

	<b>Tuesday 16<sup>th</sup> of July</b>	<b>Wednesday 17<sup>th</sup> of July</b>	<b>Thursday 18<sup>th</sup> of July</b>	<b>Friday 19<sup>th</sup> of July</b>
10:00-11:30	<b>Moodie</b> (McGill) Statistical Methods for Precision Medicine Pt1	<b>Golchi</b> (McGill) Statistical Design of Bayesian Adaptive Trial Designs Pt1	<b>De Angelis</b> (Cambridge) Statistical inference for infectious diseases evidence synthesis models Pt2	<b>Moodie</b> (McGill) Statistical Methods for Precision Medicine Pt2
11:30-12:00	Coffee	Coffee	Coffee	Coffee
12:00-13:30	<b>De Angelis</b> (Cambridge) Statistical inference for infectious diseases evidence synthesis models Pt1	<b>12:00-12:20 McLatchie</b> (UCL) Predictive performance of power posteriors	<b>12:00-12:20 Presanis</b> (Cambridge) Multi-state models of hybrid protection against SARS-CoV-2 infection: the SIREN study	<b>Golchi</b> (McGill) Statistical Design of Bayesian Adaptive Trial Designs Pt2
		<b>12:20-12:40 Dimitriou</b> (UCL) Data Fusion for Heterogeneous Treatment Effect Estimation with Multi-Task Gaussian Processes	<b>12:20-12:40 Alexopoulos</b> (AUEB), Efficient Bayesian modelling and computational framework for epidemic forecasting	
		<b>12:40-13:00 Antonelli</b> (Florida) Sensitivity analysis for multiple treatments and multiple outcomes with applications to air pollution mixtures	<b>12:40-13:00 Guzman-Rincon</b> (Warwick) Statistical framework for the nowcasting and forecasting of infectious disease growth rates	
		<b>13:00-13:20 Minas</b> (St Andrews) Efficient Bayesian inference for large biological oscillators	<b>13:00-13:20 Birrell</b> (UK HSA) Parsimonious models for $\beta_t$ and the consequences for forecasting	
13:30-14:40	Lunch	Lunch	Lunch	Lunch
14:40-16:00	<b>Stephens</b> (McGill) Semiparametric Bayesian inference for optimal dynamic treatment regimes via dynamic marginal structural models	<b>Bui</b> (BU Vietnam) Multi-state model: Statistical inference for cross-sectional data for cancer	<b>Knock</b> (Imperial) A multi-region, Bayesian hierarchical approach to epidemic modelling	<b>Stival</b> (Ca' Foscari Venice) A Bayesian approach to explain spatio-temporal heterogeneity in repeated cross-sectional health surveys
	<b>Papadogeorgou</b> (Florida) Spatial causal inference in the presence of unmeasured confounding and interference	<b>Thoma</b> (Turing/UCL) A scalable formulation of joint modelling for longitudinal and time-to-event data and its application on large electronic health record data of diabetes complications	<b>Gill</b> (Warwick) Bayesian Inference of Reproduction Number from Epidemic and Genomic Data using Particle MCMC Methods	<b>Derezea</b> (Bristol) Network meta-analysis of diagnostic test accuracy reported at multiple thresholds
	<b>Lehmann</b> (UCL) A causal debiasing framework for improving local prevalence estimates of SARS-CoV-2	<b>Seymour</b> (Birmingham) Comparative Judgement Modelling to Map Women's Health and Rights at Community Level	<b>Baguelin</b> (Imperial) Comparative Assessment of Deterministic and Stochastic Epidemic Modelling Approaches: Insights for Real-Time Outbreak Analysis	<b>Vasdekis</b> (Newcastle) Skew-symmetric sampling schemes for SDEs and where to find them
	<b>Samartsidis</b> (Cambridge) A modularized Bayesian factor analysis model for estimating heterogenous causal effects from observational time-series data	<b>Kaisaridi</b> (Sorbonne) A multivariate disease progression model. Application in identifying subtypes in CADASIL	<b>Blenkinsop</b> (Imperial) Bayesian viral phylogenetic source attribution that exploits time since infection estimates	<b>Nikoloulopoulos</b> (UEA) Vine copula mixed models for meta-analysis of diagnostic accuracy studies without a gold standard