# Sina Rüeger

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Areas of	biostatistical	and	data	analytica	l experience

Statistical modelling Regression modelling techniques (linear mixed effects, nonlinear, generalised additive,

robust and survival models), analysis of variance, model & variable selection, multiple

testing approaches.

Data mining Data dimension reduction, classification, unsupervised clustering.

Data visualisation Visual representation of concepts and results.

Data pre-processing Data storage & documentation, quality control, missing data handling.

Trackability of work Version control, documentation with knitr/rmarkdown and best-practices in program-

ming

Domain knowledge Genomics, Parkinsons disease, bariatric surgery, hepatitis C & other viral infections,

solid organ transplantation, multiple sklerosis, human height, critical care patients.

### Education

Oct-2013 – (Spring 2018) PhD student, Statistical Genetics Group, Institute of Social and Preventive Medicine,

Lausanne University Hospital (CHUV), Lausanne, Switzerland, supervised by Prof.

Zoltán Kutalik.

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

Feb-2009 - Jun-2011 Master of Science in Engineering, Zurich University of Applied Sciences, Win-

terthur, Switzerland.

Oct-2005 - Oct-2008 Dipl. Ing. Data Analysis and Process Design, Zurich University of Applied

Sciences, Winterthur, Switzerland.

## **Employment**

Feb-2013 – Sep-2013 Internship, Statistical Genetics Group, Department of Medical Genetics, University of Lausanne, Lausanne, Switzerland.

Feb-2012 – Mar–2013 **Biostatistician**, *Division of Biostatistics, ISPMZ (now EBPI)*, University of Zurich, Zurich, Switzerland.

Jan-2012 - Aug-2012 Biostatistician, Center of clinical nursing science, University Hospital Zurich, Zurich,

Mar-2009 – Jan-2012 **Research Assistant**, *Institute of Data Analysis and Process Design*, *Zurich University of Applied Sciences*, Winterthur, Switzerland.

Jul-2007 - Sep-2007 Internship, Official Statistics Zurich, Zurich, Switzerland.

# Computing skills

Statistics/Programming R [highly proficient], Matlab [familiar], Java [familiar], SQL [familiar]

Genomics tools Plink, QCTOOL, GTCA, ...

Support GitHub, LaTex, emacs, Affinity/Adobelllustrator/Inkscape

R Daily used for statistical computing and data visualisation; familiar with the architecture of R, package development and best-practices.

Computing experience Five years of experience using high performance clusters for analysis and storage of large files.

#### Other activities

R-Ladies Founding member and co-organiser of R-Ladies Lausanne. R-Ladies is a world-wide organization to promote gender diversity in the R community.

## Languages

**German** [native speaker, C2], **English** [fluent, C1], **French** [good command, B1]

#### Awards

07-Apr-2017 The 'Human Heredity' Best Student Platform Presentation Award at the European Mathematical Genetics Meeting in Tartu, Estonia.

15-Mar-2017 Poster prize at the ePerMed Workshop about 'Functional annotation of genome-wide variants' in Lausanne, Switzerland.

Lodewijk Sandkuijl Award for an outstanding talk in the field of complex genetics and statistical genetics at the European Society of Human Genetics Conference, Barcelona, Spain.

## Output list

#### Main publications

Sina Rüeger, Aaron McDaid, and Zoltán Kutalik. Improved imputation of summary statistics for realistic settings. bioRxiv, page 203927, 2017b

Sina Rüeger, Aaron McDaid, and Zoltán Kutalik. Evaluation and application of summary statistic imputation to discover new height-associated loci. bioRxiv, page 204560, 2017a

Sina Rüeger et al. Impact of common risk factors of fibrosis progression in chronic hepatitis C. Gut, 64(10):1605-1615, 2015