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), Honour 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 2021-03-18.

- 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 2021-03-18.

</> SOFTWARE

@ edibble

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

Creator and maintainer. GitHub stars: 121

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

₱ 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: 1

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: 1129

Source: https://github.com/yihui/xaringan

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 Industrial Transformation Training Centre

2019

Lead: Sally Cripps \$3,973,202

R Consortium Grant

2019

Lead: Emi Tanaka

Sydney Institute of Agriculture Research Project Grant

2018

Lead: Thomas Bishop

\$90,000

• SSA NSW Branch J. B. Douglas Award Prize

2011

\$500

Winter School Student Travel Bursaries

2010

\$250

Statistical Society of Australia Inc. Student Travel Grant

2010

\$300

Australian Postgraduate Award

2010

\$78,750

Dean's List for Academic Excellence

2008

• Tim Brown Prize II

2008

\$275

Barker Scholarship IV

2008

\$750