



Genotypes & Intermediate Phenotypes

Room Lanier Ballroom DE, Level 4
Hilton Americas Houston
Houston, TX, USA

15th October 2019

(a satellite of ASHG)

7:15 - 8:00 Registration

8:00 - 8:05 Welcome

Session 1 **Chair: Marc Greenblatt**

8:05 - 8:35 **Using Gene Expression To Interpret Common And Rare Genetic Disease**

Stephen Montgomery
Stanford University School of Medicine

8:35 - 9:05 **Trans-Regulation of Gene Expression Can Drive Omnigenic Inheritance of Human Complex Traits**

Jonathan Pritchard
Stanford University

9:05 - 9:35 Title TBA

Kaitlin Samocha
Wellcome Sanger Institute

9:35- 10:05 Title TBA

David Craig
Institute of Translational Genomics, USC Keck School of Medicine

10:05 - 10:10 **RAPID FIRE PRESENTATIONS FROM ABSTRACTS**

10:10 - 10:40 **Coffee Break**

Session 2 **Chair: Steven Brenner**

10:40 - 11:05 **Using Co-segregation Analysis and Other Family Data to Evaluate Intermediate Phenotypes**

Brian Shirts

University of Washington Medical Center

11:05 - 11:30 **Exploring The Relationship Between the Genome and the Phenome Through PheWAS**

Marylyn Ritchie

University of Pennsylvania

11:30 - 11:55 Title TBA

Stanley F. Nelson

David Geffen School of Medicine at UCLA

11:55 - 12:00 **Summary and Closing Remarks**

12:00 **MEETING END**

12:00 - 12:45 **HGVS Annual General Meeting**