BROAD INSTITUTE | ASHG 2018

BOOTH 1634 | THURSDAY, OCTOBER 18TH

San Diego Convention Center, San Diego, CA

Expo Hall Booth #1634

10:00 AM - 4:30 PM
Genomic Sequencing Services
Jane Wilkinson & Andy Hollinger

10:00 AM - 4:30 PM
Broad Institute Recruiting
Andrea Petrosino

10:00 AM -12:00 PM
Broad Institute Strategic Alliances and Partnering
Nate Kurtis

10:00 AM -12:00 PM
All of Us Genome Center Q & A
Stacey Gabriel

10:30 AM - 1:00 PM
Rare Genomes Project
Clara Williamson
Dr. Anne O'Donnel Luria

1:00 PM - 2:00 PM
Q & A with Heidi Rehm
Heidi Rehm

1:00 PM - 2:00 PM
Session Q & A - Analysis of 33,527 haploid sperm genomes from 20 individuals reveals new relationships underlying meiotic recombination and aneuploidy
Avery Davis Bell

2:00 PM - 4:00 PM
Poster Q & A - Genome-wide polygenic scores for common diseases identify individuals with risk equivalent to monogenic mutations
Amit Khera

3:30 AM - 4:30 PM
Hail Q & A
Tim Poterba

Sessions

9:00 AM - 9:15 AM
PgmNr 143: Insights into effective methods for Mendelian gene discovery from family-based genomic analysis of over 22,000 families from worldwide populations.
Ballroom 20D - Upper Level
Anne O'Donnell-Luria

9:00 AM - 9:15 AM
PgmNr 173: Using untargeted metabolomics and Mendelian randomization to dissect causal relationships in the obesity metabolome.
Room 6E - Upper Level
Yu-Han Hsu

9:15 AM - 9:30 AM
PgmNr 132: Association studies for all: A novel framework to allow for the well-calibrated genomic analysis of underrepresented admixed individuals.
Ballroom 20A - Upper Level
Elizabeth Atkinson

9:15 AM - 9:30 AM
PgmNr 144: The genetic landscape of Diamond-Blackfan anemia.
Ballroom 20D - Upper Level

Jeffrey Verboon

9:45 AM - 10:00 AM
PgmNr 176: Analysis of 33,527 haploid sperm genomes from 20 individuals reveals new relationships underlying meiotic recombination and aneuploidy.
Room 6E - Upper Level
Avery Bell

10:15 AM - 10:30 AM
PgmNr 142: Identification and characterization of adaptive regulatory variation in diverse human populations.
Ballroom 20BC - Upper Level
Joseph Vitti

10:15 AM - 10:30 AM
PgmNr 148: Diagnostic yield from WES, WGS and RNA testing among 213 neuromuscular families: Known versus novel disease genes, coding versus non-coding variants.
Ballroom 20D - Upper Level
Leigh Waddell

11:00 AM - 11:15 AM
PgmNr 203: Comparative genetic architectures of schizophrenia in East Asian and European populations.
Room 6A - Upper Level
Hailiang Huang

11:15 AM - 11:30 AM
PgmNr 204: Exome sequencing of 23,851 cases implicates novel risk genes and provides insights into the genetic architecture of schizophrenia.
Room 6A - Upper Level
Tarjinder Singh

11:30 AM - 11:45 AM
PgmNr 193: Human lineage tracing enabled by mitochondrial mutations and single cell genomics.
Ballroom 20BC - Upper Level
Leif Ludwig

11:30 AM - 11:45 AM
PgmNr 211: Tissue-specific molecular signatures associated with 16p11.2 reciprocal genomic disorder.
Room 6B - Upper Level
Parisa Razaz

12:00 PM - 12:15 PM
PgmNr 189: Analysis of 5'UTR variation in WGS from over 18,000 individuals identifies highly constrained regulatory elements and a role in human disease.
Ballroom 20A - Upper Level
Nicola Whiffin

4:14 PM - 4:30 PM
ASHG William Allan Award Presentation and Lecture: The Genetics of Common Disease: From Variants to Function
Hall C - Ground Level
Eric Lander