

# Michael Seo

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CONTACT INFORMATION	My personal website: <a href="https://mikejseo.github.io">mikejseo.github.io</a>	Email: <a href="mailto:swj8874@gmail.com">swj8874@gmail.com</a>
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	<b>Ph.D. student in Biostatistics</b> , Institute of Social and Preventive Medicine, University of Bern, Switzerland, 2019-Present  <b>Biostatistician</b> , LLX Solutions, Boston, 2018-2019  <b>Research Associate</b> , Center for Evidence Synthesis in Health, Brown University, 2015-2018  <b>Research Assistant</b> , Department of Statistics, Stanford University, 2014-2015	
PUBLICATIONS	<b>Seo M</b> , 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, <b>Seo M</b> , 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, <b>Seo M</b> , 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	<b>Michael Seo</b> and Christopher Schmid (2020). bnma: Bayesian Network Meta-Analysis using 'JAGS'. R package version 1.3.0. URL <a href="https://CRAN.R-project.org/package=bnma">https://CRAN.R-project.org/package=bnma</a> .  R. Tibshirani, <b>Michael J. Seo</b> , G. Chu, Balasubramanian Narasimhan and Jun Li (2018). samr: SAM: Significance Analysis of Microarrays. R package version 3.0. <a href="https://CRAN.R-project.org/package=samr">https://CRAN.R-project.org/package=samr</a> .	
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	<b>Teaching Assistant</b> , Evidence Synthesis Methods, University of Bern, 2019 <b>Teaching Assistant</b> , Applied Regression Analysis (PHP 2511), Brown, Spring 2016 <b>Teaching Assistant</b> , Fundamentals of Probability and Statistical Inference (PHP 2515), Brown, Fall 2015 <b>Teaching Assistant</b> , Probability and Statistics Inference (STAT 103), Duke, Spring 2011 <b>Teaching Assistant</b> , Statistics (STAT 114), Duke, Fall 2010 <b>Teaching Assistant</b> , Probability and Statistics for Engineers (STAT 113), Duke, Spring 2010 <b>Teaching Assistant</b> , Probability (STAT 104), Duke, Fall 2009	
PROGRAMMING	R (Extensive), Python, SAS	
LANGUAGES	Korean (Native), English (Fluent)	
CITIZENSHIP	U.S.A	