparasite biodiversity with age and sex of semi-captive Asian elephant hosts Carly Lynsdale

#### S23.P10

Invasions create competitors: How novel interactions among native and invasive parasites modify host parasite coevolution

K. Mathias Wegner

Thermal plasticity in immunity in association with different seasonal strategies for reproduction Yara Rodrigues

#### S23.P13

Does gut passage help keep fungus-growing termite gardens disease free?

Leandro Guimaraes

#### S23.P14

Selection on MHC class II haplotypes in a free-living ruminant

Wei Huana

#### S23.P18

Less or more? Protective microbe density and defence against parasites

Georgia C Drew

Seasonal patterns of parasite infection and larval developmental mode variation in the polychaete Pygospio elegans

Anna-Lotta Hiillos

### 25. Assortative mating for quantitative traits: mechanisms, estimation, and evolutionary consequences

#### S25.P1

Hybridization reduces the variation of male sexual phenotype in F1 hybrids: A Meta-analysis Keisuke Atsumi

#### S25.P2

Is assortative fertilisation after sperm competition driven by male environment or genetics in common bedbugs?

Jana Křemenová

#### S25.P3

Assortative mating based on circalunar and circadian timing of adult emergence in Clunio marinus Sina Schirmer

#### S25.P4

Sperm fertilization capability is shaped by the gamete-level immunological incompatibility in humans Annalaura Jokiniemi

#### S25.P5

Characterising genomic patterns of divergence underpinning reproductive isolation in the Drosophila virilis group

Leeban Yusuf

#### 31. Life history evolution: bridging theory and data

#### S31.P1

Body size variation in European common lizards: a range-wide study of a wide-ranging species Evgeny S. Roitberg

#### S31.P2

Reproductive costs in eastern grey kangaroo females, the bigger picture

Pauline Toni

#### S31.P3

Key to kangaroo siring success: be in the right place at the right time Luca Montana

#### S31.P4

Optimal germination times in unpredictable environments: the importance of dormancy for among- and within-year variation

Hanna ten Brink

#### S31.P5

The evolution of reproductive diapause facilitates insect radiation into African savannahs during the late-Miocene

Sridhar Halali

Small eggs, large clutches and parental care: unexpected life-history evolution patterns in shield bugs Shin-ichi Kudo

#### S31.P7

Individual variation and evolutionary potential of parasite traits in a songbird-tick system Gerardo Fracasso

#### S31.P8

Variation in lifetime reproductive strategies in a self-compatible hermaphrodite: combining field data with laboratory experiments

Anja Felmy

#### S31.P9

Life history evolution: the constraints of sexual plasticity Chiara Benvenuto

#### S31.P10

Experimental removal of sexual selection turns males in to couch potatoes

Martin Garlovsky

#### S31.P11

Co-evolution of maternal thyroid hormones and life history strategies: comparative and experimental tests Bin-Yan Hsu

#### S31.P12

Family dynamics and age-related patterns in marriage probability in historical human population Jenni Pettay

#### S31.P13

Seasonality as a predominant control factor of the moult dynamics in birds - a meta-analysis

Agnieszka Gudowska

#### S31.P14

Natural and laboratory competition experiments between sexual and parthenogenetic polyclonal species of brine shrimp (Artemia)

Robert Browne

### S31.P15

Within individual variance in foraging behaviour mediates the fitness consequences of broad scale climate phenomena

Samantha Patrick

#### S31.P16

Alternative reproductive strategies are associated with distinct queen and worker size in Temnothorax rugatulus ants

Marina Choppin

#### S31.P17

Thermal environment at the juvenile stage affects morph-specific offspring survival in a polymorphic damselfly

Maarit Mäenpää

The evolution of adult sex ratios in dragon- and damselflies

Martin Alejandro Serrano-Meneses

Waiting for love - Reproductive delay and assurance under fluctuating population density Chantal Stock

### S31.P20

Maternal resources, early-life care and life-history outcomes: why some mongooses are more equal than others

Emma Vitikainen

Lifespan in wild butterfly populations: genomic determinants and maintenance of standing variation Vicencio Oostra

#### S31.P22

Aiming for the moon: Maintenance of local adaptation to tidal regime

Runa Kvamme Ekrem

Survival costs of sons and daughters in Asian elephants Vérane Berger

### S31.P24

Evolutionary trade-offs in antigen-presentation mediate risk for infection and autoimmunity Jatin Arora

#### S31.P25

Dietary restriction, sexual selection, and the life history trade-off between reproduction and somatic maintenance

Elisabeth Bolund

#### S31.P26

A model for the evolution of interspecific obligate brood parasitism in fish

Pierick Mouginot

### 32. Niche width evolution and its (mal) adaptive significance

#### S32.P1

The evolution of habitat choice facilitates niche expansion

Eva Kisdi

#### S32.P2

Cumulative stress or acclimation? Thermal performance in all life stages of a thermal generalist butterfly Loke von Schmalensee

### 33. Evolutionary Ecology of Ageing: from mechanisms to life-history consequences

Deleterious mutations show increasing negative effects with age in Drosophila melanogaster Martin Brengdahl

#### S33.P2

RNAi screening of lifespan and fecundity genes in Drosophila melanogaster TOR and IIS pathways Daniel Pritchard

#### S33.P3

Reversal of the fecundity-longevity trade-off across the spectrum of sociality in the bees Andreia Teixeira

#### S33.P4

The effect of food restriction on ageing and fecundity in termite

Silu LIN

Pedigree-based estimation of germline mutation rate of Rhesus macaque (Macaca mulatta) linked to parental age

Lucie Bergeron

#### S33.P6

The evolutionary and developmental dynamics of life history

Mauricio González-Forero

#### S33.P7

The role of laying and hatching order in early-life telomere dynamics in a wild passerine Tiia Kärkkäinen

The harsh life of annual killifish - new insights from natural demography and life history Milan Vrtilek

#### S33.P9

Diversity of ageing patterns in a highly promiscuous songbird

Eve Cooper

#### S33.P10

Ant workers' susceptibility to paraquat induced oxidative stress in relation to age and reproductive status Megha Majoe

#### S33.P11

Nutrient balance as a mechanism to understand the longevity/fecundity trade-off in ants Abel Bernadou

#### S33.P12

Why do dietary restricted animals live longer? Testing the evolutionary theories

Laura Travers

### S33.P13

Trade-off between reproductive effort and oxidative status as a response to warming in marine environment Ella von Weissenberg

### S33.P14

A theoretical investigation of the effect of early-life somatic damage levels on senescence rates Matthias Galipaud

#### S33.P15

An evolutionary explanation for longevity in naked mole

Robert Noble

Is age just a number? The role of senescence in the fish populations' eco-evolutionary dynamics Silva Uusi-Heikkilä

Contributions of sperm and seminal fluid to ejaculate senescence, and its amelioration via insulin signalling Irem Sepil

### 34. Mathematical models in evolutionary biology

### S34.P1

Evolutionary rescue and dispersal: the effect of habitat choice on successful adaptation

Peter Czuppon

Microbes can explain the evolution of paternal care Yael Gurevich

#### S34.P3

Systemic consequences of reproductive strategies Laura Hildesheim

#### S34.P4

Joint evolution of dispersal and connectivity Petteri Karisto

#### S34.P5

Evolutionary and demographic consequences of temperature-induced masculinization: the role of sexual selection

Edina Nemesházi

#### S34.P6

Mutation bias in empirical genotype-phenotype land-

Alejandro V Cano

#### S34.P7

What phylodynamic skyline models can and cannot do Veronika Boskova

#### S34.P8

Bet-hedging across generations can affect the evolution of variance-sensitive strategies within generations Thomas Ray Haaland

#### S34.P9

Evolution of enzyme concentrations in metabolic pathwavs

Charlotte Coton

#### S34.P10

Horizontal transfer and phylogenetic calibration in linguistics: a Bayesian approach Luke Maurits

### S34.P11

Fitness versus risk: calculating a fitness-risk ratio that describes optimal seed heteromorphism P. William Hughes

### S34.P12

Life is not a long quiet river: modelling population genetic divergence when migration is fluctuating Vincent Calcagno

#### S34.P13

Gene flow can also lead to a U-shaped site frequency spectrum Nina Marchi

#### S34.P14

How does the occasional absence of resources for cooperation affect the evolution of direct reciprocity? Shun Kurokawa

#### S34.P15

Coevolution in a hierarchically structured host-parasite metapopulation

Robert Pascal Dünner

### S34.P16

Contrasting the impact of cytotoxic and cytostatic drug therapies on tumour progression Jani Anttila

### S34.P17

Sexual selection in stpace: The effect of spatial structure on sexual selection

Maximilian Tschol

Investigating the role of dispersal in African monarch butterflies and their male-killing endosymbiont Franziska Brenninger

Modelling the killer T-cell and cancer cell sub-population dynamics under immuno- and chemotherapies Anni S. Halkola

### 35. Evolution outreach projects: Keep SCREAMing (Science Communication **Research Empowers AMazing outreach)**

Connecting Mediterranean fishery stakeholders and scientists to resolve connectivity of fishery populations Iva Sabolić

#### S35.P2

A competition in evolution Barbora Trubenova

#### S35.P4

Evolution on the way - migrations with EvoCorner Bojan Kenig

#### S35.P5

Experimental evolution of E. coli resistance to starvation: a practical course for undergraduate students Christine Dillmann

#### S35.P6

Power of hands-on experience - DIY Biology and Bio-hacklabs in science outreach Elzbieta Iwaszkiewicz

#### S35.P7

Hidden in soil: DNA barcoding and teaching microscopic diversity of soil mites (Acari:Oribatida) Riikka Elo

#### 36e. Phenotypic Evolution

#### S36e.P2

A time series model for estimating temporal variation in phenotypic selection

Yihan Cao

#### S36e.P4

Morphological integration in a cannibalism reaction norm

Kinya Nishimura

#### S36e.P5

Quantifying evolutionary bias from comparative datasets: a parametric bootstrapping approach for evolutionary covariance matrices

Junya Watanabe

#### S36e.P6

Capture from the wild and its consequences for Asian elephant reproduction

Mirkka Lahdenperä

### S36e.P7

Local adaptation to photoperiod and the endogenous clock in Daphnia

Anke Schwarzenberger

### S36e.P8

Physiological responses to seasonal environmental variation in a long-lived mammal Sophie Reichert

### S36e.P9

The chewing machine - evolution of mouth morphology in Drosophila larvae

Nuno Silva-Soares

The evolutionary trajectory of consistency in behavioural traits across ontogeny in fast-slow life histories Will Sowersby

#### S36e.P11

**Fast Adaptive Plastic Responses to Diurnal Temperature** Variation in an Arctic Specialist Arthropod Natasja Krog Noer

### S36e.P12

Predicting ecological responses to global warming in Iris pumila: an open-topped chamber experiment Katarina Hočevar

#### S36e.P13

Assessing consequences of environmental stress on wild rodent gut health by transcriptomics, microbiomics and histology

Toni Jernfors

### S36e.P14

Age and environment (but not genetics) affect mitochondrial function in a wild bird species Coline Marciau

#### S36e.P15

Can female pheromone contribute to the co-existence of color morphs in a moth species? Chiara De Pasqual

#### S36e.P16

Decomposing phenotypic skew into genetic and environmental components reduces the predicted response to strong selection

Jarrod Hadfield

#### S36e.P17

Evolution to temperate climates in the grass subfamily **Pooideae** 

Marian Schubert

### S36e.P18

Gender-specific variation in leaf shape under environmental stress in an understorey forest perennial Dragana Cvetković

### S36e.P19

The scent of divergence: chemical communication mediates reproductive isolation of two wood tiger moth populations

Cristina Ottocento

### S36e.P21

Colour distribution in hummingbird communities results from the interplay between selection for camouflage and communication

Hugo Gruson

### S36e.P22

Pheomelanin pigmentation and oxidative balance in Asian barn swallows

Emi Hasegawa

## POSTER SESSION FRIDAY 17.20-19.20

### 1. Trans generational plasticity in animals

#### **S1.P1**

Gametic plastic responses in thermally evolving lines of Tribolium castaneum

Ramakrishnan Vasudeva

Trans-generational plasticity and bet-hedging: A framework and a meta-analysis on insect diapause reaction norms

Jens Joschinski

#### **S1.P3**

Trans-generational effects of commensal microbiota on pupal production and body weight of a polyphagous fly Binh Nguyen

#### **S1.P4**

Influence of environmental heterogeneity on the evolution of phenotypic plasticity and bet-hedging Zuzana Sekajova

Epigenetic reprogramming during gametogenesis and embryogenesis of threespine stickleback: windows for adaptation to climate change? Lisa Shama

### **S1.P6**

Effects of immune priming on honeybee pollination Matti Leponiemi

#### **S1.P7**

Prenatal programming of mitochondrial function: a potential mediator of transgenerational plasticity in animals?

Antoine Stier

### **S1.P8**

Parental age effects on offspring telomere length in a natural avian population

Hannah Dugdale

### **S1.P9**

Trans-generational effects of early developmental stress on morphology and reproductive performance in captive zebra finches

Yifan Pei

Maternal effects are the predominant source of intraspecific variation in spider foraging traits Jorge Henriques

### **S1.P11**

Adaptation to climatic differences and the role of avian volk thyroid hormones Martje Birker

### **S1.P12**

Thermal sensitivity and heat hardening capacity of Drosophila melanogaster vary during ontogeny Neda Nasiri Moghadam

### S1.P14

Phenotypic plasticity within and across generations in a polyphagous moth Axel Rösvik

#### 2. Evolution in real time: experimental evolution approaches

#### S2.P1

Natural selection drives leaf shape divergence in experimental populations of Senecio lautus under natural conditions

Thomas Richards

Evolvability of orthologous genes (effect of global suppressors)

Hind Abdalaal

#### S2.P3

The Evolution of Aggression in Response to Sexual Selection in male and female Drosophila melanogaster Danielle Edmunds

### **S2.P4**

Does sex-specific selection change mating behaviour in a hermaphrodite?

Aivars Cirulis

#### S2.P5

Rapid evolution of reproductive morphology and fitness in a model pest insect

Rebecca Lewis

#### **S2.P6**

Female-limited X chromosome evolution and its effect on sperm competitiveness

Yesbol Manat

#### **S2.P7**

Non-consumptive effects drive rapid evolution in a prey population

Chao Zhang

#### S2.P9

Experimental evolution of biological control agents Sara Magalhães

### S2.P10

Sexual selection favoured higher offspring production via evolution of both male and female traits Daisuke Kyogoku

#### S2.P11

Role of phenotypic plasticity for evolutionary adaptation: Experimental approaches using Tribolium castaneum and Bacillus thuringiensis Ana Sofia Lindeza

Experimental adaptation to malnutrition reveals tradeoff in extraction of protein versus sugar from diet Fanny Cavigliasso

### S2.P13

Sexually-selected male weapon causes gender load and increases the risk of extinction Jacek Radwan

#### S2.P14

Experimental evolution for collagen invasion in cancer cell lines Louise Johnson

#### **S2.P15**

Combined effects of toxins on non-target dung breeding flies (Diptera: Sepsidae)

Natalia Gourgoulianni

No evidence found for sexual conflict over cuticular hydrocarbons in female-limited X chromosome evolution experiment

Katrine K. Lund-Hansen

#### **S2.P17**

Evolutionary ecology of multiple-interaction networks in bacterial communities

Marie Vasse

#### S2.P19

Examining the selective potential of artificial light at night in Drosophila melanogaster Lucy McLay

#### S2.P20

Can we delimit individuals in species with blur concept of individuality?

Sundy Maurice

#### 3. Exploring the role of nongenetic inheritance in evolution

More than methylation: does pleiotropy drive the complex pattern of evolution of dnmt1?

Patricia Moore

#### S3.P2

Sex-specific social learning in juvenile zebra finches Boglárka Morvai

#### S3.P4

Eco-cultural range expansion of modern humans in **Paleolithic** 

Joe Wakano

Indirect genetic effects genetic correlation contribute to the total heritable variance in parental care Julia Schroeder

### **S3.P6**

Genetic and linguistic histories in Central Asia inferred using Approximate Bayesian Computations Frédéric Austerlitz

#### S3.P7

Offspring phenotype is shaped by the non-sperm fraction of semen

Jukka Kekäläinen

#### S3.P8

Comparative epigenomics unravels the evolutionary landscape of insect DNA methylation Panagiotis Provataris

### S3.P9

Differential maternal and paternal effects on offspring fitness traits

Valérian Zeender

Plasticity, inheritance and epigenetics in plants: Can these be linked?

Morgane Van Antro

#### S3.P11

Epigenetic contribution to phenotypic plasticity and biotic stress-induced memory in Populus nigra Cristian Peña-Ponton

Evolutionary and plastic cytosine methylation responses to embryonic rearing temperature in European grayling Tiina Sävilammi

#### 5. Aging & cancer through the lens of evolution

#### S5.P1

Predicting tumor evolution and estimating its evolutionary unpredictablity using cancer progression models Ramon Diaz-Uriarte

### S5.P2

Extreme-downregulation of chromosome Y and male disease

Alejandro Caceres

### 9. Microbial genome and community evolution in food environments

#### S9.P1

Microbial community dynamics in Gwell, a fermented milk specialty from Brittany. A participatory study Lucas von Gastrow

#### S9.P2

Triphosphate nucleotide transport by bacteria is constrained by the oxidative environment Enrique Gonzalez-Tortuero

Contributions of plasticity and evolution to trait change in a community context

Lynn Govaert

Cheese shapes its Penicillium fungi Jeanne Ropars

#### 13. Genetics and genomics of adaptation

Rapid divergence of a 'great speciator' following a human-mediated introduction Ashley Sendell-Price

Local continuous genetic Isolation-by-Environment in the threespine stickleback in the Baltic Sea following predator collapse

Casey Yanos

### S13.P3

Detecting deleterious variants in the pig Martin Johnsson

#### S13.P4

Inter- and intra-population gene expression variation in the fat body during Drosophila melanogaster development

Amanda Glaser-Schmitt

Genomics of adaptation in the Alpine whitefish radiation Rishi De-Kayne

#### S13.P8

Genomics of Microphallus parasite adaptation to its host, Potamopyrgus antipodarum Natalia Zajac

**Metabolic Efficiency Variation Across Bird Families Measured with Relative Mitochondrial Abundance** Sergio Andreu-Sánchez

Convergent expansion in gene-families and their role on the blood-feeding diet in Insecta lineages Lucas Freitas

#### S13.P11

Genomics of clinal adaptation with gene flow in parapatric lake-stream stickleback

Quiterie Haenel

#### S13.P12

Susceptibility to gapeworm parasite has both additive and dominant genetic components in house sparrows Sarah Lundregan

#### S13.P13

The genomes of Poeciliopsis retropinna and Poeciliopsis turrubarensis reflect differences in reproductive strategy Henri van Kruistum

#### S13.P14

Comparative transcriptome profiling of Triplophysa bleekeri and Triplophysa rosa, reveals potential mechanisms of eye degeneration

Qingyuan Zhao

Winter moth adaptation to climate change: genetic changes in thermal plasticity of embryonic development

Natalie E. van Dis

#### S13.P16

Discovering genetic diversity and structural variation underlying local adaptation in Scots pine Tanja Pyhäjärvi

#### S13.P17

Investigating genetic basis of geographic variation in innate immunity of a butterfly Naomi L.P. Keehnen

A Flutter of Genomes: New and revised high quality genomic resources for 50 Heliconiini species Francesco Cicconardi

### S13.P19

Gene Trf2 and the microbiome underpin the expression of dormancy in Drosophila

Manolis Lirakis

#### S13.P20

Disperse, acclimatise or adapt: seascape genomics along a thermal gradient

Anna Muir

#### S13.P21

The genetic basis of convergent adaptation to altitude in Arabidopsis thaliana

Pádraic Flood

#### S13.P22

Genomics of expanded avian sex chromosomes shows predisposition of certain chromosomes towards sex-linkage in vertebrates

Hanna Sigeman

#### S13.P23

Genome-wide effects of selection in two outcrossing plant species Tiina Mattila

### S13.P25

Nordic conquest: Post-glacial radiation and evolutionary history of lunar-rhythmic and lunar-arrhythmic reproduction in marine midges

Nico Fuhrmann

#### S13.P26

Natural Variation of defense response genes in Arabidopsis thaliana reveals evidence for balancing selection Mehmet Göktay

### S13.P27

The role of gene interactions and gene interaction networks in speciation

Ina Satokangas

#### S13.P28

Sex-specific alternative splicing in Drosophila melanogaster

Julia Raices

#### S13.P29

Linking genotype, phenotype, and environment to understand climate adaptation in the Glanville fritillary butterfly

Michelle DiLeo

### S13.P30

Non-neutral impact of synonymous mutations: example of an antibiotic resistance gene expressed in human cells Marion AL Picard

White to brown and back: circannual genic regulation of coat colour change in snowshoe hares João Pimenta

### S13.P32

Integration of proteomic data into constraint-based models reveals the molecular bases of yeast life-history trade-offs

Marianyela Petrizzelli

#### S13.P33

Muller's Ratchet and the Long-Term Fate of Chromosomal Inversions

Alexandre Blanckaert

Cross-temperature comparisons of gene expression between heat tolerant and heat sensitive Brachionus species

Sofia Paraskevopoulou

Genomic basis of rapid parallel ecological adaptation to heterogeneous environments

Hernan Morales

Chance and predictability: the genomic basis of convergent dietary specializations in an adaptive radiation Joel Vizueta

#### S13.P38

Co-speciation in bed bug Wolbachia Ondřej Balvín

#### S13.P39

The genomic basis of humic substance-driven adaptation in Eurasian perch

Mikhail Ozerov

#### S13.P40

Mito-jay-nomics: Signatures of environmental adaptation in the first assembled mitogenomes from New World Jays (Corvidae)

Katia Bougiouri

#### S13.P41

Effects of selection on haplotypes and genealogy trees of subdivided populations

Yichen Zheng

#### S13.P42

Genomics of feralization processes on Hawaiian and Bermudan chickens

Maria Luisa Martin Cerezo

#### S13.P43

Keeping pace with fast environmental changes, a science-based approach for sustainable Cork Oak forests Octávio Paulo

#### S13.P44

Detection of phylogenetically-informative SNPs in human Y-chromosome from next-generation sequencing data

Koji Ishiya

### S13.P45

Identifying the genomic and sex-specific characters underlying recombination rate variation

Suvi Ponnikas

Transposable Elements as agents of adaptation in the invasive species Drosophila suzukii?

Vincent Merel

#### S13.P47

Local adaptation of phenotypic plasticity: pupal diapause in the butterfly Pieris napi

Peter Pruisscher

#### S13.P48

Repetitive DNA: a force shaping karyotype evolution in blue butterflies (Lycaenidae, Lepidoptera)

Martina Dalikova

#### S13.P49

Transcriptomes from four Iberian Squalius fish species indicate stronger positive selection in Mediterranean climate type

Carlos Ramirez

#### S13.P50

Evolution of AT/GC content in vertebrates Radka Symonova

#### S13.P51

The role of host plant in symbiosis stability of Arbuscular mycorrhizal fungi

Shadi Eshghi Sahraei

#### S13.P52

Heritability of intra-individual variation in body temperature in the wild yellow-necked mouse, Apodemus flavicol-

Rohan Raval

#### S13.P53

Landscape genomics of the wood decay fungus Phellopilus nigrolimitatus

Jørn Henrik Sønstebø

#### S13.P54

Comparative genomics and lineage specific adaptations in Lepidoptera

Karin Näsvall

#### S13.P55

Genomic architecture of divergence between parasitic and non-parasitic lamprey ecotypes

Ahmed Souissi

#### S13.P56

New Insights into the Genetic Basis and Evolutionary History of Lactase Persistence in Africa

Alessia Ranciaro

#### S13.P57

Complete plastid genome sequence of African nightshade (Solanum scabrum) and its comparative plastomics across Solanales

Gaurav Sablok

#### S13.P58

Antagonistic coevolutionary selection patterns in the Galerucella-Asecodes host-parasitoid system Xuyue Yang

### S13.P59

Icefish genome reveals key role of mitochondria for a life without hemoglobin at sub-zero temperature Chiara Papetti

#### 15. Tracing evolution through time using ancient DNA

Distinguishing among complex evolutionary models using unphased whole-genome data through Approximate **Bayesian Computation** 

Maria Teresa Vizzari

#### S15.P2

**Ancient DNA screening from Finnish Stone Age sediments** Sanni Peltola

### S15.P3

Archaeological sediments from Finland as a source for ancient microbiomes

Enrique Rayo

#### S15.P4

Mitochondrial DNA from Iron Age to present in Eastern **Fennoscandia** 

Sanni Översti

Inferring population dynamics of the genera Oryx and Addax using modern and historical DNA Elisabeth Hempel

The first historic Treponema pallidum genomes from **Colonial Mexico** 

Aditya Kumar Lankapalli

#### S15.P7

Temporal and spatial insights into the genomic evolution of Yersinia pestis through comparative analysis Aida Andrades Valtueña

Optimization of double-stranded library preparation methods for ancient and degraded DNA Marianne Dehasque

Ancient DNA provides insights into the population history of the reindeer

Matti Heino

#### 16. Mito-nuclear interactions across levels of biological organisation

The role of selection in maintaining sympatric mito-nuclear variation in Drosophila subobscura Pavle Erić

#### S16.P2

Mitochondrial Haplotypes and Gene Expression in Laying Hens

Elisabeth HempelClara Heumann-Kiesler

#### S16.P3

A new layer of genetic regulation in the mitochondrial genome: small mitochondrial RNAs Andrea Pozzi

#### S16.P4

The mtDNA-encoded COX2 protein: bivalves have the longest

Eric Pante

#### S16.P5

Sex-specific effects of candidate Trojan Female Technique haplotype on fertility in pest species Acanthoscelides obtectus

Lea Vlajnić

#### S16.P6

How mitochondrial genetic variation affects longevity in Drosophila melanogaster Ekta

#### S16.P7

Mito-nuclear interactions in innate immunity and life-history traits

Tiina Salminen

#### S16.P8

**Mitochondrial Diseases and Compensated Pathogenic Deviations** 

Abhilesh Dhawanjewar

**Ancestry Package - Merging uniparental and autosomal** genetic histories into one picture

Vladimir Bajić

### 19. Gene-phenotype associations across evolutionary scales

#### S19.P1

Identifying host genomic regions influencing microbial traits in mice

Shauni Doms

#### S19.P2

Convergence, common ancestry and novelty: Genomics of sex chromosome diversity in cichlid fishes Astrid Böhne

#### S19.P3

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Rearrvisr: an R package to detect, classify, and visualize genome rearrangements

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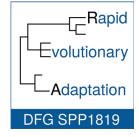


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