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 Switzerland, 2019-Present	ve Medicine, University of Bern,
	Biostatistician, LLX Solutions, Boston, 2018-2019	
	Research Associate, Center for Evidence Synthesis in Health, Bro	own University, 2015-2018
	Quantitative Analyst, In4mation Insights, Boston, 2015 Summer	
	Research Assistant, Department of Statistics, Stanford University	y, 2014-2015
Publications	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. <i>Statistics in Medicine</i> , 1-21.	
	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. <i>Research Synthesis Methods</i> , 12, 86-95.	
	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, Mo S (2020). Comparison of revascularization strategies in patients w multivessel coronary disease: A systematic review and network m cardiovascular interventions, 96, E447-E454.	with acute coronary syndrome and
R PACKAGES	Michael Seo and Christopher Schmid (2020). bnma: Bayesian Net R package version 1.4.0. https://CRAN.R-project.org/package=b	
	R. Tibshirani, Michael J. Seo , G. Chu, Balasubramanian Narasim Significance Analysis of Microarrays. R package version 3.0. https://doi.org/10.1001/j.com/	· /
Contributed Presentations	Predicting real world effectiveness of interventions, combining individual randomized and non-randomized studies. 41 st Annual Conference of Biostatistics (ISCB), Krakow; August, 2020.	
	The Kilim plot: a tool for visualizing network meta-analysis results for multiple outcomes (poster). 41 st Annual Conference of the International Society for Clinical Biostatistics (ISCB), Krakow; August, 2020.	
	Comparing methods for variable selection in individual patient data of the Austro-Swiss Region (ROeS) of the International Biometric S	*
TEACHING	Prognostic Research: from Basics to Modelling, University of Bern,	2020
Assistantship		
ASSISTANTSHIP	Evidence Synthesis Methods, University of Bern, 2019	

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