June En-Yu Lai Ph.D.

Contact Information

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Education

Ph.D. Bioinformatics, National Yang Ming University & Academia Sinica, TIGP^{*}, 2017.B.S. Biotechnology and Laboratory Science in Medicine, National Yang Ming University, 2008.

Research

TOPICS OF INTEREST

Bioinformatics — Quantitative Proteomics and Genomics, NGS Sequence Analysis especially in Structural Variation, Development of Bioinformatic Algorithms, Networks and Pathways Analysis, Annotation and Enrichment Analysis.

Biostatistics - Causal Inference and Mediation Analysis, Statistical Genetics, Clinical Meta Analysis.

Analytic Philosophy – Metaphysical Causation and Causal Models, Formal Epistemology and Modal Logics, Theology.

DOCTORAL DISSERTATION

Functional Analysis for Quantitative Proteomic Data. Advisor: Kun-Pin Wu. Committee: Hsuan-Cheng Huang, Yi-Hau Chen, Wailap Victor Ng, Hsin-Chou Yang, and Kun-Pin Wu.

Publications in Refereed Journals

Causal inference in the multiverse of hazard. Yen-Tsung Huang and En-Yu Lai. – under review. Massively parallel multi-mediator analyses under a composite null.

En-Yu Lai and Yen-Tsung Huang.

in revision.

Natural killer cells influence the clearance of SARS-CoV-2

Wan-Chen Hsieh, **En-Yu Lai**, Yu-Ting Liu, Yi-Fu Wang, Yi-Shiuan Tzeng, Lu Cui, Yun-Ju Lai, Hsiang-Chi Huang, Jia-Hsin Huang, Hung-Chih Ni, Dong-Yan Tsai, Jian-Jong Liang, Chun-Che Liao, Ya-Ting Lu, Laurence Jiang, Ming-Tsan Liu, Yao-Ming Chang, Gerlinde Wernig, Chia-Wei Li, Kuo-I Lin, Yi-Ling Lin, Huai-Kuang Tsai, Yen-Tsung Huang, and Shih-Yu Chen

Journal of Clinical Investigation (2021+) accepted.

Estimating the causal effect of embryo transfer day on clinical in vitro fertilization outcomes using propensity score matching. Han-Chih Hsieh, Chun-I Lee, **En-Yu Lai**, Jia-Ying Su, Yi-Ting Huang, Wei-Lin Zheng, Chien-Hong Chen, Chun-Chia Huang, Pin-Yao Lin, Maw-Sheng Lee, Mark Liu and Yen-Tsung Huang

BMC Pregnancy and Childbirth (2021+) accepted.

A mediator analysis for a binary non-rare outcome with sequentially ordered multiple mediators.

 ${\bf En-Yu}\;{\bf Lai}^{\dagger},$ Stephannie Shih $^{\dagger},$ Yen-T
sung Huang, and Shunping Wang.

- Statistics in Medicine (2020) Volume 39, Issue 10, Pages 1415-1428.

^{*}Taiwan International Graduate Program by Academia Sinica.

[†]Equally contributed.

A knowledge-based T^2 -statistic to perform pathway analysis for quantitative proteomic data. En-Yu Lai, Yi-Hau Chen, and Kun-Pin Wu.

- PLoS Computational Biology (2017) Volume 13, Issue 6, e1005601.

Drug class, renal elimination, and outcomes of direct oral anticoagulants in Asian patients: A meta-analysis. Kang-Ling Wang, Chun-Chih Chiu, Robert Giugliano, Doreen Tan, Chun-Yi Lin, **En-Yu Lai**, Shinya Goto, and Chern-En Chiang. – Journal of Stroke and Cerebrovascular Diseases (2018) Volume 27, Issue 4, Pages 857-864.

Once- or twice-daily mon-vitamin K antagonist oral anticoagulants in Asian patients with atrial fibrillation: A metaanalysis of randomized controlled trials.

Kang-Ling Wang, Chun-Chih Chiu, Doreen Su-Yin Tan, Chun-Yi Lin, **En-Yu Lai**, Shinya Goto, Robert P. Giugliano, and Chern-En Chiang. – Journal of the Formosan Medical Association (2017) Volume 116, Issue 8, Pages 591-598.

Extracranial arterial and venous thromboembolism in patients with atrial fibrillation: A meta-analysis of randomized controlled trials.

Kang-Ling Wang, Harry R. Büller, Shinya Goto, Chun-Yi Lin, **En-Yu Lai**, Chun-Chih Chiu, Chern-En Chiang, and Robert P. Giugliano. — Heart Rhythm (2017) Volume 14, Issue 4, Pages 599-605.

Standard versus low dose non-Vitamin K antagonist oral anticoagulants in asian patients with atrial fibrillation: A metaanalysis of contemporary randomized controlled trials.

Kang-Ling Wang, Robert P. Giugliano, Shinya Goto, Chun-Chih Chiu, Chun-Yi Lin, **En-Yu Lai**, and Chern-En Chiang. — Heart Rhythm (2016) Volume 13, Issue 12, Pages 2340-2347.

A linear-time algorithm for reconstructing zero–recombinant haplotype configuration on a pedigree. En-Yu Lai, Wei-Bung Wang, Tao Jiang, and Kun-Pin Wu.

- BMC Bioinformatics (2012) Volume 13, Supplement 17, S19.

BOOK CHAPTER

Causal models. En-Yu Lai, Wen-Fang Wang. – accepted.

INVITED TALKS / CONFERENCE TALKS

Massively parallel multi-mediator analyses under a composite null.

- South Taiwan Statistics Conference (STSC) 2021 in Kaohsiung, oral presentation
- ISMB/ECCB 2021 virtual conference, poster presentation
- South Taiwan Statistics Conference (STSC) 2020 in Chiayi, oral presentation
- A mediator analysis for a binary non-rare outcome with sequentially ordered multiple mediators.
- Chinese Statistical Association Taiwan (CSAT) Annual Meeting 2019 in Hsinchu, oral presentation

Mediation analyses of ultraviolet, air pollution, and structural variations using the Taiwan Biobank.

- ICSA International Conference 2019 in Hangzhou, oral presentation
- South Taiwan Statistics Conference (STSC) 2019 in Taichung, oral presentation
- ISI-ISM-ISSAS Joint Conference 2019 in Taipei, oral presentation
- Program in Quantitative Genomics (PQG) 2018 in Boston, poster presentation
- A genetic algorithm to locate responsive subpathways for time-course proteomic data.
- HUPO 2016 in Taipei, oral presentation
- A knowledge-based T^2 -statistic to perform pathway analysis for quantitative proteomic data.
- ISMB/ECCB 2017 in Prague, poster presentation
- Taiwan Proteomics Society (TPS) Annual Meeting 2014 in Taipei, poster presentation
- RECOMB 2013 in Beijing, poster presentation

A linear-time algorithm for reconstructing zero-recombinant haplotype configuration on a pedigree.

- InCoB 2012 in Bangkok, oral presentation
- ISMB 2010 in Boston, poster presentation

Scientific Software

PGCtest: Identifying phenome-genome functional clusters. [to submit]
https://github.com/roqe/PGCtest

MACtest: Multi-mediator analyses under a composite null. [to submit] https://github.com/roqe/MACtest

SOMM: Mediation analysis with sequentially ordered multiple mediators. [Statistics in Medicine, 2020] https://github.com/roqe/SOMM

T2GA: Pathway analysis under a self-contained null. [PLoS Computational Biology, 2017] https://github.com/roqe/T2GA

KCHI: A linear-time algorithm for haplotype inference. [BMC Bioinformatics, 2012] https://github.com/roqe/KCHI

Experiences

POSTDOCTORAL FELLOW

Variant component test on a model of causally ordered multiple mediators. – In Dr. Huang's lab at Institute of Statistical Science, Academia Sinica, 2018 – present.

Analysis of structure variation using the Taiwan Biobank NGS data.

In Dr. Huang's lab at Institute of Statistical Science, Academia Sinica, 2017 – present.

Consultant

Cooperation Center of Statistics and Data Science at Academia Sinica, Nov 2019 – present. General Clinical Research Center at Taipei Veterans General Hospital, Jan 2016 – Sept 2017.

Lecturer

TIGP-MBAS/MCB/MM 2019-2022: Biostatistics Summer 2018: Introduction to statistical learning with applications in R

TEACHING ASSISTANT

Autumn 2015: Introduction to statistical learning and data mining

Spring 2014: Bioinformatic algorithms; Machine learning and statistical inference

Autumn 2012: An overview of computer science

Autumn 2010: Basic molecular biology; Algorithms and data structures

Research Assistant

Algorithm development of gene ontology, network module and pathway analysis for quantitative proteomic data.

In Dr. Wu's lab at Institute of BioMedical Informatics, National Yang Ming University, 2015 – 2017.

In Dr. Chen's lab at Institute of Chemistry, Academia Sinica, 2012 – 2015.

Algorithmic studies of protein identification with glycosylation.

- In Dr. Wu's lab at Institute of BioMedical Informatics, National Yang Ming University, 2008 - 2009.

VISITING RESEARCHER

A linear-time algorithm for reconstructing zero–recombinant haplotype configuration on a pedigree. – In Dr. Jiang's lab at Bioinformatics Division, TNLIST[‡], Beijing, Summer 2009.

INTERNSHIP

Medical Technologist at Taipei Veterans General Hospital, July 2007 - January 2008.

[‡]Tsinghua National Laboratory for Information Science and Technology

Miscellaneous

Translator at *Research Ethics Education*, Jun 2014 – Mar 2015. Database developer at *MSI Taiwan*, Jan 2013 – Mar 2014.

Honors & Awards

Fellowships

Academia Sinica Postdoctoral Fellowship, July 2020 – June 2022 (proposal acceptance rate: \sim 30%). Academia Sinica Postdoctoral Fellowship, December 2018 – June 2020 (proposal acceptance rate: \sim 30%). NSC and ISCB Travel Grant for attending *RECOMB 2013*. NSC and APBioNET Travel Grant for attending *InCoB 2012*. NSC Travel Grant for attending *ISMB 2010*. TIGP Doctoral Fellowship, September 2009 – Augest 2012. NYMU Student Exchange Fellowship for internship at *TNLIST* during 2009 summer.

References

Kun-Pin Wu Professor Institute of BioMedical Informatics National Yang Ming Chiao Tung University +886-2-2826-7273 kpwu@nycu.edu.tw Yi-Hau Chen Research Fellow Institute of Statistical Science Academia Sinica +886-2-2787-1968 yhchen@stat.sinica.edu.tw Yen-Tsung Huang Research Fellow Institute of Statistical Science Academia Sinica +886-2-2787-5685 ythuang@stat.sinica.edu.tw