

# **AGENDA ESEB 2025 Congress**

## Barcelona, 17-22 August 2025

\*\* Modifications received after this agenda is released will not be shown in this document\*\*

\*\* Changes notified after 12 August may be reflected only in the Program located in the Congress App \*\*

## Monday, 18 August 2025

MEETING ROOM 113-117: 08:45 AM - 09:00 AM Opening Speech & Congress Guidelines

#### PLENARY SESSION (MEETING ROOM 113-117): 09:00 AM - 09:45 AM - Plenary Talk – Keynote Speaker MICHAEL LYNCH

9:00 AM 9:45 AM Drift, Mutation, and the Origin of Cellular Features Michael Lynch

EXPO AREA: 9:45 AM - 10:30 AM - Coffee Break

#### MEETING ROOM 113: 10:30 AM - 12:30 PM - S18.01 - Evolution of biotic interactions across scales

Session Ch	Session Chairs: Dr. Peter Czuppon, Prof. Lena Wilfert				
10:30 AM	11:00 AM	Mutualistic networks in the face of global change	Prof. Jordi Bascompte		
11:00 AM	11:15 AM	From mosquitoes to cuckoo bumblebees; selfish genetic elements to spotted hyenas; penguins to pathogens: can we explain how antagonists choose their victim?	Ms. Mairenn Attwood		
11:15 AM	11:30 AM	Macroevolutionary Patterns in Trees Shaped by Three Key Mutualisms: Pollination, Seed Dispersal, and Mycorrhizal Symbiosis	Prof. Akira Yamawo		
11:30 AM	11:45 AM	Competition and facilitation at the within- and between-host levels between spider mites evolving in metal-accumulating plants	Prof. Inês Fragata		
11:45 AM	12:00 PM	The Legacy of Past Heatwaves on 'Off-Host' Parasite Stages: Reduced Infection Risk and Costs in the Daphnia-Pasteuria system	Dr. Justine Boutry		
12:00 PM	12:15 PM	Bacteria-phage coevolution drives patchy distribution of plant disease through phage resistance-virulence trade-offs	Prof. Ville-Petri Friman		
12:15 PM	12:30 PM	Landscape structure as a driver of eco-evolution in host-parasite systems	Dr. Emanuel Fronhofer		

## MEETING ROOM 114: 10:30 AM - 12:30 PM - S23.01 - Evolutionary Genomics: Understanding and Adapting to Climate Change (organised by the journals Molecular Ecology & Evolutionary Applications)

Session Ch	Session Chairs: Prof. Luciano Beheregaray, Prof. Maren Wellenreuther						
10:30 AM	11:00 AM	Adaptation to Climate in European Conifers: Lessons Learned	Dr. Santiago González Martínez				
11:00 AM	11:15 AM	Spatial scales of genomic adaptation to climate and phenology in the winter moth	Dr. Andrea Estandia				
11:15 AM	11:30 AM	Environmental effects on the origin and establishment of Arabidopsis arenosa	Ms. Yu Cheng				
11:30 AM	11:45 AM	Deep-time genomes reveal the evolution of distinct genetic variants in the woolly mammoth	Ms. Kelsey Moreland				
11:45 AM	12:00 PM	Genomic bases of short term evolution in the wild revealed by long term monitoring and population-scale sequencing.	Dr. Tristan Cumer				
12:00 PM	12:15 PM	The genetics of thermal adaption in tropical butterflies	Dr. Nicola Nadeau				
12:15 PM	12:30 PM	Speciation under stress: Genomic and ecological divergence in Eastern Baltic Cod	Dr. Kwi Young Han				

#### MEETING ROOM 115: 10:30 AM - 12:30 PM - S17.01 - Evolution of behavioural diversity: from ecology to genes and neural systems

Session Ch	Session Chairs: Dr. Alexandra de Sousa, Dr. Katja Heuer, Dr. Stephen Montgomery					
10:30 AM	11:00 AM	New and old genes to generate novel neural systems: the origin of the Turbanate eyes	Dr. Isabel Almudi Cabrero			
		in mayflies.				
11:00 AM	11:15 AM	Life in dim light: The evolution of vision in deep-sea fishes	Dr. Lily Fogg			
11:15 AM	11:30 AM	Visual System Adaptations Underlying the Evolution of Flower-Visiting Behavior in	Dr. Yuki Ishikawa			
		Drosophila				
11:30 AM	11:45 AM	Modality-specific memory enhancements in Heliconius butterflies result from specific	Ms. Elizabeth Hodge			
		mushroom body expansion, rather than increased investment in the sensory pathways				



11:45 AM	12:00 PM	Evolution of Taste Neurons and Behaviour	Dr. Roman Arguello
12:00 PM	12:15 PM	Convergent evolution of pain insensitivity in a new Heliophobius species	Dr. Daniel Mendez Aranda
12:15 PM	12:30 PM	Chemosensory evolution at the origin of inquilinism in the bee louse fly (Braula coeca)	Ms. Alizée Delarue

Session Ch	Session Chairs: Dr Florent Mazel, Dr. Melissah Rowe					
10:30 AM	11:00 AM	Microbiome-mediated adaptation and niche construction	Dr. Carola Petersen			
11:00 AM	11:15 AM	Are the root microbiome and plant phenotypic traits affected when a wild perennial is transplanted across contrasting habitats?	Dr. Åsa Lankinen			
11:15 AM	11:30 AM	How the environment, genome, and microbiome influence niche width in spider mites	Dr. Karen Bisschop			
11:30 AM	11:45 AM	Elucidating interactions of the microbiome and host genomics during population decline through a temporal hologenomics approach	Dr. Annie West			
11:45 AM	12:00 PM	Behavioural Manipulation by Parasitic Nematodes is Associated with Microbiome Composition	Dr. Priscila Salloum			
12:00 PM	12:15 PM	Influence of Gut Microbiota Perturbation on Obligate Endosymbiont Abundance and Host Phenotype in the German Cockroach (Blattella germanica)	Mrs. Srijita Ray			
12:15 PM	12:30 PM	The effects of climate change on the microbial community associated with a butterfly across a 30-year period	Mr. Linyang Sun			

## MEETING ROOM 117: 10:30 AM - 12:15 PM - S51-01 - Ecology

Chairs: Pau	ı Carazo, Iva	n Gomez Mestre	
10:30 AM	10:45 AM	Continental-scale variation of Saccharomyces paradoxus	Dr. Nicolo Tellini
10:45 AM	11:00 AM	Environmental dependence of local adaptation and expansion load across the range of an annual plant	Dr. Shengman Lyu
11:00 AM	11:15 AM	Casting light on the role of light-sensitive bacteria in plant adaptation to polar light regimes	Ms. Emilia Mäkinen
11:15 AM	11:30 AM	Condition matters: First insights into alarm cue production in a freshwater gastropod.	Mr. Nhamo Mutingwende
11:30 AM	11:45 AM	Domestic dog introgression in Australian dingoes: environmental drivers and evolutionary consequences	Dr. Malgorzata Pilot
11:45 AM	12:00 PM	Breakdown of speciation processes between two genetically and ecologically divergent ecotypes of a wood decay fungus	Ms. Ingvild Myhre Ekeberg
12:00 PM	12:15 PM	$\label{thm:condition} Evolution of developmental bias explains divergent patterns of phenotypic evolution in two nematode clades$	Dr. João Picão Osório

## MEETING ROOM 118+119: 10:30 AM - 12:45 PM - S20 - Evolutionary biology meets genetic pest control

Session Chairs: Dr. Florence Débarre, Dr. Nicolas Rode					
10:30 AM	11:00 AM	Don't you know we're talking 'bout (rapid) evolution: recombination and copy-number variation in Wolbachia shift cytoplasmic incompatibility phenotypes.	Dr. Alice Namias		
11:00 AM	11:15 AM	Evaluating Wolbachia and host genetics for incompatible insect technique development in Drosophila suzukii	Dr. Svitlana Serga		
11:15 AM	11:30 AM	Assessing the role of mitonuclear and genotype-by-environment interactions for the Trojan Female Technique in pest biocontrol	Ms. Lea Vlajnić		
11:30 AM	12:00 PM	Predicting the invasiveness of threshold-dependent gene drives	Mr. Philipp Messer		
12:00 PM	12:15 PM	Mitigating resistance in gene drives using sex distortion strategies.	Ms. Oksana Vertsimakha		
12:15 PM	12:30 PM	Gene drive dynamics in plants: the role of seedbanks	Ms. Isabel Kim		
12:30 PM	12:45 PM	$\label{thm:continuous} \textbf{Out-of-locus genomic editors for efficient and localisable pest population suppression.}$	Dr. Katie Willis		

## MEETING ROOM 120+121: 10:30 AM - 12:30 PM - S05 - Aliens among us: ecological drivers, evolutionary dynamics, and rapid ecosystem reshaping by biological Invasions

Session Chairs: Dr. Oriol Lapiedra, Prof. Anna Traveset						
10:30 AM	11:00 AM	Marine Aliens: How They Impact Biodiversity and Evolution in Urban Marine	Dr. Frédérique Viard			
		Environments				
11:00 AM	11:15 AM	Bidirectional exchange of resistance genes between native Helicoverpa zea and	Dr. Henry North			
		invasive H. armigera				
11:15 AM	11:30 AM	Invasion dynamics of the house mouse in Africa	Dr. Daniel Poveda Martinez			
11:30 AM	11:45 AM	Unusual genetic structure of the alien plant Cotula coronopifolia in Europe, resulting	Mr. Raúl Sánchez García			
		from selfing, introduction history and dispersal by migratory waterbirds				
11:45 AM	12:00 PM	Rapid adaptation in an island endemic lizard in response to an invasive predator?	Dr. Adrián Talavera			
11:45 AM	12:00 PM	Rapid adaptation in an island endemic lizard in response to an invasive predator?	Dr. Adrián Talavera			



12:00 PM	12:15 PM	Short and long term effects of an invasive parasite on the physiology of host fish	Mr. Abhishek Nair Anil
12:15 PM	12:30 PM	populations. A genomically informed prediction of the overwintering range of cotton bollworm in North America	Mr. Cian Williams

MEETING ROOM 122+123: 10:30 AM - 12:30 PM - S01 - Adaptation to environmental changes in trees through the lens of common gardens and genomics

Session Ch	Session Chairs: Dr. Benjamin Brachi, Dr Domitille Coq-Etchegaray, Dr. Sofia van Moorsel					
10:30 AM	11:00 AM	Next generation experimental citizen science: when research meets the needs of forestry facing climate change	Dr. Katalin Csilléry			
11:00 AM	11:15 AM	When South meets North: the shaping of a contact zone in three boreal tree species	Ms. Maria del Pilar Herrera Egoavil			
11:15 AM	11:30 AM	A higher load of deleterious mutations has a detrimental effect on tree growth	Dr. Rosalia Piñeiro Portela			
11:30 AM	12:00 PM	Towards understanding the pangenomic basis of environmental adaptation in European beech using common gardens	Ms. Desanka Lazic			
12:00 PM	12:15 PM	Genomic signatures of climate-driven (mal)adaptation in an iconic conifer, the English yew (Taxus baccata L.)	Mr. Thomas Francisco			
12:15 PM	12:30 PM	Adaptation to ash dieback disease in natural populations of European ash	Dr. Dario Galanti			

MEETING ROOM 129+130: 10:30 AM - 12:00 PM - S03.01 - Advances in technology, mathematical and statistical models and their application in evolutionary ecology: the dawn of a new era

Session Chairs: Prof Andy Gardner, Dr. Juliano Morimoto, Prof. Hanna Kokko					
10:30 AM	11:00 AM	Which theoretical tools should be our toolbox?	Prof. Hanna Kokko		
11:00 AM	11:15 AM	The variance world: the era of location-scale models in evolutionary meta-science	Prof. Szymon Drobniak		
11:15 AM	11:30 AM	Inference of ecological and life-history traits evolution from genomic and epigenomic polymorphism data.	Prof. Aurelien Tellier		
11:30 AM	11:45 AM	Neural posterior estimation for high-dimensional simulation data from complex ecological and population genetic models	Ms. Yuxin Ning		
11:45 AM	12:00 PM	Trend-filtered support vector machine to detect natural selection from genomic autocovariation	Mr. Md Ruhul Amin		

MEETING ROOM 131: 10:30 AM - 12:30 PM - S41 - Post-phylogenomics: new and evolving molecular methods to address challenging phylogenies

Session Ch	Session Chairs: Dr. Mattia Giacomelli, Dr. Jesus Lozano-Fernandez, Dr. Anthony Redmond				
10:30 AM	11:00 AM	Animal Phylogenomics: Navigating Data, Paralogy, and Non-Tree-like Events	Prof. Mary O'connell		
11:00 AM	11:15 AM	CASTER: Direct species tree inference from whole-genome alignments	Dr. Chao Zhang		
11:15 AM	11:30 AM	Phylotranscriptomics as the Most Appropriate Tool for Resolving Complex Phylogenies: Insights from the Ophrys genus	Mr. Lucas Vandenabeele		
11:30 AM	12:00 PM	Using simulation to understand phylogenetic problems.	Prof. Max Telford		
12:00 PM	12:15 PM	Paralogy, orthology inference biases, and extreme discordance in plant phylogenomic data sets driven by assembly artifacts and sequencing strategies	Dr. Diego Morales-Briones		
12:15 PM	12:30 PM	Unraveling evolutionary complexity in a young adaptive radiation of New Caledonian persimmons	Ms. Teerna Khastgir		

## MEETING ROOM 132: 10:30 AM - 12:30 PM - S51-03 - Trait evolution

Chairs: Borja Milà, Elena Bosch					
10:30 AM	10:45 AM	Reconstructing the Evolution of Iridescence Across Birds	Dr. Michaël Nicolaï		
10:45 AM	11:00 AM	Long-fuse evolution of carnivoran skeletal phenomes through the Cenozoic	Dr. Chris Law		
11:00 AM	11:15 AM	Dressed for survival: Positive survival selection on plumage colouration revealed by	Ms. Lisa Sandmeyer		
		modelling labile trait expression			
11:15 AM	11:30 AM	Using a Meta-Omics Approach to Explore Opsin Diversity in Zooplankton	Dr. Giacinto De Vivo		
11:30 AM	11:45 AM	Investigating changing Wnt signal as a mechanism influencing differential evolutionary	Dr. Katie Pelletier		
		rates of vulval cell fates in nematodes.			
11:45 AM	12:00 PM	The genetic basis of multiple independent origins of a placenta	Dr. Margarida Cardoso-moreira		
12:00 PM	12:15 PM	Opsin gene expression plasticity and spectral sensitivity in male damselflies could	Prof. Erik Svensson		
		mediate female colour morph detection			
12:15 PM	12:30 PM	Under pressure: Adaptation of opsin proteins shaped by interacting selective factors	Mr. Peter Searle		

EXPO AREA: 12:30 PM - 2:00 AM - Lunch



#### MEETING ROOM 113: 2:00 PM - 5:00 PM - S18.02 - Evolution of biotic interactions across scales

Session Ch	nairs: Dr. Pe	ter Czuppon, Prof. Lena Wilfert, Prof. Stineke van Houte	
2:00 PM	2:30 PM	Phage warfare: Understanding the importance of prophage-encoded anti-phage	Prof. Stineke van Houte
		defence in Pseudomonas aeruginosa.	
2:30 PM	2:45 PM	I'm the captain now: Host fitness is predicted by host life-history strategy in an	Dr. Chloe Fouilloux
		emerging model host-parasite system	
2:45 PM	3:00 PM	Reversal of host-parasite dynamics following an introduction of non-native species	Prof. Martin Reichard
3:00 PM	3:15 PM	Co-circulation and co-infection: parasite evolutionary ecology across scales	Mr. Giacomo Zilio
3:15 PM	3:30 PM	Coinfection with a both a competitor and a facilitator parasite increases within-host growth rate and transmission	Dr. Alison Duncan
3:30 PM	3:45 PM	Evolution of buzz pollinated flowers in response to pollen-feeding insects: Pollen dosing	g Prof. Mario Vallejo Marin
		in response to vibrations	
3:45 PM	4:00 PM	Repeated breakdowns of mutualism into parasitism and expansions of ecological opportunity in a globally distributed plant-insect symbiosis	Prof. David Hembry
4:00 PM	4:15 PM	Tracing coevolutionary divergence in a host-parasite model system	Dr. Eva Lievens
4:15 PM	4:30 PM	Evolution of the plasticity of pathogen virulence in temporally and spatially variable environments	Dr. Ryuichi Kumata
4:30 PM	4:45 PM	Changing the script: ecological reversal of predator–prey roles in microbial communities	Dr. Marie Vasse
4:45 PM	5:00 PM	Costs of immunity avoided by immune suppression in novel terminal investment response	Dr. Nick Priest

# MEETING ROOM 114: 2:00 PM - 5:00 PM - S23.02 - Evolutionary Genomics: Understanding and Adapting to Climate Change (organised by the journals Molecular Ecology & Evolutionary Applications) Session Chairs: Prof. Luciano Reheregaray, Prof. Maren Wellenreuther, Dr. Janne Swaegers

Session Ch	airs: Prof. L	uciano Beheregaray, Prof. Maren Wellenreuther, Dr. Janne Swaegers	
2:00 PM	2:15 PM	Evolutionary trajectories of plasticity when encountering novel thermal conditions during range expansion in the wild	Dr. Janne Swaegers
2:15 PM	2:30 PM	Evolutionary genomic predicts Adaptation, Maladaptation, and Plastic Expression Responses to Climate Change in Two Closely Related Species with Contrasting Distribution Ranges and Population Sizes	Mr. Xuming Dan
2:30 PM	2:45 PM	Genetic variation in the response to multiple stresses: the case of Lymnaea stagnalis exposed to metals and global warming	Mrs. Cassandre Aimon
2:45 PM	3:00 PM	Genes that underlie photoreception and seasonal reproduction are crucial for local adaptation in the Northern krill, a keystone zooplankton for the North Atlantic Ocean	Dr. Per Unneberg
3:00 PM	3:15 PM	Cryptic diversity and phylogeographic patterns in Mediterranean acrobat ants (Crematogaster) reveal complex evolutionary histories in response to climate change	Ms. Jody Helena Voges
3:15 PM	3:30 PM	Genomic signals of adaptation during replicate range shifts of the blue-tailed damselfly, Ischnura elegans, in Scotland and Fennoscandia	Dr. Nicky Lustenhouwer
3:30 PM	3:45 PM	Unveiling future maladaptation and preadaptation in locally adapted Mediterranean beech populations under climate change	Mr. Josep Morando Mila
3:45 PM	4:00 PM	Effect of temperature on viability selection of developing hybrids revealed by whole genome pool-seq	Ms. Beatriz Portinha
4:00 PM	4:15 PM	Local adaptation and forecasting response in the endangered Scottish freshwater pearl mussel	Ms. Victoria Gillman
4:15 PM	4:30 PM	Contrasting contemporary changes in large-effect life history loci in Salmo salar across Europe	Dr. Maria Cadiz Escobar
4:30 PM	4:45 PM	Adaptation to Multiple Environmental Stressors in Tetrahymena thermophila	Dr. Laure Olazcuaga
4:45 PM	5:00 PM	Adaptive introgression enhances heat tolerance despite barriers to gene flow: a case study in Drosophila flavomontana and Drosophila montana	Dr. Noora Poikela

## MEETING ROOM 115: 2:00 PM - 5:00 PM - S17.02 - Evolution of behavioural diversity: from ecology to genes and neural systems

Session Cl	nairs: Dr. Al	exandra de Sousa, Dr. Katja Heuer, Dr. Stephen Montgomery	
2:00 PM	2:30 PM	The evolution of brain structures in mammals linked to behavioural ecology and from	Dr. Ornella Bertrand
		the perspective of the fossil record	
2:30 PM	2:45 PM	Phylogenetic insights into the emergence of brain anatomy and cognition.	Dr. Katja Heuer
2:45 PM	3:00 PM	Why did the human brain size evolve? A way forward	Dr. Mauricio González-Forero
3:00 PM	3:15 PM	Neuron Number in Fish Brains: Evolutionary Insights into Computational Power	Mr. Francesco Dionigi
3:15 PM	3:30 PM	Single-nuclei transcriptomics reveals neural cell type innovation associated with	Dr. Francesco Cicconardi
		mushroom body expansion in Heliconiini Butterflies.	
3:30 PM	3:45 PM	How to build a bigger brain?	Ms. Amaia Alcalde Anton
		The cellular basis of expanded mushroom bodies in Heliconius butterflies.	
3:45 PM	4:00 PM	Ecological divergence and post-eclosion brain development shape visual performance	Mr. Jose Borrero
		in Heliconius butterflies	



4:00 PM	4:15 PM	Heterochrony in orthodenticle expression is associated with ommatidial size variation between Drosophila species	Dr. Nico Posnien
4:15 PM	4:30 PM	The genetic basis of neural circuit evolution for Drosophila mate preferences	Dr. Emily Behrman
4:30 PM	4:45 PM	The evolution of diversity in rhythm: Developing cross-species metrics to assess the evolutionary dynamics of rhythm and syntax in birdsong	Ms. Ximena Leon
4:45 PM	5:00 PM	Genetics and environment shape vocal frequency variation in birds	Dr. Alex Kirschel

#### MEETING ROOM 116: 2:00 PM - 5:00 PM - S07.02 - Contribution of the microbiome to host adaptation and plasticity

Session Ch	Session Chairs: Dr Florent Mazel, Dr. Melissah Rowe, Dr. Hassan Salem				
2:00 PM	2:30 PM	Adaptation through symbiosis	Dr. Hassan Salem		
2:30 PM	2:45 PM	Beyond Genes: Modeling Microbiome-Mediated Responses to Selection	Dr. Bob Week		
2:45 PM	3:00 PM	The role of the host-associated microbiome in host tolerance and cross-tolerance to pollutants	Ms. Marlies Van De Maele		
3:00 PM	3:15 PM	The gut microbiome of invasive squirrels shows greater diversity, stability and functional potential than that of their native competitors	Dr. Claudia Romeo		
3:15 PM	3:30 PM	Bat gut microbiomes recapitulate host phylogeny and foraging niches	Dr. Dominik Melville		
3:30 PM	3:45 PM	Microbiome structure of French Guyana Drosophila community	Mr. Thibault Laffargue		
3:45 PM	4:00 PM	A bloody old habit: Conservation of gut-associated bacteria across blood-feeding leeches	Dr. Alejandro Manzano Marín		
4:00 PM	4:15 PM	Functional implications of changes in the rumen microbiome	Dr. Aoife Leonard		
4:15 PM	4:30 PM	Proteomic Profiling Reveals Brain Proteins Linked to Social Interactions in Gut Microbiota-Harbouring Honeybees	Mr. Gregorio Calderoni		
4:30 PM	4:45 PM	Can the gut microbiome mediate long-lasting effects of early-life environmental conditions on adult thermal physiology?	Dr. Charli Davies		
4:45 PM	5:00 PM	Of mice and microbes: the gut microbiome as a driver of thermal plasticity.	Ms. Carla Wagener		

## MEETING ROOM 117: 2:00 PM - 4:00 PM - S51-02 - Ecology

Chair: Eler	Chair: Elena Bosch					
2:00 PM	2:15 PM	The evolutionary history of a commensal species: lessons from adapting to the Anthropocene	Dr. Mark Ravinet			
2:15 PM	2:30 PM	A proposal on how to study the multifarious environmental causes of natural selection	Prof. Pim Edelaar			
2:30 PM	2:45 PM	Comparative analysis of predator-induced plastic responses in tadpoles and its regulation by corticosterone	Dr. Ivan Gomez-Mestre			
2:45 PM	3:00 PM	Gene expression profiling across the three-host life cycle of Schistocephalus solidus: how decoupled are developmental stages?	Dr. Laura Gramolini			
3:00 PM	3:15 PM	How does microbial diversity affect ecological stability?	Dr. Ming Liu			
3:15 PM	3:30 PM	The impact of population substructure on SMC-based inference tools analyzed through extensive demographic simulations on spatially complex demographic models	Mrs. Alba Nieto			
3:30 PM	3:45 PM	Symbioses between uncultivated microbes in the open ocean	Prof. Alexandra Z. Worden			
3:45 PM	4:00 PM	An overview of open science in eco-evo research and the publisher effect.	Dr. Nicolas Galtier			

## MEETING ROOM 118+119: 2:00 PM - 5:00 PM - S13 - Evolution after whole genome duplication: transformative advances and new directions

Session Chairs: Dr. Dearbhaile Casey, Dr. Ilia Leitch, Prof. Dan Macqueen						
2:00 PM	2:30 PM	SUMMARY NOT SEND	Dr James Clark			
2:30 PM	2:45 PM	Evolution of Gene Content and Expression Following Various Whole Genome	Dr. Florent Murat			
		Duplication Events in Vertebrates				
2:45 PM	3:00 PM	Evolution of Regulatory Elements Following Autopolyploidization in Salmonid Fish	Dr. MANU KUMAR GUNDAPPA			
3:00 PM	3:15 PM	Breaking the ploidy barrier: Genomic insights into allopolyploidy and cross-ploidy	Ms. Elizabeth Davies			
		hybridisation in British plant genus Euphrasia (eyebrights).				
3:15 PM	3:30 PM	Consequences of whole-genome duplication on expansion and adaptation to	Dr. Sandra Grünig			
		environmental heterogeneity in Biscutella laevigata				
3:30 PM	4:00 PM	Prof. Aoife Mclysaght's presentation	Prof. Aoife Mclysaght			
4:00 PM	4:15 PM	The genome evolution of young autopolyploids and their rediploidization	Dr. Luohao Xu			
4:15 PM	4:30 PM	Is there an advantage to polyploidy in animals? C. elegans neotetraploids resist severe	Dr. Laetitia Chauve			
		cold by escaping cold induced death as gravid adults.				
4:30 PM	4:45 PM	Structural variation and dysploidy shape drought responsive gene landscapes in	Dr. Luiz Augusto Cauz dos Santos			
		Nicotiana sect. Suaveolentes				
4:45 PM	5:00 PM	Evidence of lineage-specific rediploidisation after whole genome duplication in	Mr. Lukasz Niezabitowski			
		vertebrate ancestor				



#### MEETING ROOM 120+121: 2:00 PM - 5:00 PM - S27 - Genetic conflict: Evolutionary and Genomic consequences

Session Ch	nairs: Dr. Cé	cile Courret, Dr Beatriz Navarro Domínguez, Prof. Laura Ross	
2:00 PM	2:30 PM	The evolution of Programmed DNA elimination in insects.	Prof. Laura Ross
2:30 PM	2:45 PM	Routes to Sciarid strangeness: modeling chromosome elimination and a selfish linkage group	Dr. Kora Klein
2:45 PM	3:00 PM	Internal conflicts and evolutionary reversions in individuality	Dr. Martijn Schenkel
3:00 PM	3:15 PM	Contribution of transposable elements to misexpression of genes in Drosophila hybrids	Mr. William Vilas Boas Nunes
3:15 PM	3:30 PM	Influence of viral infections on transposition rates in Drosophila	Mrs. Chloé Garambois
3:30 PM	4:00 PM	Evolutionary Arms Races Shaping the Mammalian Epigenome	Dr. Antoine Molaro
4:00 PM	4:15 PM	Biodiversity and evolution of L chromosomes using germ-line sequencing data from single wild-caught fungus gnat males	Dr. Kamil Jaron
4:15 PM	4:30 PM	Multi-modal action of a mouse meiotic driver revealed by single nucleus sequencing	Dr. Reka Kelemen
4:30 PM	4:45 PM	The innovative germline-restricted chromosome of passerine birds	Dr. Francisco J. Ruiz-Ruano

## MEETING ROOM 122+123: 2:00 PM - 5:00 PM - S10 - Eco-evolutionary dynamics driven by mobile genetic elements

Session Ch	Session Chairs: Dr. Maliheh Mehrshad, Dr. Taylor Priest, Prof. J.Peter Gogarten				
2:00 PM	2:30 PM	The Secret Life of Inteins	Prof. J.Peter Gogarten		
2:30 PM	2:45 PM	A phage protein evolved to limit satellite phage induction and mobilization	Ms. Carlee Morency		
2:45 PM	3:00 PM	Genetic exchange networks bridge mobile DNA vehicles in the bacterial pathogen Listeria monocytogenes	Dr. Cheryl Andam		
3:00 PM	3:15 PM	Too Hot to Keep, Too Cool to Skip: The influence of thermal environment on the diversity of transposable elements in Polychaets	Mrs. Laure Lamothe		
3:15 PM	3:30 PM	A selective bottleneck during host entry drives the evolution of new plant symbionts	Dr. Philippe Remigi		
3:30 PM	4:00 PM	Mechanisms of virus-microbe adaptation, innovation, and coexistence	Dr. Adriana Lucia-Sanz		
4:00 PM	4:15 PM	Into the Evolutionary Arms Race: Role of Mobile Genetic Elements in Giant Viruses	Ms. Laura Sánchez		
4:15 PM	4:30 PM	Evolutionary dynamics of phage domestication across prokaryotes	Mr. Youn Le Cras		
4:30 PM	4:45 PM	Rapid genetic differentiation across short geographic scales in firefly Photinus pyralis	Dr. Rebekah Rogers		
4:45 PM	5:00 PM	Bacterial warfare plasmids are associated with virulence and antimicrobial resistance	Dr. Connor Sharp		

## MEETING ROOM 129+130: 2:00 PM - 5:15 PM - S14 - Evolution at species range margins

Session Ch	Session Chairs: Dr. Shengman Lyu, Prof. John Pannell, Prof. Anna Hargreaves				
2:00 PM	2:30 PM	'Who's afraid of assisted gene flow?' & other thorny questions about evolution at range edges	Prof. Anna Hargreaves		
2:30 PM	3:00 PM	Niche evolution at expanding range margins	Prof. Lesley Lancaster		
3:00 PM	3:15 PM	Eco-evolutionary feedback of thermal adaptation during range expansion	Ms. Saismit Naik		
3:15 PM	3:30 PM	Winters restrict a climate change-driven butterfly range expansion despite rapid evolution of seasonal timing traits at the range margin	Prof. Matthew Nielsen		
3:30 PM	3:45 PM	Sexual selection shapes eco-evolutionary dynamics at range margins	Dr. Greta Bocedi		
3:45 PM	4:00 PM	Adaptation to 44 years of climate change across a gradient from niche center to edge	Prof. Jake Alexander		
4:00 PM	4:15 PM	The impact of climate change on the adaptive potential of alpine plants	Dr. Hanna Nomoto		
4:15 PM	4:30 PM	Genetic variation in plasticity at different environmental scales determines fitness variation within and outside the elevational range margin of Senecio daisies.	Prof. Jon Bridle		
4:30 PM	4:45 PM	Digestive efficiency and mitochondrial phenotype underlie the rapid evolution of a faster pace-of-life at the margins of a range-expanding species	Ms. Sarah Jorissen		
4:45 PM	5:00 PM	The causes of elevational range limits: insights from population genomic and quantitative genetic approaches	Dr. Aaditya Narasimhan		
5:00 PM	5:15 PM	Flowering responses of the woodland strawberry to local climate and reduced precipitation along a European latitudinal gradient	Dr. Ivan De la Cruz		

## MEETING ROOM 131: 2:00 PM - 3:30 PM - S46 - The future meets the beginning: Synthetic biology, evolution, and the origin of life

Session Chairs: PhD Bruno Cuevas-Zuviría, Ms. Evrim Fer, Katsumi Hagino						
2:00 PM	2:30 PM	Protein Structure before LUCA	Dr. Klara Hlouchova			
2:30 PM	3:00 PM	Darwinian evolution of RNA replicators in artificial cell-like systems	Dr. Ryo Mizuuchi			
3:00 PM	3:15 PM	Order of amino acid recruitment into the genetic code resolved by last universal	Ms. Sawsan Wehbi			
		common ancestor's protein domains				



# MEETING ROOM 132: 2:00 PM - 3:30 PM - S51-04 - Trait evolution

Chairs: Isal	Chairs: Isabel Almudi, Jennifer Leonard						
2:00 PM	2:15 PM	Local trade-offs shape flower size evolution across Arabidopsis thaliana distribution.	Dr. Kevin Sartori				
2:15 PM	2:30 PM	Trade-offs beget trade-offs: Causal analysis of mammalian population dynamics	Dr. Juraj Bergman				
2:30 PM	2:45 PM	Single-Cell Dissection of Evolved Malnutrition Adaptation in Drosophila Larval Fat Body	Mr. Shrinath Narayanan				
2:45 PM	3:00 PM	Functional components of neurotransmitter synthesis and transport were present in the ancestor of Animals and Fungi	Mr. Yiannis Pyrris				
3:00 PM	3:15 PM	Phenotypic and genotypic evidence of physiological and ecological costs of tomato defence suppression in the spider mite Tetranychus evansi	Dr. Leonor R Rodrigues				
3:15 PM	3:30 PM	Morphological diversification of continental and insular fish radiations follow similar	Prof. Julia Day				

MEETING ROOM 131: 3:30 PM - 5:00 PM - S31 - Habitat-forming species and global change: a multidisciplinary perspective on their evolution and adaptive potential to improve their conservation

Session Chairs: Dr. Didier Aurelle, Dr. Jean-Baptiste Ledoux, Prof. Thorsten Reusch							
3:30 PM	4:00 PM	Magic meadows: the future of seagrasses under global change	Prof. Thorsten Reusch				
4:00 PM	4:30 PM	Life history constrains adaptation to climate change in habitat-forming species: insights	Dr. Ophélie Ronce				
		from 3 models					
4:30 PM	4:45 PM	Intraspecific variability in thermal tolerance shapes future declines in macroalgal	Dr. Jana Verdura				
		forests					

#### EXPO AREA: 5:00 PM - 7:00 PM - POSTER SESSION 1 - with Aperitif

Full list of posters can be found at the end of the program document



## Tuesday, 19 August 2025

## MEETING ROOM 113-117: 8:30 AM - 8:45 AM: Tuesday Access to Plenary Session & Announcements

#### PLENARY SESSION (MEETING ROOM 113-117): 8:45 AM - 9:30 AM - Plenary Talk – Keynote Speaker BEATRIZ VICOSO

8:45 AM 9:30 AM Sex (and lack thereof) in Artemia brine shrimp Prof. Beatriz Vicoso

## PLENARY SESSION (MEETING ROOM 113-117): 9:30 AM - 10:15 AM - Plenary Talk – Distinguished Fellow Talk DIETER EBERT

9:30 AM 10:15 AM Red Queen dynamics from months to megayears Mr. Dieter Ebert

#### EXPO AREA: 10:15 AM - 11:00 AM - Coffee Break

## MEETING ROOM 113: 11:00 AM - 1:00 PM - S29.01 - Genomic Basis of Evolutionary Innovations (organised by GEVOL)

Session Chairs: Prof. Erich Bornberg-Bauer, Dr. Barbara Feldmeyer, Dr. Eckart Stolle					
11:00 AM	11:30 AM	Chromosome chains and genome structural changes in termite sex chromosome evolution	Prof. Ann Kathrin Huylmans		
11:30 AM	11:45 AM	Exploring the Emergence of Newly-Evolved Expressed Open Reading Frames (neORFs) in Drosophila	Ms. Marie Kristin Lebherz		
11:45 AM	12:00 PM	MBD2/3 lost its methyl-CpG binding ability in multiple families of Holometabola	Ms. Elisa Israel		
12:00 PM	12:15 PM	The Role of Gene Copy Numbers, Gene- and Isoform-Expression on Cuticular Hydrocarbon Diversity across Hymenoptera species	Dr. Barbara Feldmeyer		
12:15 PM	12:30 PM	The epigenetic landscape across Lepidoptera and its impact on chromosomal rearrangements	Dr. Paula Escuer Pifarre		
12:30 PM	12:45 PM	Genomic determinants of gall formation and lifestyle transitions in Cynipoidea	Dr. Eliette Reboud		
12:45 PM	1:00 PM	Predicting the functional impact of single nucleotide variants in Drosophila melanogaster with FlyCADD	Ms. Julia Beets		

## MEETING ROOM 114: 11:00 AM - 1:00 PM - S23.03 - Evolutionary Genomics: Understanding and Adapting to Climate Change (organised by the journals Molecular Ecology & Evolutionary Applications)

journals ivi	orcealar Leo	iogy & Evolutional y Applications						
Session Cha	Session Chairs: Prof. Luciano Beheregaray, Prof. Maren Wellenreuther							
11:00 AM	11:30 AM	Pleiotropy as a constraint on physiological adaptation to changing environmental conditions	Prof. Cameron Ghalambor					
11:30 AM	11:45 AM	Plasticity, local adaptation and maladaptation to climate in a wild relative of the cultivated apples	Mr. Ronan Dadole					
11:45 AM	12:00 PM	Using Landscape Genomics to Predict Climate Change Vulnerability in a Large Mammal Grazer	Dr. Aja Tengstedt					
12:00 PM	12:15 PM	Local Adaptation to Climate in Wild and Feral Brassica rapa: Genetic Architecture and Genetic Offset	Mr. Sebastien Geneste					
12:15 PM	12:30 PM	A locally adapted neotropical bromeliad shows loss of macrosynteny and expansion of stress related gene families among individuals along an elevational gradient	Mr. Paulo Aecyo					
12:30 PM	12:45 PM	Can Mediterranean Gorgonian Populations Cope with Extreme Climatic Events? Insights from Common Garden Experiments and Population Genomics in Paramuricea clavata.	Dr. Jean-Baptiste Ledoux					
12:45 PM	1:00 PM	Strong Selection, but low repeatability: Temperature-specific effects on genomic predictions of adaptation	Dr. David Berger					

## MEETING ROOM 115: 11:00 AM - 1:15 PM - S16.01 - Evolution in small populations

Session Chairs: Ms. Bárbara Freitas, Dr. Maëva Gabrielli							
11:00 AM	11:30 AM	Evolutionary legacies of demographic collapse: Genomic time series reveal dynamics of small populations	Dr. Hernán Morales				
11:30 AM	11:45 AM	Recombination rate and efficiency of linked selection in small and large stickleback populations	Prof. Juha Merilä				
11:45 AM	12:00 PM	Developmental temperature, not inbreeding, shape life history and locomotor behaviors in juvenile guppies (Poecilia reticulata)	Mr. Md Mahmud Al Hasan				
12:00 PM	12:15 PM	Ancient DNA rewrites the evolutionary history of the Western Palearctic robins	Dr. Juan Carlos Illera				
12:15 PM	12:30 PM	Fighting Through the Heat: How Sexual Selection Shapes Small Population Demography Under Recurrent Heatwaves	Ms. Neelam Porwal				
12:30 PM	12:45 PM	Small populations and disease: Genomics of a severely bottlenecked species, the Critically Endangered Bellinger River Turtle (Myuchelys georgesi)	Dr. Holly Nelson				



12:45 PM	1:00 PM	Using proxies of genetic load to guide conservation management: do they deliver what	Dr. Diana Robledo-ruiz
		they promise?	
1:00 PM	1:15 PM	Adapting to a changing future: Is the 50/500 conservation rule adequate?	Prof. Leonard Nunney

Chairs: Isal	hel Almudi. I	van Gomez Mestre, Marina Marcet, Nicolas Galtier	
11:00 AM	11:15 AM	Can transmissible tumors manipulate their hosts?	Dr. Justine Boutry
11:15 AM	11:30 AM	Antibiotics of the future induce the evolution of hypervirulent bacterial pathogens	Prof. Csaba Pál
11:30 AM	11:45 AM	Ecological drift drives variation in the microbiome and its effects on the red flour beetle	Dr. Pratibha Sanjenbam
11:45 AM	12:00 PM	Non-hypermutator cancers access driver mutations through reversals in germline mutational bias	Dr. David Castellano
12:00 PM	12:15 PM	Domestication Reduces Plant Immune Receptor Gene Repertoires Across Lineages	Mr. Noah Bourne
12:15 PM	12:30 PM	Functional fate of duplicated genes	Dr. Irene Julca
12:30 PM	12:45 PM	Expanding the genomic landscape of Capuchino seedeater speciation using a	Dr. Maria Recuerda
		pangenome approach	
12:45 PM	1:00 PM	Parallel gene losses underly convergent evolution of the oral apparatus in ant-eating mammals	Dr. Frédéric Delsuc
1:00 PM	1:15 PM	Adaptive capacity, climatic vulnerability and reintroduction success of small populations: insights from an Australian freshwater fish	Prof. Luciano Beheregaray

## MEETING ROOM 117: 11:00 AM - 1:00 PM - S47.01 - The interplay between genetic architecture and the evolution of biodiversity

Session Chair: Dr. Claire Mérot						
11:00 AM	11:30 AM	Adaptive architecture of complex traits in Drosophila	Dr. Neda Barghi			
11:30 AM	11:45 AM	Genome reorganisation and expansion shape 3D genome architecture and define a distinct regulatory landscape in coleoid cephalopods	Dr. Thea Rogers			
11:45 AM	12:00 PM	Genomic Signatures of Coevolution and Climate in an Ant Social Parasitism System Reveal Links Between Signalling and Perception	Prof. Susanne Foitzik			
12:00 PM	12:15 PM	Deciphering Chromosomal Inversions And Their Evolutionary Dance In A Wing Polymorphic Beetle	Ms. Maria Madrid			
12:15 PM	12:30 PM	Structural mutations set an equilibrium non-coding genome fraction	Ms. Juliette Luiselli			
12:30 PM	12:45 PM	Pleiotropy promotes preservation of genes when a phenotypic trait is lost: Evidence from knockdown of multifunctional fatty acid synthase genes	Prof. Jacintha Ellers			
12:45 PM	1:00 PM	Convergent evolution through recurrent rearrangements and regulatory rewiring in the primate amylase locus	Dr. Charikleia Karageorgiou			

## MEETING ROOM 118+119: 11:00 AM - 1:00 PM - S38 - New Frontiers in Genome Diversity and Evolution: Exploring the 3D Organization and Function of Genomes

<b>Session Chairs:</b>	Dr	Thea Rogers	Prof.	<b>Aurora</b>	Ruiz-Herrera
ocoolon Chano.		i i iica nogcis,		Autotu	Maiz Hichicia

11:00 AM	11:30 AM	The Earth BioGenome Project and its Impact on our Understanding of Chromosome Evolution in Mammals	Dr. Harris Lewin
11:30 AM	11:45 AM	An in-vivo approach to population 3D genomics: age and sex affect 3D genome in a healthy controlled cohort of chickens	Dr. Juan Antonio Rodríguez
11:45 AM	12:00 PM	Chromosome-level genomics and historical museum collections reveal new insights into the evolution of waterbuck	Dr. Marta Farré
12:00 PM	12:30 PM	Structural Genomic Variants: Diversity, Functional Impact and Role in Evolution	Dr. Claire Mérot
12:30 PM	12:45 PM	New insights into chromatin organization and gene regulation in the germ line of vertebrates	Mrs. Laia Marín-Gual
12:45 PM	1:00 PM	Comparative Genomics to Unravel Chromosomal Instability in Butterflies	Mx. Camille Cornet
1:00 PM	1:15 PM	Evolution of 3D chromatin architecture of brown algal genomes	Mr. Pengfei Liu
1:15 PM	1:30 PM	The combination of population-level PacBio HiFi and transcriptome data enables the investigation of structural variants in regulatory evolution and speciation of two songhirds	Mrs. Carina Mugal

## MEETING ROOM 120+121: 11:00 AM - 1:00 PM - S26 - Gene flow to the rescue: Assessing the need, effectiveness, risks and ethical implications of manipulating gene flow to improve adaptation to climate change

<b>Session Chairs:</b>	Dr. Julie Gauzere,	Dr. Santiago	González Martínez,	Dr. Ophélie Ronce
------------------------	--------------------	--------------	--------------------	-------------------

11:00 AM	11:30 AM	Predictive ecological genomics: can we use correlative models to inform assisted gene	Dr. Thibaut Capblancq
		flow?	
11:30 AM	11:45 AM	Will enhanced gene flow rescue populations? Multiple genetic effects interact to	Prof. Jane Reid
		constrain the population benefits of increased immigration	
11:45 AM	12:00 PM	Adaptive introgression mediates vulnerability and facilitates adaptation to climate	Mr. Yupeng Sang
		change in Asian aspen species	
12:00 PM	12:30 PM	Genetic monitoring of plant translocations in practice	Dr. Fabienne Van Rossum



Dr. Nicole Walasek

12:30 PM 12:45 PM A century's wait for hybrids: investigating the legacy of assisted gene flow in European beech forests
 12:45 PM 1:00 PM Restoring adaptive potential: genomic insights from a 25-year plant translocation program

## MEETING ROOM 122+123: 11:00 AM - 1:15 PM - S49 - Time-dependency in micro- and macroevolutionary rates

Session Ch	airs: Dr. Ste	phen De Lisle, Prof. Lesley Lancaster, Prof. Erik Svensson	
11:00 AM	11:30 AM	Noise leads to the perceived increase in evolutionary rates over short time scales	Dr. Jeremy Michael Beaulieu
11:30 AM	12:00 PM	The importance of quantifying rate uncertainty	Dr. Rosana Zenil-Ferguson
12:00 PM	12:15 PM	The rise, decline and fall of clades	Dr. Ignacio Quintero
12:15 PM	12:30 PM	Measuring morphological changes in unit of nucleotide changes to disentangle neutral and adaptive processes	Dr. Thibault Latrille
12:30 PM	12:45 PM	The role of mutational neighborhoods and evolvability in shaping evolutionary tempo	Prof. Guillaume Beslon
12:45 PM	1:00 PM	Estimates of the Mutation Rate per Year Can Explain Why the Molecular Clock Depends on Generation Time	Ms. Loveday Lewin
1:00 PM	1:15 PM	Rate-time scaling in phenotypic evolution: Limitations of current models in capturing temporal dynamics	Ms. Vilde Bruhn Kinneberg

## MEETING ROOM 129+130: 11:00 AM - 1:00 PM - S03.02 - Advances in technology, mathematical and statistical models and their application in evolutionary ecology: the dawn of a new era

Session Ch	Session Chairs: Prof Andy Gardner, Dr. Juliano Morimoto					
11:00 AM		•	r Dr Callum Malaan			
11.00 AW	11:30 AM	Using machine learning to measure phenotypic evolution: butterflies as a case study for emerging technologies	Dr. Callum McLean			
11:30 AM	11:45 AM	Eco-evolutionary dynamics of finite populations from first principles and the noise-	Mr. Shikhara Bhat			
		induced reversal of natural selection				
11:45 AM	12:00 PM	The value of 'simple' models in the era of complexity: Case studies from sexual	Dr. Jussi Lehtonen			
		selection				
12:00 PM	12:15 PM	How does species diversity affect ecological stability?	Dr. Ming Liu			
12:15 PM	12:30 PM	Phylogenetic probabilistic PCA	Mx. Paola Montoya			
12:30 PM	12:45 PM	Neural ODEs for understanding population dynamics	Dr. Matishalin Patel			

#### MEETING ROOM 131: 11:00 AM - 1:00 PM - S19 - Evolution of symbioses and interactions in microbial eukaryotes

12:45 PM 1:00 PM The evolution of reversible plasticity in stable environments

Session Cha	airs: Prof. A	nna Karnkowska, Prof. Alexandra Z. Worden	
11:00 AM	11:30 AM	Unlocking the molecular complexity of endosymbiosis in diverse model systems	Dr. Ben Jenkins
11:30 AM	12:00 PM	Anaerobic protist survival in microcosms is dependent on microbiome metabolic function.	Dr. Courtney Stairs
12:00 PM	12:15 PM	Subcellular proteomics illuminates the relationship between host and symbiont in trypanosomatid Angomonas deanei	Dr. Michael Hammond
12:15 PM	12:30 PM	Interplay of HGT and targeting during the chromatophore organellogenesis in Paulinella	Mr. Moisès Bernabeu
12:30 PM	12:45 PM	Stepwise genome evolution from a facultative symbiont to an endosymbiont in the N2-fixing diatom-Richelia symbioses	Dr. Vesna Grujcic
12:45 PM	1:00 PM	An evolutionary mosaic of proteins mediates metabolic integration in transient kleptoplasts in the euglenozoan Rapaza viridis	Dr. Ivan Garcia Cunchillos

#### EXPO AREA: 12:30 PM - 2:00 AM - Lunch

#### MEETING ROOM 113: 2:00 PM - 5:00 PM - S29.02 - Genomic Basis of Evolutionary Innovations (organised by GEVOL)

Session Ch	nairs: Prof. E	rich Bornberg-Bauer, Dr. Barbara Feldmeyer, Dr. Eckart Stolle	
2:00 PM	2:30 PM	Arthropod comparative evolutionary and functional genomics	Dr. Robert Waterhouse
2:30 PM	2:45 PM	Patterns of Gain and Loss of microRNA Families Explain Divergence in Mammal	Dr. Jonathan Fenn
		Pregnancy Phenotype	
2:45 PM	3:00 PM	Template-switching and rapid emergence of novel gene regulators	Dr. Heli Mönttinen
3:00 PM	3:15 PM	Evolution of salivary protein repertoire in true bugs and its role in dietary adaptations	Dr. Maike Fischer
3:15 PM	3:30 PM	Origin of evolutionary innovations: from cell and developmental biology to trait utility and diversification.	Dr. Abderrahman Khila
3:30 PM	3:45 PM	The Escherichia coli genome is relatively devoid of cryptic promoters and repressor binding sites	Dr. Timothy Fuqua



3:45 PM	4:00 PM	Deconstructing the contribution of lineage-specific transposable element insertions to nervous and sensory system diversification in Lake Malawi cichlid fishes	Dr. Aleksandra Marconi
4:00 PM	4:15 PM	From digestion to communication: molecular evolution during the rise of a social fluid	Mr. Arthur Matte
4:15 PM 4:30 PM	4:30 PM 4:45 PM	Evolutionary novelties escape loss by recruiting life-history trade-offs  The role of transposable elements in the adaptation of amphipod crustaceans to the deep sea.	Prof. Nathan Bailey Ms. Matilda Scott
4:45 PM	5:00 PM	Olfaction evolving: a model of the evolutionary transition from deterministic to stochastic odorant receptor gene choice	Dr. Roman Zug

## MEETING ROOM 114: 2:00 PM - 5:00 PM - S36.01 - Microevolutionary processes and Macroevolutionary patterns.

Session Ch	airs: Dr. Th	éo Gaboriau, Dr. Carolin Kosiol, Dr. Thibault Latrille	
2:00 PM	2:30 PM	Phenotypic trade-offs as a bridge in micro- and macroevolution	Dr. Lee Hsiang Liow
2:30 PM	2:45 PM	Measuring the Macroevolutionary Adaptive Landscape	Dr. Masahito Tsuboi
2:45 PM	3:00 PM	The diffused evolutionary dynamics of morphological novelty	Dr. Ignacio Quintero
3:00 PM	3:15 PM	Diving depth and light environment influence countershading in pinnipeds	Prof. Changku Kang
3:15 PM	3:30 PM	A general evolutionary model for the emergence of novel characters from serial homologs	Dr. Daohan Jiang
3:30 PM	3:45 PM	Diverging selection on body size in specialist mammals	Prof. Xiang-Yi Li Richter
3:45 PM	4:00 PM	From phylogenomics to population genomics: the role of geography versus ecology in driving Coenonympha butterflies diversification	Mrs. Laurence Després
4:00 PM	4:15 PM	Evidence for Bergmann's and Allen's rules operating between species across the world's mammals	Mr. James Nathan Tang
4:15 PM	4:30 PM	Building upon a speciation theory to predict the link between microevolutionary processes and speciation time	Mr. Pierre Veron
4:30 PM	4:45 PM	Within-population variance predicts evolution in a coccolithophore lineage transitioning between adaptive zones on the macroevolutionary adaptive landscape	Dr. Kjetil Voje
4:45 PM	5:00 PM	Applications of stochastic reproduction-selection models for the study of non-equilibrium dynamics across different evolutionary time-scales	Mrs. Carina Mugal

## MEETING ROOM 115: 2:00 PM - 5:00 PM - S16.02 - Evolution in small populations

Session Chair: Ms. Bárbara Freitas					
2:00 PM	2:30 PM	Who let the frogs out? Insights from an experimental island population	Prof. Eva Ringler		
2:30 PM	2:45 PM	Dispersal evolution as a driver of island biodiversity	Mr. Siebe Van Wunnik		
2:45 PM	3:00 PM	A natural experiment in brown trout Salmo trutta demonstrates the impact of demography on linked selection	Dr. Oliver Stuart		
3:00 PM	3:15 PM	Extreme climatic event causes genetic bottlenecks across a wild butterfly metapopulation: genomic consequences for population adaptative potential	Prof. Marjo Saastamoinen		
3:15 PM	3:30 PM	Investigating mutation load associated with adaptation in threespine stickleback	Dr. Jana Nickel		
3:30 PM	3:45 PM	Rapid vs. Slow Habitat Loss: Species Traits Shape Genetic Diversity Patterns During Contraction	Mr. Ravi Vishwakarma		
3:45 PM	4:00 PM	Sky-Islands under pressure: how geography, climate, and genomic shape the history of an endemic Pyrenean species	Ms. Noèmie Collette		
4:00 PM	4:15 PM	Genetic and dental variation: The effects of small population size in the endangered Saimaa ringed seal	Mrs. Emmi Olkkonen		
4:15 PM	4:30 PM	Phenotypic and genomic impacts of using inbred versus outbred rescuers in the experimental genetic rescue of small, inbred populations	Prof. David Richardson		
4:30 PM	4:45 PM	Variation among neighbouring wild populations of Nicrophorus vespilloides in mutation load and dependence on parental care	Ms. Willow Dalehite		
4:45 PM	5:00 PM	Coping with insularly: genome-wide effects of isolation and genetic drift in highly diverse primates.	Dr. Ivo Colmonero Costeira		

## MEETING ROOM 116: 2:00 PM - 5:15 PM - S51-07 - Genetics

Chairs: Ivan Gomez Mestre, Nicolas Galtier					
2:00 PM	2:15 PM	Interplay between low-recombining regions and overdominance in a plant genome?	Dr. Marine SALSON		
2:15 PM	2:30 PM	Gene expression variability and evolutionary outcomes after whole-genome duplication in fishes	Prof. Marc Robinson-rechavi		
2:30 PM	2:45 PM	Uncovering the genetic and mechanistic basis of phenotypic polymorphisms in island populations of wall lizards	Mrs. Stéphanie Sherpa		
2:45 PM	3:00 PM	Repeated emergence of specialised ribosomes suggests a complex role for deterministic evolutionary forces across eukaryotes	Dr. Alan Beavan		



3:00 PM	3:15 PM	See you in the dark: molecular evolution of extraordinary vision in deep-sea fishes	Dr. Zuzana Musilová
3:15 PM	3:30 PM	Machine learning evaluation of evolutionary accelerated regions reveals genomic basis of adaptation in bats	Dr. Bai-wei Lo
3:30 PM	3:45 PM	Dual roles of DNA methylation in gene regulation and chromatin structure of a hemimetabolous insect	Mr. Nicholas Planidin
3:45 PM	4:00 PM	Complex genomic content allows for rapid giant virus adaptation to a novel host	Dr. Anouk Willemsen
4:00 PM	4:15 PM	Genetic architectures preventing species fusion between the Japanese sticklebacks	Takuya Hosoki
4:15 PM	4:30 PM	Junk or Dark DNA? Using a human-Arabidopsis hybrid cell line to realise the Random Genome Project at mega-base Scale	Mr. Brett Adey
4:30 PM	4:45 PM	Unicellular regulatory landscapes and the origins of animal multicellularity	Dr. Marta Alvarez Presas
4:45 PM	5:00 PM	Asymmetry in vertebrate gene repertoires: what happened to the shark HoxC cluster?	Dr. Shigehiro Kuraku
5:00 PM	5:15 PM	Genomic insights into the evolutionary potential of sharks in a changing ocean	Mrs. Juliana Sofia Alves

## MEETING ROOM 117: 2:00 PM - 5:00 PM - S47.02 - The interplay between genetic architecture and the evolution of biodiversity

Session Ch	Session Chair: Dr. Claire Mérot					
2:00 PM	2:15 PM	The Role of Sperm Competition and Genome Duplication in Cell Type Diversity in Fish Gonads	Dr. Francisca Hervas-sotomayor			
2:15 PM	2:30 PM	Gene network topology drives the mutational landscape of gene expression	Dr. Sylvain Pouzet			
2:30 PM	2:45 PM	Genomic architecture and functional constraints associated with chromosomal fusions in natural populations	Prof. Aurora Ruiz-Herrera			
2:45 PM	3:00 PM	Exploring unique genetic diversity in Dutch cattle breeds using a pangenome graph	Mr. Rensco Hogers			
3:00 PM	3:15 PM	Speciation genomics in Bryophytes: unraveling the genetic architecture of reproductive isolation in haplo-diplontic plants.	Mr. Fabien Rey-giraud			
3:15 PM	3:30 PM	Chromosomal inversions, polymorphic duplications and parallel patterns of differentiation in a marine bivalve, the European flat oyster (Ostrea edulis)	Ms. Lila COLSTON-NEPALI			
3:30 PM	3:45 PM	Detecting polygenic selection in time series data.	Dr. Margarita Takou			
3:45 PM	4:00 PM	A mosaic of modular variation at a single gene underpins convergent plumage coloration	Dr. Madeline Chase			
4:00 PM	4:15 PM	How do Atlantic salmon respond to selective fishing under various genetic architectures of a life-history trait? An individual-based demo-genetic model as a virtual laboratory	Dr. Amaïa Lamarins			
4:15 PM	4:30 PM	The evolution of sex-specific genetic architectures	Dr Peter D Price			
4:30 PM	4:45 PM	The genetic architecture of an adaptive phenotype conditions populations evolutionary response to climate change	Dr. Genís Garcia-erill			

## MEETING ROOM 118+119: 2:00 PM - 5:00 PM - S51-08 - Ethology

Chairs: Pa	u Carazo, Ai	da Verdes	
2:00 PM	2:15 PM	Individual sociability accelerates call combination learning in Western Australian magpies	Ms. Stephanie Mason
2:15 PM	2:30 PM	Mechanisms of Community Assembly through the lens of Phylogenetic Diversity: a Critical Reappraisal.	Mr. Thibault Kasprzyk
2:30 PM	2:45 PM	The evolution of masturbation in birds	Dr. Chloe Heys
2:45 PM	3:00 PM	Dietary choice and life history in D. melanogaster	Ms. Oonagh Barker
3:00 PM	3:15 PM	Phenotypic Plasticity in Predator-Induced Defenses: Species- and Genus-Level Responses to Invertebrate Predation in Cladocerans	Dr. Marjohn Baludo
3:15 PM	3:30 PM	Why do the songbirds perform tap dance?: Behavioral mechanisms and functions of mutual multimodal courtship displays	Dr. Nao Ota
3:30 PM	3:45 PM	The role of song in population divergence: Insights from the La Palma's Canary Islands Chaffinch	Ms. Bárbara Freitas
3:45 PM	4:00 PM	Plasticity and evolution of metabolic division of labour within families	Dr. Eleanor Bladon
4:00 PM	4:15 PM	The origin and spread of sorghum in Africa: a multidisciplinary perspective integrating genomics, archaeobotanical and ethnolinguistic data	Dr. Aude Gilabert
4:15 PM	4:30 PM	Genetic Patterns of Social Structuring in the Cuckoo Catfish	Mr. Lukas Koch
4:30 PM	4:45 PM	Natural variation in the gut microbiota is associated with behavioural differences in wild mice. But does it cause them?	Dr. Alexandre Figueiredo
4:45 PM	5:00 PM	Individual social bonds predict fitness in a cooperative bird	Dr. Gabriel Munar Delgado



#### MEETING ROOM 120+121: 2:00 PM - 4:00 PM - S33 - Linking recombination rates and supergene evolution with the genomics of complex traits

Session Ch	Session Chairs: Dr. Eyal Privman, Dr. Jonathan Romiguier				
2:00 PM	2:30 PM	Supergene evolution between coadaptation and degeneration	Dr. Mathieu Joron		
2:30 PM	2:45 PM	Genomic diversity and evolution in the Anopheles gambiae complex: the extreme specialisation of Anopheles bwambae to thermal springs in Uganda	Dr. Michael Fontaine		
2:45 PM	3:00 PM	Convergent evolution of inversion-based supergenes in neotropical lepidoptera	Mr. Edward Page		
3:00 PM	3:30 PM	What determines interspecific gene flow in butterflies?	Dr. Konrad Lohse		
3:30 PM	3:45 PM	Small Queens, Big Questions: Investigating the Interaction between Two Supergenes in the Ant Formica cinerea	Dr. Giulia Scarparo		
3:45 PM	4:00 PM	Supergene-associated variation in queen size and number in the ant Temnothorax longispinosus	Ms. Ina Knuf		

## MEETING ROOM 122+123: 2:00 PM - 4:00 PM - S11 - Enhancing Diversity and Transparency in Ecology and Evolution: Reliable Practices for Research and Organisations

Session Chairs: Dr. Edward Ivimey-Cook, Ms Marija Purgar Filjak					
2:00 PM	2:30 PM	Transforming Science Culture: Inclusive Meta-Research for Transparency	Dr. Malgorzata Lagisz		
2:30 PM	2:45 PM	Career breaks and care-giving: challenges and opportunities for flexible working in evolutionary ecology	Dr. Sinead English		
2:45 PM	3:00 PM	Towards an open and responsible publication model	Prof. Jacek Radwan		
3:00 PM	3:30 PM	The role of journals in promoting a diverse, credible and open research culture in ecology and evolutionary biology	Dr. Joel Pick		
3:30 PM	3:45 PM	Three species, three labs, three experiments: testing the reproducibility of evolutionary ecological studies on insect behaviour in a multi-laboratory setting	y- Prof. Joachim Kurtz		
3:45 PM	4:00 PM	Lessons learned from the COVID-19 pandemic on Open, reliable, and transparent practices	Dr. Florence Débarre		

#### MEETING ROOM 129+130: 2:00 PM - 5:15 PM - S04 - Ageing outside of the box: insights from unusual and non-model species

Session Ch	airs: Dr. Ma	rgaux Bieuville, Dr. E. Yagmur Erten	
2:00 PM	2:30 PM	'Lenses' bring ageing into focus: Senescence in a tiny, floating macrophyte	Dr. Robert Laird
2:30 PM	2:45 PM	The selection shadow measured at the single-cell type level	Dr. Mehmet Somel
2:45 PM	3:00 PM	Peto's Paradox for fungi	Dr. Ben Auxier
3:00 PM	3:15 PM	The antagonistic pleiotropy and disposable soma theories of ageing differ in their	Dr. Joost Van Den Heuvel
		predictions of age specific survival and fecundity	
3:15 PM	3:30 PM	Aging in ants: from the individual to the colony level	Dr. Luisa M. Jaimes Nino
3:30 PM	4:00 PM	Death might keep the clone alive	Prof. Thorsten Reusch
4:00 PM	4:15 PM	Divergence in allelic variation and gene expression in molecular networks that underly	Dr. Tonia Schwartz
		aging between fast- and slow-aging garter snakes.	
4:15 PM	4:30 PM	Germline mutation rate and somatic maintenance improved simultaneously by reduced	Dr. Elizabeth Duxbury
		insulin signalling	
4:30 PM	4:45 PM	Lifespan extension in Heliconius butterflies: from ultimate to proximate mechanisms	Dr. Stephen Montgomery
4:45 PM	5:00 PM	Senescence evolution under the catastrophic accumulation of deleterious mutations	Dr. Thomas Aubier
5:00 PM	5:15 PM	Senescence in an unusual invertebrate model: insights from the tsetse fly	Dr. Sinead English

## MEETING ROOM 131: 2:00 PM - 5:00 PM - S51-09 - Sex

Chairs: Jen	nifer Leonai	rd, Patricia Álvarez	
2:00 PM	2:15 PM	Genetic architecture of male reproductive success in a lekking bird: insights from	Ms. Rebecca Chen
		predicted deleterious mutations	
2:15 PM	2:30 PM	The effects of sex-specific genetic variation and combined thermal and nutritional	Mr. Aidan Stuckey
		stress on body shape	
2:30 PM	2:45 PM	Resurrecting a 100-year-old hypothesis: Effects of sheltering on sex chromosome	Dr. Andrea Mrnjavac
		degeneration and sex-biased gene content evolution	
2:45 PM	3:00 PM	Indications of different sex determination systems in two closely related Eurasian	Dr. Madlen Stange
		minnow (Phoxinus spp.) species	
3:00 PM	3:15 PM	Stage-Specific Diets Shape Sex-Specific Plasticity and Evolution of Stress Resistance	Dr. Sudipta Tung
		Across Age in Drosophila melanogaster	
3:15 PM	3:30 PM	Sex chromosome turnovers in fish	Dr. Astrid Böhne
3:30 PM	3:45 PM	Differences in responses to seminal fluid between European populations of the	Dr. Joris Koene
		simultaneous hermaphrodite Lymnaea stagnalis	
3:45 PM	4:00 PM	A comparison of selection on autosomes and the young X chromosome in Mercurialis	Mr. Suhaas Sehgal
		annua	
4:00 PM	4:15 PM	On the evolution and mechanisms of reproductive diapause in tropical insects	Mr. Marcus Hicks



4:15 PM	4:30 PM	Disentangling plasticity from genetics: reproductive effort of urban and forest great tits	Ms. Laurine Mathieu
		in a common garden experiment	
4:30 PM	4:45 PM	Diversification of stickleback sex chromosomes driven by introgression	Ms. Dandan Wang
4:45 PM	5:00 PM	The effects of costly telomere maintenance on lifespan - reproductive tradeoffs in sand	Dr. Mats Olsson
		lizards	

MEETING ROOM 120+121: 4:00 PM - 5:30 PM - S43 - The dynamics and consequences of bacteria-bacteriophage interactions and co-evolution in complex communities

Session Ch	Session Chairs: Prof. Ville-Petri Friman, Dr. Pauline Scanlan					
4:00 PM	4:30 PM	Prof. Michael Brockhurst presentation	Prof. Michael Brockhurst			
4:30 PM	5:00 PM	Exploring multi-level coevolution and evolutionary conflict in wild marine phage—bacteria systems	Mrs. Frederique Le Roux			
5:00 PM	5:15 PM	Adaptive strategies during coevolution depends on multipartite interactions between bacteria and their temperate phages.	Dr. Olaya Rendueles			
5:15 PM	5:30 PM	The phageome of apricot trees and its association with bacterial canker disease	Dr. Clara Torres Barceló			

MEETING ROOM 122+123: 5:00 PM - 6:30 PM - ESEB 2025 Satellite Symposium: Engineering Ecosystem Resilience: An interactive ARIA Workshop

EXPO AREA: 5:00 PM - 7:00 PM - POSTER SESSION 2 - with Aperitif

Full list of posters can be found at the end of the program document



## Wednesday, 20 August 2025

## MEETING ROOM 113-117: 8:45 AM - 9:00 AM: Wednesday Access to Plenary Session & Announcements

## PLENARY SESSION (MEETING ROOM 113-117): 09:00 AM - 09:45 AM - Plenary Talk – Keynote Speaker Love Dalén

9:00 AM 9:45 AM Evolutionary insights from deep-time palaeogenomes Prof. Love Dalén

EXPO AREA: 9:45 AM - 10:30 AM - Coffee Break

## MEETING ROOM 113: 10:30 AM - 12:45 PM - S28.01 - Genome Architecture and Their Role in Evolution

Session Ch	Session Chairs: Dr. Marcial Escudero, Prof. Kay Lucek, Dr. Petr Nguyen				
10:30 AM	11:00 AM	Escaping the trap! How flies evolved new sex chromosomes	Dr. Melissa Toups		
11:00 AM	11:15 AM	Non-canonical sex chromosome evolution revealed by extreme heterogeneity in homomorphic Y chromosome differentiation in the common frog	Prof. Wen-juan Ma		
11:15 AM	11:30 AM	Recombination suppression through structural variation leads to sex chromosome evolution in the plant Silene exscapa	Mr. Fabien Duez		
11:30 AM	11:45 AM	What shapes the degeneration of neo-sex chromosomes across Lepidoptera?	Mr. Thomas Decroly		
11:45 AM	12:00 PM	Do Newly Z-linked Sequences in Butterflies and Moths Show Evidence of Enhanced Protein Adaptation?	Dr. Claudia Weber		
12:00 PM	12:15 PM	Evolutionary dynamics of enlarged sex chromosomes and novel pseudoautosomal regions in Sylvioidea songbirds	Prof. Bengt Hansson		
12:15 PM	12:30 PM	Sex chromosome divergence and barriers to gene flow in brown algae with haploid–diploid life cycles	Dr. Agnieszka Lipinska		

## MEETING ROOM 114: 10:30 AM -12:30 PM - S36.02 - Microevolutionary processes and Macroevolutionary patterns

Session Ch	airs: Dr. Thé	o Gaboriau, Dr. Carolin Kosiol, Dr. Thibault Latrille	
10:30 AM	11:00 AM	Conceptual and empirical bridges between micro- and macroevolution	Dr. Jonathan Rolland
11:00 AM	11:15 AM	A computational model for the macroevolutionary bias towards simple leaf shape in flowering plants.	Mr. James Malone
11:15 AM	11:30 AM	From micro to macroevolution: Evolvability predicts skull evolution across mammals	Dr. Daniela Rossoni
11:30 AM	11:45 AM	Genetic constraints unaffected by selection predicts evolution within a fossil lineage	Dr. Meghan Balk
11:45 AM	12:00 PM	How reliable are phenotypic proxies of evolvability? Evaluating the relationships between P- and G-matrices across mammals	Dr. Barbara Costa
12:00 PM	12:15 PM	Merging evolutionary timescales to quantify adaptation	Ms. Ioanna Kotari
12:15 PM	12:30 PM	Bridging micro and macroevolution: Insights from chromosomal dynamics in holocentric true sedges	Dr. Marcial Escudero

## MEETING ROOM 115: 10:30 AM -12:30 PM - S44.01 - The ecological and evolutionary implications of climate change on reproduction

Session Ch	Session Chairs: Prof Amanda Bretman, Dr Liam Dougherty, Dr. Claudia Fricke, Prof. Rhonda Snook				
10:30 AM	11:00 AM	Elevational variation in gametophytic thermal performance and its influence on floral thermoregulatory evolution	Dr. Matthew Koski		
11:00 AM	11:15 AM	The effect of of intrasexual and intersexual selection on survival and reproduction under a heatwave.	Ms. Karendeep Sidhu		
11:15 AM	11:30 AM	Life-stage- and sex-specific fitness dynamics of thermal stress intensity and duration in Drosophila prolongata	Dr. Abhishek Meena		
11:30 AM	11:45 AM	Genotypic responses to different environments and reduced precipitation reveal signals of local adaptation and phenotypic plasticity in woodland strawberry	Dr. Ivan De la Cruz		
11:45 AM	12:00 PM	Impact of thermal stress on colony foundation in ants	Dr. Romain Libbrecht		
12:00 PM	12:15 PM	Climate change impacts on pup production, survival and recruitment in Antarctic fur seals	Mrs. Anna Paijmans		
12:15 PM	12:30 PM	Reproductive success under hypoxia: unveiling the adaptive potential of a freshwater fish	Dr. Amélie Crespel		
12:30 PM	12:45 PM	Do heatwaves affect insect reproductive behaviour?	Dr. Cristina Tuni		



#### MEETING ROOM 116: 10:30 AM -12:30 PM - S22.01 - Evolutionary Ecology of Microbial Symbioses

Session Chairs: Dr. Anne Duplouy, Prof. Wolfgang Miller					
10:30 AM	11:00 AM	The bacterial march to endosymbiosis: on-ramps and off-ramps	Ms. Marjolein Bruijning		
11:00 AM	11:15 AM	Symbiosis-mediated gene transfer in early-diverging Metazoan	Dr. Sofia Paraskevopoulou		
11:15 AM	11:30 AM	Eco-evolutionary context shapes fitness and diversity in a protective symbiosis	Dr. Georgia Drew		
11:30 AM	11:45 AM	Unraveling the mechanisms driving transitions to parthenogenesis in a Wolbachia-	Ms. Kristine Jecha		
		infected ladybug			
11:45 AM	12:00 PM	Could co-evolution explain Wolbachia-driven loss of pesticide resistance in spider mite	Dr. Flore Zélé		
		populations?			
12:00 PM	12:15 PM	Infection dynamics of endosymbionts that manipulate arthropod reproduction	Dr. Franziska A. Brenninger		
12:15 PM	12:30 PM	Spread of Wolbachia in haplodiploids is driven by extensive rewiring of female	Ms. Emma Van Reempts		
		reproductive physiology			

#### MEETING ROOM 117: 10:30 AM -12:30 PM - S47.03 - The interplay between genetic architecture and the evolution of biodiversity

Session chair: Dr. Claire Mérot						
10:30 AM	11:00 AM	The Evolution of Polymorphic Mimicry and Supergenes in Butterflies	Prof. Krushnamegh Jagannath Kunte			
11:00 AM	11:15 AM	Colour genes and clock genes: Genomic drivers of local adaptation in the wood tiger moth	Dr. Melanie Brien			
11:15 AM	11:30 AM	The evolutionary trajectory of seahorse inversions is shaped by multiple types of selection	Ms. Laura Meyer			
11:30 AM	11:45 AM	Highly modular genomic architecture underlies rapid adaptive radiation	Ms. Pooja Singh			
11:45 AM	12:00 PM	Gene copy number variations are untapped key players in adaptation along environmental gradient	Dr. Qiujie Zhou			
12:00 PM	12:15 PM	Genetics of repeated ecotype evolution in a Malawi cichlid	Mr. Luka Moritz Blumer			
12:15 PM	12:30 PM	One for all: the role of the FAR5 gene in a repeated polygenic alpine adaptation	Mrs. Alžběta Poupětová			

## MEETING ROOM 118+119: 10:30 AM -12:30 PM - S15.01 - Evolution in and of diverse genetic systems

Session Cha	Session Chairs: Dr. Thomas Hitchcock, Dr. Kora Klein, Dr. Martijn Schenkel					
10:30 AM	11:00 AM	Crazy reproduction in crazy ants	Dr. Hugo Darras			
11:00 AM	11:15 AM	Disentangling the genomic architecture of sex chromosomes in an androdioecious species with two mating types	Dr. Adrian Contreras Garrido			
11:15 AM	11:30 AM	Evolution of divergent male and female mitochondrial genomes in a basal protobranch bivalve Yoldia hyperborea	Mr. lakov Korobitsyn			
11:30 AM	11:45 AM	Evolutionary drivers and consequences of monogeny: a unusual reproductive system in fungus gnats	Ms. Melany Henot			
11:45 AM	12:00 PM	Stepwise recombination suppression around the mating-type locus associated with a diploid-like life cycle in Schizothecium fungi	Mrs. Elsa De Filippo			
12:00 PM	12:15 PM	How parasites can influence recombination in an almost fully parthenogenetic population : the case of the Bedegar gall wasp.	Dr. Antoine Branca			
12:15 PM	12:30 PM	The cheater-driven evolution of reproductive division of labour	Dr. Nobuto Takeuchi			

## MEETING ROOM 120+121: 10:30 AM -12:30 PM - S37.01 - Museomics: Challenges and Possibilities

Session cha	Session chair: Prof. Carles Lalueza-fox				
10:30 AM	11:00 AM	Museomics for understanding adaptive potential - what else do we need to develop to reach this goal?	Prof. M. Thomas P. Gilbert		
11:00 AM	11:15 AM	Temporal genomics reveals widespread but unexpected consequences of a bottleneck in the Scandinavian brown bear	Ms. Amanda Lindahl		
11:15 AM	11:30 AM	Ancient Genomes, Modern Climates: Tracking Alpine Plant Changes Through Herbarium Records	Ms. Yasaman Ranjbaran		
11:30 AM	11:45 AM	Combining eDNA and Museomics to Enhance Biodiversity Monitoring	Dr. Sarah Schmid		
11:45 AM	12:00 PM	Harnessing the potential of historical DNA in population genomic studies on insects - a case study	Mr. Michael Mitschke		
12:00 PM	12:15 PM	High-throughput palaeogenomics reveals lost bovine diversity	Mr. Alexandre Gilardet		
12:15 PM	12:30 PM	Uncovering Lost Diversity: Insights into the Genomic Structure and Population Dynamics of European Wolves using Museum Specimens	Dr. Evelyn Todd		

## MEETING ROOM 122+123: 10:30 AM -12:30 PM - S48.01 - The maintenance of adaptive polymorphisms

Session Chairs: Dr. Hannah Augustijnen, Dr. Harshavardhan Thyagarajan				
10:30 AM	11:00 AM	The maintenance and detection of sexually antagonistic genetic variation	Dr. Karl Grieshop	
11:00 AM	11:15 AM	The micro- and macroevolutionary predictability of female-limited phenotypic	Prof. Erik Svensson	
		polymorphisms		
11:15 AM	11:30 AM	Ecotype evolution in the deep sea	Prof. Rus Hoelzel	



11:30 AM	11:45 AM	Sex-specific overdominance for reproductive fitness at a large-effect maturation locus in wild Atlantic salmon	Prof. Craig Primmer
11:45 AM	12:00 PM	Genetic architecture and the maintenance of variation by sexual selection in the guppy	Dr. Wouter van der Bijl
12:00 PM	12:15 PM	Genetic interactions shape the evolution of adaptive polymorphism under resource competition	Mr. Suman Das
12:15 PM	12:30 PM	Antagonistic selection and genetic drift in eusocial ant colonies	Dr. Malvika Srivastava

MEETING ROOM 129+130: 10:30 AM -12:45 PM - S02 - Addressing new and long-standing evolutionary questions with linkage disequilibrium based approaches

Session Ch	airs: Dr. Pao	lo Franchini, Prof. Paolo Momigliano, Dr. Francesca Raffini	
10:30 AM	11:00 AM	Genomic signatures of intra-chromosomal epistasis in hybrid populations	Prof. Claudia Bank
11:00 AM	11:15 AM	Detecting genomic incompatibilities and barriers during divergence with gene flow in Formica ants	Mr. Patrick Heidbreder
11:15 AM	11:30 AM	Utilizing linkage disequilibrium (LD) to improve signal to noise ratios in outlier analyses	Dr. Petri Kemppainen
11:30 AM	12:00 PM	Inferring human evolutionary history using linkage disequilibrium	Prof. Aaron P Ragsdale
12:00 PM	12:15 PM	Inferring the evolutionary history of Caenorhabditis elegans while accounting for its self fertilising reproductive mode	· Ms. Chenxi Wang
12:15 PM	12:30 PM	Genomic incompatibilities and admixture between oviparous and viviparous lizards at a hybrid zone	Prof. Kathryn Elmer
12:30 PM	12:45 PM	Integrating LD- and Coalescent-Based Approaches Reveal Stability-Driven Genetic Diversity in the Adaptive Radiation of Lake Tanganyika Cichlids	Mrs. Daniela Souza Costa

#### MEETING ROOM 131: 10:30 AM -12:30 PM - S35 - Mechanisms, barriers, and impacts of horizontal gene transfer across the Tree of Life

#### Session Chairs: Dr. Cheryl Andam, Dr. Ben Pascoe

10:30 AM	11:00 AM	Horizontal gene transfer as an evolutionary lifeline in intracellular bacteria	Prof. Matthias Horn
11:00 AM	11:15 AM	Horizontal transfers of polydnavirus reveals new hosts of parasitoid wasps	Ms. Ines Matrougui
11:15 AM	11:30 AM	Multiple horizontal transfers of immune genes between vertebrates	Dr. Maxime Policarpo
11:30 AM	12:00 PM	Who Moves Whom and For Whose Benefit?	Dr. Eduardo Rocha
12:00 PM	12:15 PM	Introgression recapitulates an ancient hybridisation phenotype in a Passer sparrow	Dr. Melissah Rowe
12:15 PM	12:30 PM	Horizontal gene transfers can reveal ghost lineages	Mr. Enzo Marsot

EXPO AREA: 12:30 PM - 2:00 AM - Lunch

MEETING ROOM 122+123: 2 PM - 3:30 PM - Satellite Symposium: ERC SESSION - ERC grantee

Speakers: Dr. Alvaro San Millan

MEETING ROOM 120+121: 2 PM - 4 PM - ESEB 2025 Satellite Symposium: Ecological Genomics of Coevolution

MEETING ROOM 131: 2 PM - 4 PM - ESEB 2025 Satellite Symposium: EvoKE workshop on communicating evolution effectively



## Thursday, 21 August 2025

## MEETING ROOM 113-117: 8:45 AM - 9:00 AM: Thursday Access to Plenary Session & Announcements

#### PLENARY SESSION (MEETING ROOM 113-117): 09:00 AM - 09:45 AM - Plenary Talk – Keynote Speaker Charissa de Bekker

9:00 AM 9:45 AM Zombihaviour: Unraveling the zombie-making strategies of Ophiocordyceps fungi to Dr. Charissa de Bekker hijack ant behaviour

EXPO AREA: 9:45 AM - 10:30 AM - Coffee Break

#### MEETING ROOM 113: 10:30 AM - 12:15 PM - S28.02 - Genome Architecture and Their Role in Evolution

Session cha	Session chairs: Dr. Marcial Escudero, Prof. Kay Lucek, Dr. Petr Nguyen					
10:30 AM	11:00 AM	Genomic rearrangements involved in speciation and evolution of sexual system in nematodes.	Dr. Kohta Yoshida			
11:00 AM	11:15 AM	Genomic rearrangements and the stepwise evolution of asexuality	Dr. Jeremias Brand			
11:15 AM	11:30 AM	Implications of Sex-Specific Genomic Differentiation in the Great Tit (Parus major) Species Complex	Prof. Toni Gossmann			
11:30 AM	11:45 AM	Sex-specific recombination landscapes are associated with genomic and transcriptomic features in passerine birds	Ms. Lisa-Maria Ammer			
11:45 AM	12:00 PM	Holocentric Chromosome Dynamics: How Structural Rearrangements Shape Recombination and Drive Population Divergence in Carex (Cyperaceae)	Mr. Rogelio Sánchez Villegas			
12:00 PM	12:15 PM	Aneuploidy generated by chromosomal rearrangements contributes to postzygotic isolation in butterfly hybrids	Dr. Jesper Boman			
12:15 PM	12:30 PM	Frequent horizontal transfer of entire accessory chromosomes in fungi	Ms. Hanne Griem-krey			

#### MEETING ROOM 114: 10:30 AM - 12:30 PM - S42.01 - Predicting evolutionary change in ecologically relevant contexts

Session chairs: Dr. Elizabeth Mittell, Dr. Joel Pic	:k
---	----

10:30 AM	11:00 AM	Partitioning the phenotypic and genetic variances of reaction norms	Mr. Pierre de Villemereuil
11:00 AM	11:15 AM	Assessing selection on reproductive phenology in forest and urban great tits: including multiple broods matters	Mr. Jérémy Defrance
11:15 AM	11:30 AM	Effects of urbanization on selection, local adaptation, and eco-evolutionary feedbacks	Ms. Ella Martin
11:30 AM	11:45 AM	Evolutionary rescue and critical genetic variance in stage structured populations.	Mr. Julien Offresson
11:45 AM	12:00 PM	When selection doesn't add up: non-additive evolutionary responses to multiple selective pressures in Drosophila melanogaster	Mr. Vincent Montbel
12:00 PM	12:15 PM	Simulated heatwaves induce rapid and suprising multidimensional evolution in host-parasitoid communities	Prof. Jon Bridle
12:15 PM	12:30 PM	Conflicting temporal dynamics of selection, additive genetic variance and plasticity in early-life seasonal migration versus residence constrain the potential for rapid adaptive evolution	Dr. Rita Fortuna

#### MEETING ROOM 115: 10:30 AM - 12:15 PM - S44.02 - The ecological and evolutionary implications of climate change on reproduction

Session chairs: Prof Amanda Bretman, Dr Liam Dougherty, Dr. Claudia Fricke, Prof. Rhonda Snook					
10:30 AM	11:00 AM	Can evolution and plasticity buffer fertility loss under climate change?	Dr. Belinda van Heerwaarden		
11:00 AM	11:15 AM	Guardians of the germline: small-RNAs and transposable elements impact on Zebrafish fitness in thermal-stressed environments	Dr. Alice Godden		
11:15 AM	11:30 AM	How does adult age affect fertility following heat stress in male Drosophila melanogaster?	Ms. India Sutherland		
11:30 AM	11:45 AM	Recombination Plasticity in Response to Temperature Variation in Reptiles	Mrs. Laura González-Rodelas		
11:45 AM	12:00 PM	Environmental Influences on Sexual Selection and Reproductive Trait Divergence in a Sex-Role Reversed Insect	Dr. Rosalind Murray		
12:00 PM	12:15 PM	Influence of heatwaves on breeding success in the collared flycatcher (Ficedula albicollis)	Ms. Monika Gronowska		



## MEETING ROOM 116: 10:30 AM - 12:30 PM - S22.02 - Evolutionary Ecology of Microbial Symbioses

Session ch	Session chairs: Dr. Anne Duplouy, Prof. Wolfgang Miller					
10:30 AM	11:00 AM	Insect facultative symbionts: a horizontal gene pool for eukaryotes	Dr. Lee Henry			
11:00 AM	11:15 AM	Dissecting the fitness components of nitrogen-fixing rhizobia throughout their symbiotic life cycle	Ms. Margarita Granada Agudelo			
11:15 AM	11:30 AM	A genome-eroded symbiont with a transcriptional response to the host's needs	Ms. Ana Carvalho			
11:30 AM	11:45 AM	The dynamics of coral symbiosis across latitude; opportunities and challenges of coral range expansion in a warming world.	Prof. Shelby Mcilroy			
11:45 AM	12:00 PM	The role of Wolbachia in reproductive isolation and speciation in Drosophila paulistorum	Mrs. Linnéa Ekström			
12:00 PM	12:15 PM	Evolutionary dynamics of nested symbiosis in Aphrodes leafhoppers	Ms. Veronika Andriienko			
12:15 PM	12:30 PM	Pushing the limits of genome reduction: evolution of ancestral planthopper symbiont genomes	Dr. Anna Michalik			

## MEETING ROOM 117: 10:30 AM - 12:30 PM - Stearns Prize Symposium

Chair: M	ax Reuter		
10:30 AM	10:45 AM	Pronounced differentiation on the Z chromosome and parts of the autosomes in crowned sparrows contrasts with mitochondrial paraphyly: implications for speciation	Mr. Quinn McCallum
10:45 AN	1 11:00 AM	Evolution of intraspecific floral variation in a generalist–specialist pollination system	Dr. Marion Leménager
11:00 AN	1 11:15 AM	Correlated genomic patterns of introgression across space despite contrasting hybridization histories.	Dr. Matthew Farnitano
11:15 AN	1 11:30 AM	Genetic architecture of multiple mutualisms and mating system in Turnera ulmifolia	Dr. Jason Laurich
11:30 AM	1 11:45 AM	Reproductive isolation via divergent genital morphology due to cascade reinforcement in Ohomopterus ground beetles	Dr. Tian Xia
11:45 AM	12:00 PM	Why do closely related species that live together differ in colour?: Experimental examinations of the drivers of signal divergence and the forces that influence signal efficacy	Dr. Haley Kenyon
12:00 PM	12:15 PM	Single cells to microbiome perspective of bacterial adaptations	Dr. Subham Mridha
12:15 PM	12:30 PM	Challenging a host–pathogen paradigm: Susceptibility to chytridiomycosis is decoupled from genetic erosion	Dr. Donal Smith

## MEETING ROOM 118+119: 10:30 AM - 12:30 PM - S15.02 - Evolution in and of diverse genetic systems

Session chairs: Dr. Thomas Hitchcock, Dr. Kora Klein, Dr. Martijn Schenkel					
10:30 AM	11:00 AM	Evolution and function of Programmed DNA Elimination in Mesorhabditis nematodes	Dr. Brice Letcher		
11:00 AM	11:15 AM	When sex stops, genes change: ending sexual conflict alters gene expression	Dr. Julie Jaquiéry		
11:15 AM	11:30 AM	Age-dependent sex under mutation accumulation	Mr. Gaurav Athreya		
11:30 AM	11:45 AM	The evolution of non-hybrid asexual reproduction in neotropical night lizards (Lepidophyma spp.)	Mr. Roel Martijn Wouters		
11:45 AM	12:00 PM	Characterizing the germline-restricted chromosome in a hybrid-species complex of sparrows	Dr. Yifan Pei		
12:00 PM	12:15 PM	The conundrum of germline restricted chromosomes in black winged fungus gnats (Diptera)	Dr. Christina Hodson		

## MEETING ROOM 120+121: 10:30 AM -12:15 PM - S37.02 - Museomics: Challenges and Possibilities

Session chair: Prof. Carles Lalueza-fox				
10:30 AM	11:00 AM	Unlocking the temporal multi-omics potential of formalin-fixed museum specimens	Dr. Erin Hahn	
11:00 AM	11:15 AM	Museomic insights into temporal genomic change in the koala	Dr. Binia De Cahsan	
11:15 AM	11:30 AM	Temporal changes of genetic diversity in Finnish butterflies with varying population trends	Prof. Marjo Saastamoinen	
11:30 AM	11:45 AM	Identification of a male Neanderthal from Prado Vargas, Spain, via enamel paleoproteomics	Mr. Guillermo Carrillo-Martin	
11:45 AM	12:00 PM	Museomic Reconstruction of Faunal Genomic History in the Hula Valley Since the Natufian	Ms. Ksenia Juravel	
12:00 PM	12:15 PM	The Museum Genomics of Bumblebees: Collections and Collecting for DNA.	Prof. lan Barnes	



#### MEETING ROOM 122+123: 10:30 AM -12:30 PM - S48.02 - The maintenance of adaptive polymorphisms

Session cha	Session chairs: Dr. Hannah Augustijnen, Dr. Harshavardhan Thyagarajan					
10:30 AM	11:00 AM	How does plant chemodiversity evolve? Testing five hypotheses in one population genetic model	Prof. Meike Wittmann			
11:00 AM	11:15 AM	Are mitochondrial sisters their own worst enemies? Testing for negative frequency-dependent selection via resource competition in Trinidadian guppies.	Dr. Tomos Potter			
11:15 AM	11:30 AM	How do we resolve sexual antagonism? - An integrative approach to understand the genetic, developmental and regulatory basis of a sex-linked color polymorphism	Dr. Nidal Karagic			
11:30 AM	11:45 AM	Larger environmental fluctuations at low frequencies can accelerate population recovery after abrupt environmental shifts by increasing genetic diversity, but this may occur along more vulnerable trajectories	Mr. Youssef Yacine			
11:45 AM	12:00 PM	Characterizing the detectable and invisible fractions of genomic loci under balancing selection	Dr. Débora Brandt			
12:00 PM	12:15 PM	Protandry influences the maintenance of sex-limited polymorphism by antagonistic pleiotropy in the wood tiger moth	Mr. Eetu Selenius			
12:15 PM	12:30 PM	Experimental support for balancing selection maintaining sexually antagonistic genetic variation in a polygenic trait	Dr. Elina Immonen			

## MEETING ROOM 129+130: 10:30 AM -12:30 PM - S34.01 - Mechanisms of adaptation to changing conditions in microorganisms

Session cha	Session chairs: Dr. Javier De La Fuente Hidalgo, Dr. Alvaro San Millan, Dr. Macarena Toll Riera					
10:30 AM	11:00 AM	Eco-evolutionary dynamics within small bacterial communities	Prof. Sara Mitri			
11:00 AM	11:15 AM	Unravelling the evolutionary and ecological impacts of multiple agricultural stressors on soil microbial communities	Dr. Siobhan O'Brien			
11:15 AM	11:30 AM	Do heat extremes and drought predictably alter the function and composition of soil microbial communities?	Mr. Zachary Bailey			
11:30 AM	11:45 AM	Experimental evolution of thermal performance in phage communities	Mr. Samuel Greenrod			
11:45 AM	12:00 PM	Adaptation to multiple simultaneous stresses leads to a more costly and easier to reverse resistance when compared with adaptation to stresses added sequentially	Dr. Ciaran Gilchrist			
12:00 PM	12:15 PM	Adaptation to complex environments reveals pervasive trade-offs and genomic targets with pleiotropic effects	Dr. Dragan Stajic			
12:15 PM	12:30 PM	Phenotypic heterogeneity in capsule production across opportunistic pathogens	Mrs. Julie Le Bris			
			Dr. Olaya Rendueles			

## MEETING ROOM 131: 10:30 AM -12:45 PM - S08 - Cooperation, Conflict and the Evolution of Socially Transferred Materials Count

Session ch	Session chairs: Dr. Joris Koene, Dr. Steven Ramm					
10:30 AM	11:00 AM	Microbes as socially transferred materials: Origin and evolution of a defensive	Dr. Aileen Berasategui			
		symbiosis in tortoise leaf beetles.				
11:00 AM	11:15 AM	Maternally secreted protein ensures symbiont transmission in tortoise beetles	Ms. Marleny García-Lozano			
11:15 AM	11:30 AM	Predictions of sexually antagonistic co-evolution extend to the molecular level of seminal fluid and female reproductive tract proteins.	Dr. R. Axel W. Wiberg			
11:30 AM	12:00 PM	Predictions from signalling theory about the molecular properties of socially transferred materials	Dr. Jen Perry			
12:00 PM	12:15 PM	Larval Altruism and the Development of a Worker Caste - Theoretical Investigations into Primitively Eusocial Wasp Larvae as Altruists.	Mr. Dylan Thatcher			
12:15 PM	12:30 PM	Ejaculate-driven trade-offs: increased fertilization success comes at the cost of lower offspring quality	Dr. Barbara Tschirren			
12:30 PM	12:45 PM	Metabolic Division of Labour as an Evolutionary Ratchet for Social Systems	Prof. Adria Leboeuf			

## EXPO AREA: 12:30 PM - 2:00 AM - Lunch

## MEETING ROOM 113: 2:00 PM - 5:00 PM - S28.03 - Genome Architecture and Their Role in Evolution

Session Chairs: Dr. Marcial Escudero, Prof. Kay Lucek, Dr. Petr Nguyen				
2:00 PM	2:15 PM	Effects of hybridization on genomic content and expression of transposable elements in a hybrid-species complex	n Mr. Alexander Lawrence	
2:15 PM	2:30 PM	DNA on Repeat: Lineage-specific patterns of repetitive DNA accumulation across the genome size spectrum in Ferns, Gymnosperms, and Angiosperms	Mr. Pol Fernández-Mató	
2:30 PM	2:45 PM	Characterization of the transposable element landscape shaping the Ectocarpus genome	Mrs. Erica Dinatale	
2:45 PM	3:00 PM	High centromere diversity is maintained across Arabidopsis genus	Ms. Anna Glushkevich	



3:00 PM	3:15 PM	Understanding the evolutionary history of inversions and their role in speciation and adaptation across the Littorina marine snails genus	Dr. Pierre Barry
3:15 PM	3:30 PM	Structural variation underlies rapid, repeatable adaptation during experimental evolution in a seed beetle	Mr. Brian Kissmer
3:30 PM	3:45 PM	Polymorphic inversions distort selection scans: selection inference in the extremely complex genomic landscape of the spruce bark beetle (Ips typographus)	Ms. Julia Morales-García
3:45 PM	4:00 PM	Signatures of local adaptation within chromosomal inversions	Ms. Karolina Wachala
4:00 PM	4:15 PM	Conserved synteny and lineage-specific genome reshuffling in the largest animal order of Coleoptera	Mr. Arif Maulana
4:15 PM	4:30 PM	Cryptobiosis is associated with rapid chromosome restructuring in Panagrolaimus nematodes	Dr. Nadège Guiglielmoni
4:30 PM	4:45 PM	An episodic burst of massive genomic rearrangements and the origin of non-marine annelids	Dr. Carlos Vargas Chavez
4:45 PM	5:00 PM	Genome scrambling in the fast-evolving chordate Oikopleura dioica questions our understanding of the evolution of genome architecture, challenging the concepts of species and reference genome.	Dr. Cristian Cañestro

## MEETING ROOM 114: 2:00 PM - 5:00 PM - S42.02 - Predicting evolutionary change in ecologically relevant contexts

Session Ch	Session Chairs: Dr. Elizabeth Mittell, Dr. Joel Pick				
2:00 PM	2:30 PM	Adaptation to climate change in wild bird populations: Variation in evolutionary potential and plasticity in blue and great tits populations across Europe	Dr. Céline Teplitsky		
2:30 PM	2:45 PM	The disperser's gambit: playing the long game in common lizard evolution	Ms. Léa Koch		
2:45 PM	3:00 PM	Using the quantitative genetic roadmap to predict responses to selection in wild populations; from theory to empirical examples of environmental shortcuts in wild roe deers, clownfish, and snapdragon plants	Dr. Benoit Pujol		
3:00 PM	3:15 PM	The quest for life-history trade-offs continues: Building on the Y-model of resource acquisition and allocation to reveal the lifespan cost of reproduction	Dr. Erik Postma		
3:15 PM	3:30 PM	Environment-specific additive genetic variances for adult survival reveal dramatic contrasts in adaptive evolution across highly fluctuating conditions	Dr. Paul Acker		
3:30 PM	3:45 PM	Inbreeding depression throughout the growth period of wild Swiss barn owls	Dr. Anna Hewett		
3:45 PM	4:00 PM	Disentangling the demographic drivers of reproductive rate evolution in a wild bird population	Dr. Simon Evans		
4:00 PM	4:15 PM	Founders predict trait evolution and population dynamics after evolutionary rescue	Dr. Deepa Agashe		
4:15 PM	4:30 PM	Repeatability of evolution is similarly and additively affected by variation in selection and variation in drift in a heterogeneously replicated experimental evolution study.	Dr. Thomas Blankers		
4:30 PM	4:45 PM	Some evolutionary demographic consequences of pollen limitation in flowering plants	Dr. Colin Olito		
4:45 PM	5:00 PM	Not just farther, but more variable: Intraspecific variation of seed dispersal strategies in Arabidopsis thaliana as a potential adaptive trait	Mrs. Catharina Utami		

## MEETING ROOM 115: 2:00 PM - 4:45 PM - S51.10 - Sex

Session Ch	nairs: Dr. Ro	salia Piñeiro Portela, Prof. Jacek Radwan	
2:00 PM	2:15 PM	Female lineages and changing mortuary practices in Neolithic Çatalhöyük: Insights from genomic and archaeological data	Dr. Eren Yüncü
2:15 PM	2:30 PM	Genetic & evolutionary mechanisms underlying phenotypic variation in a sexually selected trait	Ms. Claudia Pruvôt
2:30 PM	2:45 PM	A pluralistic approach to sex: a combination of coevolving parasite and increased mutation rate maintain outcrossing	Dr. Michelle Mccauley
2:45 PM	3:00 PM	Can sexually selected traits influence the efficacy genetic rescue? An evolve and resequence approach	Mr. Jonathan Parrett
3:00 PM	3:15 PM	Unveiling the Evolutionary Dynamics of the X Chromosome in Aphids	Dr. Gaorui Gong
3:15 PM	3:30 PM	Adaptive evolution of viviparity resists introgression at genome regions housing genes functional for reproduction	Prof. Kathryn Elmer
3:30 PM	3:45 PM	Understanding the determinants of divergence and reproductive isolation in marine fishes using a comparative speciation genomics approach	Dr. Pierre Barry
3:45 PM	4:00 PM	Half Clone Full Complexity: Unexpected Genomic Diversity and Reproductive Strategies in Hybrid Water Frog Hemiclonal Population Systems Across Ukraine	Dr. Anna Fedorova
4:00 PM	4:15 PM	Unusual sex chromosome structure of B. pusilla and the evolution of sex chromosomes in liverworts	Dr. Yuling Yue
4:15 PM	4:30 PM	Sperm - female reproductive fluid interactions in a fish with a parasitic reproductive strategy	Dr. Alexandra Glavaschi
4:30 PM	4:45 PM	Intergenerational transfer of reproductive plasticity in fruitflies	Prof. Tracey Chapman



MEETING ROOM 116: 2:00 PM - 5:30 PM - S51-11 - Po	pulation Genetics
---	-------------------

Session Ch	Session Chairs: Dr. Jennifer Leonard, Prof. Elena Bosch					
2:00 PM	2:15 PM	Population genomics resolves the complex domestication origin of adzuki bean	Prof. Cheng-ruei Lee			
2:15 PM	2:30 PM	Genome-wide Population Structure of Lake Whitefish (Coregonus clupeaformis) in a	Mr. Philippe Hénault			
		Subarctic Great Lake.				
2:30 PM	2:45 PM	Deleterious mutations cause evolution of lifespan extension by dietary restriction	Ms. Sara Irish			
2:45 PM	3:00 PM	Are long-lived species doomed to slow adaptive evolution?	Dr. Catalina Chaparro Pedraza			
3:00 PM	3:15 PM	How models accounting for population structure can improve our understanding of the	Ms. Camille Steux			
		evolutionary history of common chimpanzees and bonobos				
3:15 PM	3:30 PM	Patterns of Duplicate Gene Retention Over Different Timescales and With Different	Prof. David Liberles			
		Selective Pressures				
3:30 PM	3:45 PM	Mitochondrial introgression affects routine metabolic rates and growth rates in	Dr. Nikhil Modak			
		tadpoles				
3:45 PM	4:00 PM	Polarizing SNPs without outgroup	Mr. Jinyang Liang			
4:00 PM	4:15 PM	Uncovering Introgression Events in Potato Domestication via TE Landscape Analysis	Ms. Zeynep Önder			
4.45.04.4	4 20 214					
4:15 PM	4:30 PM	Genomics, Epigenomics, and Transcriptomic Insights into Hybridization and the	Dr. Clarisse Palma-silva			
		Evolution of Reproductive Barriers in Pitcairnia, a Neotropical Bromeliad of adaptive				
4.20 DN4	4.4E DN4	radiation	De Martin Tanialus			
4:30 PM	4:45 PM	Not the same: genomic analysis explains variation in strength of positive selection	Dr. Martin Tesicky			
4.4E DN4	E-00 DM	across avian immune genes	Dr. Joan Pontista Ladauw			
4:45 PM	5:00 PM	Comparative Population Genomics Illuminates Species Boundaries and Symbiotic Disruption in Eunicella Octocorals	Dr. Jean-Baptiste Ledoux			
E.OO DN4	F.1 F DM	·	Dr. Luiz Augusta Couz dos Contos			
5:00 PM	5:15 PM	Dysploidy and demography: Chromosome number variation shapes population	Dr. Luiz Augusto Cauz dos Santos			
		genomics in Nicotiana section Suaveolentes				

## MEETING ROOM 117: 2:00 PM - 5:00 PM - S51-12 - Phylogenetics

Session Ch	Session Chair: Dr. Borja Milá, Dr. Patricia Álvarez					
2:00 PM	2:15 PM	A comprehensive phylogenomic view of the living xenarthran radiation	Dr. Mathilde Barthe			
2:15 PM	2:30 PM	Towards the end of the lipid divide?	Dr. Damien Devos			
2:30 PM	2:45 PM	Novel genomic approaches support Xenacoelomorpha as sister to all Bilateria	Dr. Jordi Paps			
2:45 PM	3:00 PM	Resolving Difficult Nodes in the Bat Phylogeny Using 103 Chromosome-Level Genomes	Dr. Ariadna Esthela Morales Garcia			
3:00 PM	3:15 PM	Timescale and genetic linkage explain the variable impact of defense systems on horizontal gene transfer	Dr. Jaime Iranzo			
3:15 PM	3:30 PM	Resampling multiple genomic matrices to detect challenging nodes	Dr. Marc Domènech			
3:30 PM	3:45 PM	From Molecular Promiscuity to Neural Specialization: Evolutionary paths to Nervous	Dr. Alexandros Pittis			
		System Complexity				
3:45 PM	4:00 PM	Phylogenetic inference with not-so-rare mutations and wee tiny organisms	Dr. Rui Borges			

## MEETING ROOM 118+119: 2:00 PM - 4:00 PM - S09 - Craniofacial Evolution in Vertebrates

Session Ch	air: Dr. Kévi	n Le Verger, Dr. Olivia Plateau	
2:00 PM	2:30 PM	The Metamorphic Blueprint: How life cycle type shapes salamander skulls	Prof. Anne-Claire Fabre
2:30 PM	2:45 PM	Neural crest biology shapes evolutionary dynamics of the lacertid skull across evolutionary scales	Dr. Quentin HORTA-LACUEVA
2:45 PM	3:00 PM	Reconciling stemness and specialization in neural crest cells drives avian beak diversification	Ms. Carmen Sánchez Moreno
3:00 PM	3:15 PM	Influence of Morphological Integration and Modularity on Evolutionary Rates and Disparity of Bird Beaks	Dr. Ricardo Ely
3:15 PM	3:30 PM	Prenatal Hormones Shape Maternal–Fetal Morphological Integration in a Model of Cephalopelvic Covariation	Ms. Eva Zaffarini
3:30 PM	4:00 PM	Genomic and developmental foundations of adaptive tooth shape evolution	Dr. Alexa Sadier

## MEETING ROOM 120+121: 2:00 PM - 5:00 PM - 530 - Genomic insights into evolutionary adaptation and species movements in a changing climate

Session Chairs: Dr. Ian Bradbury, Dr. Danielle Davenport					
2:00 PM 2:3	0 PM Can eco	ological genomics predict population maladaptation under climate change?	Prof. Matthew Fitzpatrick		
	Lessons	learned from 10 years of working with genomic offsets			
2:30 PM 2:4	5 PM Climate	adaptation and genomic offset in north Atlantic ptarmigan	Mr. Theodore Squires		
2:45 PM 3:0	0 PM The evo	olution of seasonal gene expression in forest trees	Mr. Shuichi Kudo		



3:00 PM	3:15 PM	Cis-regulatory divergence and heritable plasticity underlie seasonal adaptation in a tropical butterfly	Dr. Madeleine Carruthers
3:15 PM	3:30 PM	Implications of expanding hybrid wood ant populations on heat tolerance and phenology under climate change	Dr. Patrick Krapf
3:30 PM	4:00 PM	Evolution of genetic variation associated with adult migration timing in Chinook Salmon & Steelhead	Dr. Shawn Narum
4:00 PM	4:15 PM	Predicting the impacts of future climate change on Arctic Charr populations in the Canadian north using genomic offset analyses	Ms. Samantha Crowley
4:15 PM	4:30 PM	A genomic perspective on population structure and adaptation to aridity in the four- striped mouse Rhabdomys bechuanae in Southern Africa	Dr. Hamilcar Keilani
4:30 PM	4:45 PM	Evolutionarily distinct lineages of a migratory bird of prey show divergent responses to climate change	Dr. Joan Ferrer Obiol
4:45 PM	5:00 PM	Convergent adaptation to latitude in fish	Dr. Maëva Gabrielli
5:00 PM	5:15 PM	Adaptation, and genetic substructure of Atlantic Cod (Gadus morhua) in a changing climate	Dr. Tony Kess

## MEETING ROOM 122+123: 2:00 PM - 4:15 PM - S21 - Evolutionary consequences of heterokaryosis, mosaicisms, chimeras and other monsters

Session Ch	Session Chairs: Dr. Dabao Sun Lü, Dr. Miguel Angel Naranjo-Ortiz					
2:00 PM	2:30 PM	Early germline sequestration in a basidiomycete fungus	Prof. Hanna Johannesson			
2:30 PM	2:45 PM	Fragmentation of mycelia caused loss of alleles and critical symbiotic traits in the	Ms. Asta Rødsgaard-jørgensen			
		domesticated fungus farmed by leafcutter ants				
2:45 PM	3:00 PM	Nuclear cooperation as a driver of evolutionary transitions in fungi	Dr. Mattias Siljestam			
3:00 PM	3:30 PM	Choosing Cooperation: Invasive Multicellularity in Fonticula alba and Beyond	Prof. Marko Kaksonen			
3:45 PM	4:00 PM	Premeiotic endoreplication as a compensatory mechanism in hybrid amniotes: A	Prof. Lukáš Kratochvíl			
		cellular pathway to asexuality, polyploidisation and genome stabilisation				
4:00 PM	4:15 PM	Co-ordinated regulation of gene expression inside of dikaryotic Basidiomycetes	Dr. Ben Auxier			

#### MEETING ROOM 129+130: 2:00 PM - 3:30 PM - S34.02 - Mechanisms of adaptation to changing conditions in microorganisms

Session ch	Session chairs: Dr. Javier De La Fuente Hidalgo, Dr. Alvaro San Millan, Dr. Macarena Toll Riera					
2:00 PM	2:30 PM	Prof. Itzhak Mizrahi's presentation	Prof. Itzhak Mizrahi			
2:30 PM	2:45 PM	Heteroresistance as an adaptive strategy of Pseudomonas aeruginosa in fluctuating antibiotic environments	Ms. Nan Ye			
2:45 PM	3:00 PM	Genetic local adaptation of gut microbial species in metabolically and structurally distinct intestinal compartments	Mr. Russ Jasper			
3:00 PM	3:15 PM	Assessing the impact of resource availability on plasmid costs and persistence in microbial communities	Ms. Enora Marrec			
3:15 PM	3:30 PM	A competition-detoxification balance controls the evolution of resistance in simple microbial communities.	Dr. Massimo Amicone			

#### MEETING ROOM 131: 2:00 PM - 3:30 PM - S32 - Letting go: reductive evolution across the tree of life

Session chairs: Dr. Alexander Bowles, Dr James Clark						
2:00 PM	2:30 PM	The role of reductive evolution in animal terrestrialisation	Dr. Rosa Fernandez			
2:30 PM	3:00 PM	Terrestrial triumph, genetic trim: Bryophyte genome reduction and the need to serve mitochondria and plastids	Prof. Sven Gould			
3:00 PM	3:15 PM	Unraveling the reductive evolution of bdelloid rotifers, while having adapted to extreme stresses.	Prof. Karine Van Doninck			
3:15 PM	3:30 PM	Genomic signature of CAM photosynthesis loss	Dr. Pauline Raimondeau			

#### MEETING ROOM 129+130: 3:30 PM - 4:30 PM - S50 - Unraveling the origin of eukaryotes: integrating prokaryotic and eukaryotic perspectives

Session chairs: Dr. Sam von der Dunk, Dr. Julian Vosseberg					
3:30 PM	4:00 PM	The symbiotic origin of the eukaryotic cell	Dr. Puri Lopez-Garcia		
4:00 PM	4:15 PM	Tracing the Prokaryotic Origins of Membrane Trafficking in Eukaryotes	Dr. Caroline Puente-Lelievre		
4:15 PM	4:30 PM	Modeling endosymbiotic gene flow at the origin of eukaryotes: a highways approach	Dr. Tara Mahendrarajah		

MEETING ROOM 131: 3:30 PM - 5:00 PM - S39 - Novel experimental and computational approaches to understand the prevalence of reticulate evolution in eukaryotes

#### Session chairs: Dr. Carla Gonçalves, Dr. Michelle Leger, Dr. Eduard Ocana-Pallares

3:30 PM	4:00 PM	Genes encoding transporter proteins are subject to horizontal gene transfer and can alter the ecology of recipient lineages	Prof. Thomas Richards
		5	
4:00 PM	4:30 PM	A natural mechanism of eukaryotic horizontal gene transfer	Dr. Andrew Urquhart
4:30 PM	4:45 PM	$\label{lem:condition} \textbf{Reticulate evolution in a plant radiation: the role of introgression in shaping ecological}$	Dr. Clara Groot Crego
		diversity in the subgenus Tillandsia	
4:45 PM	5:00 PM	Genome polarisation for detecting and characterising admixture	Dr. Stuart Baird

EXPO AREA: 5:00 PM - 7:00 PM - POSTER SESSION 3 - with Aperitif

Full list of posters can be found at the end of the program document  $% \left( 1\right) =\left( 1\right) \left( 1\right$ 



## Friday, 22 August 2025

## MEETING ROOM 113-117: 8:45 AM - 9:00 AM: Friday Access to Plenary Session & Announcements

#### PLENARY SESSION (MEETING ROOM 113-117): 9:00 AM - 9:45 AM - Plenary Talk – Keynote Speaker JOHANNA MAPPES

9:00 AM 9:45 AM Signals of Survival: The Genetics and Ecology of Looking Dangerous Prof. Johanna Mappes

EXPO AREA: 9:45 AM - 10:30 AM - Coffee Break

## MEETING ROOM 118+119: 10:30 AM - 12:00 PM - S25.01 - Gene Content Across Genomes: Models and Genomic Data

Session Chairs: Dr. Raquel Assis, Prof. David Liberles						
10:30 AM	11:00 AM	From non-coding to coding: the de novo emerging proteome	Dr. Aaron Wacholder			
11:00 AM	11:15 AM	The march to land: unearthing how gene repertoire evolution triggered	Ms. Gemma Martínez Redondo			
		terrestrialization across the Animal Tree of Life				
11:15 AM	11:30 AM	Boom and Bust in the Parasite Genome: The adaptive benefits of massive gene family	Prof. Erich Bornberg-bauer			
		losses in slave-maker ants				
11:30 AM	11:45 AM	Understanding Evolutionary Changes in the Genome through investigation of	Dr. Marta Farré			
		Evolutionary Breakpoint Regions and Homologous Synteny Blocks				
11:45 AM	12:00 PM	Natural protein structures have evolved exceptional robustness to mutations	Dr. Sam von der Dunk			

## MEETING ROOM 120+121: 10:30 AM - 12:30 PM - S12.01 - Epigenetics and adaptation to global change: climate and biotic interactions

Session Chairs: Prof Sofia Consuegra, Dr Carlos Guerrero Bosagna					
10:30 AM	11:00 AM	Genomic and epigenomic variation contribute to local adaptation to temperature in the	e Dr. Dafni Anastasiadi		
		sea			
11:00 AM	11:15 AM	Maternal RNA variability as an adaptive feature of environmental responsiveness and heritable diversification	Prof. Irene Adrian-Kalchhauser		
11:15 AM	11:30 AM	DNA methylation: a way for fast adaptation in clonal species? An example with the plant species Fragaria vesca and adaptation to altitude.	Dr. Audrey Le Veve		
11:30 AM	12:00 PM	Taming the Genome: Epigenetic Contributions to Domestication and Sex-Specific Regulation in Chickens	Dr Carlos Guerrero Bosagna		
12:00 PM	12:15 PM	Epigenetic differences from floater to territory holding male Namibian cheetahs (Acinonyx jubatus)	Dr. Alexandra Weyrich		
12:15 PM	12:30 PM	Epimutations in Neurospora crassa	Prof. Ilkka Kronholm		

#### MEETING ROOM 122+123: 10:30 AM - 12:30 PM - S06.01 - Cancer in an evolutionary framework: across species and within individuals

Session Cha	Session Chairs: Dr. Andriy Marusyk, Prof Aurora Nedelcu, Prof. Beata Ujvari					
10:30 AM	11:00 AM	Comparative Oncology: Discovering Natural Cancer Defense Mechanisms Across	Prof. Lisa Abegglen			
		Species to Inspire Future Therapies				
11:00 AM	11:15 AM	Selection for function in cancer: a novel evolutionary principle with potentially	Prof. Frederic Thomas			
		groundbreaking therapeutic implications				
11:15 AM	11:30 AM	Selective sweep probabilities in tumours and other range expansions	Dr. Robert Noble			
11:30 AM	11:45 AM	Sublineage dynamics and mitochondrial DNA exchange in Tasmanian devils'	Ms. Sophia Belkhir			
		transmissible cancers				
11:45 AM	12:00 PM	Peristromal niches provide ecological rescue and facilitate the evolution of resistance	Dr. Andriy Marusyk			
		to targeted therapies in lung cancer				

## MEETING ROOM 129+130: 10:30 AM - 12:30 PM - S24 - Forecasting evolution in natural populations

Session Chairs: Dr. Simon Evans, Dr Erik Postma				
10:30 AM	11:00 AM	Two explanations for variation in the predictability of evolution	Mr. Patrik Nosil	
11:00 AM	11:15 AM	Quantitative prediction of population mean fitness evolution: intra- and inter-sexual genetic (co)variances measure the expected amount of adaptive change to survival and reproduction in the face of sexual conflict and competition	Dr. Matthew Wolak	
11:15 AM	11:30 AM	Divergence time and environmental similarity shape the repeatability of morphological evolution in stick and leaf insects.	Dr. Romain Boisseau	
11:30 AM	12:00 PM	Dr. Colin Garroway's presentation	Dr. Colin Garroway	
12:00 PM	12:15 PM	The use of multi-response models to improve inferences about natural selection	Ms. Sarah Dobson	
12:15 PM	12:30 PM	Jointly forecasting adaptive evolution and demography: refining tools and semantics.	Dr. Timothée Bonnet	



#### MEETING ROOM 131: 10:30 AM - 12:30 PM - S40. - Phylogenomics methodology and the deep tree of life

Session Chairs: Dr. Laura Eme, Dr. Daniel Tamarit					
10:30 AM	11:00 AM	Clarifying the origin of eukaryotic cells using new deep-time phylogenetic models.	Prof. Andrew Roger		
11:00 AM	11:15 AM	An independent phylogenomic dataset sheds new light on the eukaryotic Tree of Life	Mr. Romain B. Leroy		
11:15 AM	11:30 AM	Large-scale phylogenomics nests ten additional phyla-rank lineages in the PVC superphylum	Ms. Kassiani Panagiotou		
11:30 AM	12:00 PM	Deep roots and evolution of cellular life on Earth	Prof. Anja Spang		
12:00 PM	12:15 PM	Solarion arienae is rare microbial relict from newly described eukaryotic supergroup	Mr. Marek Valt		
12:15 PM	12:30 PM	Deciphering the origin and timing of major waves of gene acquisition unravel complex interactions underlying the origin of eukaryotes	Ms. Saioa Manzano-Morales		

## PLENARY SESSION (MEETING ROOM 113-117): 10:30 AM - 12:30 PM - JMS Prize and EUEA Award Winners Symposium

Session Chair: Dr. Josefa Gonzalez, Anne Charmantier, Noah Hensen					
10:30 AM 11:00 AM From chromosomal inversions to sex chromosomes: how deleterious mutations shape Dr. Paul	ıl Jay				
recombination landscapes					
11:00 AM 11:20 AM The path to freedom: Surviving as a trans academic present	ted by Noah Hensen				
11:20 AM 11:50 AM Evolutionary interplay between male-killing symbionts and insects Dr. Hiro	oshi Arai				
11:50 AM 12:10 PM Palestinian researchers talk May Sh	iehady				

## EXPO AREA: 12:30 PM - 2:00 AM - Lunch

#### MEETING ROOM 129+130: 1:30 PM - 3:00 PM - S45 - The evolution of microbial pangenomes

Session Chairs: Dr. Franz Baumdicker, Dr. Jaime Iranzo, Dr. Anne Kupczok				
1:30 PM	2:00 PM	Why do bacterial pangenomes vary across species?	Dr. Anna Dewar	
2:00 PM	2:30 PM	Title not Send	Dr. Jesse Shapiro	
2:30 PM	2:45 PM	Solving the pangenome paradox: detecting accessory genes under balancing selection	Dr. Cara Conradsen	
		in Klebsiella pneumoniae		
2:45 PM	3:00 PM	Emergent epistasis mediates the role of negative frequency-dependent selection in	Prof. Sonja Lehtinen	
		bacterial strain structure		

#### MEETING ROOM 118+119: 2:00 PM - 3:00 PM - S25.02 - Gene Content Across Genomes: Models and Genomic Data

Session Chairs: Dr. Raquel Assis, Prof. David Liberles				
2:00 PM	2:30 PM	Exploring Biodiversity Genomics to Reveal Soil Invertebrates' Role in the Carbon Cycle	Prof. Ingo Ebersberger	
2:30 PM	2:45 PM	Genome Size Reduction Under Non-Adaptive Evolution in an Island Radiation	Dr. Sara Guirao-Rico	
2:45 PM	3:00 PM	Evolution of genome size in self-fertilising populations	Dr. Diala Abu Awad	

## MEETING ROOM 120+121: 2:00 PM - 3:00 PM - S12.02 - Epigenetics and adaptation to global change: climate and biotic interactions

Session Chairs: Prof Sofia Consuegra, Dr Carlos Guerrero Bosagna					
2:00 PM	2:15 PM	Transgenerational transmission of epigenetic responses as a second driver of adaptive evolution	Prof. Pim Edelaar		
2:15 PM	2:30 PM	Epigenetic potential and range expansion in the house sparrow	Prof. Lynn Martin		
2:30 PM	2:45 PM	A Hardy–Weinberg Equilibrium–Based Neutral Model for Plant DNA Cytosine Methylation Variation	Dr. Arunkumar Ramesh		
2:45 PM	3:00 PM	The role of epigenetics in local adaptation in sessile oak trees: a population epigenetic association study $ \frac{1}{2} \left( \frac{1}{2} \right) = \frac{1}{2} \left( \frac{1}{2} \right) \left( \frac{1}{2} \right)$	Dr. Shannon Brandt		

## MEETING ROOM 122+123: 2:00 PM - 3:00 PM - S06.02 - Cancer in an evolutionary framework: across species and within individuals

Session Chairs: Dr. Andriy Marusyk, Prof Aurora Nedelcu, Prof. Beata Ujvari					
2:00 PM	2:15 PM	An unexplored connection between mutation, phenotypic plasticity and evolutionary	Prof Aurora Nedelcu		
		convergence in cancer			
2:15 PM	2:30 PM	Estimating excess cancer risk: an evolutionary approach	Prof. Leonard Nunney		
2:30 PM	2:45 PM	Body size and sex differences affect cancer across species in lemurs	Dr. E. Yagmur Erten		
2:45 PM	3:00 PM	The permissive binding theory of cancer: an evolutionary theory of metastasis	Dr. Caroline Weisman		

#### PLENARY SESSION (MEETING ROOM 113-117): 2 PM - 3 PM - ESEB MEMBERSHIP MEETING



PLENARY TALK (MEETING ROOM 113-117): 3 PM - 4 PM - Presidential Address by MIKE RITCHIE

Chair: Tracey Chapman Speaker: MIKE RITCHIE

PLENARY TALK (MEETING ROOM 113-117): 4 PM - 4:30 PM - CLOSING CEREMONY



## **POSTER SESSIONS**

## Monday 18 August, EXPO AREA: 5:00 PM - 7:00 PM - POSTER SESSION 1 - with Aperitif

S01 - Adaptation to environmental changes in trees through the lens of common gardens and genomics					
P01.001	Multi-Omics Insights into Seasonal Adaptation and Resilience in Prosopis cineraria	Dr. Bhumika Dubay			
P01.002	Evolutionary genomics unravels the responses and adaptation to climate change in a key alpine forest tree species	Ms. Zhiqin Long			
P01.003	Towards an integrative understanding of taxonomic diversity, climate adaptation, and introgression in Southeastern European white oak species	Dr. Clara Groot Crego			
P01.004	Estimating the demographic history of fir species (Abies ssp.) and predicting their spatial patterns of genetic diversity	Ms. Azzurra Pistone			
P01.005	Local adaptation and Genomic vulnerability to climate change in oak species	Prof. Baosheng Wang			
P01.006	FruitRescue: Large-Scale Fitness Phenotyping of Fruit Trees for Climate Resilience across multiple environments in Europe.	Mr. Mathieu Brisson			
P01.007	Islands as model systems for evolutionary divergence: a case study of Hypericum lanceolatum on Réunion Island	Ms. Romane Callarec			
P01.008	Genomic Signatures of Drought Adaptation in Eucalyptus: A Phylo-transcriptomic Approach Across 50 Million Years of Divergence	Dr. Luke Yates			
P01.009	Genotype–environment interactions underlying resistance to Hymenoscyphus fraxineus in European ash	Dr. Joanna Meger			
P01.010	Genomic assessment of existing resources and climate maladaptation risk in Scots pine: implications for forest management	Prof. Witold Wachowiak			
P01.011	Divergent early life-history strategies and climate sensitivity in Fir (Abies spp.) and Beech (Fagus spp.) provenances across Europe	Dr. Leo Zeitler			
P01.012	Landscape Genomic Tool built on genotype-environment associations of truly wild olive trees sampled along a 13-degree north-south in the Western Mediterranean to Assess the Adaptive Value of Cultivars	Dr. Bouchaib Khadari			

#### S03 - Advances in technology, mathematical and statistical models and their application in evolutionary ecology: the dawn of a new era

P01.014	Detection of eco-evolutionary dynamics in communities using Joint Species Distribution Models	Dr. Ruben Hermann
P01.016	How Green Is the Grass? Using Remote Sensing to Track Forage for Red Deer	Mr. Shane Butt
P01.017	GHIST: the Genomic History Inference Strategies Tournament	Prof. Ryan Gutenkunst
P01.018	The Power of Three: Enhancing Arthropod–Plant Interaction Monitoring with Citizen	Mr. Ivan Perez
	Science, Expert Taxonomic Criteria, and Computer Vision Models	
P01.019	High levels of mitotic gene conversion are needed to effectively purge deleterious	Dr. Matthew Hartfield
	mutations in asexual organisms	
P01.021	Estimating bacterial mutation rates with simulation-based methods (ABC)	Mr. Aurélien Tauzin
P01.022	EvAM-Tools: a tool for Evolutionary Accumulation Models	Prof. Ramón Diaz-Uriarte
P01.023	A comprehensive representation of selection at loci with multiple alleles that allows	Prof. Toni Gossmann
	complex forms of genotypic fitness	
P01.025	Predator decision shape dynamics and stability of mimicry systems.	Mr. Yi Sun
P01.026	Species trait data imputation via deep learning	Mr. Matthew Turner
P01.027	Crossovers and Consequences: Genomic Prediction Reveals Selection on Recombination	Mr. Kenneth Aase
	Rates in a Natural Population	
P01.028	Evidence that human skill level is more reactive to developmental history than to genes or	Dr. Mauricio González-Forero
	environment	
P01.029	The avian colourscape and its uncertain future	Mr. Robert MacDonald
P01.030	Semi-supervised selection target detection with positive-unlabeled learning	Mr. Sandipan Paul Arnab
P01.032	A Simulation Study Utilising REvoSim to Investigate the Impact of Predation on Species	Mr. Cameron Peacock
	Diversity Across Environmental Heterogeneity	
P01.033	Quantifying camouflage evolution: digital phenotyping and macroevolutionary modelling in	Dr. Takao K. Suzuki
	Neotropical butterflies	
P01.034	An interpretable machine learning-based alternative to genome-wide association studies	Mr. Gard W. Gravdal
	(GWAS), and its application in a wild population	
P01.035	Hues of blues: Using Comparative Genomics and Machine Learning to Uncover the	Dr. Andreas Wallberg
DO4 006	Evolutionary Ecology of Color Vision in Krill	
P01.036	Investigating the Maintenance of Alternative Reproductive Tactics in Tetranychus urticae	Mr. Koen Freerks
	males	



P01.037	The Inference Challenge: Regularization Methods Across Computational Paradigms in Evolutionary Ecology	Dr. Luke Yates
P01.038	Quantitative approaches reveal that viral cheats are highly abundant and form species- specific communities in natural infections of H5N1 avian influenza.	Dr. Asher Leeks
P01.039	SAI: A tool for Statistics for Adaptive Introgression	Prof. Martin Kuhlwilm
P01.040	Inferring the history of gene copy number evolution	Dr. Thomas Wiehe
P01.041	Quantifying the effects of antibiotic resistance and within-host competition on strain fitness in Streptococcus pneumoniae	Mr. Aswin Krishna
P01.042	The Role of Robustness in Shaping Metabolic Network Evolution Across Species	Ms. June Monge Lorenzo
P01.044	Inference of dispersal and density parameters from spatial genomics data, using simulation-based machine learning methods	
P01.045	Division of Labour in the Evolution of Group Exclusion	Mr. Wayne Liang
P01.046	Comparison of classical and Al-based computational tools for predicting deleterious mutations in plants	Ms. Eva Pardo Otero
P01.047	When barriers break: the probabilistic assembly of pathogen's host-range and the percolation of epidemics	Dr. Pedro Bolanho Mendes
P01.048	The friendship paradox and social structuring in animal societies	Ms. Eloise Newman
P01.050		
	How does ecological stability affect evolution?	Dr. Jessica King
P01.051	Modelling the evolution of division of labour in clonal groups	George Shillcock
S05 - Alien	s among us: ecological drivers, evolutionary dynamics, and rapid ecosystem reshaping by biological	ogical Invasions
P01.052	Introgression in the Anthropocene: detecting gene flow between native and alien flowering plants in the British flora	
P01.053	Evolution of the genetic load in invasive insects: insights from a genomic cross-species study	Mr. Eric Lombaert
P01.054	Hybridization at the front wave of expansion increases genomic variability within an invasive species of slug, overcoming genetic drift	Mr. Lucas de Freitas Lacerda
P01.055	Coexistence and Succession in Disequilibrium: Insights from Nematode Communities on Decaying Beetles	Dr. Ata Kalirad
P01.056	Maternal care thwarts parasitoids in the invasive brown widow spider (Latrodectus geometricus)	Dr. Monica Mowery
P01.057	Reconstructing the Origins of Contrasting Colonization Events in Littorina saxatilis	Mrs. Beatrice Sammarco
P01.058	Investigating morphological and behavioral spatial sorting in a predatory snake across a rapidly expanding invasion gradient	Mr. Marc Vez-Garzón
P01.059	The Genomic Basis of Success and Failure in Marine Snail Range Expansions	Dr. Roberto Biello
P01.060	Transgenerational Effects of Sublethal Insecticides in the Colorado Potato Beetles	Dr. Aigi Margus
P01.061	Impact of dispersal and spatial processes on niche expansion during invasions	Ms. Caitlin Miller
P01.062	Contrasting genomic signatures of hybridisation across admixed populations of native and invasive Mytilus mussels in Australia	Ms. Samantha Howitt
P01.063	Bottlenecked but booming: paradoxically strong inbreeding in highly invasive freshwater mussels	Dr. Antti Miettinen
P01.064	Transcriptomic Evidence for Enhanced Gut Function in Non-Native House Sparrows	Mrs. Kailey McCain
P01.065	Model-based demographic inference of recent invasions from genomic data	Mr. Francisco Campuzano Jiménez
P01.066	Phylogenomics and Trait Evolution in Kalanchoe (Crassulaceae): Insights from Whole- Genome Sequencing and Hybridization Analyses	Mr. Ronen Shtein
P01.067	Urbanization and the Evolution of Stoichiometric Flexibility in Trinidadian Guppies	Mr. Jeferson Amaral
SO7 - Contr	ibution of the microbiome to host adaptation and plasticity	
		Dr. Pachaol Krama
P01.069	Sex-Specific Evolution of Host-Parasite Interactions: The Interplay Between Host Genetics	Dr. Rachael Kramp
	and Microbiome in the Immune Response in Trinidadian Guppies	
P01.070	Gut Microbial Dynamics in Atlantic Salmon: A Natural Field-Based Common Garden Experiment on the Role of Host, Environment, and Stochastic Processes	Dr. Chloe Heys
P01.071	Shaping Metabolism Through the Microbiome: Genetic Constraints on Host Plasticity in a Model of Obesity	Ms. Noelle Curtis-Joseph
P01.072	Metagenomic insights into the ageing gut microbiome in a natural population	Mr. Chuen Zhang Lee
P01.073	Ancient Host-Associated Microbes obtained from Mammoth Remains	Dr. Benjamin Guinet



P01	.075	Linking phenotypic responses and advanced metagenomic techniques in Daphnia magna in response to combined environmental stresses	Ms. Kristina Yefimak
P01	.076	Heat-stressed Aspergillus terreus strain as a potential candidate for improving crop performance	Dr. Khaled Hazzouri
P01	.077	Different drivers govern the fungi and bacteria components in the dung beetle microbiota	Dr. Angela Roggero
P01	.078	Comparative microbiome analyses reveal differences between wild and captive populations of the Martsony Break Newt (Calatriton arreldi)	Mr. Sergi Tulloch Jiménez
P01	.079	of the Montseny Brook Newt (Calotriton arnoldi)  Fast-paced evolution of commensal bacteria facilitates insect host development under	Dr. Youn Henry
DO1	.080	chemical stress Diet-microbiome associations in a wild Swedish carnivore	Ms. Charlotte Enkvist
	.080	Ethanol preference as adaptive plasticity? Behavioral and microbiotic shifts in Nosema-	Mrs. Monika Ostap-Chec
		infected honeybees	·
	.083	The wild mammal oral microbiome is shaped by host diet and ecology	Ms. Markella Moraitou
P01.		Honey as a window into the honey bee microbiome: shedding light into diversity and host–environment interactions	Mrs. Julie Birgel
P01.	.087	Exploring the Effects of Habitat Modification on Rodent Gut Microbial Communities in the Yucatán Peninsula	Ms. Gabriela Borja-Martínez
P01	.088	Testing for genetic differences in fitness across an environmental gradient using germ-free hosts to incorporate "local" microbiomes	Ms. Gwendoline Acerbi
P01	.089	The gut microbiota of hybrid birds: a possible role in speciation?	Dr. Ester Martínez Renau
P01	.090	Facultative symbionts and climatic adaptation: testing thermal benefits of Hamiltonella defensa in aphids across latitudes	Mr. Dominic Stalder
P01	.091	Rapid changes in the gut microbiome revealed by repeated short-term sampling in a captive guppy population	Dr. Magdalena Herdegen-radwan
P01	.092	Integrative multi-omic analysis reveals oral microbiome-metabolome signatures of obesity	Prof. Aashish Jha
P01	.093	Leveraging metagenomics to characterize the gut microbiome of caribou (Rangifer tarandus) for conservation.	Ms. Charlotte Bourbon
P01	.094	Alternative photobiont - where did it come from? Metabarcoding exploration of algal	Dr. Magdalena Kosecka
P01	.095	diversity living on lichens in the phyllosphere framework.  Human oral microbiome and its impact on dental caries across the last 7000 years in	Ms. Şevval Aktürk
P01	.096	Anatolia Exploring the role of Zostera marina's microbiome composition under heat stress across life	Ms. Christina Bakowski
P01	.097	stages and populations Sustainable antimicrobial management in the burying beetle external microbiome	Dr. Sarah Duxbury
<b>D</b> ∩1	.098	What maintains the diversity of the gut microbiome in termite populations?	Dr. Franck Dedeine
P01		Field realistic stressors alter the bee gut microbiome	Dr. Julia Jones
P01		Reproductive Success Dependencies of Gut Microbiome: Insights from a Wild Population of	
		Blue Tits	ivis. Iviai tyria eeriarowska
P01.	.101	The Impacts of Feralization on the Horse Gut Microbiome	Ms. Madeleine Van Well Bergström
P01	.102	Egg and embryo microbiomes in the house sparrow (Passer domesticus): acquisition and consequences for embryo success	Ms. Sophia Wolfe
P01	.103	Genotype–environment interactions determine microbiota symbiosis in Syllidae polychaetes	Dr. Patricia Álvarez-Campos
P01	.104	Diet-microbiome covariation across three giraffe species in a close-contact zone	Dr. Elin Videvall
P01	.106	Impact of the presence of Varroa Mite on the Microbiota of Apis mellifera and Bombus terrestris	Dr. Fanni Borvető
D∩1	.107	Microbiome diversity in a successful urban bird	Mr. Maciej Kamiński
P01		Development and Stability of the Equine Gut Microbiome: Microbial Succession in Foals	Ms. Sofie Geck Sevatdal
. 01.	. 100	and Seasonal Variation in Adult Horses at the Spanish Riding School	Some Geek Sevatual
P01	.109	Symbiotic communities of Planococcus ficus (vine mealybug) from two vineyards varieties	Dr. Diego Santos-Garcia
P01	.110	Intestinal tissue remodeling as a key adaptation to starvation in cave-dwelling fish	Dr. Ana Santacruz
P01	.111	The role of the chicken genome in shaping gut microbiota	Mr. Matias Becker Burgos
	.112	Early life assembly of the gut microbiome and its consequences in wild Soay sheep	Dr. Amy Sweeny
P.C -	442	Comment and the Comment of the Comme	Mar Flore Cabilla Calif
P01.	.113	Can my gut save me? Heatwaves, cloacal microbiome and survival of lesser kestrel nestlings (Falco naumanni)	Ms. Elena Catelan-Carphio



P01.114	A New Strepsipteran Parasite Xenos gadagkari sp.nov and its effect on Microbial	Mr. Deepak Nain
P01.115	Communities of Primitively Eusocial wasp Polistes wattii Characterization of Microbial Communities in Winter Ticks & Moose in Maine	Ms. Monica Miles
S10 - Eco-e	evolutionary dynamics driven by mobile genetic elements	
P01.118	Diverse patterns of transposable elements expressions in Macaca mulatta and the regulation of gene expression in adjacent genes by tissue-specific TEs	Dr. Dae-Soo Kim
P01.119	Role of AluYRa1 insertion in Generating a New Isoform of the BHMT Gene in Old World	Ms. Yun-Jung Lee
P01.121	Monkeys  How TEs dynamics follow lifestyle transitions in gall wasps (Hymenoptera: Cynipoidea)	Ms. Jieying Feng
P01.123	Purifying Selection Shapes the Dynamics of P-element Invasion in Drosophila simulans Populations	Dr. Anna Maria Langmüller
P01.124	The latent cis-regulatory potential of mobile DNA in Escherichia coli	Dr. Timothy Fuqua
P01.125	Sequence composition and conservation predict the phenotypic relevance of transposable elements	Mr. Yari Cerruti
P01.126	Mechanisms of infection in freshwater bacteria–phage systems	Dr. Vesna Grujcic
P01.127	Unveiling Hidden Phage Diversity: Addressing Assembly and Identification Biases in	Dr. Vinicius Silva Kavagutti
P01.128	Metagenomic Analyses of Mutation Hotspots  Transposable element mobilization in the soma of hypomethylated Arabidopsis thaliana	Mr. Andrea Movilli
P01.129	Disentangling the complex network of phage-host interactions at genome resolution in freshwater	Ms. Lauren Davies
P01.130	Towards building a global atlas of microbial Mobile Genetic Elements	Dr. Supriya Khedkar
P01.131	Plasmid recombination: a key strategy for understanding the dynamics of plasmids and	Ms. Emma Acacia
P01.133	antibiotic resistance in bacteria Stealth plasmids: rapid evolution of deleted plasmids can displace antibiotic resistance	Dr. Tatiana Dimitriu
101.133	plasmids under selection for horizontal transmission	Dr. Tatiana Dimitria
P01.134	Repeatome analysis in an Amazonian riparian forest ground-herb reveal potential role of LTR-Retrotransposons on waterlogging adaptation mechanisms	Dr. Drielli Canal
P01.136	The co-evolutionary dynamics of CRISPR-Cas and plasmids	Dr. Berit Siedentop
	The co-evolutionary dynamics of CRISPR-Cas and plasmids attenuation after whole genome duplication: transformative advances and new directions	Dr. Berit Siedentop
		Dr. Berit Siedentop  Ms. Brenda Irene Medina Jiménez
S13 - Evolu	ution after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in	·
S13 - Evolu P01.138	ution after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara
S13 - Evolu P01.138 P01.139 P01.141	stion after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara
S13 - Evolu P01.138 P01.139 P01.141	ation after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms  How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova
S13 - Evolu P01.138 P01.139 P01.141	ation after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms  How does the common barbel (Barbus barbus) see? The effect of the whole-genome	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova
S13 - Evolu P01.138 P01.139 P01.141	Ition after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms  How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision  Dynamic Transposon Activation and Gene Regulatory Rewiring Drive Functional Divergence	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova
S13 - Evolu P01.138 P01.139 P01.141 P01.143 P01.144	Intion after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms  How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision  Dynamic Transposon Activation and Gene Regulatory Rewiring Drive Functional Divergence of Duplicated Genes During the Rediploidization Process in Orychophragmus violaceus  Polyploidy in Action: Disentangling the Consequences of Genome Doubling in Arabidopsis arenosa  Challenging reproductive barriers between and within species: the role of ploidy in the	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova Mr. Changfu Jia
S13 - Evolu P01.138 P01.139 P01.141 P01.143 P01.144	Intion after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms  How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision  Dynamic Transposon Activation and Gene Regulatory Rewiring Drive Functional Divergence of Duplicated Genes During the Rediploidization Process in Orychophragmus violaceus  Polyploidy in Action: Disentangling the Consequences of Genome Doubling in Arabidopsis arenosa  Challenging reproductive barriers between and within species: the role of ploidy in the genesis of new species  The phased genome assembly of Xenopus mellotropicalis sheds light on the pace of	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova Mr. Changfu Jia Ms. Eliška Petříková
S13 - Evolu P01.138 P01.139 P01.141 P01.143 P01.144 P01.145 P01.146	Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision Dynamic Transposon Activation and Gene Regulatory Rewiring Drive Functional Divergence of Duplicated Genes During the Rediploidization Process in Orychophragmus violaceus Polyploidy in Action: Disentangling the Consequences of Genome Doubling in Arabidopsis arenosa Challenging reproductive barriers between and within species: the role of ploidy in the genesis of new species The phased genome assembly of Xenopus mellotropicalis sheds light on the pace of subgenome dominance and diploidization in animal polyploids Sex, behavior, and giant cells: missing elements of animal polyploid evolution with the	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova Mr. Changfu Jia Ms. Eliška Petříková Ms. Diana Frazão
S13 - Evolution P01.138  P01.139  P01.141  P01.143  P01.144  P01.145  P01.146  P01.148	Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision Dynamic Transposon Activation and Gene Regulatory Rewiring Drive Functional Divergence of Duplicated Genes During the Rediploidization Process in Orychophragmus violaceus  Polyploidy in Action: Disentangling the Consequences of Genome Doubling in Arabidopsis arenosa Challenging reproductive barriers between and within species: the role of ploidy in the genesis of new species The phased genome assembly of Xenopus mellotropicalis sheds light on the pace of subgenome dominance and diploidization in animal polyploids Sex, behavior, and giant cells: missing elements of animal polyploid evolution with the parasitoid Nasonia vitripennis Chromosomes, the more the merrier? How does tetraploid C. elegans resist severe cold	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova Mr. Changfu Jia Ms. Eliška Petříková Ms. Diana Frazão Mr. Zhen Ll
S13 - Evolution Pol. 138 Pol. 139 Pol. 141 Pol. 143 Pol. 144 Pol. 145 Pol. 146 Pol. 148 Pol. 149	stion after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms  How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision  Dynamic Transposon Activation and Gene Regulatory Rewiring Drive Functional Divergence of Duplicated Genes During the Rediploidization Process in Orychophragmus violaceus  Polyploidy in Action: Disentangling the Consequences of Genome Doubling in Arabidopsis arenosa  Challenging reproductive barriers between and within species: the role of ploidy in the genesis of new species  The phased genome assembly of Xenopus mellotropicalis sheds light on the pace of subgenome dominance and diploidization in animal polyploids  Sex, behavior, and giant cells: missing elements of animal polyploid evolution with the parasitoid Nasonia vitripennis  Chromosomes, the more the merrier? How does tetraploid C. elegans resist severe cold stress at the gravid adult stage?  Metabolic and physiological responses to genome duplication in duckweed: Linking	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova Mr. Changfu Jia Ms. Eliška Petříková Ms. Diana Frazão Mr. Zhen Ll Dr. Kelley Leung
S13 - Evolution P01.138  P01.139  P01.141  P01.143  P01.144  P01.145  P01.146  P01.148  P01.149  P01.150	stion after whole genome duplication: transformative advances and new directions  Single-cell sequencing suggests a conserved function of Hedgehog-signaling in spider eye development  Ploidy-dependent differences in mutation rate, spectrum and adaptive outcomes in Chlamydomonas reinhardtii  A tektin gene family analysis of chromosome-level annelid genomes identifies an ancestral cluster of three tektin35B genes for one of the two major clades of Polynoidae/scaleworms  How does the common barbel (Barbus barbus) see? The effect of the whole-genome duplication on vision  Dynamic Transposon Activation and Gene Regulatory Rewiring Drive Functional Divergence of Duplicated Genes During the Rediploidization Process in Orychophragmus violaceus  Polyploidy in Action: Disentangling the Consequences of Genome Doubling in Arabidopsis arenosa  Challenging reproductive barriers between and within species: the role of ploidy in the genesis of new species  The phased genome assembly of Xenopus mellotropicalis sheds light on the pace of subgenome dominance and diploidization in animal polyploids  Sex, behavior, and giant cells: missing elements of animal polyploid evolution with the parasitoid Nasonia vitripennis  Chromosomes, the more the merrier? How does tetraploid C. elegans resist severe cold stress at the gravid adult stage?	Ms. Brenda Irene Medina Jiménez Mr. Antoine Van de Vloet Ms. Steffanie Mutiara Ms. Zuzana Konvickova Mr. Changfu Jia Ms. Eliška Petříková Ms. Diana Frazão Mr. Zhen Ll Dr. Kelley Leung Mr. Clement Verdier

P01.155	Population genomic evidence of inter-ploidy gene flow in natural plant populations	Dr. Jörn Frederik Gerchen
P01.156	Genomic and functional mechanisms underlying early adaptation to whole genome duplication per se.	Dr. Stella Huynh
P01.157	Consequences of induced whole genome duplication on the memory of environmental stress in tetraploid C. elegans	Ms. Emma Bazzani
P01.158	Zombie Gene Flow: Asexual Polyploid Hybrids Reintroduce Genomic Elements of Extinct Species into Contemporary Sexual Populations	Dr. Lucija Andjel
P01.159	Evidence for lineage-specific rediploidisation following the 1R whole-genome duplication at the base of the vertebrate lineage	Ms. Róisín Long

. 02.203	the base of the vertebrate lineage	
S14 - Evolut	ion at species range margins	
P01.160	Adaptive patterns of anti-predator escape behavior in a globally introduced bird species	Prof. Mark Hauber
P01.161	Dispersal evolution can only rescue a limited set of species from climate change	Mr. Peter Kamal
P01.162	Where we're going, we won't need eyes to see: Distributional range size drivers for subterranean spiders	Dr. Adrià Bellvert
P01.163	Unraveling the evolutionary history of Orestias agassii: a phylogeographic study considering its current range.	Ms. Viviana Araya
P01.164	Gradients of increased reproductive effort and increased leakiness in sex expression with range expansion in the dioecious plant Mercurialis annua	Ms. Mai Thu Nguyen
P01.166	Populations on the Edge: Range Dynamics and Conservation of Great Crested Newts Under Global Change	Mr. Charlie Towler
P01.168	Consequences of range shifts on the genetic diversity of butterflies in Finland	Ms. Lola Fernández Multigner
P01.169	An empirical test of Baker's law: Dispersal favors increased rates of self-fertilization	Dr. Michelle Mccauley
P01.170	The K-tabase: A Comprehensive Database of Plant Genetic Diversity and Population Structure Statistics for Macrogenetic Analysis.	Ms. Tinu Adenegha
P01.171	Uncovering Hybridization and Species Boundaries in Peripheral Populations of the Great Tit (Parus major) Using High-Resolution Genomics	Dr. Sahar Javaheri Tehrani
P01.172	Pan-range SNP phylogeography highlights differing effects of postglacial colonisation history on mitochondrial, autosomal, and sex-linked genome compartments	Dr. Miranda Sherlock
P01.173	Speciation between populations at the margins of Solanum chilense: Phenotypic differences and transcriptomics of maternal effects in F1 hybrids.	Ms. Sidra Tul Muntaha
P01.174	Behavioral isolation between two color morphs of the Mourning Wheatear (Oenanthe lugens)	Prof. Roi Dor
P01.175	Genomic Effects of the Successful Post-Glacial Expansion of Bombina pachypus	Dr. Victor Hugo Muñoz Mora
P01.176	Sex ratio in core, edge and refuge populations of Mercurialis annua: Home and away	Dr. Yuval Sapir
P01.178	The Evolutionary Path and Divergence of the Oncocyclus Irises Species Complex Across the Southern Levant	Dr. Yamit Bar-Lev
P01.179	Revisiting Classic Theories to Explain Global Patterns of Genetic Diversity	Dr. Alberto Garcia
P01.180	The genotype-phenotype map of repeated seasonal adaptation during range expansion of a butterfly $ \\$	Prof. Karl Gotthard
P01.181	Reconstructing the Evolutionary History of the Genus Orestias (Teleostei: Cyprinodontidae): An Integrative Phylogenomic, Morphological, and Biogeographic Approach across the Central Andes	Dr. Marco Mendez
P01.182	A systematic review of hybridization dynamics in frogs: Associations with breeding patterns and courtship behavior	Dr. Renato C. Nali
P01.183	Common Origins, Divergent Paths: Gonadal Development in the Domestic Chicken and the Zebra Finch in the Context of Avian Evolution	Ms. Paulina C. Mizia
P01.184	Divergence at the edge: genomic signatures of adaptation in marginal populations of the European mole (Talpa europaea)	Dr. Paolo Colangelo
P01.185	Genomic architecture of genetic differentiation in two hybridizing mussels from the mid- Atlantic Ridge	Dr. Coral Diaz-Recio Lorenzo
P01.186	Evolutionary suicide in the secondary contact zone	Dr. Tom Keaney
P01.187	Phylogeography of the shrimp Palaemon longirostris reveals high connectivity along the Atlantic European coast with a latitudinal gradient in diversity	Dr. Enrique Gonzalez Ortegon
P01.188	Estimating individual stress along expansion ranges: a telomere based approach	Mrs. Gaëlle Brahy



P01.189 Dr. Will Nash From Breakthrough to Breakdown? Genetic Consequences of a Rapid Range Expansion in a

Key European Pollinator.

P01.190

Genomic diversity of jaguars at the edge of their range

Dr. Patricia Saragüeta

## S17 - Evolution of behavioural diversity: from ecology to genes and neural systems

P01.015 Personality development in wild house mice: Evidence for a nutrition-dependent sensitive Dr. Nicole Walasek period early in life

	period early in life			
S18 - Evolu	S18 - Evolution of biotic interactions across scales			
P01.191 P01.193	Syntopy promotes song divergence in a Neotropical avian radiation	Dr. Vicente García-Navas Dr. Luis-miguel Chevin		
P01.193 P01.195	Evolution of plasticity and character displacement in a fluctuating environment Local adaptation of host-parasite coevolution on spatial networks	Ms. Saismit Naik		
P01.196	Experimental Removal of Niche Construction Alters the Pace and Mechanisms of Resistance			
. 02.250	Evolution			
P01.198	Towards evidence for plant-bacteria coevolution: an ecophylogenetic framework reveals selection through niche construction	Dr. Yanis Bouchenak-khelladi		
P01.199	Altruism or Selfishness? Evolution of floral behavior in response to neighboring kinship.	Mr. Haruto Tomizuka		
P01.200	From spore to infection: How the intensity and frequency of heatwaves shape bacterial infections of a crustacean	Ms. May Shehady		
P01.201	Parasite population genomics sheds light on canid evolution	Dr. Rosemonde Power		
P01.202	Making sense of floral scents: species- and population-level variation in floral scent in Mimulus section Erythranthe and its evolutionary significance	Dr. Kelsey Byers		
P01.203	Is a generalist parasitic plant locally adapted to its plant hosts?	Ms. Grace Doherty		
P01.204	Transcriptomic signatures of infection and insights into transgenerational immune priming in an invertebrate host	Mrs. Sofia Paraskevopoulou		
P01.205	The genomic basis of an obligate ant-plant symbioses	Mr. Noah Bourne		
P01.207	Ecological conditions can promote female-biased sex-ratios: a case study in haplodiploid gregarious pine sawfly	Dr. Carita Lindstedt		
P01.208	Neuropeptides mediate behavioral and lifespan changes in cestode-infected ants	Ms. Giulia Blasi		
P01.209	Examining the relationship between genomic variation and avian cholera survival in an Arctic sea duck	Ms. Shayla Kroeze		
P01.211	Enhancing plant disease resistance through intra and interspecific diversification	Mrs. Claire Neema		
P01.212	From antagonists to mutualists: the emergence of brood-site pollination in Ochyromerini weevils revealed by a new weevil specific UCE probe set.	Mr. Benjamin Zelvelder		
P01.213	The INTERACT project: Understanding evolution of common bean plant-plant interactions	Mrs. Francesca Francioni		
P01.214	Exploring Density-Dependent Shifts in Symbiotic Interactions Using Paramecium bursaria and Chlorella Species	Ms. Liwen Chu		
P01.215	Community tipping driven by predator-prey role reversal in a fishing-induced trophic cascade	Ms. Dhanya Bharath		
P01.216	Signatures of Reproductive Skew and Corresponding Multiple Merger Coalescence (MMC) in Populations of Plasmodium falciparum	Ms. Uthara Srinivasan		
P01.217	Evolution of coexistence ability in experimental Daphnia mesocosms	Mr. Sebastian Borgmann		
P01.218	Zostera marina and Labyrinthula spp.: Exploring Pathogen Virulence and Host Defense Across A Thermal Gradient	Ms. Jana Schuster		
P01.219	RangeShifter 4.0: modelling spatial eco-evolutionary dynamics for multiple interacting species	Dr. Theo Pannetier		
P01.220	Unraveling the molecular basis of host plant adaptation in a generalist herbivore	Dr. Ernesto Villacis Perez		
P01.221	Exploring the evolutionary dynamics of mating system shifts and speciation in Capsella	Mrs. Alessandra Lombardi		
P01.222	Host-parasite coevolution in complex natural systems: linking immune gene variation to infection patterns across amphibian species	Dr. Gemma Palomar		
P01.223	Divergent Defence: Multiple Evolutionary Routes of Algal Resistance and Their Costs	Ms. Piyumi Sandaruwani De Alwis		
P01.224	Twenty-years of Tuberculosis-driven selection on meerkat MHC	Ms. Vanessa Riegel		
P01.226	When Durum Wheat grows together with Alfalfa: The Genetic Dialogue of Neighboring Plants	Mr. Nicolas Salas		

A CONTRACTOR OF THE PARTY OF TH	

P01.227	Ecological aspects of coinfections: Do laboratory experiments of within-host parasite interactions reflect the reality of coinfections in nature?	Prof. Frida Ben-ami
P01.229	Questioning the early Miocene origin of grass-feeding noctuids: evidence from a phylogenomic dataset	Dr. VINEESH NEDUMPALLY
P01.230	A la carte seed harvesting: Messor barbarus selects durum wheat genotypes	Mr. Clément Plessis
P01.233	Butterfly counter-adaptations against multiple plant defences: arms race, plasticity-first	Dr. Erika Pinheiro de Castro
101.233	and convergent evolution	Dr. Erika i ililicilo de custro
S20 - Evolu	tionary biology meets genetic pest control	
P01.235	Integration of databases and High-Performance Computing in Dengue Virus research: Advances in the development for RT-LAMP detection	Profesor. Cristian E. Cadena-Caballero
P01.236	Sting Operation: Population Genetics of Yellow Legged Hornets (Vespa velutina) Using	Ms. Sarah Griffin
	Single Nucleotide Polymorphisms	
P01.237	POLYMORPHISM OF THE OVINE-MHC DRB1 EXON 2 GENE AND ITS ASSOCIATION WITH TICK	Prof. Theopoline Omagano
	INFESTATION IN DAMARA SHEEP AT NEUDAMM FARM, NAMIBIA	Amushendje
P01.238	Epistasis among fungicide resistance mutations	Mr. Luka Biočanin
P01.239	Insect-plant interactions at cell-type resolution: insights from the green-veined white	Ms. Brenda Irene Medina Jiménez
P01.240	Transcriptional profiles of the fish parasite Neoechinorhynchus agilis (Acanthocephala) emphasize energetic stress in males and high cell-division activity in females	Dr. Camille-sophie Cozzarolo
P01.242	Limited spatial propagation of quorum-drive systems through wave pinning	Dr. Guilhem Doulcier
P01.242 P01.243	Spatial soft-sweeps: a means to jointly estimate recent demography and migration in	Dr. Bhavin Khatri
101.243	Anopheles gambiae from insecticide resistance loci	DI. DIIAVIII KIIACII
P01.244	Optimizing biological pest management by investigating the genetic basis of parasitoid	Dr. Jan Buellesbach
	chemical communication systems	2.734.7 246.163246.1
P01.246	Thermal adaptation amplifies increases in insect crop damage from climate warming	Dr. Loke Von Schmalensee
COA Frank	*:	
P01.247	tionary consequences of heterokaryosis, mosaicisms, chimeras and other monsters  NLR evolution and the origins of heterokaryon incompatibility genes in Podospora	Dr. C. Lorona Amont Volásquez
FU1.247	NER evolution and the origins of neteroraryon incompatibility genes in Podospora	Dr. S. Lorena Ament-Velásquez
P01.249	Heterosis in Saccharomyces hybrids: regulation of cold-adapted proteins under heat stress	Ms. Damla Temel
S23 - Evolu	tionary Genomics: Understanding and Adapting to Climate Change (organised by the journals	Molecular Ecology & Evolutionary
Application		, , , , , , , , , , , , , , , , , , , ,
P01.250	Disentangling the genomic basis of terrestrialization across animals through a	Ms. Klara Eleftheriadi
	transcriptomics spyglass	
P01.254	Environmental adaptations in Arabidopsis lyrata	Mrs. Laura Steinmann
P01.255	Evolutionary Tuning: Understanding the Influence of Stage-Specific Diet in Shaping Developmental time in Drosophila melanogaster	Mr. Chandrakanth Mohankumar
P01.256	Fecal metagenomics as a conservation tool: Assessing pathogen loads in the critically	Mr. Román Sapino Funes
P01.257	endangered Iberian desman  Adaptive and non-adaptive responses to metal contaminants and pathogens in wild brown	Dr. Jessica Côte
FU1.237	trout Salmo trutta	DI. Jessica Cote
P01.258	An evolutionary and mechanistic study of stressor interaction patterns between pesticides,	Ms. Jue Li
101.230	fish predation and heatwaves in natural populations of Daphnia magna	Wist suc El
P01.259	Genomic signatures and demographic history of the widespread and critically endangered	Dr. Guoling Chen
	yellow-breasted bunting	
P01.260	Predicting adaptive potential from genomic data and its implications for conservation	Ms. Katie Abson
DO1 261	Convergent molecular evalution of thermal televance in re	Dr. Christopher Toylor
P01.261	Convergent molecular evolution of thermal tolerance in mammals	Dr. Christopher Taylor
P01.263	Uncovering adaptive traits in legumes through novel genotyping approaches	Dr. Giacomo Conti
P01.264	Predicting the genomic offset of fruit trees to climate change by building a reproducible	Mr. Maxime Criado
P01.265	bioinformatic pipeline  How does introduing affect local adaptation in the mountain plant Arabis aloina?	Ms Fraya Way
ru1.203	How does inbreeding affect local adaptation in the mountain plant Arabis alpina?	Ms. Freya Way
P01.266	Tracing genomic consequences of historical sealing in South American fur seal populations	Ms. Beril Yildiz
P01.267	Cold-response is altered by salinity exposure in an intertidal snail	Dr. Alice Dennis
. 01.20/	cora response is aftered by sammey exposure in an interdual small	Dirithic Denilli



DO1 360	Discourage is and adoptative history of the houteuff. Halisania, and in Nastanaiael	NAs Patrícia Mashada	
P01.268	Biogeographic and adaptative history of the butterfly Heliconius erato in Neotropical	Ms. Patrícia Machado	
P01.269	forests: understanding the past to preserve the future A Tale of Two Seas: Local Adaptation and Osmoregulation in the copepod Acartia tonsa	Ms. Alexandra Hahn	
P01.272	Selection mirage - Ecological selection led to repeated color changes during plant domestication	Mr. Tom Sönke Winkler	
P01.273	Phylogeographic and genomic insights unveil the evolutionary history and post-glacial recolonization routes of the Palmate Newt (Lissotriton helveticus) into Europe	Dr. Bernat Burriel-Carranza	
P01.274	Plasticity of mutation rates and spectra in the green algae Ostreococcus tauri - Effects of temperature and salinity?	Mrs. Lisa Mettrop	
P01.275	Genomics of Brazilian howler monkeys reveals recent inbreeding and species-specific positive selection	Ms. Katherine McVay	
P01.276	Mitochondrial Insights into the Unique Evolutionary History of Eastern Baltic Cod	Ms. Érika Endo Kokubun	
P01.277	THE EVOLUTIONARY GENOMICS OF PESTICIDE RESISTANCE IN THE WATER FLEA DAPHNIA MAGNA	Mr. Alexandros Kourtidis	
P01.278	TO HAVE OR NOT TO HAVE: Does Having a Reference Genome Make a Difference?	Mrs. Ainhoa López	
P01.279	Gene expansions in herbivorous rodents drive rapid dietary adaptation to climate change	Prof. Maria Denise Dearing	
P01.280	Selection pressures across the introgression landscape of Southeast Asian domestic cattle	Dr. Sabhrina Gita Aninta	
P01.281	Acute heat stress results in sex-specific mortality in birds	Ms. Brynleigh Payne	
P01.282	Environmental insights into sex chromosome evolution in a sex-reversing reptile (Carinacincus ocellatus)	Mr. Carles Ferre Ortega	
P01.283	Tracking Polygenic Adaptation: Phenotypic Stasis After Rapid Change in Large Populations of Drosophila	Ms. Claudia Ramirez-Lanzas	
P01.284	Time-structured genomic sampling reveals abrupt and rare genetic shifts in Eunicella singularis populations, challenging snapshot connectivity estimates.	Ms. Camille Sant	
P01.286	Evolution of facultative symbiosis in the stony coral Oculina	Dr. Xavier Grau-Bové	
P01.287	A new method for identifying local adaptation	Ms. Isabela Do O	
P01.288	Homology-Based Mapping of microRNA Evolution Across Mammals	Ms. Sarahjane Power	
P01.289	Uncovering the genetic response of Oikopleura dioica against climate change-related biotoxins: insights into evolutionary enhance resilience	Dr. Nuria Paz Torres Aguila	
P01.290	Time-Lagged Genetic Erosion from Habitat Fragmentation Undermines Evolutionary Resilience in Arthropods	Prof. Trine Bilde	
P01.291	Bridging developmental plasticity and climate adaptation through evolutionary genomics, population genetics and transcriptomics in African Bicyclus butterflies	Dr. Océane Seudre	
P01.292	Genomic analysis reveals local adaptation and vulnerability to climate change of the African orphan crop finger millet (Eleusine coracana)	Dr. Margret Veltman	
P01.293	Conservation genomics of the threatened boreal wood-decay fungus Amylocystis lapponica	Ms. Jorunn Hellekås	
P01.294	Narrow admixture zone between populations of the fungus Trichaptum abietinum in Norway	Ms. Anneli Andersen	
P01.295	Genetic Diversity and Population Structure of Bolivian Camelids Revealed by SNP Genotyping	Ms. Alejandra Del Pilar Roman Peña	
P01.297	Using admixture to understand local adaptation in the common wood decay fungus Trichaptum abietinum	Dr. Dabao Sun Lü	
P01.298	Of resilience, population growth, and genetic diversity	Dr. Ivan Scotti	
P01.299	Repeated local adaptation and genetic vulnerability to climate change in North American songbirds	Dr. Joan Ferrer Obiol	
P01.300	The Evolution of Plasticity and Heat Adaptation in Green Algae	Ms. Yeshoda Harry-Paul	
S27 - Gene	tic conflict : Evolutionary and Genomic consequences		
P01.301	Expression divergence in response to sex-biased selection	Ms. Michelle J. Liu	
P01.302	Phylogenomic Reconstruction and Taxonomic Clarification of the Ustilaginales Order	Mr. Sean Blundell	
P01.303	Evolutionary capacitance driven by Hsp90 during the de novo evolution of multicellularity	Ms. Dung Lac	
P01.304	Investigating the cause of non-Mendelian inheritance of the Robertsonian 6.16 chromosome in mouse	Dr. Frances Burden	



P01.305 P01.306	Evolution and genomics of an X-linked meiotic driver in stalk-eyed flies  The 'Jump-Corruption (Gekokujō) Evolution' hypothesis in the sex-determining gene Sry in	Mr. Benjamin Alston Ms. Honoka Okuyama	
P01.308	therian mammals  Causes and consequences of sex-biased reproduction in the flea beetle Altica lythri	Dr. Kim Rohlfing	
P01.309	Following the genomic traces of introgressed mitochondria in Drosophila paulistorum	Mr. Konstantinos Papachristos	
P01.310	How sexes diversify: Molecular drivers underlying sexual color dimorphism in cichlids	Ms. Muktai Kuwalekar	
P01.312 P01.313 P01.314	Red Queen dynamics in the determination of recombination hotspots Selfish male reproduction in stick insects Selfish Elements and Selective Sperm: impact of haploid selection on zebrafish fitness	Dr. Diego Hartasánchez Mr. Luca Soldini Dr. Alice Godden	
P01.316 P01.317	Genetic drift influences levels of sex-biased expression in metazoans Sexual tension over intragenomic conflict: Bacillus subtilis bacteria and pB32 plasmid	Dr. Carolina Barata Dr. Ana - Hermina Ghenu	
P01.318 P01.319	Adaptive potential of hybridization despite initial hybrid breakdown On the roles of transposable elements in East African cichlid evolution	Ms. Marit Kuijt Dr. Miguel Almeida	
S31 - Habit	cat-forming species and global change: a multidisciplinary perspective on their evolution and a	adaptive potential to improve their	
P01.322	Hidden Diversity: Genomic Evidence of Cryptic Species and Habitat Specialization in the Brooding Coral Seriatopora hystrix Along the Great Barrier Reef	Ms. Ilha Byrne	
P01.324	'Green tide' on the Korean coast: the major green tide forming species, biomass, and ecological impacts on benthic invertebrate ecosystems	Prof. Hyuk Je Lee	
P01.325	Maximising adaptive capacity and climate resilience in structural rainforest trees: a versatile population genetics workflow for both restoration and conservation outcomes	Mr. Richard Dimon	
S41 - Post-	phylogenomics: new and evolving molecular methods to address challenging phylogenies		
P01.326 P01.327	A timetree of Fungi dated with fossils and horizontal gene transfers SARS-CoV-2 Zoonotic and Reverse-Zoonotic Transmissions Between Humans and Mink	Dr. Eduard Ocana-pallares Ms. Marlies Jilles Francine Goedknegt	
P01.328	Population genomic divergence reveals uncertainties in species identification of cave- dwelling groundwater isopods	Mr. Stefano Lapadula	
P01.329	Amino Acid Exchangeabilities Differ More Between Surface and Buried Sites than Among Species	Mr. Peter Goodman	
P01.330	PhyloFusion - Fast and easy fusion of rooted phylogenetic trees into rooted phylogenetic networks	Ms. Banu Cetinkaya	
P01.331	Transcriptomic insights into the enigmatic origins of monotypic genera Mankyua and Helminthostachys in Ophioglossaceae	Ms. Jaeseo Shin	
P01.332 P01.333	Diversification of allotetraploid species with identical genomic composition  Resolving taxonomic uncertainties in genus of the edible chayote (Sechium P. Browne (Cucurbitaceae)) and exploring potential center of domestication of chayote through genomics analysis	Mr. Benneth Nass Ms. Cécile Truchot Taillefer	
P01.334	Inferring gene flow from phylogenies with too many genomes	Mr. Diogo Ribeiro	
S42 - Predi P01.024	cting evolutionary change in ecologically relevant contexts  Rapid evolution reveals trade-offs across ecological interactions in a host-parasitoid system	Mr. Juan Esteves	
S43 - The dynamics and consequences of bacteria-bacteriophage interactions and co-evolution in complex communities			

## S43 - The dynamics and consequences of bacteria-bacteriophage interactions and co-evolution in complex communities

P01.335	Understanding phage-host dynamics during the course of a cyanobacterial bloom in the Baltic Sea	Ms. Zahra Goodarzi
P01.336	Spacer Chronicles: Reconstructing ancestral CRISPR Spacer Arrays to Reveal Phage-Host	Mr. Axel Fehrenbach
P01.337	Dynamics  Eco-evolutionary dynamics of phage-bacteria interactions in the human gut through the	Dr. Jaime Iranzo
P01.338	analysis of CRISPR arrays  Temperature extremes alter phage predation dynamics and the capacity for bacterial	Dr. Zachary Bailey
PU1.338	coexistence	Dr. Zachary Balley
P01.339	Surprising morphological and genetic diversification during experimental evolution of phage M13	Ms. Liya Miksovsky



Disentangling the evolutionary impacts of relatedness and facultative/obligate life cycles during the transition to multicellularity	Ms. Autumn Peterson
The Race Between Entropy and Maximum Entropy: Balancing Innovation and Survival	Mr. Jiří Nedomlel
Applying Statistical Models to the Family Resemblance Definition of 'Life'	Prof. Jessica Abbott
Symposium	
Introgression and divergence in a young species group	Dr. Ina Satokangas
Urban warriors have bigger shields – how urbanization affects ornament expression and individual quality in a common waterbird	Dr. Amelia Chyb
Ecological and developmental mechanisms of floral diversification in selected South	Dr. Marina Strelin
One species, many shells: Heritable habitat-associated divergence in the sole lymnaeid snail	Ms. Micaela Müller Baigorria
	Mr. Basanta Khakurel
A formal theory of group-level adaptation for obligate eusociality	Ms. Kalyani Twyman
	Mr. Harry Ames
RECUR: Identifying recurrent amino acid substitutions from multiple sequence alignments	Dr. Yi Liu
Evolutionary origin of vertebrate hematopoiesis: insights from lamprey development	Mrs. Ana Hernandez-Martinez
Using ddRADseq to explore the genomic landscape of serpentine habitation in three diploid plant species from the Iberian Peninsula	Mr. Pablo Arrufat
Monogamy, Parental Care and the Emergence of Eusociality in Insects	Dr. Rosa Bonifacii
Genomics and evolution of the expanding pine pathogen Lecanosticta acicola	Dr. Marina Marcet Houben
The interplay of sexual selection and hybridization can drive sexual radiation	Dr. Kotaro Kagawa
Socially responsive crickets: insights into the evolutionary consequences of behavioural interactions	Dr. Tom Ratz
Host-shift changes patterns of morphological integration in a sex-specific manner: an experimental evolution approach in a seed beetle	Dr. Sanja Budečević
	Dr. Carola Greve
Mitochondrial Haplotype Distribution and Association with Reproductive Mode in Wild New Zealand Mud Snail	Ms. Giulia Lin
Patterns of Soil Nematode Biodiversity in the Atacama Desert	Mrs. Laura Villegas
Crossing the biggest ocean in the world – the phylogeography of how plant genus Wikstroemia colonised the islands of the Pacific	Mr. Ruben Cousins-Westerberg
Evolutionary Plasticity of Sex Determination: Multi-Omic Insights from a Rudimentary Hermaphroditic Sparid Teleost	Mr. Sam Modern
A new model of background selection improves our understanding of human genetic diversity	Dr. Gustavo Valadares Barroso
Adaptive physiological responses of Daphnia pulicaria to elevated salinity	Ms. Varvara Paida
Post-copulatory sexual selection intensity is associated with vagina length but not sperm storage traits in Galliformes	Dr. Katherine Assersohn
Quantity over quality? Parental senescence effects in an annual killifish	Dr. Milan Vrtílek
Butterfly adaptations to seasonal tropical environments: links between, polyphenism, reproductive diapause, and migrations	Prof. Freerk Molleman
DNA barcoding of passerine birds in Iran	Dr. Sahar Javaheri Tehrani
	Prof. Hisashi Ohtsuki
Evaluation of simulation-based supervised machine learning methods for inferring	Dr. Frédéric Austerlitz
Intra-specific variation in aging and its life history implications in African annual	Dr. Radim Blazek
Hidden Strength: Mutational robustness in changing environments	Ms. Ronja Hulst
Demographic history inferred from an inversion-rich spruce bark beetle genome	Dr. Piotr Zieliński
Emergence and evolution of angiosperm forests	Mrs. Laura Pajot
Spawning patterns and symbiotic relationships of Korean bitterling fishes (Pisces:	Dr. Hee-kyu Choi
Divergent visual system adaptations in two sturgeon species revealed by single-nuclei analysis	Mrs. Kristýna Eliášová
	The Race Between Entropy and Maximum Entropy: Balancing Innovation and Survival Applying Statistical Models to the Family Resemblance Definition of 'Life'  Symposium Introgression and divergence in a young species group Urban warriors have bigger shields – how urbanization affects ornament expression and individual quality in a common waterbird Ecological and developmental mechanisms of floral diversification in selected South American angiosperms One species, many shells: Heritable habitat-associated divergence in the sole lymnaeid snall species of southern Patagonia Characters changing speed: the Covariomorph model and its impact on phylogenetic trees A formal theory of group-level adaptation for obligate eusociality Exploring the inbreeding dynamics in a cooperatively breeding mongoose. RECUR: Identifying recurrent amino acid substitutions from multiple sequence alignments Evolutionary origin of vertebrate hematopoiesis: insights from lamprey development Using ddRADseq to explore the genomic landscape of serpentine habitation in three diploid plant species from the Iberian Peninsula Mongamy, Parental Care and the Emergence of Eusociality in Insects Genomics and evolution of the expanding pine pathogen Lecanosticta acicola The interplay of sexual selection and hybridization can drive sexual radiation Socially responsive crickets: insights into the evolutionary consequences of behavioural interactions Host-shift changes patterns of morphological integration in a sex-specific manner: an experimental evolution approach in a seed beetle InUrFaCE - Initiative of Urban Facades Creature Exposition Mitochondrial Haplotype Distribution and Association with Reproductive Mode in Wild New Zealand Mud Snail Patterns of Soil Nematode Biodiversity in the Atacama Desert Crossing the biggest ocean in the world – the phylogeography of how plant genus Wikstroemia colonised the islands of the Pacific Evolutionary Plasticity of Sex Determination: Multi-Omic Insights from a Rudimentary Hermaphroditic Sparid Teleost A new model of



			Control of the contro
P01.234	Using moulting cycle time series transcriptome analysis to pinpoint host candidate genes for the Daphnia magna – Pasteuria ramosa host-parasite interaction.	Mr. Luca Pecalli	
P01.241	Estimating Seagrass Meadow Persistence via Clone Aging	Ms. Clara Winguth	
P01.245	Unraveling the migratory puzzle: population genetics of the least flycatcher (Empidonax minimus)	Ms. Sara Castro García	
P01.248	Study of coloration diversity with iPACS: A new methodology to link morphology and transcriptomes at the single cell level.	Dr. Dimitri Meistermann	
P01.251	Hot stuff! Large brains buffer against heatwave induced cognitive decay	Dr. Annika Boussard	
P01.252	Unraveling the gene regulatory networks of predator-induced developmental plasticity within and across generations in Daphnia magna	Ms. Zorimar Vilella-pacheco	
P01.253	Methylome and transcriptome responses to hydrocarbon pollution: acute and priming effects	Mr. Hamish Williams	
P01.262	Cytogenetic analysis of satellite content of five species of arvicolid rodents (Arvicolinae, Rodentia)	Ms. Alona Yurchenko	
P01.270	The evolutionary impact of migration on genetic and cultural differentiation	Mrs. Niccole Porras	
P01.271	Co-diversification of sperm and female sperm storage organs cause reproductive isolation in diverging populations	Mr. Dick Moberg	
P01.285	Exploring plumage polymorphism in birds of prey – a systematic map	Dr. Pawel Podkowa	
P01.296	Selfing VS Outcrossing: Evolutionary Consequences in a Haploid World	Mr. Timothee Fichant	
P01.307	Gestation length both shapes and is shaped by other life history traits in eutherian mammals.	Mr. Theodoros Danis	
P01.311	Elastic solution for damage avoidance in genital lock-and-key evolution	Prof. Yasuoki Takami	
P01.315	Genome-wide assessment of three diverged lineages of common seadragon in the face of climate change	Dr. Laura Tensen	
P01.320	The adaptive dynamics of a trait operating over multiple interaction networks	Mr. Lewis Flintham	
P01.321	Does the evolution for predatory aggression alter conspecific aggression? Insights from a selection experiment in bank voles	Mr. Gokul Bhaskaran	
P01.323	Parental age effects on offspring telomere length across vertebrates: a meta-analysis	Prof. Hannah Dugdale	
P01.343	Active predators do not necessarily specialize in sedentary prey: a simulation model	Prof. Inon Scharf	
P01.344	Do social bird species have more stable demographic histories?	Mr. Louis Bliard	
P01.345	Plugging into Innovation: Spatial Transcriptomics of Tissue Organization and Gene Expression in the Ricefish Plug	Ms. Alina Schüller	
P01.346	Insights into the genome of dromedary camels: signatures of positive selection and copy number variations	Dr. Hussain Bahbahani	
P01.347	Extraordinary adaptations: Functional and evolutionary synergy of trait components can explain the existence of leaf masquerade	Dr. Benito Wainwright	
P01.348	Co-evolutionary dynamics of dietary breadth and body size in moths	Mr. Stenio Foerster	
P01.349	Heavy metal tolerance and bioaccumulation potential in fungi isolated from Kiruna Iron Ore mines $ \\$	Mx. Noah Hensen	
P01.350	Conservation genetics in ectomycorrhizal fungi: estimating the effective population size (Ne)	Ms. Anouck Champion	
P01.351	The evolution of chemical defence in the mint family - investigating the functional divergence of terpene synthases	Ms. Violette Pepper	
P01.352	Impact of landscape structure on trait variation	Ms. Ulla Riihimäki	
P01.353	Inferring genome-wide patterns of ancestral divergence and introgression in genus Epinephelus	Dr. Yan Chi Arthur Chung	
P01.354	From Policy to Practice: Progress Towards Data-and Code-Sharing in Ecology and Evolution	Dr. Edward Ivimey-Cook	
P01.355	Impact of geography, seasonality and experimental selection on Chironomus riparius recombination rates	Dr. Maria Esther Nieto Blazq	uez
P01.356	Is there a limit to the adaptive potential of chromosomal inversions with regard to global warming?	Prof. Frances Mestres	
P01.357	Gene flow between marine crab (Liocarcinus depurator) populations from the Atlantic- Mediterranean transition	Prof. Frances Mestres	
P01.358	Holobiont plasticity and transcriptome stability in a gorgonian facing local anthropic pressures	Dr. Didier Aurelle	
P01.359	Potential Application of Disease Associated SNPs for Forensics: A study on the Kuwaiti population	Prof. Suzanne Albustan	
P01.360 P01.361	The genetic dynamics of Red Queen coevolution extracted from sediment cores Fluctuating selection among years in a wild insect	Mr. Jahn Ringger Prof. Tom Tregenza	

		And the second s
P01.362	THE HORMONAL CONTROL OF REGENERATION AND REPRODUCTION IN THE MARINE ANNELID SYLLIS PROLIFERA	Ms. ANDREA Martín-Salas
P01.363	Mutations conferring resistance to second-line drug treatments in the Mexican population affected by tuberculosis	Dr. Ikuri Alvarez-Maya
P01.364	Genomic tools for identifying the origin of Pinus pinaster reproductive material	Dr. Sanna Olsson
P01.365	Super-shedder vs. super-spreader: from physiology to behaviour	Dr. Luis Silva
P01.366	lodine deficiency-induced goiter constrains inland colonization of fish	Prof. Jun Kitano
P01.367	The evolutionary predictability of avian colouration under natural and sexual selection	Ms. Jasmine Hardie
P01.368	Diversity and adaptation in avian wing bone inner structure: insights from extant and extinct Strisores	Dr. Fabio Alfieri
P01.369	Timing is everything: Evolution of ponerine ants highlights how dispersal history shapes modern biodiversity	Dr. Maël Doré
P01.370	Transition from monogamy to polygamy in human-modified environments: the case of free-ranging dogs	Dr. Clément Car
P01.371	Sexual conflict over floral behaviour: pollination-induced flower closure is costly to the pollen recipients in a sexually reproducing diploid dandelion	Dr. Daisuke Kyogoku

## Tuesday 19 August, EXPO AREA: 5:00 PM - 7:00 PM - POSTER SESSION 2 - with Aperitif

research on species interactions

S04 - Ageir	ng outside of the box: insights from unusual and non-model species	
P02.001	Surviving on limited resources: effects of caloric restriction on growth, gene expression and	Dr. Freya Pappert
	gut microbiota in a species with male pregnancy (Hippocampus erectus).	,
P02.002	The role of selection pressure in the evolution of asynchrony of aging	Ms. Claire Tsui
P02.003	Molecular and structural diversity of multi-ciliated cell types during larval development in the marine annelid Platynereis dumerilii	Ms. Steffanie Mutiara
P02.004	Nutritional trade-offs in model and non-model insects and its consequences for lifespan, reproduction and sexual conflict	Dr. Juliano Morimoto
P02.007	Exploring the causes of lifespan differences among dog breeds: a bourgeoning model for longevity research	Dr. Alejandro Gonzalez Voyer
P02.008	Epigenetic Clocks of Soay Sheep	Ms. Lucy Barnard
P02.009	Age-related changes and selective disappearance shape variation in bold-shy continuum in guppies	Dr. Magdalena Herdegen Radwan
P02.010	Environmental influences on sex-specific senescence in Soay sheep: insights from a longitudinal study	Ms. Elizabeth Drake
P02.011	A phylogenetic analysis of sporocarp longevity and alternate life history strategies in Polypore mushrooms	Ms. Gayathri Venkatraman
P02.012	A life history model of continuous growth and negligible senescence under resource allocation trade-offs	Mr. Arttu Soukainen
P02.013	Investigating the Plastic and Evolutionary Effects of Isocaloric Macronutrient Composition on Aging and Organismal Performance in Outbred Drosophila melanogaster Populations	Mr. Devashish Kumar
P02.014	Connecting moth flight timing, ageing and predation risk – a citizen science approach	Dr. Jelle Zandveld
P02.015	Asymmetric life history trade-offs shape sex-biased longevity patterns	Ms. Ella Rees-baylis
P02.016	An Evolutionary Origin of Ageing: Jensen's Inequality and Asymmetric Division in Unicellular Life	Mr. William Singleton
P02.017	The ageing methylome of the King penguin	Dr. Flávia Nitta Fernandes
P02.018	Nonhuman primate species share Alzheimer-like neuropathology with humans	Ms. Brier A. Rigby Dames
P02.019	The relationship between phylogeny and age alignments in mammals	Dr. Alexandra de Sousa
P02.020	Extreme Sexual Size Dimorphism: Sex-Specific Growth Pattern in a Spider	Mr. Tim Prezelj
P02.021	The Mammal That Rewired Insulin Signaling to Live Long and Prosper on an Extreme High-	Dr. Jasmin Camacho
	Sugar Diet	
S11 - Enha	ncing Diversity and Transparency in Ecology and Evolution: Reliable Practices for Research and C	Organisations
P02.022	Psychological insights for building accessible R Shiny apps in ecological and evolutionary modeling	Dr. Ana - Hermina Ghenu
P02.023	SPI-Birds: increasing open, reliable and transparent analyses based on avian long-term monitoring in the wild	Dr. Amélie Fargevieille
P02.024	From predation to mutualism: authors' gender identities influence the focus of ecological	Dr. Camille Thomas-Bulle



Do artificial intelligence tools help students with their evolutionary ecology homework? And are they willing to disclose it?

P02.025

Prof. Joanna Rutkowska

S15 - Evolu	ition in and of diverse genetic systems	
P02.026	The evolution of reproductive mode in the Carassius auratus complex	Mrs. Reem Alhaidose
P02.027	Mitochondrial RNA interference in the blue mussel: barriers to overcome	Mx. Marianne Hubert
P02.028	Mitonuclear coevolution as a driver of latitudinal avian speciation trends	Ms. Ellen Nikelski
	·	
P02.029	Hermaphroditism in Insects; The Unbelievable Case of Icerya purchasi	Ms. Elpida Skarlou
P02.030	Evolution of Translation GTPases: From LUCA to Extant Diversity	Ms. Evrim Fer
P02.032	Breeding system influences mating-type proteins evolution in Microbotryum fungi	Dr. Alice Namias
P02.033	Sharks and rays have the oldest vertebrate sex chromosome with unique sex determination mechanisms	Mr. TAIKI NIWA
P02.034	Faster-X evolution in globular springtails	Dr. Sam Ebdon
P02.037	Mutational Biases and Selection in Mitochondrial Genomes: Insights from a Comparative	Mr. Alexandre Schifano
F02.037	Analysis of Natural and Experimental Populations of Caenorhabditis elegans	Wil. Alexandre Scillano
P02.038	Disentangling the complex evolutionary history of sex chromosomes in snakes	Mr. Tomáš Pšenička
P02.039	Mechanisms of transition to parthenogenesis in hybrid stick insects	Ms. Morgane Massy
P02.041	Repeated evolution of all-female populations in facultatively parthenogenetic stick insects	Mr. Luca Soldini
P02.042	Massive genome expansion and rearrangement in an androdioecious scale insect	Dr. James Galbraith
P02.043	DNA methylation patterns underlying paternal genome silencing in the mealybug Planococcus citri	Ms. Tamsin Woodman
P02.044	From Genome to Function: Evidence of Transcriptional activity in Sciaridae Germline- Restricted Chromosomes	Mr. Riccardo Kyriacou
P02.045	Elucidating molecular mechanisms of sex determination in stick insects	Ms. Emelyne Gaudichau
P02.046	Sexual conflict and the evolution of unconventional sex-roles and male pregnancy in pipefish	Mr. Kevin Hsiung
P02.047	Insights into inheritance and genomic variation patterns following hybridisation between two divergent cichlid lineages	Ms. Jaysmita Saha
P02.048	The genomic predictability of hybridization: insights from pine contact zones across contrasting environments	Dr. Bartosz Łabiszak
P02.050	Recurrent centromere repositioning in an otherwise stable karyotype of birds	Prof. Alexander Suh
P02.051	IT'S A MATCH: using modelling to understand the evolution of fungal mating and	Dr. Dabao Sun Lü
	incompatibility systems	
P02.052	Unexpected diversity of telomeric repeats in Andrena bees	Dr. Sven Findeiss
P02.053	Diversity of sex chromosome gene regulation in squamate reptiles	Dr. Michail Rovatsos
P02.054	Escape from a conserved system: Sex chromosome turnover in the Hispaniolan giant	Dr. Michail Rovatsos
	galliwasp (Caribicus warreni, Diploglossidae)	
S16 - Evolu	ition in small populations	
P02.055	Mutation bias influences adaptive evolution, even at large population sizes.	Dr. Jake Barber
P02.056	Evolutionary dynamics of invasive Vespula germanica: The role of dispersal and climate in a restricted population	
P02.057	Turnover shapes evolution of birth and death rates	Mr. Teemu Kuosmanen
P02.058	Estimating realized relatedness in a wild baboon population and its implications for kin- biased behavior	Ms. Carlota Galán-Plana
P02.059	Population Genomics of Laboratory Zebrafish: Insights into Domestication and genetic diversity	Prof. Irene Adrian-Kalchhauser
P02.061	Range of trait variation in prey determines evolutionary contributions to predator growth rates	Dr. Ruben Hermann
P02.062	DNA methylation: a way for fast-renewed inbreeding depression in animals? An example with the mollusc species Physa Acuta.	Dr. Audrey Le Veve
P02.063	A century of isolation in contrasting environments drives evolutionary divergence in a small population of salmonid.	Mr. Hervé Rogissart
P02.065	Genomic Insights into Anthropogenic Impacts on Small Mammal Populations Since the Natufian Period	Ms. Ksenia Juravel
P02.066	Reframing the Himalaya: Genomic Insights from 1,217 individuals Into Migration, Isolation, and Malaria Adaptation in a Complex Human Landscape	Prof. Aashish Jha
P02.067	Effects of long-term inbreeding on assortative mating preferences and dynamic colour-	Dr. Timo Thünken

ornament expression in a cichlid fish



P02.068 P02.069	Genetic Diversity and Population Structure of Spanish Goat Breeds  Manning extinction rick in scalary and policy	Dr. Antonia Noce Dr. Francesca Raffini
	Mapping extinction risk in seahorses: integrating genomics, phylogeny, ecology and policy towards evolution-guided conservation	
P02.070	Genetic surveillance reveals low, sustained malaria transmission with clonal replacement in Sao Tome and Principe	Dr. Hsiao-Han Chang
P02.071	From hives to genomes: inferring human impacts and genomic consequences of evolution in small bee populations	Dr. Thibault Leroy
P02.072	Colonisation costs drive weak purifying selection in island rails	Dr. Kees Wanders
P02.073	Mitochondrial Divergence and Population Genomics in a Dwarf Insular Snake	Ms. Alexis Lindsey
P02.074	Genomic Insights into the Evolution of Inbred Loach Populations: Phylogeny and Runs of Homozygosity	Ms. SIZHE Hu
P02.075	Ecological genetics of isolated loach populations indicate compromised adaptive potential	Ms. Xi Wang
P02.076	Purging and Accumulation of Genetic Load in an Expanding Population of Trinidadian Guppies	Dr. Mateusz Konczal
P02.077	First Insights into Population Structure of the Endangered Darwin's Fox (Lycalopex fulvipes).	Mr. Cristobal Valenzuela-Turner
P02.078	Purging of highly deleterious mutations through an extreme bottleneck	Dr. Oliver Stuart
P02.079	Evolutionary rescue in small populations: latitudinal variation in wing polyphenism modifies genetic diversity and connectivity across natural populations in a water strider.	Mr. Aleix Palahi I Torres
P02.080	Exploring the Genomic Signatures of Convergent Dietary Specialization in the Adaptive Radiation of Canary Island Dysdera Spiders	Ms. Marta Olivé Muñiz
P02.081	Loss of genetic variation and establishment of damaging variants in effectively small populations	Dr. Martin Johnsson
P02.082	Microevolutionary changes of Daphnia galeata during the oligotrophication of Lake Constance	Dr. Pelita Octorina
P02.083	Pleistocene climate and human impact shaped the demographic history of Mediterranean gray wolves (Canis lupus)	Dr. Carlos Sarabia
P02.084	Evolutionary potential for pollutant tolerance in urban Daphnia	Mr. Felix Mende
P02.085	Genomics of caribou (Rangifer tarandus) in Western Canada: an assessment of inbreeding and demographic history	Ms. Charlotte Bourbon
P02.086	Behavioral adaptations at key genes in a small and isolated bear population facilitate coexistence with humans.	Dr. Giulia Fabbri
P02.087	Differentiation in host plant performance among beetle populations – adaptation to local plant community?	Mrs. Erica Winslott
P02.088	Unraveling the basis of adaptation to divergent habitats in chaffinch populations on the small island of La Palma	Mr. Miguel Fernández-Janoher
P02.089	Population genomics of the genus Megasyllis (Annelida, Syllidae) reveals hidden connectivity in the Canary Islands	Mrs. Irene Del Olmo
P02.090	Exploring the effects of first year cold-dormancy on the phenotypic, molecular and metabolic responses in head-started Gopher tortoises (Gopherus polyphemus)	Ms. Anet Filipova
P02.091	Conservation genomics in action: Advancing conservation of the small populations of threatened Australian flora	Dr. Jia-yee Yap
P02.092	Toads in Town: Conservation genetics of increasingly fragmented Western Leopard Toad populations in Urban Cape Town	Dr. Holly Nelson
P02.093	Understanding the social and genetic intergenerational impact of inbreeding depression in a wild population.	Mr. Alessandro Vincenzo Pinto
P02.094	A genomic toolkit reveals severe inbreeding depression in Seychelles warblers	Mr. Kiran Gok Lune Lee
P02.095	Sexual selection and population spatial structure interact to shape sex-specific evolutionary responses in physiology	Dr. Maider Iglesias
P02.096	Topi or not topi: population structure of Damaliscus lunatus in Africa and its implication for conservation	Dr. Sabhrina Gita Aninta
P02.097	Speciation from Small Beginnings: Population Size and Environmental Heterogeneity Shape Speciation Dynamics in Individual-Based Simulations	Mr. Cameron Peacock
P02.098	Drivers of adaptive evolution in a relict brown bear population	Dr. Paolo Colangelo
P02.099	Genome Scans of Adaptive Introgression in the genomically eroded Iberian Lynx (Lynx pardinus)	Ms. Laia Pérez Sorribes
P02.100	Patterns of inbreeding and purging selection in sand lizard populations in Sweden	Ms. Lucia Caroli
P02.101	Evaluating genotype imputation in the Iberian lynx: a conservation genomics approach for small and endangered populations	Mrs. Lucía Mayor-Fidalgo

P02.102	Long-term genomic and morphological evolution of island vertebrates	Dr. Margarida Barcelo-Serra
P02.103	Habitat-Driven Divergence: Early Development Variation in Two Ecotypes of the Marine Gastropod Littorina fabalis	Ms. Luisa Kumpitsch
P02.104	Genomic consequences of recent population decline in the endangered butterfly Hipparchia sbordonii	Dr. Sebastiano Fava
P02.105	Comparative Genomics in an Ex Situ Gorilla Population	Ms. Elena Asensi Martínez
P02.106	Beyond panmixia: Disentangling structure and recent declines in a threatened humpback whale population using SFS- and LD-based demographic inferences	Mr. Maël Le Gouellec Le Gouellec
P02.107	Landscape driven isolation and inbreeding in specialist grassland butterflies	Mr. Zachary Nolen
P02.108	On the edge of extinction: genetic monitoring of the Aeolian wall lizard in a fragmented island system	Dr. Josephine Paris
S19 - Evolut	cion of symbioses and interactions in microbial eukaryotes	
P02.110	Genomic signatures of adaptation to stress reveal shared evolutionary trends between Tetrahymena utriculariae and its algal endosymbiont, Micractinium tetrahymenae	Dr. Joseph Kelly
P02.111	Genomic signatures and evolutionary consequences of symbiosis in a beetle-associated fungus	Ms. Frances Alice Pitsillides
P02.112	Intracellular signaling in proto-eukaryotes evolves to alleviate regulatory conflicts of endosymbiosis	Dr. Sam von der Dunk
P02.113	A genomic approach to investigate the specificity of symbiont-conferred defence in the tripartite system aphid – Hamiltonella - parasitoid wasp	Dr. Giacomo Moggioli
P02.114	Genetic variation among progeny shapes symbiosis in a basidiomycete with poplar	Dr. Benjamin Dauphin
P02.115	Host species identity and phylosymbiosis as the main evolutionary and ecological drivers of gut microbiota in wild rodent species in the Yucatán peninsula	Ms. Gabriela Borja Martínez
P02.116	Few Targets but Broad Effects: Molecular Mechanisms of Host Phenotype Modulation by a Cestode Parasite	Dr. Tom Sistermans
P02.117	Eco-Evo of Symbiosis: Environmental Heterogeneity Meets Density-Dependent Feedback	Dr. Matishalin Patel
S22 - Evolut	ionary Ecology of Microbial Symbioses	
P02.118	Pathogen Discovery in Ancient Cattle Remains: Possible Cause of Taurine Extinction in Thailand	Dr. Wunrada Surat
P02.119	Social transmission of the microbiome in wild baboons	Dr. Reena Debray
P02.120	Investigating Past Human Mobility and Interactions in the Southern Cone of the Americas using Phylogenomics of Ancient Commensal Microbes.	Mr. Arve Lee Willingham Grijalba
P02.121	Plant conservation hologenomics: combining population genomics with microbiont community assessments for two orchid species in Denmark	Dr. Ida Hartvig
P02.122	The microbiome of suprapubic and sternal scent glands of wild tamarins suggests microbes play a key role in olfactory communication	Mrs. Silvia Carboni
P02.123	Investigating signatures of local adaptation in the ubiquitous forest ectomycorrhizal fungus Cenococcum geophilum	Mr. Felix Zimmermann
P02.124	Beyond abundance: Symbiont transcriptional plasticity meets host's nutritional requirements	Mr. Aftab Mahmood Ayas
P02.125	Eco-evolutionary dynamics of Pisum-Bacteria interactions mediated by reciprocal selection	Mr. Victor Angot
P02.126	Temperature mediates coevolution between parasitoids and symbiont-protected hosts	Prof. Christoph Vorburger
P02.127	Symbiopectobacterium in Rhodnius prolixus: evidence for a transitional intracellular symbiosis	Ms. Tanisha Moons
P02.128	Foregut microbial communities and the possibility of endogenous ethanol production in honeybees	Dr. Weronika Antoł
P02.129	Ruling the kingdoms: coordinated stress responses in a lichen-forming fungus partnered with algal or cyanobacterial symbionts	Ms. Andrea Goss
P02.130	Micromanaging: The architecture and evolution of host control in a microbial symbiosis	Dr. Erika Hansson
P02.131 P02.132	The influence of gut microbes on the social brain  Host-gut microbiota coevolution in insular populations: A case study of the Balearic lizard	Prof. Joanito Liberti Ms. Caterina Marquès Gomila
P02.133	Podarcis lilfordi Expanding the P. bursaria-algal model for endosymbiosis	Mr. Daniel Malumphy Montesdeoca



P02.134	Phylosymbiosis and other factors shaping gastrointestinal microbiota composition of captive parrots: interspecific and intraspecific variation revealed by large-scale profiling	Mrs. Kateřina Marková	
P02.136	Contextualizing symbiosis: microbial interactions, host environments, and the dynamics of association	Dr. Marie Vasse	
P02.137	The rat gut microbiome is affected by indirect genetic effects from social partners, which could drive rapid evolutionary change	Dr. Amelie Baud	
P02.138	Bee gut bacteria and their arsenal: Evolution and Diversity of secretion systems	Mr. Samuel Akwasi Acheampong	
P02.139	Experimental evolution within the bee gut	Dr. Ramya Ganesan	
P02.140	The Living Fossils of Symbiosis: Wolbachia Genes in Rhodnius prolixus	Prof. Eva Novakova	
P02.141	Quantifying the impact of insect host genotype on Wolbachia titer	Ms. Mountassira MATIN	
P02.142	Breaking eco-evolutionary feedbacks leads to extinction in a bacterial mutualism	Mrs. Yael Baranovitch (Sorokin)	
P02.143	Evolution of gut microbiome during ageing in short-lived African turquoise killifish	Mr. Ulas Isildak	
P02.145	Characterization of Klebsiella pneumoniae-Targeting Phages in an In Vitro FUT2-Dependent Mucosal Model: Implications for Host–Phage Evolution	Ms. Anna Ewertz	
P02.146	Eco-evolutionary feedbacks in C. elegans-microbe interactions link community dynamics to genetic evolution and back	Dr. Thomas Blankers	
P02.147	Exploring the evolutionary ecology of microbes associated with the common garden snail	Dr. Andrew Matthews	
P02.148	Composition of gut microbiota is mostly driven by developmental stage and substrate in an invasive insect pest	Ms. Sido Dunis	
P02.149	Is bird egg sterile? Challenges of metabarcoding in a low-biomass microbiome study	Dr. Martin Tesicky	
P02.150	Persistence of an Influential Passenger: The Endosymbiotic Association of Wolbachia with the Neotropical Fly Drosophila willistoni	Mr. Marcos Aurélio Martins Oliveira Da Silva	
P02.151	Tissue Type drives plant-associated bacterial community structure more than geographic location: Insights from Antarctica and Patagonia	Dr. Suni Mathew	
P02.153	Disentangling Ecological Variability: The Role of Richness and Environment in Gut Microbiota Dynamics	Ms. Maria Virginia Ramirez-Montoya	
P02.154	Mitochondrial adaptation regarding different chloroplast retention capabilities in Sacoglossa	Dr. Carola Greve	
P02.155	Patterns of correlation between mycorrhizal specificity and host genomics in Platanthera orchids across Eastern North America	Dr. Ida Hartvig	
S26 Cono	flow to the rescue: Assessing the need, effectiveness, risks and ethical implications of manipula	ating gone flow to improve adaptation to	
climate cha		ating gene flow to improve adaptation to	
P02.156	Evolutionary potential in a plant at the brink of extinction: Centaurea Corymbosa	Dr. Sudeeptha Yainna	
P02.157	Environment and forest management: How forest context drives gene flow and demographic dynamics in the rainforest timber tree Dicorynia guianensis	Mr. Julien Bonnier	
P02.158	Pervasive and recurrent hybridisation mitigates inbreeding in Europe's most threatened seabirds	Mr. Guillem Izquierdo Arànega	
P02.159	Impact of Habitat Fragmentation on Genetic Health in European Hamster Populations: A Microsatellite and Genomic Analysis	Ms. Juan Manuel Ortega	
P02.160	Population cross-breeding improves adaptive potential in the sand lizard (Lacerta agilis)	Dr. Seraina E. Bracamonte	
P02.161	Using novel selection strategies to bolster climate resilience in restoration planting stock	Mr. Daniel Krumm	
P02.162	Harnessing Genetic Rescue: A Case for Marsh Fritillary Butterflies in a Fragmented Habitat	Dr. Aja Tengstedt	
P02.164	Genetic rescue from the brink of extinction: genomic consequences of a severe bottleneck and intense conservation management in the pink pigeon	Dr. Carolina Pacheco	
P02.165	Soil characteristics shape optimal migration scenarios for assisted gene flow of crops vulnerable to climate change	Dr. Margret Veltman	
P02.167	Hybrid regeneration dynamics in a mixed plantation of interfertile species	Ms. Caroline SCOTTI-SAINTAGNE	
S28 - Genome Architecture and Their Role in Evolution			
P02.168	Repeat Driven Biases in Nanopore Sequencing Based Structural Variant Detection Reveal Clues to Chromosomal Rearrangements Linked to Infertility	Mrs. Kammy Howe	



		V**
P02.169	Sex-specific chromosome number and sex-determining mechanisms in the rhizocephalan Peltogasterella gracilis (Cirripedia: Rhizocephala: Peltogasterellidae)	Dr. ASAMI KAJIMOTO
P02.170	Telomere Position Effect and the Evolutionary Dynamics of Gene Expression Regulation	Prof. Wanil Kim
P02.171	Refining Sex Chromosome Strata Annotation Using X-Y Allelic Topologies	Ms. Suzie Tallon
P02.172	A high-quality reference genome for the Ural Owl (Strix uralensis) enables investigations of	Mr. Ioannis Chrysostomakis
. 02.272	cell cultures as a genomic resource for endangered species.	viii rodiiiis ciii ysostoiiidkis
P02.173	Genome defence and expansion – Repeat-Induced Point mutations shaping genome-wide TE activity in a ubiquitous ectomycorrhizal fungus	Dr. Benjamin Dauphin
P02.175	Supergene polymorphism and its role in male morph differentiation in the bulb mite,	Dr. Sebastian Chmielewski
102.173	Rhizoglyphus robini.	Dr. Sebastian emmelewski
P02.176	The effect of chromosomal inversions on gene expression in Mimulus guttatus	Dr. Paris Veltsos
P02.177	Impact of gene family expansions, gene flow and a recent TE burst on the evolution of a	Dr. Julia Schwarzer
FUZ.177	unique reproductive strategy in fishes	Dr. Julia Scriwarzer
DO2 170		Dr. Chuya Kata
P02.178	Genomic architecture imposes common constraints on hybrid genome composition across	Dr. Shuya Kato
D02.470	replicated hybridization events in an intertidal goby	
P02.179	Mutating the mutators: mutational processes acting on transposable elements	Ms. Svitlana Sushko
P02.180	A novel genomic structure organising the centromeres of the haplochromine cichlid	Mr. Pío Sierra
	radiation	
P02.181	Inversions Associated with Local Genomic Structure in Culex pipiens Mosquitoes from	Mrs. Sonia Cebrián-Camisón
	Southern Spain	
P02.183	Chromosomal rearrangements and their role in rapid sex chromosome turnovers in African	Ms. Sophie Smith
	cichlid fishes	
P02.185	Investigating the role of indels in allele specific expression in insects	Ms. Frances Swift
P02.186	Genome evolution in a karyotypically diverse primate group and the role of chromosomal	Mr. Axel Jensen
	fissions as reproductive barriers	
P02.187	Genome Structure Evolution in Ithomiini Butterflies	Dr. Karin Näsvall
P02.188	The understanding of structural complexity requires evolutionary context: Lessons from	Dr. Magdalena Migalska
	MHC architecture in Salamanders	
P02.189	Characterizing sex-chromosomes and the sex-linked region in Wikstroemia sandwicensis	Ms. Lila Maladesky
	from Hawai'i island	
P02.190	Genomic architecture of the self-incompatibility locus in the apple tree provides insights	Dr. Aureli Mesnil
	into the evolution of collaborative non-self-recognition	
P02.191	Genomic Architecture of the Major Histocompatibility Complex Genes in the Neotropical	Dr. Marta Barluenga
	Midas Cichlid	
P02.192	Unravelling Sex Chromosome Evolution and Dosage Compensation Mechanisms in	Mr. Vincent Bett
	Cameraria ohridella (horse-chestnut leaf miner moth)	
P02.193	Sloths Harbor the Highest Retrocopy Abundance Across Mammalian Genomes	Prof. Marcela Uliano Da Silva
P02.194	Reconstruction of the human amylase locus reveals ancient duplications seeding modern-	Dr. Charikleia Karageorgiou
	day variation	
P02.195	Understanding the role and evolutionary dynamics of "dark matter of the genome" of the	Dr. Eva Šatović Vukšić
	economically important beetle Tenebrio molitor	
P02.197	Genome wide identification of polymorphic inversions in a wild bird population and their	Mr. Hugo Corval
	possible phenotypic and fitness consequences	-
P02.198	The genomic landscape of speciation in an Amazonian songbird	Ms. Else Mikkelsen
P02.199	Comparative genomics illuminates karyotype and sex chromosome evolution of sharks and	Ms. Jiahong Wu
	rays	-
P02.200	Stepwise recombination suppression and degeneration in the mating-type chromosomes of	Dr. Loreleï Boyer
	two anther-smut fungi	,
P02.201	Genome evolution in holocentric organisms: insights from reconstructing the ancestral	Mr. James McCulloch
	linkage groups of sedges (Cyperaceae)	
P02.202	Layers of Adaptation: Transcriptomic divergence in the mantle tissue of Littorina saxatilis	Ms. Gabriella Malmqvist
		4
P02.203	Divergent sex chromosome systems in Australian marsupials – implications for evolution	Dr. Laia Marín-Gual
. 02.200	2.10. School of the months of	211 2010 11101111 0001
P02.204	Bringing light into the enigmatic genome architecture of B chromosomes: germ line	Ms. Agda Maria Bernegossi
. 52.207	dynamics in Neotropical deer	1500 Maria Delliegossi
P02.205	Evolution of Gene Regulatory Networks under Negative Frequency-Dependent Selection	Mr. Ritam Das
. 52.203	2.5.4.5 or defic hegalatory rections and contesting requestly dependent delection	Main Dus
P02.206	From Genome to Chromosomes: Satellite DNA Mapping and Comparative Cytogenomics in	Ms. Jennifer Pompeo
. 52.200	Engystomops pustulosus (Anura, Leptodactylidae)	Tompeo
P02.207	The rise and fall of insecticide resistance: the role of genetic architecture	Dr. Aude Caizergues
1 02.207	The rise and rail of insectione resistance, the role of genetic architecture	Dr. Adde Calzergues



P02.209	Microsynteny and functional relatedness: insights into eukaryotic genome evolution	Ms. Silvia Prieto-Baños
P02.210	Chromosomal structure and parallel evolution in the largest known avian sex chromosomes	Mr. Simon Jacobsen Ellerstrand
P02.211	Genome assembly and annotation of the parasitoid jewel wasp Nasonia oneida, with insights into the genomic evolution of Nasonia species	Ms. Anjali Rana
P02.212	Chromosome-Level Genome Assemblies Reveal Sex Chromosome Dynamics in Rapidly Evolving Spined Loaches (Cobitis)	Ms. Lucija Andjel
P02.213	The role of gene duplications in speciation of Lake Baikal Amphipods	Mr. Iakov Korobitsyn
S29 - Genor	nic Basis of Evolutionary Innovations	
P02.214	Comparative genome analysis of giant clams and non-photosymbiotic mollusks	Ms. Gabriella Juliane Maala
P02.216	Host use drives convergent evolution in clownfish	Dr. Théo Gaboriau
P02.217	Lineage-specific diversification of AluSg and AluSx3-derived alternative splicing events of exonization in RPS29 gene during primate evolution	Ms. Min-gyeong Ko
P02.218	The Emergence of Diatom Heterotrophy on Brown Algal Polysaccharides through Horizontal Gene Transfer, Gene Duplication and Neofunctionalization	Mr. Zeng Hao Lim
P02.219	De novo birth of transmembrane yeast proteins from intergenic polyA/T tracts	Dr. Nikolaos Vakirlis
P02.220	Investigating the Cellular and Molecular Basis of Regeneration	Mrs. Irene Del Olmo
P02.221	The expanded Bostrychia moritziana genome unveils evolution in the most diverse and	Mrs. Romy Petroll
	complex order of red algae	,
P02.222	A High-Quality Reference Genome Assembly of Oophaga pumilio, the Strawberry Poison Frog	Ms. Paula Mora Rojas
P02.223	The genomic basis of Lepidoptera diversity and evolution	Dr. Peter Mulhair
P02.224	Multiomics reveal associations between CpG methylation, histone modifications and	Mrs. Zoe Marie Länger
	transcription in a species that has lost DNMT3, the Colorado potato beetle	ŭ
P02.225	Mapping the shape of the helmet onto the genomic phylogeny reveals that this trait is	Dr. Marisol De La Mora Curiel
	phylogenetically conserved across the tribes of treehoppers (Hemiptera: Membracidae).	
P02.226	Stingless and Superorganismal: Multi-omic analysis reveals population differentiation and	Dr. Benjamin Taylor
	signatures of social evolution in Tetragonula bees	- ,-
P02.227	The genomic and developmental basis of an exaggerated sexual signal in the stalk-eyed fly Teleopsis dalmanni	Dr. Victoria Lloyd
P02.228	Parallel Ranges, Divergent Histories: A Comparative Genomic Study of Two Brassicaceae Species	Dr. Rachita Srivastava
P02.229	Evolution of chemosensory genes in the Chagas disease vectors of the genus Rhodnius	Dr. Marie Merle
P02.230	Exploring the genetic signatures of convergent evolution in termitophilous rove beetles	Ms. Caroline Høegh-guldberg
P02.231	Testis Transcriptome Evolution in the Absence of Sperm Competition: Insights from Pregnant Males	Dr. Bernadette Johnson
P02.232	Comparative Genomics of Syngnathiform Fishes: Linking Genome Evolution to Ecological and Phenotypic Diversity	Ms. Jule Drewalowski
P02.233	Support for the ovarian ground plan hypothesis in the evolution of eusociality from transcriptome analyses of halictid bees	Ms. Mozhgan Khodadadi
P02.234	Incomplete Lineage Sorting and Its Role in the Recurrent Evolution of Parthenogenesis in Stick Insects	Mr. João Souto
P02.235	Multiple Origins, Singular Success: Genomic Evidence for Recurrent Hybridization in Chenopodium album s. str.	Dr. Farzaneh Habibi
P02.236	Ecological divergence and retinal specialization in crater lake cichlids: a multiomic view of early sensory adaptation	Dr. Pau Balart i García
P02.237	Genomic mechanisms and evolution of a novel hemolymph neurotoxin in the Colorado Potato Beetle	Ms. Himrekha Agarwal
P02.238	Milk transcriptomes provide insights into the evolution of pinniped lactation traits	Ms. Jeni Sidwell
P02.239	Waddington revisited: Standing genetic variation drives the evolution of acquired traits	Dr. Gonzalo Sabarís
P02.240	Modeling Sex-biased Gene Expression Evolution Across Firefly Phylogeny	Mr. Haoqing Du
P02.241	Genetic parallelism underpins convergent mimicry coloration across Lepidoptera	Dr. Kanchon Dasmahapatra
P02.242 P02.243	Multivalent Mysteries: How Termite Chromosome Chains Shape Evolution The transcriptomic basis of phenotypic diversity in stick insects	Mrs. Mara Jean Julseth Dr Iulia Darolti



P02.244	Conserved and Lineage-Specific Gene Regulation Underlie Daphnia's Fish Predation	Mr. Aaron Kiggen
	Responses: A Cross-Species Meta-Analysis	
P02.245	Recurrent genomic dynamics linked to parallel evolution of secondary phytophagy in	Ms. Ronja Reinisch
	Hymenoptera	
P02.247	The Genomic Basis of Convergent Fusiform Body Evolution in Lake Tanganyika Cichlid Fishes	Dr. Emanuell Duarte-ribeiro
P02.251	ANTSR is an ancient sex-determining locus in bees	Dr. Tilman Rönneburg
P02.252	Assessing the phylogenetic position of sun spiders within chelicerates using complete	Ms. Inés Galán-Luque
	genomes	
P02.253	Convergent Recruitments of Venom Genes in Rear-Fanged Snakes	Dr. Inacio Azevedo
P02.256	Decoding the evolutionary developmental genetic basis of novel animal signal traits	Ms. Jae Walker
P02.257	Sex-biased gene expression evolution in extreme sexually dimorphic fireflies	Dr. Nicolás Lichilín
P02.258	Recently evolved, stage-specific genes are enriched at life stage transitions in flies	Mr. Andreas Remmel
P02.259	Transposable elements contribute to Anopheles coluzzii and An. gambiae adaptation to	Dr. Marta Coronado-Zamora
	urban environments	
P02.261	"Tinkering loci" rapidly generate new genes in Drosophila	Dr. Caroline Weisman
S32 - Lettir	ng go: reductive evolution across the tree of life	
P02.263	Neural and Molecular Signatures of Olfactory Specialization in Ant Social Parasites	Mr. Amirhossein Karimizadeh
P02.264	Increasing mutation rates and population sizes both lead to a reduction in the genome, but	Mrs. Juliette Luiselli
	not in the same genome compartments	
P02.265	Illuminating the mystery of thylacine extinction: a role for relaxed selection and gene loss	Mr. Buddhabhushan Salve
P02.266	Tracing genome reduction among prokaryotic lineages to reveal shared patterns and	Dr. Fabian van Beveren
	drivers of gene loss	
P02.268	Molecular basis of convergent evolution: parallel reduction of the sporophyte phase in	Mr. Huaxing Huang
	mosses	
S36 - Micro	pevolutionary processes and Macroevolutionary patterns	
S36 - Micro P02.269	vevolutionary processes and Macroevolutionary patterns  Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858):	Dra. Kelly Souza
	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858):	Dra. Kelly Souza
P02.269	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America	·
	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858):	Dra. Kelly Souza Mr. Jules Ferreira
P02.269	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae)	Mr. Jules Ferreira
P02.269 P02.270	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution	·
P02.269 P02.270 P02.271	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution	Mr. Jules Ferreira Mr. Oscar García Miranda
P02.269 P02.270	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution	Mr. Jules Ferreira
P02.269 P02.270 P02.271 P02.272	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae	Mr. Jules Ferreira Mr. Oscar García Miranda Dr. mariella Baratti
P02.269 P02.270 P02.271	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species	Mr. Jules Ferreira Mr. Oscar García Miranda
P02.269 P02.270 P02.271 P02.272 P02.273	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill
P02.269 P02.270 P02.271 P02.272	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel	Mr. Jules Ferreira Mr. Oscar García Miranda Dr. mariella Baratti
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez
P02.269 P02.270 P02.271 P02.272 P02.273	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly,	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and morphological analyses of the Korean dark chub (Nipponocypris koreanus) compared with	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282 P02.283	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and morphological analyses of the Korean dark chub (Nipponocypris koreanus) compared with its close relative (Nipponocypris temminckii)	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang  Ms. Yu Rim Kim
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and morphological analyses of the Korean dark chub (Nipponocypris koreanus) compared with its close relative (Nipponocypris temminckii) Life-history trade-offs and lifetime reproductive success under high vs. low food availability	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang  Ms. Yu Rim Kim
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282 P02.283	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and morphological analyses of the Korean dark chub (Nipponocypris koreanus) compared with its close relative (Nipponocypris temminckii) Life-history trade-offs and lifetime reproductive success under high vs. low food availability in a spider	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang  Ms. Yu Rim Kim
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282 P02.283 P02.283	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and morphological analyses of the Korean dark chub (Nipponocypris koreanus) compared with its close relative (Nipponocypris temminckii) Life-history trade-offs and lifetime reproductive success under high vs. low food availability in a spider The role of temperature and life cycle variation in Caudata diversification	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang  Ms. Yu Rim Kim  Prof. Simona Kralj-Fišer  Dr. Vivien Louppe
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282 P02.283 P02.283	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and morphological analyses of the Korean dark chub (Nipponocypris koreanus) compared with its close relative (Nipponocypris temminckii) Life-history trade-offs and lifetime reproductive success under high vs. low food availability in a spider The role of temperature and life cycle variation in Caudata diversification Developing an experimental toolbox for entomopathogenic nematodes	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang  Ms. Yu Rim Kim  Prof. Simona Kralj-Fišer  Dr. Vivien Louppe  Dr. Janneke Wit
P02.269 P02.270 P02.271 P02.272 P02.273 P02.274 P02.275 P02.276 P02.278 P02.279 P02.282 P02.283 P02.283	Landscape influences on gene flow in Cochliomyia hominivorax (Coquerel, 1858): implications for population connectivity in South America Fossil Placement in Dated Phylogenies: A Case Study on Bark Beetles (Coleoptera: Curculionidae) Sex roles, breeding systems and sexual dimorphism: combining studies of microevolution and macroevolution Illuminating firefly diversity: molecular taxonomy and conservation of Italian Lampyridae  Eco-evolutionary processes can explain morphological similarities in cryptic species complexes Phylogeography and Emergence of the Southernmost Orthohantavirus Lineage: A Novel Clade in the Magellanic Rainforests of Chile Why do kangaroos hop? Inferring ancestral foraging and locomotion traits to trace the evolution of hopping among marsupials Extreme Morphological Divergence Without Visual Compromise: Conserved Visual Field Geometry in Sexually Dimorphic Spiders From microevolution to macroevolution: sexual selection in plants shapes speciation in a 1000-species phylogeny of Brassicaceae Experimental evolution of metabolic fluxes in a nutritionally challenging context  Population genomic structure and geographical differentiation of a spiny crawler mayfly, Drunella ishiyamana, in erosive and flashy streams in Mongolia Cryptic species complex in a Korean endemic freshwater fish: genetic structure and morphological analyses of the Korean dark chub (Nipponocypris koreanus) compared with its close relative (Nipponocypris temminckii) Life-history trade-offs and lifetime reproductive success under high vs. low food availability in a spider The role of temperature and life cycle variation in Caudata diversification	Mr. Jules Ferreira  Mr. Oscar García Miranda  Dr. mariella Baratti  Dr. Abigail Cahill  Prof. Fernando Torres-Pérez  Mr. Lucas Le Gall  Mr. Atal Pande  Mr. Mohammadjavad Haghighatnia  Dr. Samuel Charberet  Dr. Ji Hyoun Kang  Ms. Yu Rim Kim  Prof. Simona Kralj-Fišer  Dr. Vivien Louppe  Dr. Janneke Wit



P02.289	Detecting evolutionary shifts in primates through model-based comparative analyses.	Dr. Fabio Barteri
P02.290	Non-random loss of ecological roles following extinction and extirpation on island bird communities	Ms. Natàlia Martínez-Rubio
P02.291	Evolution of nest architecture in tyrant flycatchers and allies	Mr. David Ocampo
P02.292	Disentangling effective population size and mutation rate: amphioxus at the edge of panmixia	Dr. Diego Hartasánchez
P02.294	Comparative establishment of species barriers in plants and animals by measuring introgression among species pairs from phylogenies	Mr. Arthur Boddaert
P02.296	Independent paths, shared landscapes: morphological and lineage diversification in Gymnophthalmidae (Reptilia: Squamata)	Mr. Juan Daniel Vasquez-restrepo
P02.297	The Evolutionary Dynamics of Transposable Elements in the Bengal Slow Loris (Nycticebus bengalensis)	Mr. Charles Michie
P02.298	Using contact zones to inform species delimitation in the gray zone of speciation: delimitation of species within the Acanthodactylus erythrurus complex	Ms. Carolina Farhat
P02.299	Scaling the Tree: How Clade Age and Size Govern Species Richness Gradients	Dr. Antonin Machac
P02.300	Adaptation to whole genome duplication across scales : convergent evolution among Angiosperm plant species	Dr. Violette Doublet
P02.302	Using Morphological Traits for the Species Delimitation under a Polymorphism- Aware Phylogenetic Models approach	Mr. Omar Ruelas
P02.303	Effects of cycles of range expansion and contraction on macroevolutionary patterns	Ms. Izabel Salvi
P02.304	Transposable Element Diversity in The Mountain Chicken Frog (Leptodactylus fallax)	Ms. Hayley Free
P02.305	Repetitive element density in micro and macro avian chromosomes is shaped by the interplay of 3D architecture constraints and recombination rate	Dr. Valentina Peona
P02.310	Assessing the population history of crop infecting pathogenic fungi and their relatives infecting wild grasses	Dr. Lucas Bonometti
P02.311	Phylogenetic Oscillating Ornstein-Uhlenbeck process: analysis and application.	Mr. Bayu Brahmantio
P02.312	Phylogenetic inference from an incomplete fossil record	Mr. Niklas Hohmann
P02.314	A Ghost in the Stream: Mitochondrial Evidence for a Cryptic Lamprey Lineage in the Upper Po Valley	Ms. Elena Catelan-Carphio
P02.315	Integration Constrains Visual System Diversification in Ecologically Diverse Spiders	Mr. Atal Pande
P02.319	The speciation continuum in bloom: Incomplete lineage sorting, gene flow, and reticulate evolution in rapidly diverging plant lineages	Mrs. Luana Soares
P02.320	Multilayer thin film produces recurrent iridescence in mammals	Ms. Jessica Dobson
S38 - New	Frontiers in Genome Diversity and Evolution: Exploring the 3D Organization and Function of Genc	omes
P02.322	Comparative Genomics of Araneoidea Spiders reveal insights into their 90-million-year evolutionary history.	Ms. Caitlin Price
P02.323	3D Genome Constrains Breakpoints of Inversions that Can Act as Barriers to Gene Flow in the Stickleback	Dr. Yo Yamasaki
P02.324	The structure of mitochondrial genomes across the plant tree of life	Dr. Max Carter-Brown
P02.325	Ancestral origin and structural characteristics of non-syntenic homologous chromosomes in abalones	Dr. Shotaro Hirase
P02.326	Divergent patterns of 3D genome organization across the Animal Tree of Life	Ms. Gala Pujol
P02.327	Understanding extreme heterochiasmy: cellular and genomic insights in frog systems	Ms. Ezgi Unal
P02.328	On the genomic underpinnings of speciation in holocentric sedges	Dr. Ashwini V Mohan
P02.329	Interspecies Functional Equivalence of Pancreatic Enhancers: Zebrafish as a Model to Study the Genetic Basis of Type 2 Diabetes	Mr. Diogo Pimenta-Sousa
P02.330	Dynamic reorganization and evolution of three-dimensional genome architecture during the species diversification of Populus	Prof. Jing Wang
P02.331	Patterns of Genome Reshuffling and 3D Chromatin Architecture in Turtles	EM. Laura González-Rodelas
P02.332	Long-term inversion recurrence and segmental duplication conservation in mammal genomes	Mrs. Maria Diaz Ros
P02.335	Unraveling the Phenotypic Evolution and Global Breeding History of Domestic Chickens	Dr. Cheng Ma



P02.337	Beyond vertical inheritance: the dynamic evolution of transposable elements in Drosophila	Mr. Riccardo Pianezza
P02.338 P02.339	Hybrid Speciation? A Case Study in Grasshopper Species Phylogeographic signatures of lineage fusion: a simulation study assessing the incorporation of geo-spatial information, and long-read nuclear haplotype data	Ms. Dörte S. Neumeister Dr. Ryan Garrick
	cological and evolutionary implications of climate change on reproduction	
P02.340	To self or to clone? Mechanisms driving reproductive assurance strategies across wild strawberry genotypes.	Dr. Carolina Diller
P02.341	Phenotypic Selection on Reproductive Traits Under Autonomous Self-Fertilization in Fragaria vesca	Dr. Sebastian Arenas
P02.342	Thermal genotype-by-environment interactions and sexual selection in Drosophila melanogaster	Mr. Ignacio López Pardo
P02.343	Stochastic reproductive plasticity in Artemia brine shrimps: bet-hedging against unpredictable environmental change?	Dr. Clementine Lasne
P02.344	What doesn't kill me makes me stronger: How naturally occurring insecticide resistance alleles lead to broader colonising opportunities in D. melanogaster	Dr. Andre Nogueira Alves
P02.345	Stage-Specific Nutritional Environments Drive Plastic and Evolutionary Shifts in Reproduction	Dr. Sudipta Tung
P02.346	Early-life social stress rewrites strategic seminal fluid investment in Drosophila males	Dr. Juliano Morimoto
P02.347	Predicting reproductive fitness under climate change: developmental effects of nutritional and thermal stress in locally-adapted populations of Drosophila melanogaster	Dr. Brooke Zanco
P02.348	Birds in the Grey Zone: High-Resolution Satellite Imagery Reveals Continental Patterns in Urban Great Tits (Parus major)	Mr. Nicolas Bekka
P02.349	Breeding phenology drives variation in reproductive output, reproductive costs and offspring fitness in a viviparous ectotherm	Dr. Théo Bodineau
P02.350	Phenological fluctuations but no advance: exploring the drivers and consequences of breeding phenology in the Soay sheep	Ms. Ellis Wiersma
P02.351	Hybridisation and Climate Change in Alpine Orchids	Ms. Hannah Gunn
P02.352	Loss of Photoperiodic Regulation of a Key Pleiotropic Gene Underlying the Evolutionary Diversification of Reproductive Seasonality in Sticklebacks	Prof. Asano Ishikawa
P02.353	ShareTrait: towards interoperable and reusable trait data in Ectotherms	Mr. Raimon Cuxart-Erruz
P02.354	Dark waters, strong beginnings: investigating maternal and environmental influences on early-life traits in Eurasian perch (Perca fluviatilis)	Mrs. Meruyert Zhumasseit
P02.355	Dynamics of Mutation Load in the Antarctic Fur Seal (Arctocephalus gazella)	Dr. Kosmas Hench
P02.356	Couch potato chipmunks: negative effects of supplemental feeding on breeding activity in eastern chipmunks	Prof. Patrick Bergeron
P02.357	Reproductive plasticity in response to short-term polystyrene nanoplastic exposure in aquatic ecosystems	Ms. Gala Pujol
P02.358	Resource availability as a key environmental constraint on maturation in the ectothermic vertebrate rainbow trout (Oncorhynchus mykiss)	Dr. Ana Lindeza
P02.361	Does temperature affect the adaptive value of male harm in Drosophila melanogaster?	Ms. Soumya Panyam
P02.362	How does turbidity and male harassment mediate the female reproductive behaviour and stress?	Mrs. Md Mahmud Al Hasan
P02.363	Impacts of thermal and salinity stress on flatworm reproduction: sex-specificity, genetic variation and mechanisms	Dr. Steven Ramm
P02.364	Comparison of reproduction-related parameters between free-ranging sex-reversed and sex-concordant agile frog males	Dr. Edina Nemesházi
P02.366	Is the incubation stage a bottleneck for avian fitness in a warming world? Evidence from an experimental study on collared flycatchers.	Ms. Karolina Skorb
P02.367	Masting modeling: evolution towards an early phenology to control seed consumers.	Dr. Sabrina Gastebois
P02.368	Environmental O₂ and the C. elegans germline	Ms. Emma Haxen
P02.369	Evolution of phenotypic plasticity during environmental fluctuations	Dr. Martin Lind
P02.370	Genomic signatures of reproductive fitness in the King penguin across two decades of climate change	Dr. Josephine Paris
P02.371	How thermal stress alters reproductive traits and foraging behavior in Trinidadian Guppies	Ms. Stephanie Tran
P02.372 P02.373	Mitonuclear interactions constrain thermal responses Environmental drivers of parental care diversity in mammals	Dr. Florencia Camus Mrs. Naomi Narzissenfeld



		·
S47 - The in	terplay between genetic architecture and the evolution of biodiversity	
P02.035	Dissecting genes of color pattern in birds and their role in maintaining species boundaries in a moving hybrid zone	Mr. Louis Hausner
P02.184	Sexually antagonism and the mating system determine the rate and distribution of recombination suppression across sex chromosomes	Dr. Ewan Flintham
P02.196	Genomic impact of asexuality in oribatid mites	Mr. Karim Gueddach
P02.130	·	
PU2.215	Evolution and divergent genetic mechanisms of annual and perennial Hordeum species	Dr. Timo Hellwig
P02.246	Constituted and malecular discostion of hotographory in Colonosco	Dr. Miguel Canta Deminas
	Genetic and molecular dissection of heteranthery in Solanaceae	Dr. Miguel Santo Domingo
P02.248	Gene copy number variations in Brassicaceae	Ms. Freja Lindstedt
P02.249	Distinct dynamics in a key herbivore detoxification gene family reveal mechanisms of convergent host plant shifts	Ms. Paula Fernandez Begne
P02.254	Population genomics and structural variation underlying local adaptation to elevational gradient in a bromeliad from the Brazilian Atlantic Forest	Mr. Paulo Aecyo
P02.255	Uncovering the Genomic Architecture of Tooth Development Across a Dietary Radiation of Bats	Dr. Ariadna Morales
P02.260	Ecological shifts and genomic architecture divergence drive sexually selected weapon size	Dr. Camille Thomas-Bulle
102.200	reduction and reproductive isolation in island populations of rhinoceros beetles	Dr. carrine mornas bunc
P02.262	Squalomix: genomic exploration of shark and ray evolution	Dr. Shigehiro Kuraku
P02.267	Genome Size: Did Flight Loss Relax Genome Size Constraints in Birds?	Ms. Zeynep Oguzhan
P02.281	Evolving inversions: the genomic architecture of parallel snail ecotypes in Sweden and	Mr. Basile Pajot
	France	
P02.284	Genome Architecture-Aware Investigation of Earthworm Terrestrialization Amidst Extensive Genomic Reshaping, Rearrangements and Polyploidy	Dr. Dearbhaile Casey
P02.295	Comparative repeatome evolution in diploid and allotetraploid Trifolium (clover)	Ms. Katie Herron
. 02.255	Comparation operations of order or an appear and another appear a fine and (correct)	
P02.301	Avian structural coloration tuning explained by changes in feather keratinization	Ms. Rita Afonso
P02.306	Evolutionary analysis of diet adaptations in anatomically modern humans using ancient	Ms. Carla Casanova Suarez
	genomes	
P02.307	Modeling the co-evolution of genetic individual condition and condition-dependent traits	Ms. Julie Roux
	involved in sexual selection	
P02.308	Quantifying structural variants in chromosomes using landmark-based disparity	Dr. Ashwini V Mohan
P02.313	A single theory for the evolution of sex chromosomes and the two rules of speciation	Dr. Thomas Lenormand
P02.317	Changes in the cell migratory environment as a driver of irregular pigment cell migration in	Ms. Olivia Dreilinger
DO2 240	the blotch color polymorphism of African cichlid fishes	Du Caulinath Chattanadh
P02.318	An empirical fitness landscape of gene regulation in E.coli	Dr. Gopinath Chattopadhyay
P02.321	Understanding the costs and benefits of pleiotropy for adaptive evolution in an in silico	Dr. Théotime Grohens
D02 222	model of serial organ evolution	De Althorfor Alocat Koldala
P02.333	A moving avian hybrid zone provides insights into how the genomic architecture of traits	Dr. Niloofar Alaei Kakhki
DO2 224	under natural and sexual selection shapes speciation	Du Katia Haadiaa
P02.334	Distinct functional effects of individual nucleotide variants in Eip75B on life history	Dr. Katja Hoedjes
DO2 226	adaptation in the fruit fly	Dr. Hannah Weller
P02.336	Self-organization on a tight leash? Identifying the mechanisms of phenotypic robustness	Dr. Hannan Weiler
DO2 250	using fluctuating asymmetry in cichlid fish color patterns	Dr. Freih Helen
P02.359	Ancestral Hybridisation Events in Snapdragons	Dr. Emily Haley
P02.360	Genomic drivers of persimmon adaptive radiation in New Caledonia, a biodiversity hotspot	Mr. Amin Ghane
DO2 265	The good, the had as the leafe 2 Hay TE driven he had degeneric interests with a globally	Dr. Jonathan Wilson
P02.365	The good, the bad or the lucky? How TE-driven hybrid dysgenesis interacts with a globally	Dr. Jonathan Wilson
DO2 274	adaptive insecticide resistance locus	Ma Changui Vian
P02.374	Haplotype structure – an overlooked key factor shaping the genomic selection response	Mr. Changyi Xiao
DO2 275	Diet qualities and concession adoptations to avera facilities to the best facilities to the	Dr. Vuoling Vi
P02.375	Diet evolution and genomic adaptations to sugar-feeding in the bat family Phyllostomidae	Dr. Xueling Yi
D02 276	technical collection of the control	May Variable Of
P02.376	Linked selection of insertions and deletions in coding regions of the great tit genome	Ms. Yu-chi Chen
DO2 277	Fundation of Tunner could be Flowerste in the Countrie Countries of Learning and Countries of Tunner count	Du Malavi Canqueta
P02.377	Evolution of Transposable Elements in the Swedish Sand Lizard, Lacerta agilis	Dr. Malavi Sengupta
P02.378	Genomic bases of a pollinator-driven speciation in sexually deceptive orchids of the Ophrys	ivirs. Pascaline Salvado
	insectifera clade	

P02.379	Identification of genes associated with domestication by using three different statistics	Mr. Diego Alonso Vargas Donayre
P02.380	The complex genomic landscape of inversion polymorphisms in the spruce bark beetle: genomic patterns, evolutionary forces, and implications for evolutionary inference	Dr. Krystyna Nadachowska-Brzyska
P02.381	Adapting complex phenotypes despites pleiotropy: the importance of regulatory networks structure	Dr. Maud Fagny
P02.382	Functional analysis of yellow gene's cis-regulation in Pieris canidia butterflies	Ms. Ler Shan Ang
P02.383	Translocations and climate glacial refugia shaped the genetic structure of the largest Italian wild ungulate, the red deer Cervus elaphus	Dr. mariella Baratti
P02.384	The genomic landscape of historical and ongoing gene flow in hybridising Geum (Rosaceae)	Dr. Meng Lu
P02.385 P02.386	The joint evolution of separate sexes and sexual dimorphism  Pangenomic variations and differentiation between the two African malaria vector sister	Dr. Thomas Lesaffre Prof. Michael Fontaine
	species Anopheles gambiae and Anopheles coluzzii	
S49 - Time	-dependency in micro- and macroevolutionary rates	
P02.040	Introducing the early high disparity phylogenetic comparative model, with applications to ichthyosaur macroevolution and implications of sedimentary sequence biases	Dr. Ricardo Ely
P02.049	Numerous karyotype rearrangements of diptera evolution on various timescales.	Dr. Kamil Jaron
P02.064	Multiple processes contribute to time-dependency in contemporary rates of genome-wide evolutionary change	Dr. Zachariah Gompert
P02.135	The origin and diversification of turmerics (Curcuma: Zingiberaceae) in paleotropical biodiversity hotspots: the role of ancient hybridization and historical climate change	Dr. Marcos V. Dantas-Queiroz
P02.138	Uncovering dynamic adaptive landscapes on a phylogeny using the Ornstein-Uhlenbeck model	Ms. Priscilla Lau
P02.144	Bioinformatic bias in pedigree-based mutation rate estimates: insights from sharks and whales	Ms. Nisha Dwivedi
P02.152	Conflicting Timelines: Exploring patterns of divergence discordance across data types among tetrapod groups	Mr. Praveen Karanth
P02.163	Speciation completion rates have limited impact on macroevolutionary diversification	Mr. Pierre Veron
P02.174	Temporal and non-temporal patterns in rates of diversification and phenotypic evolution in a hyperdiverse fish clade	Prof. Carmelo Fruciano
P02.182	Idiosyncrasies unveiled: examining the pace, patterns and predictors of biotic diversification in peninsular India	Mr. Pragyadeep Roy
S51 - Oper	n Symposium	
P02.005	Are fish on land to escape an intolerable aquatic environment?	Dr. Iker Irisarri
P02.006	New mutation, gene flow or plasticity? Using genomic variation to test the origins of seasonal allochrony in storm-petrels	Prof. Vicki Friesen
P02.031	Comparison of molecular diversity of spring fen macrozoobenthos from two geomorphological regions	Ms. Magdalena Gajdosova
P02.036	The consequences of constrained sex allocation under local mate competition	Mr. Chedhawat Chokechaipaisarn
P02.060	Saving the locals; conservation genomics for the Spanish Toothcarp (Aphanius iberus).	Ms. Maria Estarellas

## Thursday 21 August, EXPO AREA: 5:00 PM - 7:00 PM - POSTER SESSION 3 - with Aperitif

Multidimensional perspective of flower color polymorphism in Iris pumila

**Evolutionary Biology Meets Artistic Research Across Scales** 

P02.109

P02.138

S02 - Addr	essing new and long-standing evolutionary questions with linkage disequilibrium based approa	ches
P03.003	Identification of novel toxin resistance genes in Drosophila via Experimental Evolution,	Mr. Michele Marconcini
	GWAS, and CRISPR screening	
P03.004	Investigating the role of meiotic recombination in social insects - Termites adding an	Ms. Turid Everitt
	important piece to the puzzle	
P03.005	A Linkage-Based Method to Detect Introgression in Polyploid Genomes	Dr. Sergio Tusso

Mr. Ivan Perez

Dr. Nataša Barišić Klisarić



P03.006	Linkage disequilibrium based inference method reveals stable recombination hotspots in the house sparrow	Dr. Marie Raynaud	W.
P03.007	Ecological speciation in North Atlantic Redfish (Sebastes): Insights from whole-genome data	Dr. Danielle Davenport	
P03.008	Filtering linked selection to improve LD-based demographic inference in heavily exploited Baltic cod (Gadus morhua) populations	Ms. Lingfeng Meng	
P03.009	Inferring population size history from both modern and ancient genomes using approximate bayesian computation	Mr. Patrick Jacques	
P03.010	Demographic inference given rampant background selection under a high deleterious mutation rate	Ms. Micaila Marcelle	
P03.011	Estimating very recent population split times using decay of IBD (identity by descent) sharing	Prof. Jeff Wall	
P03.013	Historical divergence and secondary contact shape the distribution of genetic variation in a key glacial refugium	Dr. Giulia Gentile	
P03.014	Asymmetric gene flow and phenotypic clines across a European butterfly hybrid zone	Dr. Bruna Cama	
S06 - Cance	er in an evolutionary framework: across species and within individuals		
P03.016	Evolution of Mutational Susceptibility to Genetic Diseases	Ms. Maria Kelly	
		•	
P03.024	Using evolutionary accumulation models to identify therapeutic targets in cancer	Prof. Ramón Diaz-Uriarte	!
P03.182	The cross-talk between ER+ breast cancer cells during progression	Ms. Katarzyna Wierzbicka	a
P03.184	Intra-tumor evolution can favour inter-clonal cooperation and facilitate metastasis	Mrs. Caroline Carneiro	
P03.187	The role of the microbiome in transmissible cancers, a case study on Tasmanian devil facial tumour diseases	Mr. Jeremy Jeanjean	
P03.191	Organ-specific variability of tumor prevalence across wild mammals	Ms. Crystal Morin	
P03.196	Investigating the metastatic potential of two transmissible cancers in Tasmanian devils	Mrs. Florence Pirard	
P03.199	Preventing evolutionary rescue in cancer using two-strike therapy	Dr. Robert Noble	
P03.202	No evidence that oestrogen positive breast cancer risk is genetically correlated with human	Prof. Hannan Dugdale	
P03.202	life-history traits Pan-Mammalian genomic insights into cancer resistance and therapeutic target discovery	Ms. Giada Padovani	
P03.204	When rest fails, tumors rise: sleep-like state deprivation reveals an ancestral cancer- protective function in Hydra	Mr. Jordan Meliani	
S08 - Coop	eration, Conflict and the Evolution of Socially Transferred Materials		
P03.140	Substrate hardness and social environment drive egg clustering behaviour in Drosophila melanogaster	Dr. Emily Fowler	
P03.146	The socio-sexual environment influences diet and oviposition site choice in female fruit flies	Dr. Mabel Sydney	
P03.150	Fecundity help may be more common than survival help in realistic ecological scenarios	Ms. Margaret Bolton	
P03.155	Studying the evolutionary impact of parental influence on mate choice in human societies	Mr. Kartikey Awasthi	
P03.157	Evolutionary Adaptation to Juvenile Malnutrition and Investment in Sexual Conflict	Dr. Berta Canal Domenec	ch
P03.159	Sexual Selection and the Evolution of the Mammalian Brain and Social Cognition: A Proposal on the Evolution of Selfishness and the 'Male Brain Meme' Concept	Dr. Michihiko Ito	
P03.162	The cooperation inside us: investigating bacterial cooperation in the human gut	Mrs. Zohar Katz	
P03.179	Reciprocal influence on viral load and effects on virulence in co-infections with latent ALSV and phylogenetically distinct viruses	Dr. Guillaume Lafforgue	
S09 - Crani	ofacial Evolution in Vertebrates		
P03.098	Evolutionary dynamics of gene and isoform regulation underlying jaw diversification in	Ms. Pooja Singh	
P03.102	parallel cichlid adaptive radiations  Heterochronic shift in gene expression contributed to the evolution of vertebrate sensory	Dr. Shigeru Sato	
	system	-	
P03.106	Exploring the genetic correlations of craniodental size dimensions in a pedigreed baboon colony	Mr. Mario Modesto Mata	a



P03.110	Applying ancient DNA extraction methods to improve the recovery of genomic material from modern bone samples	Dr. Alexandra Schuh
P03.115 P03.129	Biomechanical adaptation to an anthropogenic diet in a human-commensal bird  Ecomorphological diversification of Caviomorph rodents and the role of environmental	Ms. Ruth Fawthrop Ms. María Cristina Aparicio De Soto
P03.130	change as a driving force  A matched genome-phenome dataset of the world's wolves (Canis lupus) and dogs (Canis familiaris)	Dr. Sarah Du Plessis
P03.133	Evidence for tonotopic organization in zebrafish otolith organ: Implications for the evolution of vertebrate auditory systems	Dr. Masashi Tanimoto
P03.136	Parallel adaptation of tooth classes to dietary niches in Chiropterans	Ms. Fanny Gagliardi
S12 - Epige	netics and adaptation to global change: climate and biotic interactions	
P03.001	Identification of Fish Epigenetic Biomarkers of Temperature Response and parasite resistance shaped by Early Rearing Conditions	Dr. Sofia Consuegra
P03.002	Epigenetic response of mountain hare (Lepus timidus) in different climate zones	Ms. Lara Marinangeli
P03.012	DNA methylation footprint of fisheries in genetically homogeneous marine metapopulations	Dr. Margarida Barcelo-Serra
P03.015	The epigenetic dimension of effects of climate change on alpine plants	Mr. Florian Brück
P03.022	Experimental evidence for short term directional selection of epigenetic trait variation	Dr. Benoit Pujol
P03.033	Epigenomic profiles underlying induced defenses in a freshwater gastropod	Mr. Dominik Periša
P03.044	Developmental plasticities within and across generations in Austrolebias annual killifish	Dr. Tom JM Van Dooren
P03.045	Evolutionary Trade-offs Between Intergenerational and Transgenerational Fitness Effects	Mr. Isaac Harris
P03.054	Genetic and epigenetic responses of a keystone forest tree species to global climate change revealed through the integration of population-scale pangenomics and multi-omics analyses	Ms. Jiajun Feng
P03.059	Temperature dependent DNA methylation in the Swedish sand lizard (Lacerta agilis)	Mr. Joshua Hufton
P03.061	Epigenetic factors associated with diel vertical migration and diapause in the marine copepod Calanus finmarchicus.	Ms. Katarzyna Zejc
P03.078	Unveiling the genetic basis of floral seasonal phenotypic plasticity in the Moricandia genus	Dr. Saloni Sharma
P03.085	Exploring the link between epigenetics and deleterious mutations	Ms. Rebecca Chen
P03.089	Evolved transcriptional responses and their regulation after long-term adaptation of	Ms. Ella Tadmor
	Bemisia tabaci to a marginally-suitable host	
P03.354	Using patterns of DNA methylation to explore the environmental drivers of senescence in the wild wood mouse	Dr. Sarah Wolf
0.17 5 1		
	tion of behavioural diversity: from ecology to genes and neural systems	
P03.017	Testing the Darwin-Bateman Paradigm: Anisogamy as a Fundamental Driver of Sex Role Evolution in Birds	Mr. Oscar García Miranda
P03.020	On the evolution of sound-related sexual dimorphism in dung beetles	Dr. Angela Roggero
P03.021	To migrate or not to migrate? Exploring the genomic basis of the partial migratory behavior in bats	Dr. Diego Matías Peralta
P03.023	Genetic and Environmental Drivers of Multiple Cognitive Abilities in Great Tits Living Along an Altitudinal Gradient	Dr. Laura Gervais
P03.025	Warmer climate disrupts pace-of-life syndromes through ontogenic changes of activity levels in the common lizard Zootoca vivipara	Mr. Nicolas Mouret
P03.026	The effects of dietary innovation on metabolism during the evolution of cognitive enhancement in Heliconius butterflies.	Mr. Rami Kersh-Mellor
P03.027	Rapid color change in cichlid fishes: Using computer vision to study behavioral diversity and evolution	Mr. Jan Häge
P03.028	The role of social learning in speciation by sexual selection: the case of a poison frog	Mr. Marco González-Santoro
P03.029	Immunohistochemical mapping of diapause in the aphid brain	Mr. José Ricardo Morales Poole
P03.030	Transcriptomic mapping of diapause in the aphid brain	Mr. José Ricardo Morales Poole
P03.031	Nest weave pattern in weaverbirds: a sexual signal selected through sensory drive?	Dr. Erwan Harscouet
P03.032	Arms-race dynamics drive rapid behavioural coevolution in a parasitoid fly and its host in Hawaii.	Dr. Leeban Yusuf



P03.034	Adaptive plasticity in escape behaviour: Crickets don't run as fast as they can when facing predator threat	Ms. Ruonan Li
P03.035	ROLE OF NICHE CHOICE IN RAPID HOST ADAPTATION OF A GLOBAL PEST	Ms. Philippa Musiolik
P03.036	Sex differences in parental cognition and response to offspring signals in birds	Dr. Shana Caro
P03.037	Functional adaptation and evolution of mandibular tusks in burrowing mayflies (Insecta:	Prof. Yeon Jae Bae
	Ephemeroptera) explored via micro-CT and 3D morphometrics	
P03.038	Alpha and Beta Diversity of Cultural Traits in Global Jukebox Cantometrics Data	Mr. Jiří Nedomlel
P03.039	Facing hypoxia/reoxygenation stress in the nervous system of the Mediterranean mussel	Dr. Serena Mirra
	Mytilus galloprovincialis	
P03.040	Identification of novel miRNAs in the deep-sea cephalopod Pteroctopus tetracirrhus with	Dr. Tosca Van Gelderen
	potential to rescue mitochondrial dysfunction in human diseases.	
P03.041	Evolutionary origin and maintenance of distinct song types that may promote reproductive	Ms. Athena Syarifa
	isolation in the willow tit	
P03.042	The effects of migration-related genotype on swimming behavior in rainbow trout	Mr. Lilian Redon
	Oncorhynchus mykiss	
P03.043	Convergent patterns of visual adaptation in a community of mimetic butterflies	Dr. J Benito Wainwright
P03.046	Dissecting the genomics of migration in the Painted Lady butterfly	Ms. Aurora García-Berro
P03.047	Non-canonical organisation of the olfactory system in the ant Temnothorax longispinosus	Ms. Clarita Mendes
P03.048	Sex-specific voluntary fasting in the ontogeny of size dimorphism in snakes	Prof. Lukas Kratochvil
P03.049	Dynamic Genomes, Dynamic Behaviors: Multi-Omic Mapping of Predator-Induced Gene	Mr. Aaron Kiggen
	Regulatory Networks in Daphnia magna	
P03.050	The evolution of socially-mediated dispersal favours the emergence and maintenance of	Mrs. Iris Prigent
	distinct social and dispersal strategies	
P03.051	Comparing the sex-specific demographic history of primates with contrasting social systems	Dr. Léa Guyon
P03.052	Linking Changes in Pollinator Behaviour with Their Effect on Male Fitness in Buzz-Pollinated	Dr. Lilian Melo
D02.0E2	Flowers	De Claudia Casadhan
P03.053	Does it pay to be picky? How local adaptation in nesting behavior influences offspring	Dr. Claudia Crowther
P03.055	phenotypes in a reptile with environmental sex determination.	Ms. Katherine Henson
FU3.U33	Patterns and Colors Just Don't Mix: Plumage Color Evolution Trends in a Songbird Clade	ivis. Ratiferine Herison
S20 - Evolut	onary biology meets genetic pest control	
P03.033	Implications of intraspecific variation in genes determining host plant use in a generalist	Dr. Ernesto Villacis Perez
	herbivore	
S24 - Foreca	sting evolution in natural populations	
P03.056	Adaptive Evolution in Brassicaceae: Unraveling Mechanisms of Genomic Convergence in	Mr. Vít Bureš
	Serpentine Adaptation Across a Broad Range of Genetic Divergence	
P03.057	Fluctuation of population abundance over decade in one amphibian assemblage	Dr. Bogdan Jovanović
D02.050		
P03.058	Local Adaptation in Baltic Cod? Insights from Early Development	Ms. Maddi Garate Olaizola
P03.060	omniCADD: prediction variant deleteriousness beyond model organisms	Dr. Julia Höglund
P03.062	Where have all the flowers gone? What could we start learning about possible	Dr. Aleksej Tarasjev
	evolutionary patterns from multi-year study of population dynamic of sexual and	
	vegetative reproduction in clonal Iris pumila natural population in Deliblato Sand	
P03.063	Evolution of a marine invertebrate in urban coastal habitats	Mr. Alan Le Moan
P03.063	The role of developmental plasticity in the resilience of spionid populations	Mr. Lukas Edwards
P03.065	How the common evening brown butterfly conquered the world	Mr. Nhat Tan Pham
P03.065	The link between Doubly Uniparental Inheritance (DUI) of mtDNA and sex determination in	
. 03.000	marine mussels: a unique case of sexual system evolution from dioecy to trioecy	Diri abio Oyaizaii
S25 - Gene	Content Across Genomes: Models and Genomic Data	
P03.067	Evolutionary diversity of CXCL16-CXCR6: Convergent substitutions and recurrent gene loss	Mr. Buddhabhushan Salve
	in sauropsids	

A phylogeny aware analysis of function for the biodiversity genomics  $\mbox{\it era}$ 

Comparative genomics provides insights into early bryophyte evolution

P03.068

P03.069

Dr. Gaurav Diwan

Dr. Alexander Bowles

P03.070	Does Metabolite Toxicity Impact Gene Order in Metabolic Operons through Selection for Robust Gene Expression?	Mr. Quentin Fernandez De Grado
P03.071		Dr. Alexander Bowles
	Comparative genomics provides insights into early angiosperm evolution	
P03.072	Studying the evolution of gene repertoire in Fisher's Geometric Model	Prof. Guillaume Beslon
P03.073	A high-quality genome for Pagodroma nivea facilitates understanding the evolution of highly adapted 'tube-nosed' (Procellariiformes) seabirds	Dr. Anna Rix
P03.074	Exploring the transcriptional landscape of environmental samples using metatranscriptomics long reads	Mrs. Carmen LAFUENTE SANZ
P03.075	Convergent evolution of cosexuality in liverworts	Dr. Peter Szoevenyi
P03.076	OrthoFinder: scalable phylogenetic orthology inference for comparative genomics	Dr. Laurence Belcher
P03.077	Genome of the Amazon Guppy (Poecilia bifurca) reveals conservation of sex chromosomes and dosage compensation	Ms. Lydia Fong
P03.079	Antibiotic Resistance Prediction in Staphylococcus aureus Using Gene Content Outperforms SNV-Based Approaches	Ms. Bruna Fernanda Fistarol
DO2 000		Ms. Luana landira Duana Landau
P03.080	Evolution of amylase copy number variation in Nile rats and functional impact	Ms. Luane Jandira Bueno Landau
P03.081	Uniform annotations reveal the connections between genome size, repeat content and	Ms. Milena Trabert
	gene family evolution	
P03.082	Widespread intra- and inter-domain horizontal transfer and bursts of gene duplication	Ms. Saioa Manzano-morales
	shape the size and content of Asgard archaeal genomes	
P03.083	Lineage-specific selection and gene family dynamics in the genus Lynx	Ms. Lorena Lorenzo Fernández
P03.084	Thymelaeaceae genome annotations: where are we?	Dr. Nikos Tsardakas Renhuldt
P03.086	The genome distribution of structural mutations in wild strains of Neurospora crassa	Dr. April Snøfrid Kleppe
P03.087	Comparative analysis of Genes encoding Ribosomal Proteins across Metazoa	Ms. Megha Suresh
P03.088	Archaeorhizomycetes, a diverse group of ancient root endophytes	Mr. Kevin Nielsen
S30 - Geno	mic insights into evolutionary adaptation and species movements in a changing climate	
P03.090	Genomics of northern adaptation: Prevalence and mode of selection in northern	Dr. Jana M. Flury
	Arabidopsis lyrata populations in Eurasia	•
P03.091	Genome assembly of pulmonate snail Ellobium chinense illuminates genomic adaptation	Prof. Joong-Ki Park
. 00.001	and contrasting historical demography	Tron Joong Kirank
P03.092	Twenty years of monitoring and common garden experiment revealed a possible difference	Dr. Vu Endo
F03.032		DI. Ta Eliao
	in heat tolerance under climate changes and robust species barriers between sympatric	
D02 002	sticklebacks	May January Character
P03.093	Analysing the Evidence for a Seasonal Migration in Panacea prola Using Population Genomics	Ms. Jenny Stewart
P03.094	When Isolation echoes: Tracing Temporal Genetic Drift and the Loss of Genetic Diversity in	Mr. Alfonso Lopez
	Baetican Toothcarps (Aphanius baeticus Doadrio, Carmona & Fernández-Delgado, 2002)	·
P03.096	Genomic Legacy of the Ice Ages	Mr. José Costa
P03.097	Evolutionary history of the rapid post glacial diversification in the songbird genus Junco	Mr. Borja Mila
P03.204	Polar bear evolutionary adaptation: insights from non-coding accelerated regions	Ms. Giada Padovani
S33 - Linkir	ng recombination rates and supergene evolution with the genomics of complex traits	
P03.099	Palms, a new study system for sex chromosome evolution in plants	Mr. Hugo Tessarotto
P03.100	Species delimitation for a male-only clonal lineage of ants	Ms. Alice Ha
P03.101	Complex supergene system and fine-scale dispersal in a socially polymorphic ant	Mr. Joshua Ducancel
. 03.101	Complex supersence system and interstate dispersal in a socially polymorphic and	Will Joshua Ducancei
DO2 102	Invarcione in the hackground: Conomic invarcions facilitate rapid adaptation by modulating	Mr. Shangaha 7hang
P03.103	Inversions in the background: Genomic inversions facilitate rapid adaptation by modulating	ivii. Silaligzlie Zilalig
D00 101	sexual conflict and regulatory interactions in Hawaiian field crickets	
P03.104	Degeneration and gene expression patterns in the morpho-social supergene in the ant	Dr. Romuald Laso-Jadart
	Myrmecina graminicola	
P03.105	Is genomic compatibility compromised after a bottleneck in Asian elephants?	Dr. Mirte Bosse
P03.107	Positive and negative selection on the supergene controlling male mating strategies in Ruff	Ms. Leyi Su

Universal bacterial clade dynamics dominate under predation despite altered phenotypes Mr. Dovydas Kiciatovas

S34 - Mechanisms of adaptation to changing conditions in microorganisms

and mutation targets

P03.108



P03.109	How turnover shapes the eco-evolutionary dynamics of complex bacterial communities	Dr. Jacob Wilde
P03.111	Genetic Adaptation to Environmental Change in Evolving Populations	Mr. Salvatore Bannò
P03.112	The role of evolving niche choice in herbivore adaptation to host plants	Dr. Peter Nabutanyi
P03.113	Stress-Driven MicroRNA Dynamics in Phaeodactylum tricornutum: Unveiling the Molecular Mechanisms of Microalgal Plasticity	Dr. Sabrina Carrella
P03.114	Evolution and evolvability of rifampicin resistance across the bacterial tree of life	Ms. Negin Bolourchi
P03.116	Role of $\beta\mbox{-lactamase}$ in cephalosporin resistance evolution in structured environments	Ms. Chang Cui
P03.117	Emergence and persistence: Structural evolution of early Nitrogenase enzymes	PhD Bruno Cuevas-Zuviría
P03.118	Does the lipid metabolic toolbox constrain the evolution of cellular biological processes?	Mr. Quentin Saintain
103.110	boes the lipid metabolic toolbox constrain the evolution of cellular biological processes.	Wil. Queritin Suntain
502.440	The sale of constant and of the sale is	Ma Bardina Communica
P03.119	The rate of spontaneous m6A changes in bacteria	Ms. Pauliina Summanen
P03.120	Nutrient-rich spatial refuges buffer against extinction and promote evolutionary rescue in	Dr. Siobhan O'Brien
	evolving microbial populations	
P03.121	Experimental evolution of antimicrobial peptide resistance, and increased sensitivity in the	Dr. Sarah Duxbury
	presence of microbial interactions	·
P03.122	Hypermutation and Fluctuating Stress Levels Can Enable Evolutionary Rescue	Mr. Gleb Ebert
P03.123	Long-term survival of microbial viruses suggests an evolutionary bet-hedging strategy	Dr. Eva Lievens
P03.124	Nitrogen availability drives ecological dynamics and parallel evolution in synthetic root-	Mr. Atharva Bhide
	associated bacterial communities.	
P03.125	Can diatoms use nutrient stores to affect temperature responses over several generations?	Mr. Shravan Raghu
		G
P03.126	Mutation bias predicts rapid emergence of antibiotic resistance	Ms. Adrita Chakraborty
		•
P03.127	Effect of multiple global change factors on microbial communities in the field	Dr. Alvaro Rodriguez Del Rio
P03.128	Multi-layered ecological interactions determine growth of clinical antibiotic-resistant	Dr. Ricardo Leon Sampedro
	strains within human microbiomes	
P03.131	World Enough and Time: Mapping the Martian Adaptive Landscape with a Terran	Dr. Peter Conlin
	Bacterium	
P03.132	Linked evolution of gene expression and fitness in fungal populations	Dr. Anne Genissel
S35 - Mech	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life	
S35 - Mecha P03.134	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life Most laterally acquired genes in grasses degrade due to the loss of utility	Ms. Catherine Collins
S35 - Mech	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De
S35 - Mecha P03.134 P03.135	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin
S35 - Mecha P03.134 P03.135	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life Most laterally acquired genes in grasses degrade due to the loss of utility	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De
S35 - Mecha P03.134 P03.135	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal
S35 - Mecha P03.134 P03.135 P03.137	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal
S35 - Mecha P03.134 P03.135 P03.137	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal
S35 - Mecha P03.134 P03.135 P03.137 P03.138	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere
S35 - Mecha P03.134 P03.135 P03.137 P03.138	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen  Ms. Flora Gaudilliere  Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation — how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere  Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze  Ms. Priya Gordon
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen  Ms. Flora Gaudilliere  Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.144	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere  Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze  Ms. Priya Gordon
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.144	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere  Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze  Ms. Priya Gordon
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.018	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti Mr. Franco Nery Liñán Vigo
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.018	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti Mr. Franco Nery Liñán Vigo
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.018	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti Mr. Franco Nery Liñán Vigo
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.019	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti Mr. Franco Nery Liñán Vigo
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.019	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range Plasmids in isolation vs plasmids in congregation — how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)  Flower traits and their relationship with evolution of tolerance to herbivory in Datura plants  omics: Challenges and Possibilities	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti Mr. Franco Nery Liñán Vigo Mr. Franco Nery Liñán Vigo
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.019 S37 - Muser P03.147	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)  Flower traits and their relationship with evolution of tolerance to herbivory in Datura plants  omics: Challenges and Possibilities  Museomic challenges to delimit the nine-banded armadillo species complex	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti Mr. Franco Nery Liñán Vigo Dr. Mathilde Barthe
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.019	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life Most laterally acquired genes in grasses degrade due to the loss of utility Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range Plasmids in isolation vs plasmids in congregation — how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)  Flower traits and their relationship with evolution of tolerance to herbivory in Datura plants  omics: Challenges and Possibilities  Museomic challenges to delimit the nine-banded armadillo species complex  From almost extinct to rescued? Temporal dynamics of genomic erosion after demographic	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti Mr. Franco Nery Liñán Vigo Dr. Mathilde Barthe
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.018 P03.019	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation — how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)  Flower traits and their relationship with evolution of tolerance to herbivory in Datura plants  omics: Challenges and Possibilities  Museomic challenges to delimit the nine-banded armadillo species complex  From almost extinct to rescued? Temporal dynamics of genomic erosion after demographic recovery.	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti  Mr. Franco Nery Liñán Vigo  Dr. Mathilde Barthe Dr. Claudia Fontsere
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.019 S37 - Muser P03.147	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation – how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)  Flower traits and their relationship with evolution of tolerance to herbivory in Datura plants  omics: Challenges and Possibilities  Museomic challenges to delimit the nine-banded armadillo species complex  From almost extinct to rescued? Temporal dynamics of genomic erosion after demographic recovery.  Genomic Insights into Biodiversity Decline: Integrating Reference Genomes and Population	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti  Mr. Franco Nery Liñán Vigo  Dr. Mathilde Barthe Dr. Claudia Fontsere
S35 - Mecha P03.134 P03.135 P03.137 P03.138 P03.139 P03.141 P03.142 P03.143 P03.144 P03.145 S36 - Micro P03.018 P03.019	anisms, barriers, and impacts of horizontal gene transfer across the Tree of Life  Most laterally acquired genes in grasses degrade due to the loss of utility  Addressing the role of horizontal gene transfer in plant-insect interactions  Host-parasitoid relationship as a driver of horizontal transfer  Long-term selection on mitocoding genes buried in mammalian and avian nuclear genomes  Unwelcome guests: characterizing the genomic niche of insertion sequence elements in prokaryotic genomes  Giant Virus Heredity is Surprisingly Vertical: Reconciliation of Nucleocytoviricota Phylogeny  Plasmid distribution reveals genomic determinants for their host range  Plasmids in isolation vs plasmids in congregation — how do evolutionary dynamics differ?  Ctrl+Alt+Conserve: Transformation Reduces Genomic Disparity in Digital Organisms  Detecting non-vertical inheritance across eukaryotes  evolutionary processes and Macroevolutionary patterns  Evolutionary patterns of tolerance to defoliation in the genus Datura (Solanaceae)  Flower traits and their relationship with evolution of tolerance to herbivory in Datura plants  omics: Challenges and Possibilities  Museomic challenges to delimit the nine-banded armadillo species complex  From almost extinct to rescued? Temporal dynamics of genomic erosion after demographic recovery.	Ms. Catherine Collins Mrs. Alexandra Jalaber Dupont De Dinechin Mrs. Audrey Portal Ms. Yu-chi Chen Ms. Flora Gaudilliere Dr. Joao Henrique Diniz Brandao Gervasio Dr. Yiqing Wang Ms. Ekaterine Kikodze Ms. Priya Gordon Mr. Giacomo Mutti  Mr. Franco Nery Liñán Vigo  Dr. Mathilde Barthe Dr. Claudia Fontsere



P03.151	Genomic evidence for West Antarctic Ice Sheet collapse during the Last Interglacial	Dr. Sally Lau
P03.152	Ancient leaves, modern trees: Phylogenomics in Amaranthaceae sensu stricto using	Ms. Tina Kiedaisch
	herbarium material	
P03.153	A legacy of low diversity: museum genomics uncovers persistent genomic erosion in the	Dr. Xuejing Wang
	Mauritius kestrel	
P03.154	Museomics reveals the true extend of horizontal transfer in Drosophila	Dr. Robert Kofler
P03.156	Speciation around a mountain chain: Hybridisation dynamics in a group of lek-mating Birds-	Mr. Ingo Müller
	of-paradise (Paradisaea)	
P03.158	Reconstructing the evolutionary history of the Tyto owls' diversification through	Dr. Angélica Pulido
	museomics	
P03.160	Shelf Life: Optimizing Ancient DNA Recovery from Museum-Curated Pleistocene Specimens	Dr. Hannah Moots
P03.161	Limits and best practices for local ancestry inference in imputed ancient genomes	Dr. Katia Bougiouri
P03.163	Genomic insight into the population history of Norwegian lemmings	Ms. Isabelle Feinauer
P03.164	DNAharvester: A Nextflow Pipeline for Processing and Analyzing Highly Degraded DNA from	Dr. Muhammad Bilal Sharif
	Ancient and Historical Specimens	
P03.165	Harnessing a Nearly Complete Herbarium Collection to Unravel the Evolutionary Radiation	Mr. Daniele Buono
	of Astragalus (Fabaceae)	
P03.166	The Genomic Revolution and Its Limitations: A Summary of Genomic Data Available for	Ms. Hayley Free
	Eukaryotic Species	, ,
P03.167	Ancient genomic reconstruction of Late Pleistocene North America Canis origins and	Dr. Paul Wilson
	distribution	
P03.168	Developing an accessible multi-locus barcoding approach for the field and museum	Mr. Yannis Schöneberg
P03.169	Past and future anthropogenic impacts on Atlantic bluefin tuna through the eyes of	Mr. Piergiorgio Massa
103.103	demographic and adaptation analyses on range-wide whole genomes	Will Fleigher Wassa
	demographic and adaptation analyses on range wide whole genomes	
S40 - Phylo	genomics methodology and the deep tree of life	
		D 4 6 6'I
PN3 170	I no (antrian knot of internavillm Salralian Relationships	I)r Ana Serra Silva
P03.170	The Gordian Knot of Interphylum Spiralian Relationships  Machine learning phylogenetics	Dr. Ana Serra Silva Mr. Nikita Kulikov
P03.171	Machine learning phylogenetics	Mr. Nikita Kulikov
	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome	
P03.171 P03.172	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome Intestine	Mr. Nikita Kulikov Ms. Mingyue Sun
P03.171	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome	Mr. Nikita Kulikov
P03.171 P03.172 P03.173	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility	Mr. Nikita Kulikov Ms. Mingyue Sun Dr. Caroline Puente-Lelievre
P03.171 P03.172 P03.173 P03.174	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility Fast Species Tree: Fast and Scalable Species tree Inference	Mr. Nikita Kulikov Ms. Mingyue Sun Dr. Caroline Puente-Lelievre Dr. Jonathan Holmes
P03.171 P03.172 P03.173	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid	Mr. Nikita Kulikov Ms. Mingyue Sun Dr. Caroline Puente-Lelievre
P03.171 P03.172 P03.173 P03.174 P03.175	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution	Mr. Nikita Kulikov Ms. Mingyue Sun Dr. Caroline Puente-Lelievre Dr. Jonathan Holmes Dr. Mahwash Jamy
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predip P03.180	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predic P03.180	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates?	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predip P03.180	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predic P03.180	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates?	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predic P03.180	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predic P03.180	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predip P03.180 P03.181 P03.183	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predip P03.180 P03.181 P03.183	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predip P03.180 P03.181 P03.183	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod Eurytemora affinis	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento Schulze
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predip P03.180 P03.181 P03.183	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod Eurytemora affinis Inter-island genetic connectivity and selection in the Avian Vampire Fly across the	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento Schulze
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predict P03.180 P03.181 P03.183 P03.185	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod Eurytemora affinis Inter-island genetic connectivity and selection in the Avian Vampire Fly across the Galápagos archipelago' Spatial patterns of intraspecific genetic diversity in herptiles: a machine learning approach	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento Schulze Ms. Abbie Hay
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178 S42 - Predict P03.180 P03.181 P03.183 P03.185	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod Eurytemora affinis Inter-island genetic connectivity and selection in the Avian Vampire Fly across the Galápagos archipelago' Spatial patterns of intraspecific genetic diversity in herptiles: a machine learning approach to macrogenetics	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento Schulze Ms. Abbie Hay  Dr. Matthew Moreira
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178  S42 - Predir P03.180 P03.181 P03.183 P03.185 P03.186 P03.188 P03.189	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod Eurytemora affinis Inter-island genetic connectivity and selection in the Avian Vampire Fly across the Galápagos archipelago' Spatial patterns of intraspecific genetic diversity in herptiles: a machine learning approach to macrogenetics Evolution to salinity in communities	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento Schulze Ms. Abbie Hay  Dr. Matthew Moreira  Ms. Emily M. Booms
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178  S42 - Predict P03.180 P03.181 P03.183 P03.185 P03.186 P03.188	Machine learning phylogenetics Co-evolution of the Myosin Vb—Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod Eurytemora affinis Inter-island genetic connectivity and selection in the Avian Vampire Fly across the Galápagos archipelago' Spatial patterns of intraspecific genetic diversity in herptiles: a machine learning approach to macrogenetics	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento Schulze Ms. Abbie Hay  Dr. Matthew Moreira
P03.171 P03.172 P03.173 P03.174 P03.175 P03.176 P03.177 P03.178  S42 - Predir P03.180 P03.181 P03.183 P03.185 P03.186 P03.188 P03.189	Machine learning phylogenetics Co-evolution of the Myosin Vb–Rab11 Interface and Villification of the Gnathostome Intestine From Sequence to Structure: Inferring the Origin and Evolution of Bacterial Motility  Fast Species Tree: Fast and Scalable Species tree Inference New Deep-Branching Environmental Plastid Genomes and Their Implications for Plastid Evolution Inferring the deep past of Asgard archaea Testing the adequacy of amino acid substitution models at different time scales DBuilder: Automated Building of Proteome Database for Phylome Reconstruction  cting evolutionary change in ecologically relevant contexts Escaping from a stressor: evolution of larval development under nutritional shortage  Is Bergmann's rule valid for terrestrial vertebrates? Does Mum always know best? Predicting the consequences of rapid host-preference evolution under climate change in Brown Argus butterfly populations  Genomic and physiological mechanisms underpinning seasonal adaptation in the copepod Eurytemora affinis Inter-island genetic connectivity and selection in the Avian Vampire Fly across the Galápagos archipelago' Spatial patterns of intraspecific genetic diversity in herptiles: a machine learning approach to macrogenetics Evolution to salinity in communities	Mr. Nikita Kulikov Ms. Mingyue Sun  Dr. Caroline Puente-Lelievre  Dr. Jonathan Holmes Dr. Mahwash Jamy  Dr. Julian Vosseberg Dr. Mattia Giacomelli Ms. Patricia Saragüeta  Ms. A. Carolina Varela-pereira  Ms. Oleksandra Oskyrko Dr. Brooke Zanco  Dr. Jennifer Catherine Nascimento Schulze Ms. Abbie Hay  Dr. Matthew Moreira  Ms. Emily M. Booms



P03.193	Host shift changes population dynamics and phenotypic plasticity in seed beetles	Dr. Uroš Savković
P03.195	Genomic Insights into Kaimanawa Horses: Demography, Diversity, and Conservation Strategies for a Feral Population	Mr. Arne Bielke
P03.197	Effect of temperature fluctuations on life history traits and strategies of phytophagous arthropods - a systematic review	Ms. Gulsamal Askarova
P03.198	Assessing the genetic vulnerability of Myotis velifer to white-nose syndrome	Dr. Flora Whiting-Fawcett
P03.200	Life-history trade-offs explain local adaptation in Arabidopsis thaliana	_
		Dr. Benjamin Brachi
P03.201	Breastfeeding difficulties and maternal trade-offs in rural southern Poland	Ms. Joanna Żyrek
P03.203	Complex ecological strategies drive genetic distribution of Arabidopsis thaliana in natural urban habitats.	Ms. Justine Floret
P03.205	The response of the life history traits of Daphnia magna to the deteriorating and fluctuating temperatures	Dr. Nikola Petkovic
P03.206	Predators in the eco-evolutionary dynamics of arthropod communities	Mr. Henri Truchassout
P03.207	Correlated genomic patterns of introgression across space despite contrasting hybridization histories.	Dr. Matthew Farnitano
P03.208	Understanding the Ecology of Sexual Size Dimorphism Through Resource Competition in a Sex- and Size-structured Model	Ms. Danyang Shi
P03.209	Statistical models of eco-evolutionary dynamics to understand the past and predict future states of ecological communities	Prof. Jelena Pantel
P03.210	Evolution of the seasonal preference of pathogens	Dr. Ryuichi Kumata
P03.211	Predicting community response to warming and heat waves using thermal performance of component species	Mr. Paresh Nath Das
P03.212	Predictability in coevolution under hierarchical competition	Prof. Inês Fragata
P03.213	Repeatability of evolution in interacting species	Ms. Loraine Hablützel
P03.214	Cryptic evolution of body size in response to climate change in the Alpine marmot (Marmota marmota)	Mr. Pierre-alexandre Quittet
P03.216	Ecological context drives introgression outcomes in experimental Drosophila hybridization	Ms. Martina Pernigotti
SAE Thor	evolution of microbial pangenomes	
		Mc Marina Mota Morlo
P03.217	Recombination enforces diversity of secreted proteins in a symbiont of honeybees	Ms. Marina Mota Merlo
P03.218	Bacteriocin competition: ecological and evolutionary dynamics in colonization	Mr. Elliott Greene
P03.219	Phylogeny, evolution and virulence of Coxiella burnetii the agent of the Q fever	Dr. Theo Tricou
S48 - The r	naintenance of adaptive polymorphisms	
P03.220	Associative overdominance can drive recombination suppression around loci under	Ms. Lou Guyot
P03.221	balancing selection	,
	The genomic basis and evolution of immune trait variation in Soay sheep	Ms. Gina Henderson
P03.223	Survival, sex, and insecticides: decoding the mechanisms behind resistance polymorphism in Drosophila melanogaster	Dr. Felipe Martelli
P03.224	Intramolecular trade-offs promote the maintenance of polymorphism in plant-pathogen coevolution	Dr. Anja Hörger
P03.225	Genetic basis and selection maintaining wing polymorphism in the evening brown butterfly, Melanitis leda	Ms. Yuqian Huang
P03.226	Gene conversion as a key contributor to adaptive polymorphism at the avian MHC	Prof. Piotr Minias
P03.227	Molecular evolution in Bryophytes: How ploidy drives patterns and efficacy of natural selection.	Ms. Muskaan Muskaan
P03.228	Differing adaptations in regulation of conjunctival immune response to a virulent pathogen across house finch populations	Dr. Michal Vinkler
P03.229	A key protein of missing-self-induced innate immunity is polymorphic and driven by diversifying selection in a wild rodent	Prof. Jacek Radwan
P03.230	Comparison of the the genetic structures of Borrelia virulence factor, the outer surface protein C, and the putatively neutral intergenic spacer	Mrs. Jozefina Wasilewska
P03.231	Deleterious hitchhiking under balancing selection as a potential fitness cost of balancing selection.	Dr. Roman Stetsenko
P03.232	Different Modes of Balancing Selection Leave Highly Similar Signatures in Time-Series	Dr. Ozgur Taskent
P03.234	Genetic Data  Maintenance of Social and Wing Polymorphism in the ant Myrmecina graminicola through the selfish transmission of a haplotype	Mr. Brandon Duquenoy
	the selfish transmission of a haplotype	

W. C.	

P03.235	Trans-species polymorphisms and long-term balancing selection across the primate radiation.	Dr. Sebastian Cuadros
P03.236	Genomic Signatures of Sex-Specific Selection in the seed beetle Callosobruchus maculatus	Dr. Alexandre Rego
P03.237	Harnessing Simulations to Identify Balancing Selection in Evolve-and-Resequence Experiments	Mr. Baron Koylass
P03.239	Balancing, Linkage and Effects of Selection: a Deep Learning Approach	Dr. Carolin Kosiol
P03.241	Using experimental evolution to explore patterns of dominance in Drosophila melanogaster	Dr. Katrine K. Lund-Hansen
S50 - Unray	reling the origin of eukaryotes: integrating prokaryotic and eukaryotic perspectives	
P03.242	Diverse origins of the last eukaryotic common ancestor proteins points to multiple	Mr. Moisès Bernabeu
P03.244	interactions during eukaryogenesis Cyanobacterial Contributions to the Evolutionary Origins of Animals	Prof. Paul Taylor
P03.245	Pluralistic Red Queen and Court Jester Interactions Drive Unisexual Reproduction in Digital Populations	Ms. Priya Gordon
P03.246	Inferring the ancestral set of eukaryotic genes - comparative genomics of jakobid and malawimonad flagellates	Mr. Samuel Prince-drouin
P03.247	Orthology of eukaryotic protein complexes	Dr. Ore Francis
S51 - Open	Symposium	
P03.194	Evolutionarily distinct Saccharomyces cerevisiae lineages coexist within a single fruit habitat (marula fruits)	Tawanda Proceed Makopa
P03.215	Integrating thermal physiology and evolutionary ecology to understand biodiversity responses to climate change in Brazilian ecosystems	Luisa Maria Diele-Viegas
P03.222	Chromosome-level genome assembly and annotation of Salvelinus malma	Meijun Wang
P03.233	Exploration of gene presence/absence variations in Oncorhynchus mykiss and their differentiation between wild and selection population	Dr. Yuan Tian
P03.238	Genomic basis of copy number variations and codon substitution provided the evidence for convergent evolution of salinity adaptation in teleosts	Qinfeng Gao
P03.243	A curated great ape genome diversity panel and a selection screen	Dr. Sojung Han
P03.248	Triploidy in parasitic worms: an overlooked aspect in host-parasite dynamics and evolution of drug resistance?	
P03.249	Sweeps in the IM model	Dr. Derek Setter
P03.250	Reproductive success of dispersers depends on the population of origin in Atlantic salmon	Mr. Emilio Egal
P03.251	Adaptive Laboratory Evolution of Aspergillus niger Using Single-Conidium Sorting for Enhanced Citric Acid Production	Ms. Adéla Schandl
P03.252	Biologically inspired warning patterns deter birds from wind turbines	Dr. Sandra Winters
P03.253	Development of a Thermophilic PURE System for Cell-Free Paleobiology: Exploring Protein Evolution at the Origin of Life	Katsumi Hagino
P03.254	Postglacial divergence and local adaptation in the wood-decay fungus Trichaptum abietinum	Ms. Kathleen Theresia Helleland
P03.255	Latitude Matters: A Global Phylogeographic Perspective on Climate-Driven Demographic Responses in Tarantulas	Mr. Praveen Karanth
P03.256	Being plastic is fantastic? How climate influences morphology across developmental strategies in European salamanders and newts	Ms. Morgane Fournier
P03.257	Assessment of Transparency Across Articles in Ecology and Evolutionary Biology Journals, 2000–2024	Ms Marija Purgar Filjak
P03.258	Intraspecific variation in mate attraction shapes hybridization in a young species group	Ms. Margarita Spirina
P03.259	One Sea, Different Whales: Genomics Sheds Light on the Puzzling Population of Mediterranean Fin Whales	Dr. Elisa Desiato
P03.260	A temporal analysis of sockeye-kokanee hybridization, behaviour, and fitness following a re- introduction program	Ms. Olivia Boven
P03.261	Distinct sex-specific patterns in hematological and biochemical aging of Macaca monkeys	Dr. Min-Gyeong Ko
P03.262	Co-Evolution between Phages and Bacterial Pangenomes in the Light of Horizontal Gene Transfer	Dr. Franz Baumdicker
P03.263	The joint influence of incompatibilities and heterosis on hybrid population genetics	Mr. Julio Ayala Lopez
P03.264	Environmental heterogeneity changes the genomic signature of adaptation to cadmium	Ms. Marta Ferreira



P03.265	Identifying candidate genes for extreme growth variation in response to nutrition in the water strider Microvelia longipes	Ms. Ingrid Dourlens
P03.266	Cracking the transcriptomic hieroglyphics code: comparative transcriptomic analysis of respiratory organ evolution across arthropods	Mr. Polychronis Tatsis
P03.267	Modeling the effect of risk tolerance on biopesticide resistance evolution & crop damage	Ms. Blair Matarlo
P03.268	Recent Positive Selection in a Wild Rodent: Investigating the Genomic Basis of Adaptation	Mrs. Lucia Ximena Alva Caballero
P03.269	Origin, evolution and biogeography of Atlantic Forest Nymphalidae (Lepidoptera)	Ms. Mar Repullés
P03.270 P03.271	Vocal divergence and incipient speciation in the Reunion Grey White-eye Would the real Ophioplocus incipiens please stand up?	Ms. Bárbara Freitas Ms. Beatrice Salgarella
P03.272	Ecological disruption and evolutionary signals from transgene spread in a wild cotton	Ms. Valeria Vázquez-barrios
500 070	metapopulation within its center of origin	
P03.273	The Population Genetic History of the Hittite Capital Hattusa	Ms. Kayra Canpolat
P03.274	Low Genetic Diversity of Key Immune Genes in the Critically Endangered Burrunan Dolphin (Tursiops australis)	Ms. Grace Day
P03.275	Parental impact on genetic inheritance: patterns of accumulating mutations across generations in Macaca fascicularis	Ms. Hyeri Park
P03.276	Biogeography of Southeast Asian Melastomes - insights from a phylogenomic study of Dissochaeteae	Ms. Linde Wieringa
P03.277	Repeated sex chromosome turnovers in African annual killifishes	Dr. Petr Nguyen
P03.278	Molecular mechanisms of colouration using salamanders as a model	Mr. Nicholas Strowbridge
P03.279	White lupin's genomic diversity and domestication history through pangenome graphs	Prof. Andrea Benazzo
103.273	white tupin's genomic diversity and domestication history through pungenome graphs	Tron. Andrea Benazzo
P03.280	The SLC30A9 gene case: Testing Denisovan Adaptive Introgression using Approximate Bayesian Computation	Mr. Jorge Garcia-calleja
P03.281	Birdlife in the Anthropocene: Understanding convergent adaptation to urbanisation	Ms. Marina Martín Maroto
P03.282	Donlife Denulation Conomics serves the Tree of Life	Dr. Sònia Casillas
	PopLife: Population Genomics across the Tree of Life	
P03.283	Comparison of genetic effective population size estimates in species across a large range of	Ms. Marie-Gabrielle Harribey
	life-history strategies	
P03.284	Evolution of evolution: how do mechanisms of evolution change over time?	Dr. Oleksandr Maistrenko
P03.285	Does the social and sexual environment impact ageing through nutrient-sensing pathways?	Dr. Emily Churchill
P03.286	Multipartite Parasitic Interactions in the Aquatic Food Web: Modulating Gene Exchange and Carbon Flow	Dr. Maliheh Mehrshad
P03.287	Genomic and Transcriptomic Insights into Salinity Adaptation and Immune Trade-offs in the Spanish Toothcarp (Aphanius iberus, Valenciennes, 1846)	Dr. Silvia Perea
P03.288	How subtle environmental differences influence the predictability of evolution?	Ms. Neetika Ahlawat
P03.289	Chronic heat tolerance reveals overestimated thermal safety margins and increased vulnerability in marine fish populations	Mr. Andrés N. Molina
P03.290	Universality and Cultural Specificity in Artifact Evolution: A Cross-Cultural Transmission Chain Experiment	Mr. Petr Chlup
P03.291	Host-Driven Divergence in the Fish Parasite Ligula intestinalis: Evidence of Parallel Evolution in Two Species Lineages	Dr. Masoud Nazarizadeh
P03.292	Genetic Basis of Immunity in Drosophila melanogaster Populations Selected for Improved Post-Infection Survivorship	Ms. Tsering Choton
P03.293	Revisiting genetic load across populations using protein language models	Ms. Ece Zeynep Atici
P03.294	Chromosome-Level Genome Assemblies Across Five Brassicaceae Spesies: New Resources for Comparative Evolutionary Studies	Dr. Mahnaz Nezamivand-chegini
P03.295	How do expression plasticity and pleiotropic gene function impact the evolution of gene	Ms. Julia Hagauer
DOC 05 -	regulation? A case study applying single cell multi-omics and environmental manipulation to embryos of divergently adapted fish	
P03.296	Genomic Insights of the Migratory Divide: Investigating Population Structure in the Red- necked Phalarope	Ms. Vishakha Gupta
P03.297	Microsatellite identification of ramet genotypes in a clonal plant Iris pumila	Dr. Stevan Avramov
P03.298	Associative Learning in Plants	Ms. Muriel Saidenberg
P03.299	The link between pace and shape of ageing: a comparative analysis in birds and mammals	Dr. Victor Ronget
P03.300	Using a computational model to explore possible eco-evolutionary drivers of swarming behavior in locusts	Mr. Dan Amichai



		· · · · · · · · · · · · · · · · · · ·
P03.301 P03.302	Multivariate selection and the evolution of Drosophila wings Individual heterogeneity in bovine tuberculosis (bTB) disease dynamics in European	Dr. Stephen De Lisle Dr. Barbara Tschirren
P03.303	badgers: genetics, senescence and social networks  Adaptive laboratory evolution of Saccharomyces and non-Saccharomyces wine strains under oxidative stress: phenotypic shifts and genomic perspectives	Mrs. Andrea Irene Silva Claros
P03.304	Experimental evidence for cryptic female choice in the plant species Brassica rapa	Mr. Timothée Chenin
P03.305	Population genomics of coastal dune building species European marram grass (Calamagrostis arenaria)	Ms. Emily Baker
P03.306	Human-commensal songbirds and the genomics of convergent adaptation	Mr. Erik Sandertun Røed
P03.307	Evolution of plasticity and evolvability in a model of bacterial surface exploration	Mr. Alger Jorritsma
P03.308	Integrating gene expression and biochemical approaches for understanding the interactions between the antibiotic enrofloxacin and environmental stressors in the Mediterranean mussel (Mytilus galloprovincialis)	Prof. Carlo Pretti
P03.309	Investigating mitonuclear interactions in hybridizing newt species Triturus ivanbureschi and T. macedonicus	Ms. Joanna Jakobik
P03.310	Nutrient-Driven Forms: Evolved Allometry in Drosophila adapted to malnutrition conditions	Mr. Shrinath Narayanan
P03.311	The tempo and mode of toxicant sensitivity evolution	Mr. Iain R. Moodie
P03.312	Identification of loci underlying local adaptation in European Drosophila melanogaster	Mr. Siddharth Murali
	populations	
P03.313	The performance of phenotypic evolutionary rate inference methods using continuous multivariate data: a simulation approach	Ms. Ruby Redlich
P03.314	Beyond Genetic Analogy: Variance Proportionality in Cultural Evolution	Mr. Peter Kutsos
P03.315	TreeProfiler: Computing Phylogenetic Profiles and Visualizing Metadata on Large Trees	Dr. Ziqi Deng
P03.316	Building Bridges in Evolution Education: A Teacher Training Experience in Argentina	Ms. Pía Pacheco
P03.317	Sexual conflict and sexual pleiotropy	Dr. Thomas Hitchcock
P03.318	Biogeographic resilience in Patagonian spiders: Glacial refugia and distribution shifts of	Ms. Pía Pacheco
103.310		ivis. Fla Facilico
P03.319	Philisca species during Pleistocene climatic oscillations It all began in the center: genomic phylogeny and ancestral range estimates provide new	Dr. Laura Hagemann
P03.320	insights into Sulawesi tarsier diversification  Context-dependent mate preferences influence the stability of premating isolation in early	Ms. Sofie Nilén
500 004	population divergence	
P03.321	Expansive pangenomes in Agaricomycetes	Dr. Miguel Angel Naranjo-Ortiz
P03.322	Using drawings to illustrate evolution	Dr. Jelle Zandveld
P03.323	Plasticity in rapidly changing nightscapes: Effects of Artificial Light at Night on color dynamism in chameleons from Madagascar	Ms. Udita Bansal
P03.324	Reconstructing the morphological and chemical evolution of the prothoracic repellent glands in stick and leaf insects (Phasmatodea)	Prof. Thomas Schmitt
P03.325	Strong sexual selection reduces population size, and limits population recovery after heat stress	Dr. Neha Pandey
P03.326	Impacts of management on the genomics of perennial crops: a case study of two mezcal Agave species with different selection histories.	Mrs. Irene Martínez Velasco
P03.327	Coevolution of ploidy and sex	Dr. Carl Mackintosh
P03.328	Genomic basis of recurring phenotypic loss of bee scopae	Dr. Eckart Stolle
P03.329	Adaptive significance of the effects of population density on life history traits in a moth	Ms. Kerli Kuusk
P03.330	Genomic Insights into the Evolutionary History and Cryptic Diversity of Two-Toed Sloths (Choloepus) in Amazonia	Ms. Linda Hagberg
P03.331	Monogamy may remove constraints on reproductive tissue investment imposed by intense	Dr. Maider Iglesias
P03.332	sexual selection  Minimal effects of starvation on pre- and post-copulatory male sexual traits in Drosophila	Mr. Ashwin Suryanarayanan
P03.333	melanogaster  Non-invasive samples reveal genetic connectivity and fine-scale population structure in	Ms. Irune Ruiz Gartzia
	western lowland gorillas	
P03.334	Glowing and seeing red in deep-sea dragonfish (Stomiiformes)	Mrs. Vit Kaufman
P03.335	Testing the link between DNA methylation and phenotypic variation in the water strider, Microvelia longipes	Dr. Mirjam Urb
P03.336	Communicating science with art – you don't need to be an artist!	Ms. Matilda Scott



P03.337	Assessing genotype imputation methods for low-coverage sequencing data in populations with differing relatedness and inbreeding levels	Dr. Audald Lloret-Villas
P03.338	Inferring the Dry Season Ecology of Malaria Vectors from Allele Frequency Dynamics	Dr. Tin-Yu Hui
P03.339	Discovering animal diversity through phylogenetic placement of 18S metabarcoding data	Mr. Javier Arañó-Ansola
P03.340	The role of ongoing introgression on diversification in an adaptive radiation of sailfin silverside fishes from Sulawesi	Dr. Els De Keyzer
P03.341	Transposable element copy number variation reveals their evolutionary dynamics in apple domestication	Mr. Anthony Venon
P03.342	Leveraging macrogenetics to understand global patterns of intraspecific genetic variation on genes under purifying selection	Dr. Ivo Colmonero Costeira
P03.343	Interplay between sex determination and sex chromosome turnover in Sepsid flies	Ms. Dunja Rokvić
P03.344	Philopatry, Dispersal, and Distance: Disentangling Genetic Structure in a Highly Mobile Terrestrial Bird	Ms. Juliana Fonseca-Tellez
P03.345	Inner ear asymmetry in birds and mammals: Clues to developmental instability and evolvability	Ms. Anna Pyttlik
P03.346 P03.347	Floral developmental constraints and pollinator dependency in soybean Identifying the most stable endogenous controls for RT-qPCR analysis of genes of interest in larvae and adult male and female fruit flies (Drosophila melanogaster) exposed to different concentrations of cadmium chloride using available statistical tools	Dr. Marina Strelin Ms. Mina Rakić
P03.348	Conservation genetics of Ambystoma dumerilii, a narrow-endemic axolotl species critically endangered	Mr. Pablo Lechuga Paredes
P03.349	Origins of androgenetic lineages and sperm competition in the clam genus Corbicula	Mr. Mohammed Benbachir
P03.350	Loud environment and noisy neighbors: impact of urbanization and interspecific competition on song characteristics of Blue and Great tits	Ms. Jeanne Legros
P03.351	On the origin of the parthenogenetic, polyploid insect Saga pedo (Orthoptera, Tetiigonidae)	Dr Beatriz Navarro Domínguez
P03.352	Environmental DNA illuminates evolutionary patterns in tardigrade communities across urban and elevational zones	Ms. Payal Dash
P03.353	Metabolic Cross-Feeding Shapes Genome Evolution and Community Assembly in Deep Groundwaters	Dr. Maryam Rezaei Somee