## Agenda – October 31<sup>st</sup> 2025



Time	Session title
09:00 - 09:30	Registration
09:30 - 09:40	Opening remarks
09:40 – 10:30	Keynote presentation James Hay: A quantitative world: should we stop binarizing our epidemiological data?
10:30 – 11:10	Scientific session
	<b>Alexis Martin</b> : Coupling models together! The love story of two mathematical models for malaria
	Annabelle Wong: Assessing the effect of social contact structure on the impact of pneumococcal conjugate vaccines
11:10 – 11:40	Coffee Break and Posters
11:40 – 12:20	Scientific session
	Matthieu Domenech de Cellés: Immune boosting and the perils of interpreting pertussis seroprevalence studies
	<b>Louis Colliot</b> : How virus genomics can help to understand the past (and present) of HIV epidemics in France?
12:20 – 13:00	Speed talks
	Baptiste Ellie: Impact of viral blips and low level viremia on HIV-specific antibody responses
	Cana Kussmaul: Modeling the effect of weather on infectious diseases: why causal structure matters
	<b>Daniella Figueroa-Downing</b> : How well does this vaccine work? Navigating efficacy, effectiveness, and modelled impact
	<b>Dennis PartI:</b> Compartmental Model on the Introduction of Zoliflodacin in South Africa
	<b>Adrian Lison</b> : Real-time estimation of pathogen transmission dynamics from wastewater
13:00 – 14:00	Lunch (Served at ground floor)
14:00 – 14:50	Keynote presentation Eva Bons: Forecasting and Scenario Modelling of Respiratory Infections in Europe: Leveraging the Ensemble Approach

14:50 – 15:30	Scientific Session
	<b>Martin Wohlfender</b> : Machine learning-based short-term forecasting of COVID-19 hospital admissions using routine hospital patient data
	<b>Branwen Owen</b> : Cost-effectiveness of typhoid conjugate vaccine across sub- Saharan Africa: Random forest analysis reveals primacy of cost over epidemiological drivers
15:30 – 15:50	Break
15:50 – 16:50	Speed Talks
	<b>Diane Duroux</b> : Early antifungal resistance identification using MALDI TOF mass spectrometry and machine learning
	<b>Luzia Nora Felber:</b> Simulating Mosquito Flight Paths with a Random Walk Model
	Javier Perez-Saez: Evolving infectious disease dynamics shape school-based intervention effectiveness
	Elizabeth Lee: Improving the timeliness of cholera outbreak response with case-based early warning signals
	<b>Sarah Kramer:</b> Purely statistical methods of inferring causality fail to accurately characterize pathogen-pathogen interactions
	Adrian Denz: Inferring a novel insecticide resistance metric and exposure variability in mosquito bioassays across Africa
	<b>Pietro Gemo:</b> Are time series regressions reliable for estimating the effects of weather on infectious diseases?
	<b>Nicolas Banholzer</b> : The relative contribution of close-proximity contacts, shared classroom exposure and indoor air quality to respiratory virus transmission in schools
16:50 – 17:00	Final Words
17:00	Apéro and Networking