Sina Rüeger Data Analyst

Education

PhD in Life Sciences in 2018

Supervision by Prof. Zoltán Kutalik, <u>Statistical Genetics Group</u> at the Lausanne University Hospital.

Thesis: Integrative statistical analysis of -omics and genome-wide association studies data.

- Improvement of summary statistics imputation for application in more realistic scenarios and helping in building an easy to use software.
- Identification of genetic risk factors in different disease areas using GWAS methodology.
- Validation of gene expression-trait associations by applying text processing to a drugbank database, extracting drug target information and matching the results with gene expression-trait associations.
- Master of Science in Engineering in 2011

Zurich University of Applied Sciences, Winterthur, average mark: 86%. Development of a statistical model to predict stages of Parkinson's disease from brain perfusion data, leading to a better understanding of disease progression. New Zealand Brain Institute, Christchurch, NZ.

Engineering degree in Data Analysis & Process Design in 2008
Zurich University of Applied Sciences, Winterthur, average mark: 82%.

Experience

- Fellay Lab, EPFL

Postdoc: since Sept 2018

Analysis of human genomic data in the context of infectious diseases.

Division of Biostatistics, University of Zurich

Biostatistician: Feb 2012 - Mar 2013

- Provided statistical consulting for medical staff and students.
- Analysed medical data for over eight research groups.
- Institute of Data Analysis & Process Design, ZHAW

Research Assistant: Mar 2009 - Jan 2012

- Provided statistical data analysis for various in-house projects.
- Built an application to automate cluster analysis on consumer testing data.
- Predicted vehicle type using sensor data and random forests.

Publications

See full list <u>here</u>

- Sina Rüeger et al. (2015) Impact of common risk factors of fibrosis progression in chronic hepatitis C. <u>Gut</u>.
- Sina Rüeger et al. (2017) Improved imputation of summary statistics for realistic settings. <u>BioRxiv</u>.
- Sina Rüeger et al. (2018) Evaluation and application of summary statistic imputation to discover new height-associated loci. <u>PLOS</u> Genetics.

Contact

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Skills

Statistical data analysis

Visualising data

R programming

Used daily since 10 years

Git

Knitr/RMarkdown

Genomics tools

High performance computing

Presentation & Communication

Adobe Illustrator

German native speaker

English fluent

French good command

Awards & Activities

Awards for best student presentation at conferences (ESHG, EMGM, ePerMed).

Founding member and co-organiser of <u>R-Ladies Lausanne</u>.