VIIth Bay Area Population Genomics Conference

Sponsored by Ancestry.com, Stanford Center for Computational, Evolutionary, and Human Genomics, and EcoEvo group of the Stanford Biology Department

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Registration is FREE but REQUIRED

Date: February 2, 2013 (Saturday),

Place: <u>Stanford University</u>, Herrin Hall T175

Sign up RSVP by <u>Attendee List</u> January 15

9:00 AM

CONTACT: Dmitri Petrov; dpetrov@stanford.edu

Breakfast Buffet

SCHEDULE

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	9:30 AM	Kirk Lohmueller, Nielsen/Lohmueller Labs, Berkeley/UCLA	Exome sequencing and heterogeneity of complex trait		
	9:50 AM	Ben Callahan, Fisher Lab, Stanford	Experimental evolution of niche construction		
	10:10 AM	Joshua Schraiber, Slatkin Lab, Berkeley	Theory and application of Wright-Fisher diffusion bridges		
	10:30 AM	Coffee Break	reak		
	10:50 AM	Pleuni Pennings, Petrov Lab, Stanford	Soft and hard selective sweeps in HIV		
	11:10 AM	Anand Bhaskar, Song Lab, Berkeley	Exact comparison between Kingman's coalescent and discrete Wright- Fisher model for large sample sizes		
	11:30 AM	Qi Zhou, Bachtrog Lab, Berkeley	Atlas of avian sex chromosome evolution		
	11:50 AM	Lunch (<u>register</u> if you would like to get food!) and Posters			
	1:00 PM	Nandita Garud, Petrov Lab, Stanford	Evidence that recent adaptation in Drosophila is primarily driven by soft sweeps		
	1:20 PM	Jeremy Roop, Brem Lab, Berkeley	Rare variants drive common traits in S. paradoxus		
	1:20 PM	Hunter Fraser, Fraser Lab, Stanford	Gene expression drives local adaptation in humans		
POSTERS:					
		Scott Roy, Roy Lab, SFSU	Many different solutions to the same problem: subcellular targeting of MDH genes in fungi		
		Joel Thompson, Fisher Lab, Stanford	Evolutionary Dynamics and Diversity in Microbial Populations		
		Rajiv McCoy, Boggs/Petrov	Evaluating methods of demographic inference using genomic data from the checkerspot butterfly Euphydryas gillettii		
		Alan Bergland, Petrov Lab, Stanford	Genomic evidence of rapid and stable adaptive oscillations over seasonal time scales in Drosophila.		
		David Lawrie, Petrov Lab, Stanford	Strong Purifying Selection at Synonymous Sites in D. melanogaster		

Sandeep Venkataram, Petrov Lab, Stanford Epistasis and Repeatability of evolution in Fisher's Geometric Model

<u>Eco / Evo</u>	Ancestry.com	BAPG Google Group	<u>Map</u>	<u>Stanford</u> <u>Biology</u> <u>Department</u>
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