

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

| Date | Topic | Presenter | Note |
|------------|---|------------------------|---------|
| 9/9/2011 | Brainstorm | Everyone | |
| 9/23/2011 | Rare variant analysis and expression data | Han Chen | |
| 10/7/2011 | Analysis of sequencing data (rare variants) | Adrienne Cupples | |
| 10/21/2011 | Replication analysis and the implication of related samples | Stephen Hartley | |
| 11/4/2011 | QC of sequencing data | Mayetri | |
| 11/18/2011 | MRI analysis | Scott Melville | |
| 12/2/2011 | Analysis of whole genome data or genetic risk prediction | Paola | |
| 12/16/2011 | Mitochondrial analysis | Chunyu Liu | NHLBI |
| 12/30/2011 | Winter Break | Winter Break | |
| 1/13/2012 | Latent variable models for assessing individual drug sensitivity and for predicting hereditary disease risk | Evan Johnson | |
| 1/27/2012 | To be determined | Xihong Lin | Harvard |
| 2/10/2012 | Joint analysis of family and unrelated | Chuanhua Xing | |
| 2/24/2012 | Contrasting associations in different ethnic groups | Ching-Ti | |
| 3/9/2012 | Bayesian method in GWAS | Luis Carvalho | |
| 3/23/2012 | Bivariate genome-wide association analysis to identify shared genetic risk factors related to Alzheimer's Disease | Gyungah Jun | |
| 4/6/2012 | Methylation | Dawn Demeo | Harvard |
| 4/20/2012 | G x E Interaction and CVD risk | Chao-Qiang Lai | Tufts |
| 5/4/2012 | To be determined | Harold Bae | |
| 5/18/2012 | Mendelian Randomization | Julius Ngwa | |
| 6/1/2012 | Gene-gene & gene-environment interaction approaches | Chen Lu & Josee Dupuis | |
| 6/15/2012 | Multiple trait analysis | Jacqui | |