Michael Seo

Email: swj8874@gmail.com Website: mikejseo.github.io Mobile: +41 - 79 - 835 - 0405Github: github.com/mikejseo

EDUCATION

University of Bern Bern, Switzerland Ph.D. Biostatistics 2019 - 2022 (Expected)

Stanford University Stanford, CA M.S. Statistics; GPA: 3.7/4.0 2012 - 2014

Duke University Durham, NC B.S. Statistics, Graduation with High Distinction; GPA: 3.7/4.0 2007 - 2011

EXPERIENCE

Institute of Social and Preventive Medicine, University of Bern

Bern, Switzerland 2019 - Present

Ph.D. Student in Biostatistics

- o Real-world effectiveness: Developed models that combine individual patient data from randomized controlled trials and non-randomized studies when aiming to predict outcomes for a set of treatments.
- o Variable selection and shrinkage: Compared variable selection and shrinkage methods for estimating patient-specific treatment effects in individual patient data meta-analysis.
- Multiple imputation: Explored methods of addressing the systematically missing predictors problem, when the aim is to build a prediction model using data from multiple studies.

LLX Solutions Boston, MA Biostatistician2018 - 2019

- o Experimental design: Drafted statistical analysis plans which address study designs and methodologies on how to evaluate efficacy and safety of the new drug in development.
- Data manipulation: Transformed raw clinical data into datasets that meet FDA standards using SQL.

Center for Evidence Synthesis in Health, Brown University Research Associate

Providence, RI 2015 - 2018

o Network meta-analysis: Developed an R package for Bayesian network meta-analysis which allows simultaneous comparison of multiple treatments.

- $\circ \ \textbf{N-of-1 clinical trial} : \ Developed \ a \ Bayesian \ statistical \ tool \ to \ analyze \ single \ patient \ trials \ with \ crossover \ design \ and$ applied it to give individualized recommendations of carbohydrate diet for patients with inflammatory bowel disease.
- o Bayesian Calibration: Developed a Markov cohort simulation on survival outcomes after contralateral prophylactic mastectomy and used Bayesian calibration to estimate parameters such as probability of metastasis.

Selected Publications

- Research Publication: Seo M, Debray TPA, Ruffieux Y, Gsteiger S, Bujkiewicz S, Finckh A, Egger M, Efthimiou O (Under Revision). Combining individual patient data from randomized and non-randomized studies to predict real-world effectiveness of interventions.
- Research Publication: Seo M, White IR, Furukawa TA, Imai H, Valgimigli M, Egger M, Zwahlen M, Efthimiou O (2021). Comparing methods for estimating patient-specific treatment effects in individual patient data meta-analysis. Statistics in Medicine, 40, 1553-1573.
- Research Publication: Seo M, Furukawa TA, Veroniki AA, Pillinger T, Tomlinson A, Salanti G, Cipriani A, Efthimiou O (2021). The Kilim plot: A tool for visualizing network meta-analysis results for multiple outcomes. Research Synthesis Methods, 12, 86-95.
- Research Publication: Furukawa TA, Debray T, Akechi T, Yamada M, Kato T, Seo M, Efthimiou O (2020). Can personalized treatment prediction improve the outcomes, compared with the group average approach, in a randomized trial? Developing and validating a multivariable prediction model in a pragmatic megatrial of acute treatment for major depression. Journal of Affective Disorders, 274, 690-697.
- R Package: Michael Seo and Christopher Schmid (2020). bnma: Bayesian Network Meta-Analysis using 'JAGS'. R package version 1.4.0. https://CRAN.R-project.org/package=bnma.
- R Package: Robert Tibshirani, Michael Seo, Gil Chu, Balasubramanian Narasimhan, and Jun Li (2018). samr: Significance Analysis of Microarrays. R package version 3.0. https://CRAN.R-project.org/package=samr.

SKILLS

• Programming: R, Python, SAS, SQL, C++, Java, LaTeX

• Statistics: Bayesian Methods, (Network) Meta Analysis, Machine Learning, Causal Inference