

Programme

Israeli Society for Evolutionary Biology 2019

Wednesday 11 December 2019

Hour		Hall	Room 1	Room2
08:00	09:00	Registration		
09:00	09:15	Opening remarks		
09:15	10:00	PLENARY 1 = DOUGLAS ERWIN Developmental novelties in the early history of complex life		
10:00	10:20	Coffee Break		
		Genome Evolution (Chair: Tal Pupko)	Population Genetics (Chair: Yoav Ram)	Experimental Evolution (Chair: Tzachi Pilpel)
10:20	10:45	EINAT HAZKANI-COVO The evolution of short inverted repeats	ILAN GRONAU The case for genetic genealogies in population genomics	RUTH HERSHBERG Recovery from costs associated with adaptation will often lead to loss of adaptive alleles once selection in their favor stops
10:45	11:00	JOACHIM SURM The draft genome of <i>Actinia tenebrosa</i> reveals insights into toxin evolution	SHIRLI BAR-DAVID Impact of human-dominated sites on the fine-scale genetic structure of the Nubian Ibex in Israel	ORNA DAHAN Exploring the balance between mutations and horizontal gene transfer during bacterial adaptation
11:00	11:15	RONI RAK A shift in tRNA modifications and abundance during T-cell activation	AYELET SALMAN-MINKOV Inference of complex scenarios of adaptive evolution during rapid species radiation	ABRAHAM KOROL Evolution of recombination features: numerical models and experiments with <i>Drosophila</i>
11:15	11:30	HADAR MEDINI Metabolic heterogeneity in pancreatic Beta cells: Single cells Mitochondrial transcript heterogeneity as a model	PNINA COHEN Demographic history of invasive fire ants	SARAH LEDUC An ecological tragedy: The collection of pre-parasitized prey by a potter wasp
11:30	11:45	TATIANA ORLI SANDBERG RAILEANU Insights into the evolution of Myxozoa mitochondrial genomes	GAD DEGANI Genetic variation of the distribution of various species of amphibians in Israel	ODED REHAVI Transgenerationally inherited information regulates reproductive strategies in <i>Caenorhabditis elegans</i>
11:45	12:00	TALIA KUSTIN Biased Evolution in RNA Viruses	OREN AVRAM Bacterial adaptations – NOT what you thought	ANNA MELLUL Establishing an experimental-evolution-based method for understanding an incomplete penetrance mechanism in a model of <i>C. elegans</i>

12:00	13:00	Lunch Break
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Hour		Hall	Room 1	Room2
13:00	13:45	PLENARY 2 = DIETER EBERT Explaining species wide variation: What shapes phenotypic and genotypic diversity?		
13:45	14:05	Coffee Break		
		Human Evolution & Paleobiology (Chair: Dan Mishmar)	Evolution of Behaviour (Chair: Lilach Hadany)	Virus & Mobile Elements (Chair: Yehu Moran)
14:05	14:30	OREN KOLODNY Why did Moderns replace Neanderthals and not vice versa? And what took them so long?!?	ARNON LOTEM Crowd wisdom enhanced by costly signaling in a virtual rating system	ADI STERN Mechanistic insights into the evolutionary dynamics of competing cheater RNA viruses
14:30	14:45	YOEL RAK The Neandertal's Place in Nature	ERAN TAUBER Nocturnal and diurnal Drosophila: insights into the genetics of diurnal preference	MAGDA LEWANDOWSKA dsRNA sensing and antiviral immune response in a non-bilaterian model animal <i>Nematostella vectensis</i>
14:45	15:00	DOR SHILTON Human Social Evolution: Self-domestication or self-control?	Yael GUREVICH The evolution of paternal care: a role for microbes?	SHAI PILOSOF Network structure of CRISPR immunity shapes the co-evolutionary dynamics between hosts and viruses
15:00	15:15	YUVAL TABACH Comparative genomics of 1600 species reveal new insights about "super-traits", Breast cancer, and human diseases	KEITH D HARRIS Using cellular automata to study genome evolution	TZACHI HAGAI Caught between pathogens and autoimmunity: The evolution of the mammalian immune system
15:15	15:30	YOAV RAM Evolution of non-vertical social learning	SIMCHA LEV-YADUN Anti-herbivory plant coloration	EREZ LEVANON Purifying selection of long dsRNA bounds the layout of mobile elements in metazoan genomes
15:30	16:45	Poster session		
16:45	17:30	PLENARY 3 = HANNA KOKKO Gentlemanly males? Always, sometimes, never?		
17:30		End		

Thursday 12 December 2019

Hour		Hall	Room 1	Room2
08:00	09:00	Registration		
09:00	09:45	PLENARY 4 = DAN GRAUR Mutational load and the functional fraction of the human genome: It's complicated		
09:45	10:05	Coffee Break		
		Evolutionary Ecology 1 (Chair: Eran Tauber)	Systems Biology & EvoDevo 1 (Chair: Ariel Chipman)	Microbial Evolution (Chair: Adi Stern)
10:05	10:30	ARIEL NOVOPLANSKY Metaplasticity: fine tuning and curbing the costs of plasticity	NAAMA BARKAI Regulatory evolution: massive variations in gene abundances contrasts a largely invariant gene-regulatory network	SARIT AVRANI Micro and macro diversity are both important for bloom-forming cyanobacteria interactions with their phages
10:30	10:45	MICHAL GRUNTMAN Evolution of plant defenses along an invasion chronosequence	TAMAR LOTAN Signaling via GABAB receptors regulates early development and neurogenesis in the basal metazoan <i>Nematostella vectensis</i>	URI OBOLSKI Vaccination affects selection for antibiotic resistance in <i>Streptococcus pneumoniae</i>
10:45	11:00	ACHIK DORCHIN The evolution of floral host preference in longhorn bees of the genus <i>Eucera</i> (Hymenoptera, Apidae, Eucerini): is association with pollen from bee-flowers advantageous?	SMADAR BEN-TABOU DE-LEON Hypoxic conditions distort sea urchin skeletal patterning through the BMP and VEGF signaling but not Hif1 α , demonstrating divergent and convergent evolution modes	EUGENE ROSENBERG & ILANA ZILBER-ROSENBERG The hologenome concept of evolution after 12 Years
11:00	11:15	EYAL ZINGER Less fit <i>Lamium amplexicaule</i> plants produce more dispersible seeds	ARSENIY MOROV Asymmetrical distribution of maternal nodal mRNA in amphioxus embryos by means of Arp2/3 complex	
11:15	11:30	NETTA DORCHIN Evolution of feeding modes and diversification rates in the megadiverse gall midges (Diptera: Cecidomyiidae)	NOA AHARON-HEFETZ The role of tRNAs in the cellular proliferation state regulation	EHUD LAMM How can holobionts evolve?
11:30	11:45	MONICA A MOWERY Newly-established brown widow spiders are more dispersive and	ARIE FRIDRICH Evolution and recurrent subfunctionalization of Argonautes in Metazoa revealed	OHAD LEWIN-EPSTEIN Host-microbiome coevolution and cooperative behavior

		less fecund across multiple invasion fronts	by a functional study in sea anemones	
11:45	12:45	Lunch Break		

Hour		Hall	Room 1	Room2
12:45	13:30	PLENARY 5 = JOANNA MASEL De novo genes and the long-term direction of their subsequent evolution		
13:30	13:50	Coffee Break		
		Evolutionary Ecology 2 (Chair: Oren Kolodny)	Systems Biology & EvoDevo 2 (Chair: Smadar Ben Tabou de Leon)	Phylogeny (Chair: Dorothee Huchon)
13:50	14:05	ALLY HARARI Male effects on the evolution of monandrous mating strategy in moths	HAGAI SHPIGLER The evolution of extreme fertility in advanced eusocial bees and modifications in gonadotropic hormone signaling to reduce brain cost	KSENIA JURAVEL Sequence-independent phylogeny for resolving the non-bilaterian animals relationships
14:05	14:20	MICHAL SEGOLI The combined effect of host and food availability on optimized parasitoid life history traits based on a three-dimensional trade-off surface	GAT KRIEGER Comparative analysis of yeast species reveals transcriptional rewiring in cis and in trans	KEREN HALABI A codon model for associating phenotypic traits with altered selective patterns of sequence evolution
14:20	14:35	FRANKLIN SARGUNARAJ Stress as an adaptation: Does experimental cortisol supplementation affect predation risk assessment in foraging gerbils?	JUDY WEXLER Hemimetabolous insects elucidate the origin of sexual development via alternative splicing	SHIRAN ABADI ModelTeller: machine-learning based model selection for optimal phylogenetic reconstruction
14:35	14:50	LEE KOREN Testosterone is associated with male and female copulation success in opposite ways in wild rock hyrax	AMIR SAPIR Metabolic reconfiguration facilitates widespread sterol auxotrophy during the evolution of the animal kingdom	ANNA RICE Model adequacy for likelihood models of chromosome-number evolution
14:50	15:05	AMIYAAL ILANY Social inheritance shapes spotted hyena social networks and influences offspring longevity	ARIEL CHIPMAN Developing an integrated understanding of the evolution of arthropod segmentation using fossils and evo-devo	ASHER MOSHE Probabilistic models for genome rearrangements
15:05	15:20	ROI HOLZMAN Using performance landscapes to understand adaptive diversification within fish	OREN LEV Anterior Segmentation in Arthropods	DANA AZOURI The phylogenetic tree reconstruction game: developing reinforcement-learning algorithms for fast and accurate inference of evolutionary trees
15:20	15:35	VICTOR CHINA Hydrodynamic constraints shape the performance landscape of the feeding mechanism of fish larvae during the critical period	YONAT GURVICH Evolution through reverse transcription: a potential means to employ Lamarckian-like adaptations	BENJAMIN YAKIR Estimating relatedness in ancient DNA samples
15:35	16:50	Poster session		
16:50	17:30	Society meeting + Poster prizes		
17:30		End		