

# #BIBC2025 Schedule

## Tuesday

Start	Arc Cinema	Theatrette
08:15	Conference Registration	
08:45	Opening remarks and housekeeping	
09:05	<b>Keynote – Chair: Chris Brien</b> <b>On Finding Good Experiments</b> by <i>Cheng Soon Ong</i>	
10:00	<b>Session 2A – Chair: Garth Tarr</b>  <b>Data-Adaptive Automatic Threshold Calibration for Stability Selection</b> by <i>Martin Huang</i>  <b>Variable Selection in a Joint Model for Huntington’s Disease Data</b> by <i>Rajan Shankar</i>  <b>StableMate: a regression framework for selecting stable predictors across heterogeneous data environments</b> by <i>Yidi Deng</i>	<b>Session 2B – Chair: Scott Foster</b>  <b>Estimating abundance in small populations using pedigree reconstruction</b> by <i>Sarah Croft</i>  <b>Accounting for heterogeneous detection rates when inferring eradication of an invasive species</b> by <i>Sean A. Martin</i>  <b>Zero-inflated Tweedie distribution for abundance of rare ecological species</b> by <i>Nokuthaba Sibanda</i>
11:00	Morning Tea	
11:30	<b>Invited Session: A Cluster of Modern Clustering Methods for Biometrics – Chair: Alan Welsh</b>  <b>Extending Spatial Capture-Recapture with the Hawkes Process</b> by <i>Alec B. M. van Helsdingen</i>  <b>A Test for Detecting Multiple Clusters with Hotspot Spatial Properties</b> by <i>Kunihiko Takahashi</i>  <b>Outlier-robust estimation of state-space models using a penalised approach</b> by <i>Garth Tarr</i>	<b>Session 3B – Chair: James Curran</b>  <b>Group Sampling with Imperfect Testing for Biosecurity Applications</b> by <i>Adele Jackson</i>  <b>Koala Distribution and Abundance</b> by <i>Scott D. Foster</i>  <b>Speed: An R package for Spatially Efficient Experimental Designs</b> by <i>Sam Rogers</i>

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Start	Arc Cinema	Theatrette
	<b>Disease cluster detection via functional additive models incorporating spatial correlation</b> by <i>Michio Yamamoto</i>	<b>Running Human Subject Experiments via Online Crowdsourcing</b> by <i>Patrick Li</i>
12:50	Lunch	
13:50	<b>Session 4A – Chair: Francis Hui</b>	<b>Session 4B – Chair: Ruth Butler</b>
	<b>Bayesian clustered ensemble prediction for multivariate time series</b> by <i>Shonosuke Sugawara</i>	<b>Building Trust Without Peer Review: Establishing Reproducibility Standards in Industrial Statistical Consulting</b> by <i>Dean Marchiori</i>
	<b>The difficulties of clustering categorical or mixed data</b> by <i>Louise McMillan</i>	<b>Teaching Meta-Analysis for Systematic Reviewers with Mixed Statistical Training</b> by <i>Xu Ning</i>
	<b>Species archetype models for presence-only data</b> by <i>Skipton N.C. Woolley and Scott Woolley</i>	<b>Tales from the jungle: a personal perspective of statistical consulting since COVID</b> by <i>Alice Richardson</i>
		<b>Better Conversations, Better Support: Strengthening Consulting through Practical Education and Community</b> by <i>Sharon G. Nielsen</i>
15:20	Afternoon Tea	
15:50	<b>Session 5A – Chair: Zhanglong Cao</b>	<b>Session 5B – Chair: Graham Hepworth</b>
	<b>Nested-factorial treatment models: types, their uses and examples</b> by <i>Chris Brien</i>	<b>Enhancing Fraud Detection in Banking through Random Survival Forests: Addressing Data Imbalance and Model Transparency</b> by <i>Arjun Sekhar</i>
	<b>Integrating Spatial Data and On-Farm Experimentation to Understand Wheat Variety Performance Across Western Australia</b> by <i>Sandra K. Tanz</i>	<b>Using point cloud data to discover genomic regions associated with dynamic height</b> by <i>Colleen H Hunt</i>
	<b>Multi-environment trial analysis of count data with complex variance structures using generalised linear mixed models</b> by <i>Michael H. Mumford</i>	<b>Automatic debiased machine learning (autoDML) for causal inference: implementation and evaluation in real-world observational studies</b> by <i>Tong Chen</i>

## Tuesday Poster Session

- Location: Gallery
  - Time: 17:00-19:00
1. **Using atmospheric transport models to predict species incursions in northern Australia** by *Zhenhua (Iris) Hao Dr*
  2. **Visualisation of multinomial multilevel time-series modelling with application to current smoking status** by *Alice Richardson*
  3. **Paired Comparison with Cyclic Dominance: An Extension of the BradleyTerry Model** by *Yuki Ohno*
  4. **Cleaning Text Data with Large Language Models** by *Jiajia Li*
  5. **ggincerta: An R Package for Uncertainty Visualisation with a Layered Grammar of Graphics** by *Xueqi Ma*
  6. **The Theory of Sampling (ToS) - have statisticians missed the boat?** by *Damian Collins*
  7. **Evaluating Diagnostic Performance via Bayesian  $F_1$  Score Estimation without a Gold Standard** by *Jun Tamura*
  8. **Evaluating Remote Monitoring in Automated Peritoneal Dialysis: A Difference-in-Differences Analysis** by *Annie Conway*
  9. **Judgement Post-Stratification for Covariate Adjustment in Pairwise Comparisons in Block Designs** by *Sam Rogers*
  10. **Bayesian inference for sparse Gaussian copula graphical model** by *Tomotaka Momozaki*
  11. **Visualization for departures from symmetry with the power-divergence-type measure in square contingency tables** by *Wataru Urasaki*
  12. **Performance of Factor Analytic Mixed Models and PlackettLuce Models for Genotype Ranking in Multi-Environment Trials** by *Jiazhe Lin*

## Wednesday

Start	Arc Cinema	Theatrette
08:30	Conference Registration	
08:50	Housekeeping	
08:55	<b>Keynote – Chair: Thomas Lumley</b> <b>Uses of gnm for Generalized (Non)linear Modelling</b> by <i>Heather L. Turner</i>	
09:50	<b>Session 7A – Chair: David Warton</b> <b>Scalable finite mixture of regression models for clustering species responses in ecology</b> by <i>Francis KC Hui</i>	<b>Session 7B – Chair: Sam Rogers</b> <b>Using a linear mixed model based wavelet transform to model non-smooth trends arising from designed experiments</b> by <i>Clayton Forknall</i>

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Start	Arc Cinema	Theatrette
	<b>Elastic Net Regularization for Vector Generalized Linear Models: A Flexible Framework for High-Dimensional Biomedical Data</b> by <i>Wenqi Zhao</i>	<b>Functional Data Analysis for the Australian Grains Industry</b> by <i>Braden J. Thorne</i>
	<b>Fitting Generalised Linear Mixed Models using Sequential Quadratic Programming</b> by <i>Peter Green</i>	<b>Bayesian Ordinal Regression for Crop Development and Disease Assessment</b> by <i>Zhanglong Cao</i>
10:50	Morning Tea	
11:20	<b>Invited Session: Statistics for Biosecurity Surveillance – Chair:</b> <i>Robert Clark</i>	<b>Session 8B – Chair:</b> <i>Linh Nghiem</i>
	<b>Optimal allocation of resources between control and surveillance for complex eradication scenarios</b> by <i>Mahdi Parsa</i>	<b>Estimating extinction time from the fossil record using regression inversion</b> by <i>David I. Warton</i>
	<b>Inferring the rate of undetected contamination using random effects modelling of biosecurity screening histories</b> by <i>Sumonkanti Das and Robert Clark</i>	<b>The performance of Yu and Hoff's confidence intervals for treatment means in a one-way layout</b> by <i>Paul Kabaila</i>
	<b>Optimal sampling in border biosecurity: Application to skip-lot sampling</b> by <i>Raphael Trouve</i>	<b>Rate-optimal sparse gamma scale mixture detection</b> by <i>Michael Stewart</i>
		<b>Extension of the corrected score estimator in a Poisson regression model with a measurement error</b> by <i>Kentarou Wada</i>
12:50	Lunch	
13:00	Social Activities	
18:30	Young (at heart) Biometrician Social Event	

## Thursday

Start	Arc Cinema	Theatrette
08:30	Conference Registration	
08:50	Housekeeping	
08:55	<b>Keynote – Chair: Warren Muller</b> <b>Modularizing Biometric Models Facilitates Multistage Computing</b> by Mevin B. Hooten	
09:50	<b>Session 10A – Chair: Vanessa Cave</b> <b>Visualize your fitted non-linear dimension reduction model in the high-dimensional data space</b> by P. G. Jayani Lakshika <b>The geometry of diet: using projections to quantify the similarity between sets of dietary patterns</b> by Beatrix Jones <b>Multivariate meta-analysis methods for high-dimensional data</b> by Alysha M. De Livera	<b>Session 10B – Chair: Alice Richardson</b> <b>Reporting Odds Ratios under Fluctuating Reporting Rates in Spontaneous Reporting Systems</b> by Tatsuhiko Anzai <b>Handling Missingness in Prevalence Estimates from National Surveys</b> by Oyelola Adegboye <b>Pooled testing with penalized regression models</b> by Christopher Bilder
10:50	Morning Tea	
11:20	<b>Session 11A – Chair: Matthew Schofield</b> <b>Integrated Species Distribution Models: A Single-Index Approach</b> by Quan Vu <b>Simultaneous Inference for Latent Variable Predictions in Factor Analytic Models</b> by Zhining Wang <b>Model-based assessment of functional and phylogenetic diversity</b> by Shaoqian Huang <b>Fitting integrated species distribution models using mgcv</b> by Elliot Dovers	<b>Session 11B – Chair: Chris Triggs</b> <b>Crossvalidation for predictive models in complex survey data</b> by Thomas Lumley <b>A Set of Precise Asymptotics for Gaussian Variational Approximation in Generalised Linear Mixed Models</b> by Nelson J. Y. Chua <b>An allometric differential equation model quantifies energy trade-offs between growth and reproduction under temperature variation</b> by Hideyasu Shimadzu <b>A circular hidden Markov model for directional time series data</b> by A.A.P.N.M. Perera
12:40	Lunch	
13:00	AGM	
13:40	<b>Invited Session: Methods and Practice in Agricultural Analytics – Chair: Emi Tanaka</b>	<b>Session 12B – Chair: Sam Mason</b>

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Start	Arc Cinema	Theatrette
	<p><b>Evaluating the impact of trait measurement error on genetic analysis of computer vision-based phenotypes</b> by <i>Gota Morota</i></p> <p><b>Predication of Daily Weight Gain with Cattle Behaviour and Daily Activity Using Triaxial Accelerometer Data</b> by <i>Shuwen Hu and Antonio Hu</i></p> <p><b>Fishing for Heritability in the Gill Microbiome: Why Statisticians Should get out into the field</b> by <i>Elle Saber</i></p>	<p><b>Do Mice Matter? The Impact of Mice on a New Zealand Ecosanctuary</b> by <i>Vanessa Cave</i></p> <p><b>Modelling Species Diversity from Citizen-Science Bird Counts</b> by <i>Graham Hepworth</i></p> <p><b>Continental-Scale Bayesian Analysis of Acacia Flowering Phenology: A Novel Framework Integrating Phylogenetic Signal and Circular Statistics</b> by <i>Owen Forbes</i></p> <p><b>Spatio-Temporal Species Distribution Modelling</b> by <i>Sam Mason</i></p>
15:10	Afternoon Tea	
15:40	<b>Session 13A – Chair: Julian Taylor</b>	<b>Session 13B – Chair: Oyelola Adegboye</b>
	<p><b>The *equalto* covariance structure for meta-analysis using the ‘glmmTMB’ R package</b> by <i>Coralie C. Williams</i></p> <p><b>Simulation-based Inference about Variance Components</b> by <i>Farwa Saleem</i></p> <p><b>A Proportional Random Effect Block Bootstrap for Highly Unbalanced Clustered Data</b> by <i>Zhi Yang Tho</i></p>	<p><b>Extending diagnostic validity meta-analysis to several diagnostic guidelines</b> by <i>Alain C. Vandal</i></p> <p><b>Risk of Guillain-Barré Syndrome after COVID-19 vaccination and SARS-CoV-2 infection: A multinational self-controlled case series study</b> by <i>Han Lu</i></p> <p><b>Childhood Risk and Resilience Factors for Pasifika Youth Respiratory Health: Accounting for Attrition and Missingness</b> by <i>Siwei Zhai</i></p> <p><b>A missing data detective story how I navigated through a perfect storm of drop-outs, COVID and informative missingness.</b> by <i>Eve Slavich</i></p>
18:30	Conference Dinner	

## Friday

Start	Arc Cinema	Theatrette
08:40	Conference Registration	
09:00	Housekeeping	
09:05	<b>Keynote – Chair: Louise McMillan</b>	
	<b>Saddlepoint approximations for likelihoods</b> by <i>Jesse Goodman</i>	
10:00	<b>Session 15A – Chair: Yidi Deng</b>	<b>Session 15B – Chair: David Baird</b>
	<b>False Discovery Rate Controlled Robust Variable Selection under Cellwise Contamination</b> by <i>Xiaoya Sun</i>	<b>Genstat Markdown: Reproducible Research with Genstat</b> by <i>James M. Curran</i>
	<b>A covariate-adaptive test for replicability across multiple studies with false discovery rate control</b> by <i>Dennis Leung</i>	<b>The 4S method for the longitudinal analysis of multidimensional questionnaires: application to Parkinsons disease progression from patient perception</b> by <i>Tiphaine Saulnier</i>
	<b>A semi-supervised framework for diverse multiple hypothesis testing scenarios</b> by <i>Jack Freestone</i>	<b>‘heritable’ An R package for heritability calculations for plant breeding trials</b> by <i>Fonti Kar</i>
11:00	Morning Tea	
11:30	<b>Keynote – Chair: James Curran</b>	
	<b>Optimizing Research Impact Through Interdisciplinary and Collaborative Research</b> by <i>Charmaine B. Dean</i>	
12:20	Closing Ceremony	
12:40	Lunch	