

Congress of the European Society for Evolutionary Biology

August 14–19, 2022 Prague Congress Centre

Book of Abstracts



CONTENTS

TALKS	4
- / 	

- **S01.** Tug of war between the sexes: The transcriptomic architecture of sex-linked traits
- **S02.** Sex chromosome evolution: the canonical model and so much beyond
- **S03.** Diversity and evolution in sperm, ova, and other primary reproductive traits
- **S04.** The evolutionary ecology of mating systems
- **\$05.** A combinatorial view on rapid speciation the role of ancient genetic variants and hybridisation
- **S06.** Revisiting chromosomal speciation in the genomic era
- **\$07.** Chromosome rearrangements in evolution
- **S08.** Integrative biogeography: Past, present, future
- **\$09.** Parallel and repeated evolution in adaptive radiation
- **\$10.** Eco-evolutionary dynamics in changing environments: insights from models, experiments and case studies
- **S11.** Adaptation and evolution across environmental gradients
- **\$12.** Resurrection ecology as a tool for the study of rapid evolution
- **S13.** Evolutionary ecology of chemically-mediated species interactions in plants
- **\$14.** Ecological drivers and evolutionary consequences of within-population colour variation
- **\$15.** Rapid evolution of color patterns
- **\$16.** Predator cognition and the evolution of prey defence strategies
- **\$17.** Brain, behaviour and cognitive evolution
- \$18. The evolution of behavioural adaptations: Genes, neurons and ecology
- **\$19.** Eco-evolutionary dynamics and feedbacks in invasive species
- **\$20.** Unravelling the interplay between plasticity and evolution during rapid global change
- **S21.** Epigenetics goes wild! Epigenetic diversity and the evolutionary potential of wild populations.
- **S22.** Phenotypic plasticity's importance in evolution: Same old dog or new tricks?
- **\$23.** The evolution and consequences of non-mendelian inheritance
- **S24.** Progress and prospects in adaptation genomics
- **S25.** The positives and negatives of whole genome duplication: synthesizing polyploid evolution across organisms and disciplines
- \$26. The biological meaning of SNPs
- **\$27.** Tandem repeats: their role in molecular evolution and methods
- **S28.** Beyond transcription: the role of post-transcriptional gene regulation in adaptation and evolution
- **\$29.** Comparative genomics: a powerful tool for exploring broad evolutionary questions
- **\$30.** Characterizing genomic landscapes of recombination and their evolution
- **\$31.** Limits to adaptation: linking evolution, ecology, and genetics
- **S32.** Inferring macroevolutionary patterns from microevolutionary processes: methods and practices
- **\$33.** Domestication: Fresh insights from ancient genomics
- **S34.** How have biomarkers improved our understanding of health and the evolution of senescence?
- **\$35.** The art of microscopic war: interference competition in microbes
- **\$36.** Evolution of antibiotic resistance: from lab to clinic



- **S37.** Microbiomes in the wild: the drivers and evolutionary consequences of microbiome variation
- **\$38.** Molecular evolution and trade-offs in host-pathogen interactions and host immunity
- **\$39.** Mechanisms of host-symbiont coevolution: from genotype to phenotype
- **\$40.** OPEN SYMPOSIUM

POSTERS.......356

- **S01.** Tug of war between the sexes: The transcriptomic architecture of sex-linked traits
- **S02.** Sex chromosome evolution: the canonical model and so much beyond
- **S03.** Diversity and evolution in sperm, ova, and other primary reproductive traits
- **S04.** The evolutionary ecology of mating systems
- **\$05.** A combinatorial view on rapid speciation the role of ancient genetic variants and hybridisation
- **S06.** Revisiting chromosomal speciation in the genomic era
- **\$07.** Chromosome rearrangements in evolution
- **S08.** Integrative biogeography: Past, present, future
- **\$09.** Parallel and repeated evolution in adaptive radiation
- **\$10.** Eco-evolutionary dynamics in changing environments: insights from models, experiments and case studies
- **\$11.** Adaptation and evolution across environmental gradients
- **\$12.** Resurrection ecology as a tool for the study of rapid evolution
- **\$13.** Evolutionary ecology of chemically-mediated species interactions in plants
- **\$14.** Ecological drivers and evolutionary consequences of within-population colour variation
- **\$15.** Rapid evolution of color patterns
- **\$16.** Predator cognition and the evolution of prey defence strategies
- **\$17.** Brain, behaviour and cognitive evolution
- \$18. The evolution of behavioural adaptations: Genes, neurons and ecology
- **\$19.** Eco-evolutionary dynamics and feedbacks in invasive species
- **\$20.** Unravelling the interplay between plasticity and evolution during rapid global change
- **S21.** Epigenetics goes wild! Epigenetic diversity and the evolutionary potential of wild populations.
- **S22.** Phenotypic plasticity's importance in evolution: Same old dog or new tricks?
- **S23.** The evolution and consequences of non-mendelian inheritance
- **\$24.** Progress and prospects in adaptation genomics
- **S25.** The positives and negatives of whole genome duplication: synthesizing polyploid evolution across organisms and disciplines
- **\$26.** The biological meaning of SNPs
- **\$27.** Tandem repeats: their role in molecular evolution and methods
- **\$28.** Beyond transcription: the role of post-transcriptional gene regulation in adaptation and evolution
- **\$29.** Comparative genomics: a powerful tool for exploring broad evolutionary questions
- **\$30.** Characterizing genomic landscapes of recombination and their evolution
- **\$31.** Limits to adaptation: linking evolution, ecology, and genetics
- **S32.** Inferring macroevolutionary patterns from microevolutionary processes: methods and practices
- **\$33.** Domestication: Fresh insights from ancient genomics
- **S34.** How have biomarkers improved our understanding of health and the evolution of senescence?



- **\$35.** The art of microscopic war: interference competition in microbes
- **S36.** Evolution of antibiotic resistance: from lab to clinic
- **S37.** Microbiomes in the wild: the drivers and evolutionary consequences of microbiome variation
- \$38. Molecular evolution and trade-offs in host-pathogen interactions and host immunity
- **S39.** Mechanisms of host-symbiont coevolution: from genotype to phenotype
- **\$40.** OPEN SYMPOSIUM



affects prophage persistence. When prophages are subjected to induction, we observed fast selection against prophages, driven by loss of prophages from the bacterial genome. In contrast, prophages subject to counteracting selective pressures from antibiotics and an inducing agent remain persistent in their bacterial host through two different mechanisms. Prophages that provided a selective advantage with antibiotic resistance that acts as a public good facilitate the emergence of phage free bacteria. Therefore, prophage containing bacteria persist at an equilibrium frequency with phage free bacteria to counteract the antibiotics. Lysogens without this beneficial resistance gene also maintain their prophage by going through successive rounds of beneficial mutations against the presence of antibiotics and inducing agent. Our study suggests that prophage persistence and the frequency of carriage within populations is environmentally dependent, and the pressure of induction can lead to rapid loss of prophages from a population.

Abstract ID: 1684
Poster board number:

P646

Temperature shapes chromosome inversion polymorphism of *Drosophila subobscura* through generations

Katarina Erić¹⁾, Marija Savić Veselinović²⁾, Aleksandra Patenković¹⁾, Pavle Erić¹⁾, Mihailo Jelić²⁾, Marina Stamenković - Radak^{1, 2)}, Marija Tanasković¹⁾

¹⁾Department of Genetics of Populations and Ecogenotoxicology, Institute for Biological Research"Siniša Stanković" – National Institute of the Republic of Serbia, Belgrade, Serbia

²⁾Faculty of Biology, University of Belgrade, Belgrade, Serbia

Drosophila subobscura is a wide spread Palearctic species, which possesses very rich inversion polymorphism in all of the five acrocentric chromosomes. As populations rapid response to fast changes of environmental conditions can be seen through changes in inversions frequency, this polymorphism is good candidate to measure the effect of temperature on the genetic structure of population. Frequencies of particular with chromosomal arrangements are correlated temperature, with arrangements increasing in frequency at lower temperatures, and more complex arrangements like J_1 , U_{1+2} , U_{1+2+6} , E_{1+2+9} and O_{3+4} increasing in frequency at higher temperatures. In order to explore the effect of optimal and suboptimal temperatures on the arrangements frequencies, samples of D. subobscura population from a high altitude were reared in laboratory at suboptimal (16°C) and optimal (19°C). Chromosome arrangements frequencies were analyzed and compared through generations (F₀, F₅ and F₁₆). The increasing frequency of complex chromosome arrangements and the decreasing frequency of standard chromosomal arrangements were observed. There are significant differences in some of the chromosome arrangements between suboptimal and optimal conditions and also between generations, indicating laboratory adaptation through generations.

Abstract ID: 1722
Poster board number:

P647

Context- and sex- dependent links between father's sexual success and offspring pathogen resistance