PHD STUDENT

BioQuant, Heidelberg University

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Data scientist tackling biomedical challenges

Education

Bielefeld UniversityBielefeld, Germany

BACHELOR OF SCIENCE - MOLECULAR BIOTECHNOLOGY

2011-14

• Thesis: Effect of overexpression of xanA on the growth and xanthan production of Xanthomonas campestris pv. Campestris in the laboratory and in the model

Bielefeld University

Bielefeld, Germany

MASTER OF SCIENCE - GENOME BASED SYSTEMS BIOLOGY

2014-17

• Thesis: Coupling of Metabolism and Gene Regulation to identify Gene Targets in Living Cells

RWTH Aachen Aachen Aachen, Germany

PHD 2017-19

• From gene expression to pathway and transcription factor activities to acquire functional and mechanistic insight into chronic liver diseases

Heidelberg University

Heidelberg, Germany

CONTINUATION OF PHD 2019-now

· From gene expression to pathway and transcription factor activities to acquire functional and mechanistic insight into chronic liver diseases

Working Experience

Reseach Center Jülich

INTERNSHIP 2015 (3 month)

• Studying Escherichia coli metabolism with focus on the uncertainties of the biomass equation

Alacris Theranostics GmbH Berlin, Germany

INTERNSHIP

2016 (3 month)

 Application of a mechanistic model and an artificial neural network for simulation and parameter optimization of the drug response model in the context of personalized medicine

Insilico Biotechnology AG Stuttgart, Germany

MASTER STUDENT 2016-2017 (9 month)

• Coupling of Metabolism and Gene Regulation to identify Gene Targets in Living Cells

Teaching Experience

Bielefeld University

Bielefeld, Germany

STUDENT ASSISTANT JOB - MATHEMATICS FOR BIOLOGISTS

2013-15

Bielefeld University

Bielefeld, Germany

STUDENT ASSISTANT JOB - STATISTICS/INFORMATICS FOR BIOLOGISTS 2013-15

Skills

Programming

R, Python, Matlab, Bash

Packages

TIDYVERSE, SHINY, BOOKDOWN, PKGDOWN, TESTTHAT, PANDAS, NUMPY

Tools

MAY 2020

GIT, TRAVIS CI, LATEX

Conferences Attended

SBMC Bremen, Germany

TALK: Unveiling of conserved transcriptomics perturbation responses in Mice and Human

2018

RECOMB/ISCB New York City, USA

POSTER: UNVEILING OF CONSERVED TRANSCRIPTOMICS PERTURBATION RESPONSES IN MICE AND HUMAN

Basel, Switzerland

POSTER: ROBUSTNESS AND APPLICABILITY OF FUNCTIONAL GENOMIC TOOLS ON SCRNA-SEO DATA

2019

Selected Publications

ISBM/ECCB

- 1. Holland, C, J Tanevski, J Perales-Patón, J Gleixner, M Kumar, E Mereu, and ... (2020). Robustness and applicability of transcription factor and pathway analysis tools on single-cell RNA-seq data. *Genome Biology*.
- 2. Tajti, F, C Kuppe, A Antoranz, M Ibrahim, H Kim, F Ceccarelli, and ... (2020). A functional landscape of CKD entities from public transcriptomic data. *Kidney International Reports*.
- 3. Garcia-Alonso, L, C Holland, M Ibrahim, D Turei, and J Saez-Rodriguez (2019). Benchmark and integration of resources for the estimation of human transcription factor activities. *Genome research*.
- 4. Holland, C, B Szalai, and J Saez-Rodriguez (2019). Transfer of regulatory knowledge from human to mouse for functional genomics analysis. *Biochimica et Biophysica Acta (BBA)-Gene Regulatory Mechanisms*.
- 5. Ghallab, A, M Myllys, CH Holland, A Zaza, W Murad, R Hassan, and ... (2019). Influence of Liver Fibrosis on Lobular Zonation. *Cells*.
- 6. Szalai, B, V Subramanian, C Holland, R Alföldi, L Puskás, and ... (2019). Signatures of cell death and proliferation in perturbation transcriptomics data—from confounding factor to effective prediction. *Nucleic acids research*.