TIME (BST)	ON-SITE AGENDA WEDNESDAY 20 MAY
08:30–09:30	**Registration with breakfast**
09:45–11:45	**Session 1: Auditorium**
09:45–10:10	Welcome to London Calling 2026
10:10–10:35	From innovation to global impact: adaptive genome sequencing for childhood leukemia | **Thomas Alexander** | University of North Carolina, USA
10:35–11:00	N-Care Project: using long-read sequencing for timely genetic diagnosis in critically ill infants across Asia-Pacific **Ni-Chung Lee** | National Taiwan University Hospital, Taiwan
11:00–11:20	Lightning talks
11:20–11:45	Embedding clinical metagenomics into NHS diagnostic pathways **Adela Alcolea-Medina** | Synnovis, Guy's and St Thomas' NHS Foundation Trust, UK | **Luke Blagdon Snell** | King's College London, UK
11:45–13:00	**Lunch***
	Democratising access to the future of paediatric leukaemia diagnostics | Showcase Stage Infectious disease networking | The Jam Networking Area
13:15–14:15	**Session 2: Breakouts**
	Metagenomics — insights without culture | Blondie Bioinformatic tools and insights for cancer research | Arctic Monkeys Underexplored molecules and gene regions | The Clash Population data and human diversity | The Jam Non-invasive biomarker detection | Bowie
14:25–15:25	**Break***
	Poster networking | Poster Hall Biodiversity genomics | Secret Cinema Technique talks | Techniques Theatre
15:35–16:40	**Session 3: Auditorium**
15:35–15:50	Spotlight session
15:50–16:15	Genetic and epigenetic landscape of self-identified Hispanics in All of Us | **Fritz Sedlazeck** | Baylor College of Medicine, USA
16:15–16:40	Improving diagnosis in Li-Fraumeni syndrome using long-read whole-genome and integrated multiomic sequencing **David Thomas** | University of New South Wales, Australia
16:50–17:40	**Break***
	Human translational research | Secret Cinema Plant & animal networking | The Jam Networking Area Technique talks | Techniques Theatre Beyond the BAM — supercharge your bioinformatics deployment | Showcase Stage
17:50–19:25	**Session 4: Auditorium**
17:50–19:20	Tech talk
19:20–19:25	Closing remarks
19:25–22:00	**Evening networking with food & drink**
22:00–01:00	**London Calling After Party!**

* Visit the Live Lounge and Data Analysis Lounge for drop-in support on Adaptive Sampling, transcriptomics, whole-genome sequencing, and data analysis. You can also book expert one-to-one support with the Oxford Nanopore team or Compatible Product Partners at the Tech Hub, visit Customer Services, and explore Oxford Nanopore posters.

TIME (BST)	ON-SITE AGENDA THURSDAY 21 MAY
08:15–08:50	**Registration with breakfast**
09:00–10:30	**Session 5: Auditorium**
09:00–09:20	Welcome back to London Calling 2026
09:20–09:45	Cancer clinical genomic testing using Oxford Nanopore whole-genome sequencing | **Rowan Howell** | Genomics England, UK
09:45–10:05	Lightning talks
10:05–10:30	Genomic integrity profiling of autologous iPSC-derived therapeutics for Parkinson's disease | **Roy Williams** | Aspen Neuroscience, USA
10:40–11:30	**Break***
	Poster networking | Poster Hall Clinical research | Secret Cinema Advanced analytics for biopharma using nanopore sequencing | Showcase Stage Technique talks | Techniques Theatre Early-career researcher networking | The Jam Networking Area
11:40–12:40	**Session 6: Breakouts**
	Profiling and classification of cancer | Blondie Pathogen surveillance from in-house to in the field | Arctic Monkeys Shedding light on biological regulation with multiomics | The Clash Pipelines to enable clinical scientists | The Jam Preparing for future clinical implementation | Bowie
12:50–14:00	**Lunch***
	Data for lunch | Arctic Monkeys & Bowie Animal genome assemblies | Secret Cinema Clinical networking | The Jam Networking Area
14:10–14:50	**Session 7: Auditorium**
14:10–14:25	Spotlight winner
14:25–14:50	Characterising the complete transcriptome and proteome with GenomeProt | **Mike Clark** | The University of Melbourne, Australia
15:00–15:50	**Break***
	Technique talks | Techniques Theatre Spotlight runners-up | Arctic Monkeys Oncology networking | The Jam Networking Area Beyond the BAM — revealing new dimensions in clinical research data | Showcase stage
16:00–17:35	**Session 8: Auditorium**
16:00–16:05	Poster & Sustainability awards
16:05–16:30	Rapid and accurate childhood cancer diagnosis with nanopore long-read RNA sequencing | **Sandy Fong** | SickKids Research Institute, Canada
16:30–17:30	Panel plenary: Methylation matters
17:30–17:35	Closing remarks
17:35–18:35	**Evening networking with drinks**

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