TIME (BST)	ONLINE AGENDA WEDNESDAY 20 MAY
09:30–13:05	**Session 1**
09:30–09:45	LC26 Studio: Where are our hidden secrets in genomics, and why should we care? | Live and exclusive speaker interviews
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
11:45–11:55	LC26 Studio: From urgent answers to equitable care in APAC | Live and exclusive speaker interviews
12:00–12:45	Knowledge exchange: bacterial and fungal isolate sequencing with NO-MISS
12:45–12:55	From London Calling 2025: Rapid-response mobile laboratory for disease elimination, outbreak investigation, and pathogen X **Arianna Ceruti** | Leipzig University, Germany
12:55–13:05	From London Calling 2025: Nanopore sequencing captures first in-patient evolution of vancomycin-resistant *E. faecium* in New Zealand **Rhys White** | Institute of Environmental Science and Research, New Zealand
13:15–15:35	**Session 2**
13:15–14:15	Breakout sessions: Metagenomics — insights without culture | Population data and human diversity | Bioinformatic tools and insights for cancer research
14:25–15:25	Poster networking
15:25–15:35	LC26 Studio: What can the St. Jude Global Alliance teach us about scaling paediatric cancer diagnostics? | Live and exclusive speaker interviews
15:35–17:50	**Session 3**
15:35–15:50	Spotlight talks
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
16:40–16:50	LC26 Studio: From representation to patient benefit in the Americas | Live and exclusive speaker interviews
16:55–17:35	Knowledge exchange: how to sequence full-length 16S and ITS microbial amplicons with Oxford Nanopore
17:40–17:50	LC26 Studio: From metagenomics to surveillance: what earlier pathogen detection changes | Live and exclusive speaker interviews
17:50–19:35	**Session 4**
17:50–19:20	Tech talk
19:20–19:25	Closing remarks
19:25–19:35	LC26 Studio | Live and exclusive speaker interviews

TIME (BST)	ONLINE AGENDA THURSDAY 21 MAY
08:45–11:40	**Session 5**
08:45–09:00	LC26 Studio: What's stopping rapid genomic insight becoming routine care — and how we fix it? | Live and exclusive speaker interviews
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
10:30–10:40	LC26 Studio: Can genomics change clinical decision making? | Live and exclusive speaker interviews
10:45–11:30	Democratising access to the future of paediatric leukaemia diagnostics: a St. Jude Global Alliance showcase
10:45–11:40	Poster networking
11:40–14:10	**Session 6**
11:40–12:40	Breakout sessions: Profiling and classification of cancer | Pipelines to enable clinical scientists | Pathogen surveillance from in-house to in the field
13:00–14:00	Data for lunch
14:00–14:10	LC26 Studio: From pilots to public services: how health systems can adopt genomics | Live and exclusive speaker interviews
14:10–16:00	**Session 7**
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
14:50–15:00	LC26 Studio: Leveraging AI and multiomics to reveal more biology | Live and exclusive speaker interviews
15:20–15:45	Spotlight runners-up
15:50–16:00	LC26 Studio: From disease biology to delivering advanced therapies | Live and exclusive speaker interviews
16:00–17:45	**Session 8**
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
17:35–17:45	LC26 Studio: Can methylation move the needle in clinical settings? | Live and exclusive speaker interviews

In addition to the above schedule, in the Auditorium you can also find on-demand Secret Cinema talks covering a variety of topics.