

# Genomics England Research Conference Agenda

4 November 2019

**08:00** Room: Entrance Hall

Registration opens

**09:00–09:15** Location: Great Hall

Welcome

*Chris Wigley* (CEO, Genomics England)

**09:15–09:20** Location: Great Hall

Opening address

*Matt Hancock* (Secretary of State for Health and Social Care)

**09:20–10:05** Location: Great Hall

Keynote plenary: Drug Development in the Era of Precision Medicine

*Professor David Goldstein* (Director of the Institute for Genomic Medicine, Columbia University)

**10:05–10:20** Location: Great Hall

Keynote sponsor

*Hans Cobben* (CEO, Bluebee)

**10:20–10:50** Coffee/tea break (with posters) Location: Library & Lecture Hall

## Parallel session 1: Beyond the Coding Regions

Chairs: *Professor Tim Hubbard* (Genomics England) and *Dr. Freya Boardman-Pretty* (Genomics England)  
Location: *Robert Perks*

**10:50–11:05**

Investigating the Microbiome of the 100,000 Genomes Colorectal Cancer Cohort

*Dr. Henry Wood* (University of Leeds)

**11:05–11:20**

Filtering Artefacts in Somatic Single Nucleotide Variant Calling Using a Panel of Normals

*Dr. Boris Noyvert* (University of Birmingham)

**11:20–11:35**

Real World Experience of Whole Genome Sequencing (WGS) in Haematological Malignancies

*Dr. Angela Hamblin* (Oxford Genomic Medicine Centre)

**11:35–11:45**

Xdrop™ - Targeted Sequencing into the Dark and Unknown

*Dr. Marie Mikkelsen* (Samplix)

## Parallel session 2: Diagnostic Discovery

Chair: *Dr. Anna Need* (Genomics England)

Location: *George Thomas*

**10:50–12:25**

Discussion of how to identify novel potential diagnoses in the Research Environment with case studies and examples

*Professor Krishna Chatterjee* (Institute of Metabolic Science, University of Cambridge)

*Dr. Jana Vandrovцова* (UCL Queen Square Institute of Neurology Department of Neurodegenerative Disease)

*Dr. Jenny Taylor* (Wellcome Centre for Human Genetics, Oxford)

*Dr. Alistair Pagnamenta* (Wellcome Centre for Human Genetics, Oxford)

*Dr. Hywel Williams* (Cardiff University)

*Dr. Guillermo del Angel* (Alexion)

**12:25–12:30**

How Tos & low-hanging fruit: Inspire and encourage Discovery Forum and GeCIP researchers to do diagnostic discovery work

*Dr. Anna Need* (Genomics England)

## Parallel session 3: Precision Clinical Studies in Cancer

Chairs: *Dr. Maria Antonietta Cerone* (Cancer Research UK) and *Dr. Lea Lahnstein* (Genomics England)

Location: *Great Hall*

**10:50–11:05**

Patient Stratification in Chronic Lymphocytic Leukaemia - Using Whole Genome Sequencing and Machine Learning to Predict Patient Outcomes in Cancer

*Dr. Kate Ridout* (University of Oxford)

**11:00–11:25**

The frontline of molecular tumour characterisation for precision medicine in cancer: How Individum developed its IndivUType, a truly multiomics and global cancer database for R&D

*Dr. Hartmut Juhl* (Indivumed)

**11:25–11:40**

Identifying fusion protein variants in 100,000 Genomes Project cancer participants for clinical trials

*Dr. John Bridgewater* (GeCIP PI)

**11:40–11:50**

Genomics England and precision clinical studies: short introduction and discussion

*Frank Nankivell* (Genomics England)

## Bioinformatics drop-in clinic

Location: *Library*

Chair: *Dr. Loukas Moutsianas* (Genomics England)

Drop-in clinic for questions on bioinformatics and tools for working in the Genomics England Research Environment



<b>Parallel session 1: Beyond the Coding Regions</b> <i>Location: Robert Perks</i>	<b>Parallel session 2: Diagnostic Discovery</b> <i>Location: George Thomas</i>	<b>Parallel session 3: Precision Clinical Studies</b> <i>Location: Great Hall</i>	<b>Bioinformatics drop-in clinic</b> <i>Location: Library</i>  <i>Chair: Dr. Loukas Moutsianas (Genomics England)</i>  <b>Drop-in clinic for questions on bioinformatics and tools for working in the Genomics England Research Environment</b>	
<b>11:45–11:55</b> <b>Genome-Wide Map of Selective Constraint from 28,000 Deep Whole Genomes Informs New Model for Non-Coding Pathogenicity</b> <i>Patrick Short (Wellcome Sanger Institute)</i>		<b>11:50–12:05</b> <b>Germline Pharmacogenomics: Analysis and feedback of variants to avoid drug toxicity</b> <i>Dr. Ellen McDonagh (Genomics England)</i>		
<b>11:55–12:15</b> <b>Whole Genome Sequencing of Primary Immunodeficiency Reveals a Role for Common and Rare Variants in Coding and Non-Coding Regions</b> <i>Professor Ken Smith (University of Cambridge)</i>		<b>12:05–12:20</b> <b>Validity of Whole Genomes Sequencing Results in Neoplasms with a High Tumour Mutational Burden in Precision Medicine</b> <i>Dr. Marc Ooft (King's College Hospital)</i>		
<b>12:15–12:30</b> <b>Functional and <i>In-Silico</i> Interrogation of Rare Genomic Variants Impacting RNA Splicing for the Diagnosis of Genomic Disorders</b> <i>Dr. Jamie Ellingford (University of Manchester)</i>		<b>12:20–12:30</b> <b>General discussion on precision studies in cancer and the role of whole genome sequencing</b> <i>Dr. Maria Antonietta Cerone (Cancer Research UK) &amp; Dr. Lea Lahnstein (Genomics England)</i>		
<b>12:30–13:20 Lunch (with posters)</b> <i>Location: Library &amp; Lecture Hall</i>				

<b>Connecting patients, participants and researchers as partners</b> <i>Location: Great Hall</i>	
<i>Chair: Vivienne Parry (Genomics England)</i>	
<b>13:20–13:35</b>	
<b>1. A parent's patient journey</b> <i>Emma Walters (100,000 Genomes Project participant)</i>	
<b>2. Recent PPI activity between a patients' group and the relevant GeCIP domain</b> <i>Fiona Copeland (GeCIP PPI representative)</i>	
<b>3. Building a multidisciplinary community for better research – where the future might lie</b> <i>Jillian Hastings Ward (Chair of the Genomics England National Participant Panel)</i>	
<b>13:35–14:20</b>	
<b>Panel discussion and open Q&amp;A</b> <i>Professor Claire Shovlin (Respiratory GeCIP domain lead)</i> <i>Dr David Church (Endometrial Cancer GeCIP domain lead)</i> <i>Fiona Copeland (GeCIP PPI representative)</i> <i>Helen White (Genomics England National Participant Panel member)</i>	



**Parallel session 1: Cancer Landscapes**

Chair: *Tahrima Rahim (Genomics England)*  
Location: *Robert Perks*

**14:25–14:40**

**The Genetic Immune Landscape of Colorectal Cancers**

*Dr. Eszter Lakatos (Queen Mary University of London)*

**14:40–14:55**

**Assessing the Mutational Landscape of Epithelial Ovarian Cancer Using Whole Genome Sequencing Data from the 100,000 Genomes Project**

*Dr. Mark Evans (University of Sheffield)*

**14:55–15:10**

**Characterisation of the Somatic Landscape of Adult Glioma**

*Dr. Ben Kinnersley (Institute of Cancer Research)*

**15:10–15:25**

**Identifying Clinically Actionable Genetic Abnormalities in Paediatric Acute Lymphoblastic Leukaemia**

*Professor Anthony Moorman (Newcastle University)*

**Parallel session 2: Diagnostic Discovery - Research Talks**

Chair: *Dr. Anna Need (Genomics England)*  
Location: *George Thomas*

**14:25–14:40**

**Interrogation of the 100,000 Genomes Project Ophthalmic Disease Cohort Reveals Novel Genes, New Associations and Previously Undetectable Mutations**

*Dr. Gavin Arno (University College London)*

**14:40–14:55**

**From the Rare Disease Pilot to the Main Program: How Many of the Newly Discovered Causal Variants and Genes can be Identified in the 100,000 Genomes Project?**

*Dr. Karyn Megy (University of Cambridge – NIHR BioResource)*

**14:55–15:10**

**Diagnosis of Rare Disease Using Whole Genome Sequencing in the 100,000 Genomes Project: Experience of the First 2000 Cases**

*Dr. Andrea Haworth (Congenica)*

**15:10–15:25**

**Increasing the diagnostic yield for craniosynostosis inside Genomics England by systematic biased “digging”**

*Dr. Eduardo Calpena (University of Oxford)*

**Parallel session 3: New Technologies & the Genomics England Research Environment**

Chair: *Dr. James Holman (Genomics England)*  
Location: *Great Hall*

**14:25–14:30**

**Introduction: Overview of work being done in the Research Environment**

*Dr. James Holman (Genomics England)*

**14:30–14:50**

**Lightning talks explaining the commercial/academic utility of the Research Environment**

*Dr. Helen Griffin (Newcastle University)*

*Dr. Mark Collins (Helomics)*

*Dr. Salih Tuna (NIHR BioResource, Cambridge University Hospitals)*

*Dr. Amy Hawarden (University of Manchester)*

**14:50–15:05**

**Overview of News, Tools and Features**

*Georgia Chan, Dr. Chris Odhams and Dr. Loukas Moutsianas (Genomics England)*

**15:05–15:15**

**UK Health Data Research Alliance: Implementation through trusted research environments**

*Professor Tim Hubbard (Genomics England)*

**15:15–15:25**

**Q&A**

*Dr. James Holman & all speakers*

**15:25–15:55 Coffee/tea break (with posters) Location: Library & Lecture Hall**





Parallel session 1: Patient Data Ecosystems	Parallel session 2: Collaborating on Research	Parallel session 3: Diagnostic Discovery – Research Talks	Research Environment User & Bioinformatics Panel
<p><i>Chair: Professor Tim Hubbard (Genomics England)</i> <i>Location: Robert Perks</i></p>	<p><i>Chair: Vivienne Parry (Genomics England)</i> <i>Location: George Thomas</i></p>	<p><i>Chair: Dr. Anna Need</i> <i>Location: Great Hall</i></p>	<p><i>Chair: Dr. Loukas Moutsianos</i> <i>Location: Library</i></p>
<p><b>15:55–16:10</b> <b>What Can We Learn About Secondary Findings from the 100,000 Genomes Project to Inform Policy and Practice? Findings from a Workshop with Patients, Clinicians, Policymakers, and Researchers</b> <i>Dr. Saskia Sanderson (University College London)</i></p>	<p><b>15:55–16:05</b> <b>Introduction</b> <i>Vivienne Parry (Genomics England)</i></p>	<p><b>15:55–16:10</b> <b>The 100,000 Genomes Project: Moorfields Eye Hospital NHS Foundation Trust Experience in Uncovering Pathogenic Variants in Inherited Optic Neuropathies</b> <i>Dr. Neringa Jurkute (UCL Institute of Ophthalmology and Moorfields Eye Hospital NHS Foundation Trust)</i></p>	<p><b>15:55–16:40</b> <b>Research Environment users and Genomics England experts discuss Slido questions from the Research Environment session and open questions from other users</b> <i>Dr. Chris Odhams (Genomics England)</i> <i>Dr. Georgia Chan (Genomics England)</i> <i>Dr. Alona Sosinsky (Genomics England)</i> <i>Dr. Mark Collins (Helomics)</i> <i>Dr. Hywel Williams (Cardiff University)</i> <i>Dr. Patrick Short (SanoGenetics)</i></p>
<p><b>16:10–16:25</b> <b>Assessing the Diagnostic Accuracy of Quantitative Phenotypic Risk Scores Based on Electronic Health Care Records for Patients with Rare Genetic Disorders</b> <i>Dr. Stefanie Mueller (Institute for Health Informatics, University College London)</i></p>	<p><b>16:05–16:40</b> <b>Interactive session on expectations and joint problem solving in groups</b></p>	<p><b>16:10–16:25</b> <b>Bronchiectasis: Developing Early Insights Into An Old Disease</b> <i>Dr. Helen Griffin (Newcastle University)</i></p>	
<p><b>16:25–16:40</b> <b>Generating Health Economic Evidence for Genomic Medicine: Opportunities Arising from the 100,000 Genomes Project</b> <i>Patrick Fahr (University of Oxford)</i></p>		<p><b>16:25–16:40</b> <b>Whole Genome Sequencing Through the 100,000 Genomes Project Identifies a Novel Cause of Pulmonary Arteriovenous Malformations Associated with Hereditary Haemorrhagic Telangiectasia</b> <i>Professor Claire Shovlin (Imperial College London)</i></p>	

**Closing plenary**  
*Room: Great Hall*

<b>16:50–17:00</b>	<b>Recap of the day</b>
<b>17:00–17:15</b>	<b>The National Genomic Healthcare Strategy</b> <i>Professor Sir John Bell (Genomics England Board Member)</i>
<b>17:15–17:30</b>	<b>Genomics England – the future: Update Towards 5 Million Genome Analyses</b> <i>Professor Sir Mark Caulfield (Chief Scientist, Genomics England)</i>
<b>17:30–17:35</b>	<b>Towards 5 Million</b> <i>UCB Industry Cohort</i>
<b>17:35–17:50</b>	<b>Genomics England – the future: Closing remarks by Chris Wigley, CEO of Genomics England</b>

**17:50–19:30 Drinks reception** *Room: Library & Lecture Hall*

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