

Room: Crosstown 305 unless change is announced			
Time: 9:30 ~11:00 AM unless change is announced			
Date	Topic	Presenter	Note
9/10/2010	Brainstorm	Everyone	
9/24/2010	Determining MicroRNA-Gene Interactions Via Evolutionary Monte Carlo-Based Variable Selection	Julius Ngwa	
	Analysis of micro RNAs in the Framingham Study	Martin Larson	
10/8/2010	Finding rare variants in family studies: for better or worse?	Cornelia van Duijn	Netherlands
10/22/2010	Gene x Environment Interaction	Alisa Manning	
11/5/2010	ASHG meeting in DC		ASHG
11/19/2010	Genetic Analysis Workshop (GAW) debrief	All GAW first authors	
12/3/2010	Analysis of DNA methylation data	Xiaoyan Yin	
12/10/2010			
2:00PM at CT462	Leveraging age information to increase power in association studies	Alkes Prices	Harvard
12/17/2010	Quality control for sequence data	Ron Do and Nathan Stitzel	Broad
12/31/2010	Happy New Year Eve		
1/14/2011	Mapping/aligning sequence data	John Brother/Becky Kusko	
1/28/2011	Analysis of Exome Sequencing Data	Honghuang Lin	
2/11/2011	CHARGE Meeting		
2/25/2011	Cancel for School Vacation week		
3/11/2011	Reanalysis strategy using published dbGAP datasets in age-related macular degeneration	Gyungah Jun	
3/25/2011	Network analysis defines the impact of gene-environment interactions on metabolic diseases	Larry Parnell	Tufts
4/8/2011	New phenotypes in Alzheimer's disease & homozygosity	Rick Sherva	
4/22/2011	Analysis of RNA-seq	Jacqui Milton	
5/6/2011	mitochondrial DNA mutation and metabolic traits	Chunyu Liu	NHLBI
5/20/2011	Topic to be determined	Shamil Sunyaev	Harvard
6/3/2011	Incorporating phenotyped but ungenotyped individuals into genetic association tests with family data	Vivian Zhuang	