

33rd International Dynamics & Evolution of Human Viruses

Wednesday, May 20, 2026

Evolutionary Dynamics of HIV: part 1 (2:00 PM - 2:40 PM)

-Conveners: Matthew Hall; Abayomi Olabode

time	[id] title	presenter
2:00 PM	[69] TOGGLING OF BOTH SYNONYMOUS AND NONSYNONYMOUS MUTATIONS CAN EXPLAIN WHY MEASURED RATES OF HIV EVOLUTION ARE FASTER WITHIN- THAN BETWEEN-HOST	BACK, Lucy
2:20 PM	[91] Phylodynamic Evaluation of Progress Towards the "Ending the HIV Epidemic" Goals in the New York City	WERTHEIM, Joel

Friday, May 22, 2026

Evolutionary Dynamics of HIV: Part 2 (10:50 AM - 12:50 PM)

-Conveners: Matthew Hall; Abayomi Olabode

time	[id] title	presenter
10:50 AM	[56] NETWORK-BASED DETECTION OF HIV-1 RECOMBINATION REVEALS UNDERLYING SUBTYPE ARCHITECTURE	OLABODE, Abayomi
11:10 AM	[33] MUTATIONAL SUPPLY SHAPES HIV EVOLUTION	NITIDO, Adam
11:30 AM	[89] THREE NEW BAYESIAN AND FIDUCIAL ANALYSIS METHODOLOGIES FOR HIV INTEGRATION SITE ANALYSIS	Dr EDLEFSEN, Paul. T
11:50 AM	[92] DYNAMICS OF VIRAL REPLICATION AND SPREAD IN TISSUES DURING EARLY SIV INFECTION	Dr IMMONEN, Taina
12:10 PM	[26] Substantial Deceleration of Adaptation of HIV-1 Within 1,500 Generations in an Experimental Evolution: A Genomic Perspective	MOVASATI, Ali
12:30 PM	[73] Suppression and fragmentation of HIV transmission following scale-up of TasP and PrEP in British Columbia, Canada: a population level phylogenetic study	JOY, Jeffrey