

# MICHAEL SEO

☎ +41 798350405 ✉ [mikejseo7\[at\]gmail.com](mailto:mikejseo7[at]gmail.com)  [LinkedIn](#)  [Personal Website](#)

## EDUCATION

---

### University of Bern

*Ph.D. Biostatistics and Epidemiology*

**Bern, Switzerland**

*2019 - 2022*

### Stanford University

*M.S. Statistics; GPA: 3.7/4.0*

**California, USA**

*2012 - 2014*

### Duke University

*B.S. Statistics, Graduation with High Distinction; GPA: 3.7/4.0*

**North Carolina, USA**

*2007 - 2011*

## WORK EXPERIENCE

---

### Roche

**2022 - Present**

*Access Evidence Lead (HTA Statistician)*

*Basel, Switzerland*

- Drafted indirect treatment comparisons statistical analysis plans and reports needed for the HTA reimbursement submissions.
- Worked with Phase II/III clinical trials data to implement inverse probability weights methods and network meta-analysis.
- Developed an R package (maicplus) for matching-adjusted indirect comparison which adjusts for differences in baseline characteristics between treatment groups when only aggregate data is available for the comparator study.

### Institute of Social and Preventive Medicine, University of Bern

**2019 - 2022**

*Ph.D. Student in Biostatistics and Epidemiology*

*Bern, Switzerland*

- Compared variable selection and shrinkage methods for estimating patient-specific treatment effects in individual patient data meta-analysis.
- Developed models that combine individual patient data from randomized controlled trials and observational studies when aiming to predict outcomes for a set of treatments.
- 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

**2018 - 2019**

*Biostatistician*

*Massachusetts, USA*

- Drafted statistical analysis plans for Phase I trials to evaluate safety of the new drug in development.
- Transformed clinical data into datasets that meet FDA standards using clinical SAS programming.

### Department of Biostatistics, Brown University

**2015 - 2017**

*Research Associate*

*Rhode Island, USA*

- Developed an R package (bnma) for Bayesian network meta-analysis which allows simultaneous comparison of multiple treatments.
- 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.

## SKILLS

---

**Programming:** R, Python, SAS

**Statistics:** indirect treatment comparison, network meta-analysis, causal inference