Michael Seo

CONTACT INFORMATION	My personal website: mikejseo.github.io	Email: swj8874@gmail.com
Research Interests	Bayesian methods; (network) meta-analysis; machine learning	
EDUCATION	Ph.D. Biostatistics, University of Bern, 2019-2022 (Expected)	
	M.S. Statistics, Stanford University (GPA: 3.7), 2012-2014	
	B.S. Statistics, Graduation with High Distinction, Duke University (GPA: 3.7), 2007-2011	
EMPLOYMENT	Ph.D. student in Biostatistics , Institute of Social and Preventive Medicine, University of Bern, Switzerland, 2019-Present	
	Biostatistician, LLX Solutions, Boston, 2018-2019	
	Research Associate, Center for Evidence Synthesis in Health, Brown University, 2015-2018	
	Quantitative Analyst, In4mation Insights, Boston, 2015 Summer	
	Research Assistant, Department of Statistics, Stanford University, 2014-2015	
Publications	Seo M, White IR, Furukawa TA, Imai Hissei, Valgimigli M, Egger M, Zwahlen M, Efthimiou O (Under revision). Comparing methods for estimating patient-specific treatment effects in individual patient data meta-analysis.	
	Seo M , Furukawa TA, Veroniki AA, Pillinger T, Tomlinson A, Salanti G, Cipriani A, Efthimiou O (2020). The Kilim plot: A tool for visualizing network meta-analysis results for multiple outcomes. <i>Research Synthesis Methods</i> , 1-10.	
	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. <i>Journal of Affective Disorders</i> , 274, 690-697.	
	Khan MS, Khan AR, Khan AI, Seo M , Yasmin F, Usman MS, Moustafa A, Schmid CH, Kalra A, Ikram S (2020). Comparison of revascularization strategies in patients with acute coronary syndrome and multivessel coronary disease: A systematic review and network meta-analysis. <i>Catheterization and cardiovascular interventions</i> , 1-8.	
R PACKAGES	Michael Seo and Christopher Schmid (2020). bnma: Bayesian R package version 1.3.0. https://CRAN.R-project.org/package	
	R. Tibshirani, Michael J. Seo , G. Chu, Balasubramanian Nar Significance Analysis of Microarrays. R package version 3.0.	
Contributed Presentations	Predicting real world effectiveness of interventions, combining individual patient data from multiple randomized and non-randomized studies. 41st Annual Conference of the International Society for Clinical Biostatistics (ISCB), Krakow; August, 2020.	
	Comparing methods for variable selection in individual patient data meta-analysis. XXXIst Conference of the Austro-Swiss Region (ROeS) of the International Biometric Society, Lausanne; September, 2019.	
TEACHING EXPERIENCE	Prognostic Research: from Basics to Modelling, University of Bern, 2020	
	Evidence Synthesis Methods, University of Bern, 2019	
	Applied Regression Analysis (PHP 2511), Brown, Spring 2016	
	Fundamentals of Probability and Statistical Inference (PHP 2515), Brown, Fall 2015	
	Probability and Statistics Inference (STAT 103), Duke, Spring 2011	

Statistics (STAT 114), Duke, Fall 2010

Probability and Statistics for Engineers (STAT 113), Duke, Spring 2010 $\,$

Probability (STAT 104), Duke, Fall 2009

PROGRAMMING R, Python, SAS

Languages Korean (Native), English (Fluent)

CITIZENSHIP U.S.A