

Thursday, 23 June

09:00-10:15

Session 3: Evolving guidelines/resources to support variant assessment continued

Chair: Matt Hurles

09:00 Integration of diverse data sources to aid clinical genome interpretation

Michael Simpson

Genomics plc. UK

09:15 Comprehensive interpretation guidelines for clinical genomics; a UK clinical science perspective Dominic McMullan Birmingham Womens Hospital, UK

09:30 Classification and clinical management of Variants of Uncertain Significance in high penetrance cancer predisposition genes

Christi van Asperen

Leiden University Medical Center, The Netherlands

09:45 Identification of novel pathways linked to Bardet-Biedl Syndrome (BBS) using a multi-omic strategy (HIGH5 Project) Phil Beales UCL, UK

10:15-10:45

Morning Coffee

10:45-12:45

Session 4: Patient reported data

Chair: Beverly Searle

- 10:45 Prospective Registry of Multiplex Testing (PROMPT): a web-based platform for patients, families, and researchers seeking to assess cancer risk of genomic variants

 Mark Robson

 Prompt, USA
- 11:15 Attitudes towards genomic data sharing: Your DNA, Your Say Anna Middleton Wellcome Trust Sanger Institute, UK
- 11:45 GenIDA (Genetics of Intellectual Disabilities and Autism spectrum disorders): an online registry and clinical database for patients, families and professionals Florent Colin IGBMC, France
- 12:15 Smart phone phenotyping Euan Ashley Stanford University, USA

12:45-14:00

Lunch