

| Wednesday 9 July | | |
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| Time | Title | Presenter |
| G.41 | | |
| 13:00-13:30 | Registration, Lunch & Networking | |
| 13:30-14:30 | Keynotes | Chair: Nick Loman |
| 13:30 | Francis Crick, the father of genomics and phylogenetics | Matthew Cobb - University of Manchester |
| 14:00 | Rapid intraoperative diagnostics of CSF tumours | Matt Loose - University of Nottingham |
| 14:30-16:00 | Evolving Technologies 1 | Chair: Michael Quail |
| 14:30 | Driving the multiomics revolution: The latest innovations from Illumina | Oliver Miller, Illumina |
| 14:50 | Element Biosciences | Rachel Piddock |
| 15:10 | From big dreams to real possibilities... | Dan Fordham |
| 15:30 | Innovation across the technology spectrum | Neil Ward |
| 16:00-16:30 | Coffee Break | |
| 16:30-18:15 | Evolving Technologies 2 | Chair: Michael Quail |
| 16:30 | Access the Full Richness of Biological Complexity with New Single Cell and Spatial tools from 10X Genomics | Nicola Cahill |
| 16:50 | Ultima | Alex Watkins |
| 17:10 | Single Technologies | Johan Stroemqvist |
| 17:30 | Roche | Bernd Timmermann |
| 18:05-19:30 | Drinks Reception and Posters | |

| Thursday 10 July | | | | |
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| | Parallel A - G.56 | | Parallel B - G.06 | |
| 08:15-09:15 | Genomics Community UK Networking Breakfast - G.06 | | | |
| 08:30-09:20 | Coffee & Pastries | | | |
| 09:20-11:00 | Plant and Animal Genomics | Chair: Al Darby | Evolutionary Genomics | Chair: Matt Loose |
| 09:20 | Decoding multiple layers of Genomics using Twist Bioscience' | Sponsor Talk - Twist | The Future of Exponentially Scalable Single Cell-omics | Sponsor Talk - ParseBio Ryan Wall |
| 09:30 | Wheat Pan Genomics | Anthony Hall | Cycles of centromere evolution | Ian Henderson |
| 09:55 | Fly genomes and transposable element invasions | Professor Andrea Betancourt | Hanging on the Tree of Life: Sequencing All Life and the Genomic Impact of Jumping Elements in Sloths. | Marcela Uliano-Silva |
| 10:20 | 3. Artificial Lighting Induces Gene Expression Responses Linked to Oxidative Stress in Bats | Tom Jenkins | 2. High-Resolution Mapping of Bacterial Evolution Under Antibiotic Stress Using Single-Cell Whole-Genome Sequencing | Matt Bawn |
| 10:35 | 20. Performance assessment of a rapid library preparation kit for short-read RNA sequencing | Thomas Brabbs | 22. Testing the regulatory effects of indels and structural variants using massively parallel reporter assays | Lindsey Plenderleith |
| 10:50 | Exploring the power of 10X spatial applications: A focus on Visium HD and Xenium In situ tools in plant and animal studies | Sponsor Talk - 10x Genomics Nicola Cahill | | |
| 11:00-11:30 | Coffee Break | | | |
| 11:30-13:00 | Spatial Genomics | Chair: Iain Macaulay | Microbial Genomics | Chair: Lauren Cowley |
| 11:30 | Enabling Spatial Omics at true single-cell resolution | Sponsor Talk - Takara Speaker: Matthieu Pesant, Takara Bio Europe | Accurate detection of small non-coding RNAs using NEBNext Low-bias Small RNA Library Prep Kit | Sponsor Talk - New England BioLabs Speaker Name: Louise Williams |
| 11:40 | Spatial transcriptomics of developing flowers across time | Christobal Uauy | Spatially-resolved host-pathogen interactions across the lung | Josie Bryant - Sanger |
| 12:05 | A Longer Look: Benchmarking Long-Read Sequencing for Single-cell Tanscriptomics | Anita Scoones | Pathogen Genomics in Maternal and Neonatal Health: Informing Targeted Interventions for Better Outcomes | Elita Jauneikaite - Imperial |
| 12:30 | 27. Spatiotemporal and Single-cell Sequencing of chicken trachea reveals differences in host Responses after Vaccination with Pathogenic and Attenuated Infectious Bronchitis Virus (IBV) | Dr Ge Wu | 17. Genomic epidemiology of plasmids carrying the iuc5 locus and antimicrobial resistance genes in Escherichia coli and Klebsiella pneumoniae from One Health settings | Keira Cozens |
| 12:45 | Introducing Illumina Spatial Technology | Sponsor Talk - Illumina Andrew Slatter | 37. 'PyamilySeq: Exposing the fragility of conventional gene (re) clustering and pangenomic inference methods' | Nicholas Dimonaco |
| 13:00-14:00 | Photo, Lunch and Poster Session II | | | |
| 14:00-15:30 | Epigenetics/Human & Clinical Genomics | Chair: Ania Piskorz | Microbial Communities & Microbiome | Chair: Nick Loman |
| 14:00 | "NGS: What are you missing?" | Sponsor Talk - Dovetail Speaker Name: Miles Collier | Empowering Discovery: Agilent's Integrated Genomics Solutions for the Future of Precision Science | Sponsor Talk - Agilent Rahna Ayub |
| 14:10 | Towards multi-modal liquid biopsies | Florent Mouliere | Resolving MAGs from complex microbiomes with long reads | Chris Quince (Earlham Institute) |

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| 14:35 | 12. Barcoded ultra-long read sequencing for genetic exploration of cancer samples | Joe Fenwick | 47. Making the most of microbiome biobanks: Enabling Cross-Cohort and Multi-Kingdom Analyses | Greg Young |
| 14:50 | 36. Understanding gene expression involved in immune response to SARS-CoV-2 Infection | Lee Murphy | 38. Beyond Taxonomic Composition: Metagenomic Evidence for Functional Niche Displacement in the Rumen Microbiome. | Katie Lawther |
| 15:05 | European Clinical Implementation of HiFi | Sponsor Talk - PacBio Riki Aydeniz | Metagenomics opportunities | Justin O'Grady |
| 15:20-16:00 | Coffee Break | | | |
| 16:00-17:30 | Computational Biology | Chair: Sion Bayliss | Human & Clinical Genomics | Chair: Katherine Johnson |
| 16:00 | Using rearrangement distances and historical samples to study plasmid evolution and epidemiology | Zamin Iqbal | Rapid genomic sequencing: Where are we and where are we going | Andy Parrish |
| 16:25 | Machine Learning and AI approaches for understanding Prokaryotic Pangenomes | James McInerney - University of Liverpool | The Generation Study: Insights from the first year | Katie Handley - Genomics England |
| 16:50 | 49. Accurate bioinformatic host pangenome depletion using fast minimizer search | Bede Constantinides, University of Birmingham | 19. Investigating mitochondrial DNA replication dynamics using thymidine analogues and ONT long read sequencing. | Katja Menger |
| 17:05 | 10. Ultra-long Sequencing Reveals Structural Heterogeneity of Common Dicentric Chromosomes in Childhood Cancer | Eleanor Woodward | Size Matters - Enhance your long-read sequencing data and yields | Sponsor Talk - Yourgene Health Speaker: Darren Marjenberg |
| 17:20 | Technology Platforms at the Earlham Institute: Research, Collaboration and Service | Alex McNeil, Earlham | | |

Friday 11 July

| Time | Title | Presenter |
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| G.41 | | |
| 08:30-09:15 | Coffee & Pastries | |
| 09:15-11:05 | New Methods in Genomics | Co-chairs: Sally James, Iain Macaulay |
| 09:15 | | Sponsor Talk - Element |
| 09:30 | Towards nanopore proteomics: Single-molecule detection of post-translational modifications on full-length proteins | Yujia Qing |
| 09:55 | Leveraging single-cell long-read sequencing and machine learning for therapeutic target identification | Adam Cribbs - University of Oxford |
| 10:20 | Revealing mechanisms of chemo-resistance and sensitisation with next generation lineage tracing technologies | Kirsty Sawicka - CRUK Cambridge |
| 10:45 | Reveal the power of the 6-base genome | Sponsor Talk - Biomodal |
| 11:00-11:30 | Coffee break | |
| 11:30-12:30 | Keynotes | Chair: Michael Quail |
| 11:30 | Shining a light on the dark genome | Wendy Bickmore |
| 12:15 | Lunch & end | |