

TIME (BST)		MASTERCLASS ONLINE AGENDA TUESDAY 19 MAY	
11:00–11:30	How do I get started with Oxford Nanopore sequencing?		
11:35–11:40	Multiomics with one platform: how do I see more from my samples with Oxford Nanopore sequencing?		
11:45–12:15	How do I extract high-quality DNA and RNA for my Oxford Nanopore sequencing experiment?		
12:15–12:30	Break		
12:30–12:35	How do I perform real-time targeted Oxford Nanopore sequencing?		
12:35–12:40	How do I enrich a digital panel in real time?		
12:45–13:15	Which library prep workflow is right for my experiment?		
13:20–13:30	Start the timer: how do I prepare a sequencing library in ten minutes?		
13:30–14:00	Break		
14:00–14:10	How do I load a PromethION Flow Cell?	How do I load a MinION Flow Cell?	
14:15–15:15	How do I analyse my Oxford Nanopore sequencing data without specialist training?		
15:15–15:30	Break		
15:30–15:35	Epigenetics without bisulfite: how do I analyse methylation and variants in one go?		
15:40–16:10	From speed to scale: how do I choose the right human whole-genome sequencing workflow for my experiment?	From full-length 16S to high-resolution metagenomics: how do I choose the right microbial community workflow for my experiment?	