

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	<a href="#">Welcome to London Calling 2026</a>		
10:10–10:35	<a href="#">From innovation to global impact: adaptive genome sequencing for childhood leukemia</a> Thomas Alexander   University of North Carolina, USA		
10:35–11:00	<a href="#">N-Care Project: using long-read sequencing for timely genetic diagnosis in critically ill infants across Asia-Pacific</a> Ni-Chung Lee   National Taiwan University Hospital, Taiwan		
11:00–11:20	<a href="#">Lightning talks</a>		
11:20–13:00	Lunch*		
	<a href="#">Democratising access to the future of paediatric leukaemia diagnostics</a>   Showcase Stage	<a href="#">Infectious disease networking</a>   The Jam Networking Area	
13:15–14:15	Session 2: Breakouts		
	<a href="#">Metagenomics — insights without culture</a>   Blondie <a href="#">Bioinformatic tools and insights for cancer research</a>   Arctic Monkeys <a href="#">Underexplored molecules and gene regions</a>   The Clash	<a href="#">Population data and human diversity</a>   The Jam <a href="#">Non-invasive biomarker detection</a>   Bowie	
14:25–15:25	Break*		
	<a href="#">Poster networking</a>   Poster Hall	<a href="#">Biodiversity genomics</a>   Secret Cinema	
15:35–16:40	Session 3: Auditorium		
15:35–15:50	<a href="#">Spotlight session</a>		
15:50–16:15	<a href="#">Genetic and epigenetic landscape of self-identified Hispanics in All of Us</a> Fritz Sedlazeck   Baylor College of Medicine, USA		
16:15–16:40	<a href="#">Improving diagnosis in Li-Fraumeni syndrome using long-read whole-genome and integrated multiomic sequencing</a> David Thomas   University of New South Wales, Australia		
16:50–17:40	Break*		
	<a href="#">Bioinformatics in practice</a>   Showcase Stage	<a href="#">Human translational research</a>   Secret Cinema	<a href="#">Plant &amp; animal networking</a>   The Jam Networking Area
17:50–19:25	Session 4: Auditorium		
17:50–19:20	<a href="#">Tech talk</a>		
19:20–19:25	<a href="#">Closing remarks</a>		
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*		
	Advanced analytics for biopharma using nanopore sequencing   Showcase Stage Poster networking   Poster Hall	Clinical research   Secret Cinema 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*		
	Animal genome assemblies   Secret Cinema	Clinical networking   The Jam Networking Area	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 Jam Networking Area
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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