#### **DECEMBER 2014 - JUNE 2015**

# Jan Gleixner Curriculum Vitae

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#### **EDUCATION**

2018 – PRESENT **Biosciences** PHD – TBD

Heidelberg University

2013 - 2018 Molecular Biotechnology

**Major: Bioinformatics**Master of Science – 1.1

Heidelberg University

 $2010-2013 \quad \textbf{Molecular} \quad \textbf{Biotechnology}$ 

**Major: Bioinformatics** 

Bachelor of Science – 1.9 *Heidelberg University* 

2009 Secondary School

Major: Math & Informatics

Abitur – 1.8

Carl-Zeiss-Gymnasium, Jena

#### WORK EXPERIENCE

MAR 2018 - EXP. 2024

German Cancer Research center (DKFZ) – Michael Boutros' lab and European molecular biology laboratory (EMBL) – Oliver Stegle's group Computational Biology

Improvement of experimental and computational methods to quantify effects on gene expression on different levels in high throughput  $^{\rm l}$ 

JUN 2017 - FEB 2018

### European molecular biology laboratory (EMBL) – Oliver Stegle's group Causal Inference

Application of Invariant Causal Prediction (ICP) to single cell RNA expression CRISPR perturbation  ${\rm data}^2$ 

OCTOBER 2015 - DECEMBER 2015

Max-Planck-Institute for Empirical Inference – Jonas Peters' group

#### Causal Inference

Development a likelihood score based bootstrap hypothesis test for the existence of a total causal effect in the framework of causal additive models

### Heidelberg University – Labs of Barbara Di Ventura and Dirk Grimm Synthetic Biology

Improving gene therapy by engineering a split Cas9 enzyme with improved expression from self-complementary Adeno-associated virus (scAAV)

February 2014 - November 2014

#### iGEM team Heidelberg 2014 Synthetic Biology

Development of a standard for cloning of Intein fusion proteins and use of Intein mediated circularization for stabilization of enzymes in a team of eleven students working full time with acquired funds over a hundred thousand euro<sup>5</sup>

February 2011 - October 2013

## Max Planck Institutes for Neurobiology and for Medical Research — Moritz Helmstaedter's group Application of Machine Learning

Programming of artificial neural networks to automatically segment 3D-electron microscope images of brain tissue and porting of those to GPUs; Development of features for and use of Random Forests for synapse detection

AUGUST & SEPTEMBER 2012

### Duke University — Ute Hochgeschwender's lab Neuro-optogenetics

Internship to learn cell culture, patch clamping and other neurobiology skills by analyzing and improving a Channelrhodopsin-Aequorin fusion protein $^4$ 

 $May - Juli \ 2010$ 

## Leibniz Institute Natural Product Research and Infection Biology — Hans Knoell Institute Image Analysis Automation

Development of scripts in R and MATLAB for image analysis and statistical evaluation

Juni 2009

### Max Planck Institute for Biogeochemistry Rustle Classifier Development

Literature search, preliminary experiments and their analysis by use of MATLAB and R

#### LANGUAGE SKILLS

ADVANCED R. data.table

INTERMEDIATE Python, Theano/TensorFlow,

Stan, Java, C++, Lar, bash

BASIC LEVEL Perl, MATLAB, Haskell,

JavaScript, Powershell, C#, CUDA and Assembler

- <sup>1</sup>MC Funk et al., *Aged intestinal stem cells propagate cell-intrinsic sources of inflammaging in mice*, Developmental Cell 58, 2914–2929.e7 (2023).
- <sup>2</sup>CH Holland et al., Robustness and applicability of transcription factor and pathway analysis tools on single-cell RNA-seq data, Genome Biol 21, 36 (2020).
- <sup>3</sup>M Goethe, J Gleixner, I Fita, and JM Rubi, *Prediction of Protein Configurational Entropy (Popcoen)*, J. Chem. Theory Comput. 14, 1811–1819 (2018).
- <sup>4</sup>K Berglund et al., *Luminopsins integrate opto- and chemogenetics by using physical and biological light sources for opsin activation*, Proc Natl Acad Sci U S A 113, E358–367 (2016).
- <sup>5</sup>MC Waldhauer, SN Schmitz, C Ahlmann-Eltze, JG Gleixner, CC Schmelas, AG Huhn, C Bunne, M Büscher, M Horn, N Klughammer, et al., Backbone circularization of Bacillus subtilis family 11 xylanase increases its thermostability and its resistance against aggregation, Molecular BioSystems 11, 3231–3243 (2015).

- 2014 **iGEM International Genetically Engineered Machines competition**Winner team "Ring of fire" (Heidelberg)
- 2012 **DAAD's (German academic exchange service) RISE worldwide program**Full schoolarship to carry out a research internship at Duke University (NC, USA)
- 2011 SYNtheSYS Student competition on Synthetic & Systems Biology
  Winner team "Faster than life"
- 2009 **Jugend forscht** State level, 2. award