GLAM-Evogen

Syracuse University, August 24, 2024

8:30am - 9:25am Registration - Breakfast/coffee			
9:30am - 10:45am - Talk Session 1			
9:30 am - 9:45 am	<i>Charikleia Karageorgiou</i> (University at Buffalo) Paleolithic gene duplications primed adaptive evolution of human amylase locus upon agriculture		
9:45 am - 10:00 am	<i>Kwondo Kim</i> (The Jackson Laboratory) An Alu insertion in OCA2 may have contributed to light pigmentation in Denisovans and present-day Melanesians		
10:00 am - 10:15 am	Tharindu Premachandra (McMaster University) Sex chromosome conservation through multiple allotetraploidization events in the African clawed frog subgenus Silurana, genus Xenopus		
10:15 am - 10:30 am	Brian Kirz (University of Rochester) Insights into rapid genome expansion in the North American red squirrel		
10:30 am - 10:45 am	<i>Jiarun Chen</i> (Columbia University) Polygenic Adaptation under Recurrent Changes in Environment		
10:45am - 11:15am Coffee Break			
11:15am - 12:15pm - Talk Session 2			
11:15am - 11:30am	Joy Zhichun Zhang (Cornell University) Bayesian phylodynamic inference of population dynamics with dormancy		
11:30am - 11:45am	Daniel Seidman (University of Rochester) Investigating mutation rates in pedigreed populations using low coverage data		
11:45am - 12:00pm	<i>Jeremy Summers</i> (University of Rochester) The multigenerational impact of gene flow in a natural population		

12:00am - 12:15pm	Peiyu (Perry) Xu (Cornell University) e3SIM: A novel epidemiological-ecological-evolut framework for genomic epidemiology	ionary simulation	
12:15pm – 1:15pm Lunch			
1:15pm – 2:30pi	m Poster session	Avantor	
2:45pm -3:45pm - Talk Session 3			
2:30pm - 2:45pm	Jolie Carlisle (Cornell University) Evolutionary and Genetic Investigation of Male ar Control of the <i>Drosophila melanogaster</i> Mating P		
2:45pm - 3:00pm	<i>Martin Knytl</i> (McMaster University) Synthetic sex chromosomes generated by CRISF Haplosufficiency mediates the origin of a new Y of concomitant extinction of an ancestral W chromos	hromosome and	
3:00pm - 3:15pm	Antonio Gomez (Syracuse University) Evolution of ejaculate structures		
3:15pm - 3:30pm	Sushant Potdar (Cornell University) Neurogenomic causes and proteomic consequen preference learning in two butterflies	ces of male mate	
3:30pm - 3:45pm Break			
3:45pm - 4:00pm	You Chen (Cornell University) Directionality of transcriptional regulatory element	ts	
4:00pm – 4:15pm	<i>Zhaozhi Li</i> (Cornell University) Multi-drug Cancer Adaptive Therapy with Deep R	einforcement Learning	
4:15pm - 5:00pm Keynote Shaila Musharoff (Cornell University) Admixture as a lens into human complex trait architecture			
5:30pm - Adjourn + Happy hour at Harvey's Garden (<u>https://www.harveysgardensyr.com</u>)			

POSTER PRESENTATIONS:

1	Benjamin C. Haller (Cornell)	SLiM: evolutionary simulation software
2	Bryan Guo (Toronto)	Functional Impact of Extracellular Loop 2 Motif in Cetacean Rhodopsin
3	Sarah Kassem (Buffalo)	Deciphering the copy number variation in the Amylase Locus in Nile Rats shed light on diet-induced diabetes
4	Alycia Johnson (Rochester)	Genome-wide scans for selection reveal the impact of climate in a widely-distributed social wasp
5	Dominic Hockenbury (ESF)	Characterizing BGLUC in Drosophila melanogaster
6	Emily Yarbrough (Binghamton)	An atrazine gradient differentially affects patterns of sexual dimorphism for Drosophila melanogaster development time
7	Lindsey Siao (Buffalo)	Unraveling global patterns of pepsinogen gene copy number variation provides insights into human dietary evolution
8	Amanda Ivanoff (Buffalo)	Investigating the mechanism of evolution of an alternative parturition signal
9	Freddie Xu (Cornell)	A screen for modifiers in the sex determination pathway of Drosophila melanogaster
10	Henna Domian (Cornell)	The Impact of Socioeconomic Status on Breast Cancer Polygenic Score Performance
11	Jaclyn Liquori (Cornell)	Sexual Dimorphism in LDL-C Measurements and Gender Bias in Statin Treatment Rates
12	Sylvia Han (Cornell)	A novel identity-by-descent approach for detecting genotype errors in large sequencing datasets
13	Luane Jandira Bueno Landau (Buffalo)	Ancestral Mammalian Salivary Gene Expression: A Comparative Transcriptomic Analysis
14	Logan Edvalson (Rochester)	Satellite DNA regulation & the mechanism of Segregation Distorter
15	Tyler Handler (Rochester)	Genomic mapping of the suppressor of Segregation Distorter in D. melanogaster
16	Ben McCormick (Cornell)	Evidence consistent with the Stellate array of Drosophila melanogaster being an active meiotic drive system
17	Ryan Chaffee (Cornell)	Developing a model for conservation of animal populations experiencing human- induced habitat fragmentation
18	Oona Shigeno Risse- Adams (Cornell)	The environmental effect of social determinants of health and race/ethnicity on human trait and disease outcomes