EMI TANAKA

WORK EXPERIENCE

Lecturer

Monash University, Department of Econometrics and Business Statistics 2020/01 - current

Lecturer

The University of Sydney, School of Mathematics and Statistics 2017/01 - 2019/12

Research fellow

University of Wollongong, School of Mathematics and Applied Statistics 2014/01 - 2017/01

EDUCATION

PhD, Statistics

Statistical Methods for Improving Motif Evaluation

Supervisor: Dr. Uri Keich

School of Mathematics and Statistics

The University of Sydney, Sydney, Australia, 2015

Bachelor of Science (Advanced Mathematics), Honours I

Major in Mathematics and Statistics

The University of Sydney, Sydney, Australia, 2010

PUBLICATIONS

- Morota, G, Cheng, H, Cook, D, & Tanaka, E (2021) ASAS-NANP SYMPOSIUM: Prospects for interactive and dynamic graphics in the era of data-rich animal science. *Journal of animal science*, 99(2). *Citations:* 1.
- 2. Tanaka, E (2020) Simple outlier detection for a multienvironmental field trial. *Biometrics*, 76(4), 1374-1382. *Citations:* 4.
- 3. Tanaka, E & Hui, F (2019) Symbolic formulae for linear mixed models. *Statistics and data science*, 3-21
- 4. Hui, F, Tanaka, E, & Warton, D (2018) Order selection and sparsity in latent variable models via the ordered factor LASSO. *Biometrics*, 74(4), 1311-1319. *Citations:* 6.
- 5. Norman, A, Taylor, J, Tanaka, E, Telfer, P, Edwards, J, Martinant, J, & Kuchel, H (2017) Increased genomic prediction accuracy in

Contact

- github.com/emitanaka

For more information, please contact me via email.

Skills

Expert: R, HTML/CSS, LaTeX **Intermediate**: Git/GitHub, Python,

Bash, JS

Languages: English (fluent) and Japanese (conversational)

Interests

Research: experimental design, mixed models, data visualisation, bioinformatics, statistical genetics, selective breeding

Non-Research: drawing (but not good at it), reading (manga, manhwa and non-fiction books)

This resume was made with the R package pagedown.

Last updated on 2021-03-21.

- wheat breeding using a large australian panel. *Theoretical and applied genetics*, 130(12), 2543-2555. *Citations: 16*.
- 6. Tanaka, E, Ral, J, Li, S, Gaire, R, Cavanagh, C, Cullis, B, & Whan, A (2017) Increased accuracy of starch granule type quantification using mixture distributions. *Plant methods*, 13, 107. *Citations:* 5.
- 7. Tanaka, E (2014) Statistical methods for improving motif evaluation. *PhD Thesis*
- 8. Tanaka, E, Bailey, T, & Keich, U (2014) Improving MEME via a two-tiered significance analysis. *Bioinformatics*, 30(14), 1965-1973. *Citations: 17*.
- Liachko, I, Tanaka, E, Cox, K, Chung, S, Yang, L, Seher, A, Hallas, L, Cha, E, Kang, G, Pace, H, Barrow, J, Inada, M, Tye, B, & Keich, U (2011) Novel features of ARS selection in budding yeast lachancea kluyveri. *BMC genomics*, 12, 633. *Citations: 22*.
- 10. Tanaka, E, Bailey, T, Grant, C, Noble, W, & Keich, U (2011) Improved similarity scores for comparing motifs. *Bioinformatics*, 27(12), 1603-1609. *Citations:* 49.

Citation counts are sourced from Google Scholar at 2021-03-21.

</> SOFTWARE

@ edibble

An R-package that implements the grammar of experimental design.

Creator and maintainer. GitHub stars: 121

Source: https://github.com/emitanaka/edibble

Ø Documentation: https://edibble.emitanaka.org

anicon

An R-package to insert animated icons for R markdown and Shiny apps.

Creator. GitHub stars: 109

Source: https://github.com/emitanaka/anicon

nestr

An R-package to build hierarchical or nested sturctures.

Creator and maintainer. GitHub stars: 3

Source: https://github.com/emitanaka/nestr

O Documentation: https://nestr.emitanaka.org

monash 🗭

A utility R-package with consolidated tools and templates for staffs at Monash University.

Creator and maintainer. GitHub stars: 3

Source: https://github.com/numbats/monash

xaringan

An R package for creating slideshows with remark.js through R Markdown.

Contributor. My contribution is the adaptation of the ninja-theme shown in the Documentation. GitHub stars: 1131

Opcumentation: https://github.com/emitanaka/ninja-theme

Q gghdr

Plots of highest density regions (HDR) for ggplot2.

Listed as author but my contribution is small. GitHub stars: 34

Source: https://github.com/ropenscilabs/gghdr

Open Documentation: https://ropenscilabs.github.io/gghdr

shinycustomloader

Add a custom loader for R shiny.

Creator. GitHub stars: 97

Source: https://github.com/emitanaka/shinycustomloader

datalegreyar

Datalegreya, the typeface that melts text and data visualisation, for R markdown.

Creator. GitHub stars: 38

Source: https://github.com/emitanaka/datalegreyar

</> Tomtom

A motif comparison tool part of the MEME Suite.

The -incomplete-scores is an implementation as a result of Tanaka et al. (2011).

■ TALKS

List of talks (and links to the slides if available) are at https://emitanaka.org/talks.html. Below show the last 10 talks.

The Grammar of Experimental Design

TokyoR

2021-01-23

 Advent of "Grammar": Bridging Statistics and Data Science for the Design of Experiments

Statistical Society of Australia 2020-11-27

 Rethinking the framework to specify the design of experiments

Department of Agriculture and Fisheries, Queensland 2020-11-11

What every young statistician should know:
 Developing your identity, networking and marketing

Statistical Society of Australia Western Australian Branch 2020-09-30

 Advent of "Grammar": Bridging Statistics and Data Science for the Design of Experiments

Monash Bioinformatics Seminar 2020-07-15

 Beyond Beamer: Modern and Dynamic Presentations with R Markdown

Statististical Society of Australia Canberra & NSW Branch 2020-06-23

 Software design, selection and estimation for latent variable models

WOMBAT 2019-11-28

 Symbolic model formulae for linear mixed models illustrated with the analysis of agricultural data
 Department of Mathematics and Statistics, Macquarie University 2019-10-15

 Symbolic model formulae for linear mixed models illustrated with the analysis of agricultural data

Research School of Finance, Actuarial Studies and Statistics, Australian National University 2019-09-16

 Statistical modelling for plant breeding trials for accurate genotype-by-environment effect predictions

Sydney Institute of Agriculture 2019-09-06

♣ WORKSHOPS

Data Wrangling with R

2020/12

Hosted by Statistical Society of Australia NSW Branch.

Data Visualisation with R

2020/11

Co-presented with Di Cook. Hosted by Statistical Society of Australia VIC Branch.

Tidyverse and R Markdown Workshop

2019/12

Hosted by International Biometrics Society Australasia Region.

R Package & R Markdown Workshop

2019/11

Co-presented with Damjan Vukcevic.

Hosted by Statistical Society of Australia VIC Branch.

Statistical Methods for Omics Assisted Breeding

2018/11

Co-presented with Gota Morota, Diego Jarquin, Malachy Campbell, Jessica Tressou, Hiroyoshi Iwata. Hosted by Univeristy of Tokyo.

▲ SERVICE

Vice President

Statistical Society of Australia, Victoria Branch 2020/03-2021/03

President

Statistical Society of Australia, Victoria Branch 2021/03-2023/03

Vice President

Statistical Society of Australia, Victoria Branch 2023/03-2024/03

Assistant Secretary

Statistical Society of Australia, NSW Branch 2017/03-2018/03

Secretary

Statistical Society of Australia, NSW Branch 2018/03-2020/03

Social Media Coordinator

International Biometrics Society, Australaisan Region 2018/01-ongoing

Member

useR! 2021, Program Committee 2021/03-2021/03

PROFESSIONAL MEMBERSHIPS

- Statistical Society of Australia
- International Biometrics Society

Q AWARDS & DISTINCTIONS

ARC Centre of Excellence

Associate Investigator
ARC Centre of Excellence for Mathematical and Statistical
Frontiers (ACEMS)
2020

ARC Industrial Transformation Training Centre

Chief Investigator
Data Analytics for Resources and Environments (DARE)
2019
\$3,973,202

R Consortium Grant

Lead Investigator
Symbolic Formulae for Linear Mixed Models
2019
\$6,000

Sydney Institute of Agriculture Research Project Grant

Chief Investigator
Land surface models of carbon and water do not work in agricultural landscapes where it actually matters they work 2018
\$90,000