Statistical Genetics Seminar Series

Room: Crosstown 305 unless change is announced Time: 9:30 ~11:00 AM unless change is announced

Date	Торіс	Presenter	Note
9/9/2016	Brainstorm Session	Everyone	
	Dissecting Genetic Architecture of Complex Diseases Through Integrated Genomic Analysis	Hongyu Zhao	Yale U.
10/7/2016	CHARGE debrief	Sam & Virginia	
10/21/2016	ASHG meeting - no statistical genetics seminar		
11/4/2016	Obesity RNA-seq differential expression analysis within post- mortem human brain	Chris Wake	
11/18/2016	Exploring gene-environment interaction by a twin structure equation model	Biqi Wang	
12/2/2016	analysis of data from single cell sequencing	Josh Campbell	
12/16/2016	metabolism and microbiome	Daniel Segre	
12/30/2016	Break - no statistical genetics seminar		
1/13/2017	Proteomic analysis of CVD associated proteins	Chen Yao	NHLBI
1/27/2017	Analysis of metabolite data	Martin Larson and Qiong Yang	
2/10/2017	Interrogation of Lipoprotein Lipase pathway	Gina Peloso	
2/24/2017	Parent of origin analyses	Chloé Sarnowski	
3/10/2017	Genetic Analysis Workshop debrief	ALL	
3/24/2017	CHARGE Meeting - no statistical genetics seminar		
4/7/2017	Deep sequencing of the mitochondrial genome	Chunyu Liu	NHLBI
4/21/2017	High-throughput path analysis based on network models to explore disease mechanism	Yuanjia Wang	Columbia U.
5/5/2017	Integration of function annotation	Virginia Fisher	
5/19/2017	Do mouse models mimic genomic response in human transcriptomic system? - An answer from Bayesian perspective	George Tseng	U. of Pittsburg