

**Genomics of Rare Disease  
27-29 March 2019**

**Wellcome Genome Campus,  
Hinxton, Cambridge, UK**

**Draft Conference Programme**

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**Wednesday 27 March 2019**

12:00-13:30	<b>Registration with lunch</b>
13:30-13:40	<b>Welcome and introduction</b> <i>Programme Committee</i>
13:40-14:40	<b>Lupski lecture</b>  <i>Nancy Cox</i> <i>Vanderbilt University Medical Center, USA</i>
14:40-16:00	<b>Session I: Solving the unsolved</b>  14:40 Harnessing transmission genetics to build models of complex disease from the ground up <i>Jennifer Posey</i> <i>Baylor College of Medicine, USA</i>  15:20 The NIH undiagnosed diseases program: expansion to a national and international networks <i>William Gahl</i> <i>NIH, USA</i>
16:00-16:40	<b>Afternoon tea</b>
16:40-17:10	<b>Session I continued: Solving the unsolved</b>  16:40 Short talk selected from submitted abstract  16:55 Short talk selected from submitted abstract
17:10-17:45	<b>Lightning talks</b>
17:45-19:15	<b>Poster session I (odd numbers) with drinks reception</b>
19:15	<b>Dinner</b>

## Thursday 28 March 2019

09:00-10:30

### Session 2: Informatics

- 09:00 Regulatory variant in rare disease: causal and modifier effects  
*Tuuli Lappalainen*  
*Genome Center, USA*
- 09:40 Short talk selected from submitted abstract
- 09:55 Short talk selected from submitted abstract
- 10:10 Short talk selected from submitted abstract
- 10:25 Short talk selected from submitted abstract

10:25-11:00

### Morning coffee

11:00-12:25

### Session 3: Developmental genetics

- 11:00 Rare disorders caused by mutations in genes encoding epigenetic regulators: novel diagnostic opportunities  
*Rosanna Weksberg*  
*The Hospital for Sick Children, Canada*
- 11:40 Short talk selected from submitted abstract
- 11:55 Short talk selected from submitted abstract
- 12:10 Short talk selected from submitted abstract

12:25-14:00

### Lunch

14:00-15:30

### Session 4: New therapeutic approaches

- 14:00 P4 medicine for intellectual disabilities: integrated -omics the way forward?  
*Clara van Karnebeek*  
*Amsterdam University Medical Centres, The Netherlands*
- 14:40 Hope for escape from a prison of bone  
*Frederick Kaplan*  
*University of Pennsylvania, USA*
- 15:20 Short talk selected from submitted abstract
- 15:35 Short talk selected from submitted abstract

15:50-16:30

### Afternoon tea

16:30-17:30

**Session 5: Electronic health records**

- 16:00 Assessing the pathogenicity, penetrance and expressivity of apparently monogenic disease variants in UK Biobank  
*Caroline Wright*  
*University of Exeter, UK*
- 16:40 Genomic disorders in the general population. What can we learn and how can we help?  
*Katrin Mannik*  
*University of Lausanne, Switzerland*
- 17:20 Short talk selected from submitted abstract
- 17:35 Short talk selected from submitted abstract

17:50-18:20

**Lightning talks**

18:20-19:50

**Poster session 2 (even numbers) with drinks reception**

19:50

**Conference dinner**

**Friday 29 March 2019**

08:30-10:20

**Session 6: Data sharing – legal and ethical issues**

- 08:30 Advancing medicine through data sharing and collaboration on a global scale  
*Heidi Rehm*  
*Broad Institute, USA*
- 09:10 Talk title TBC  
*Ewan Birney*  
*EMBL-EBI, UK*
- 09:50 Short talk selected from submitted abstract
- 10:05 Short talk selected from submitted abstract

10:20-11:00

**Morning coffee**

11:00-12:50

**Session 7: Functional genomics**

- 11:00 Development and disease at single cell resolution  
*Malte Spielmann*  
*Max Planck Institute for Molecular Genetics, Germany*
- 11:40 Functional characterization and therapeutic targeting of gene regulatory elements  
*Nadav Ahituv*  
*UCSF, USA*

- 12:20 Short talk selected from submitted abstract
- 12:35 Short talk selected from submitted abstract
- 12:50-14:15 **Lunch**
- 14:15-16:05 **Session 8: Intersection of polygenic and monogenic**
- 14:15 The Finnish population isolate: examples of polygenic and rare variants in brain diseases  
*Aarno Palotie*  
*FIMM, University of Helsinki, Finland*
- 14:55 Short talk selected from submitted abstract
- 15:10 Short talk selected from submitted abstract
- 15:25 Using polygenic scores to define lifetime trajectories of disease  
*Michael Inouye*  
*University of Cambridge, UK*
- 16:05-16:15 **Closing remarks**  
*Programme Committee*
- 16:25 **Coaches depart to Cambridge city centre via train station, and Heathrow airport via Stansted airport**