

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
11:20–13:00	Lunch*
	Democratising access to the future of paediatric leukaemia diagnostics Showcase Stage Infectious disease networking The Underground
13:15–14:15	Session 2: Breakouts
	Bioinformatic tools and insights for cancer research Arctic Monkeys Underexplored molecules and gene regions The Clash Metagenomics — insights without culture Blondie Population data and human diversity The Jam Non-invasive biomarker detection Bowie
14:25–15:25	Break*
	Poster networking Poster Area Biodiversity genomics programmes Secret Cinema
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	Plenary talk Name Institution, country
16:50–17:40	Break*
	Bioinformatics in practice Showcase Stage Animal genome assemblies Secret Cinema Plant & animal networking The Underground
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!

*Live Lounge features demo zones and ONT expert support, including digital panels powered by Adaptive Sampling, dRNA & cDNA, methylation, data analysis, project planning, flow cell loading, and Customer Services.

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*		
	Advanced analytics for biopharma using nanopore sequencing Showcase Stage Poster networking Poster Area	Clinical research Secret Cinema Early career researcher networking The Underground	
11:40–12:40	Session 6: Breakouts		
	Pathogen surveillance from in-house to in the field Arctic Monkeys Profiling and classification of cancer Blondie Preparing for future clinical implementation Bowie	Shedding light on biological regulation with multiomics The Clash Pipelines to enable clinical scientists The Jam	
12:50–14:00	Lunch*		
	Human translational research Secret Cinema	Clinical networking The Underground	Data for lunch Arctic Monkeys & Bowie
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*		
	Bioinformatics in practice Showcase stage	Spotlight runners-up Arctic Monkeys	Oncology networking The Underground
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		

*Live Lounge features demo zones and ONT expert support, including digital panels powered by Adaptive Sampling, dRNA & cDNA, methylation, data analysis, project planning, flow cell loading, and Customer Services.