Michael Seo

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EDUCATION

University of Bern Bern, Switzerland Ph.D. Biostatistics and Epidemiology 2019 - 2022

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

Duke University North Carolina, USA

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

EXPERIENCE

Institute of Social and Preventive Medicine, University of Bern

Ph.D. Student in Biostatistics and Epidemiology 2019 - 2022

Bern, Switzerland

o Variable selection and shrinkage: Compared variable selection and shrinkage methods for estimating patient-specific treatment effects in individual patient data meta-analysis.

• Real-world effectiveness: Developed models that combine individual patient data from randomized controlled trials and observational studies when aiming to predict outcomes for a set of treatments.

o Missing data: 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 Massachusetts, USA Biostatistician2018 - 2019

• Study design: Developed statistical analysis plans and TFL shells for Phase I trials.

• Statistical programming: Generated and reviewed CDISC SDTM and ADaM datasets using SAS.

Center for Evidence Synthesis in Health, Brown University Rhode Island, USA Ph.D. Student in Biostatistics 2015 - 2017

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

PUBLICATIONS

- Research Publication: Seo M, Furukawa TA, Karyotaki E, Efthimiou O (2023). Developing prediction models when there are systematically missing predictors in individual patient data meta-analysis, Research Synthesis Methods, 14, 455-467.
- Research Publication: Seo M, Debray TPA, Ruffieux Y, Gsteiger S, Bujkiewicz S, Finckh A, Egger M, Efthimiou O (2022). Combining individual patient data from randomized and non-randomized studies to predict real-world effectiveness of interventions. Statistical Methods in Medical Research, 31, 1355-1373.
- 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.
- R Package: Michael Seo (2022). bipd: Bayesian Individual Patient Data Meta-Analysis using 'JAGS'. R package version 0.3. https://CRAN.R-project.org/package=bipd.
- R Package: Michael Seo and Christopher Schmid (2020). bnma: Bayesian Network Meta-Analysis using 'JAGS'. R package version 1.5.0. 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

Statistics: Bayesian methods, (network) meta-analysis, causal inference