

# Agenda 2016

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**9:00 a.m. Breakfast**

**9:15 a.m. Welcome**

Jeff Leek, Ph.D.  
Associate Professor  
Department of Biostatistics  
Johns Hopkins Bloomberg School of Public Health  
*Speaker at 3<sup>rd</sup> symposium*

**9:30 a.m.**

Jonathan Pevsner, Ph.D.  
Professor  
Department of Neurology,  
Kennedy Krieger Institute  
Department of Psychiatry and Behavioral Sciences  
Johns Hopkins University School of Medicine  
Somatic mutation in disease:  
from Sturge-Weber syndrome to autism

**10:30 a.m.**

Hongkai Ji, Ph.D.  
Associate Professor  
Department of Biostatistics  
Johns Hopkins Bloomberg School of Public Health  
Genome-wide Prediction of DNase I  
Hypersensitivity Using Gene Expression

**11:00 a.m. Break**

**11:10 a.m.**

Luigi Marchionni, M.D., Ph.D.  
Assistant Professor  
Oncology Cancer Biology  
Sidney Kimmel Comprehensive Cancer Center  
Johns Hopkins University School of Medicine  
The restaurant at the end of the "omics"  
universe – What to order and what not...

**11:40 a.m.**

Margaret A. Taub, Ph.D.  
Assistant Scientist  
Department of Biostatistics  
Johns Hopkins Bloomberg School of Public Health  
Megakaryocytes derived from iPSCs show  
concordant eQTLs across racial groups and  
platelet-related signature of CD41+CD42a+  
percentage

**12:10 p.m.**

Lightning Poster Session presentations

**12:20 p.m.**

Lunch (provided)

**1:00 p.m.**

Andrew Jaffe, Ph.D.  
Lead Investigator  
Lieber Institute for Brain Development  
Developmental and genetic regulation of  
the human frontal cortex transcriptome  
in schizophrenia

**1:30 p.m.**

Héctor Corrada Bravo, Ph.D.  
Associate Professor  
Center for Bioinformatics and Computational Biology  
UMIACS  
Department of Computer Science  
University of Maryland  
Statistics and visualization for metagenomic  
data analysis

**2:00 p.m. Break**

**2:20 p.m.**

Carlo Colantuoni, Ph.D.  
Investigator, Genome Informatics,  
Lieber Institute for Brain Development  
Assistant Professor,  
Department of Neurology  
Johns Hopkins University School of Medicine  
Genome Dynamics in Pluripotent Stem Cells:  
Can we define stable variation in individual  
human cell biology?

**2:50 p.m. Talk Wrap-Up**

**3:00 p.m.**

Giovanni Parmigiani, Ph.D.  
Professor  
Department of Biostatistics  
Dana-Farber Cancer Institute  
Cross-study Analysis of Prediction Algorithms  
in Genomics

**4-5 p.m. Poster Session - Greenhouse Cafe**