

Virtual Conference Agenda

| Start (GMT) | Finish (GMT) | Time allocated | Format | Presenter details |
|--------------------------------|--------------|----------------|--------------|--|
| Monday, 31 January 2022 | | | | |
| 09:00 | 09:10 | | | Welcome |
| 09:00 | 09:10 | 10 mins | Live | Scientific Programme Committee: Gemma Chandratilake, University of Cambridge, UK Catherine Houghton, Liverpool Women's Hospital, UK Nicki Taverner, Cardiff University, UK |
| 09:10 | 10:45 | | | Session 1: The role of genomics in healthcare |
| 09:10 | 09:15 | 5 mins | Live | Introduction to the session <i>Chair: Nicki Taverner</i> |
| 09:15 | 09:30 | 15 mins | Pre-recorded | Development of Genomic Testing <i>Gemma Chandratilake</i> |
| 09:30 | 09:45 | 15 mins | Pre-recorded | The role in the NHS <i>Catherine Houghton</i> |
| 09:45 | 10:00 | 15 mins | Pre-recorded | Genomics in Africa: Health and Research Tina Wessels, University of Cape Town, South Africa |
| 10:00 | 10:15 | 15 mins | Pre-recorded | Genetic Counselling Landscape in the Middle East Tamam Khalaf, Igenomix Middle East, United Aab Emirates |
| 10:15 | 10:45 | 30 mins | Live | Q&A <i>Chair: Nicki Taverner</i> <i>Moderator: Gemma Chandratilake</i> |
| 10:45 | 11:05 | 20 mins | | Break |
| 11:05 | 14:55 | | | Session 2: Cancer Genomics |
| 11:05 | 11:10 | 5 mins | Live | Introduction to the session <i>Chair: Nicki Taverner</i> |
| 11:10 | 12:40 | 90 mins | Live | Workshop 1: Hereditary Cancer Jonathan Roberts, Cambridge University Hospitals, UK <i>Interactive workshop</i> |
| 12:40 | 13:25 | 45 mins | | Break |
| 13:25 | 13:55 | 30 mins | Pre-recorded | A million to one: how we find (and miss) meaningful variants in genomic sequencing <i>Gemma Chandratilake and Catherine Houghton</i> |
| 13:55 | 14:25 | 30 mins | Pre-recorded | Cancer Genomics: Bridging from the tumour to the germline in variant interpretation Clare Turnbull, The Institute of Cancer Research, UK |
| 14:25 | 14:55 | 30 mins | Live | Q&A <i>Chair: Nicki Taverner</i> <i>Moderator: Catherine Houghton</i> |
| 14:55 | 15:55 | 60 mins | Live | Networking/discussion |

Tuesday, 1 February 2022

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| 09:00 | | 11:20 | | Session 3: Variant Interpretation | |
| 09:00 | 09:05 | | Live | Introduction to the session <i>Chair: Catherine Houghton</i> | |
| 09:05 | 09:20 | 15 mins | Pre-recorded | Introduction to a genome browser <i>Gemma Chandratilake</i> | |
| 09:20 | 09:50 | 30 mins | Pre-recorded | Functional studies - When is a variant pathogenic? <i>Nicki Taverner</i> | |
| 09:50 | 11:20 | 90 mins | Live | Workshop 2: Variant interpretation using DECIPHER Julia Foreman, Wellcome Trust Sanger Institute, UK Group work /interactive elements | |
| 11:20 | 11:40 | 20 mins | | Break | |
| 11:40 | | 13:15 | | Session 4: Broadening access to genomics I | |
| 11:40 | 11:45 | | Live | Introduction to the session <i>Chair: Gemma Chandratilake</i> | |
| 11:45 | 12:15 | 30 mins | Pre-recorded | Polygenic Risk Scores in Genomic Healthcare: what, why, how and when Cathryn Lewis, King's College London, UK | |
| 12:15 | 12:45 | 30 mins | Pre-recorded | West Midlands FH Service - Cascading an innovative model of delivery. <i>Elaine George</i> | |
| 12:45 | 13:15 | 30 mins | Live | Q&A <i>Chair: Gemma Chandratilake</i> <i>Moderator: Nicki Taverner</i> | |
| 13:15 | 14:00 | 45 mins | | Break | |
| 14:00 | 15:00 | 60 mins | Live | Networking/Case study discussion | |

Wednesday, 2 February 2022

| Start | End | Duration | Type | Topic |
|--|-------|----------|--------------|---|
| Session 5: Broadening access to genomics II | | | | |
| 10:30 | 12:35 | | | |
| 10:30 | 10:35 | 5 mins | Live | Introduction to the session <i>Chair: Nicki Taverner</i> |
| 10:35 | 11:05 | 30 mins | Live | The role of GCs in the genomic era - MDTs, practice development <i>Led by programme committee</i> |
| 11:05 | 11:35 | 30 mins | Pre-recorded | Developing a national newborn screening programme using whole genome sequencing Amanda Pichini, Genomics England, UK |
| 11:35 | 12:05 | 30 mins | Pre-recorded | RDNOW: bridging the gap to diagnosis in rare disease <i>Lyndon Gallacher, Victorian Clinical Genetics Service, Australia</i> |
| 12:05 | 12:35 | 30 mins | Live | Q&A <i>Chair: Nicki Taverner</i> <i>Moderator: Catherine Houghton</i> |
| 12:35 | 12:55 | 20 mins | | Break |
| Session 6: Cardiac genomics | | | | |
| 12:55 | 15:15 | | | |
| 12:55 | 13:00 | 5 mins | Live | Introduction to the session <i>Chair: Catherine Houghton</i> |
| 13:00 | 13:30 | 30 mins | Pre-recorded | Cardiac genomics Chloe Reuter, Stanford University, USA |
| 13:30 | 13:45 | 15 mins | | Break |
| 13:45 | 15:15 | 90 mins | Live | Workshop 3: Cardiac genomics - Variant interpretation <i>Chloe Reuter, Stanford University, USA</i> <i>Group work /interactive elements</i> |
| 15:15 | 15:30 | 15 mins | | Break |
| Closing remarks | | | | |
| 15:30 | 16:10 | | | |
| 15:30 | 16:00 | 30 mins | Live | Next steps: applying what you have learned on this course - Panel discussion <i>Led by programme committee</i> |
| 16:00 | 16:10 | 10 mins | Live | Closing remarks Scientific Programme Committee: Gemma Chandratilake Catherine Houghton Nicki Taverner |