

# CAROLINE VIKTORIA WEIS

*Curriculum vitae*

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🔗 weis.ml

## EDUCATION

- 2017–present **Ph.D. candidate** in Machine Learning and Computational Biology at ETH Zurich, Switzerland  
*Machine Learning and Computational Biology*  
Adviser: Prof. Dr. Karsten Borgwardt
- 2014–2016 **M.Sc.** in Biotechnology at ETH Zurich, Switzerland, final grade **5.55** (very good)<sup>1</sup>  
Thesis: *Assessing the Potential of Feature-pairs in Predicting the Impact of Missense Variants*  
Advisers: Dr. L. Folkman, Dr. D. Grimm, Prof. Dr. Karsten Borgwardt
- 2010–2014 **B.Sc.** in Integrated Life Sciences at FAU Erlangen, Germany, final grade **1.6** (excellent)<sup>2</sup>  
Thesis: *Analysis of zinc-oxide particle growth under influence of triethylamine by using Small Angle X-Ray Scattering and UV/Vis Spectroscopy*  
Adviser: Prof. Dr. Tobias Unruh
- 2001–2010 *Abitur*<sup>3</sup>, Alexander-von-Humboldt Gymnasium Schweinfurt<sup>4</sup>, Germany, final grade **1.8** (very good)

## PROFESSIONAL EXPERIENCE

- 10/2015–03/2016 **Industrial research intern** at *Genedata AG*, Basel, Switzerland.  
As a Machine Learning research intern in the Screener Business Unit I assessed different algorithms for the task of classifying screening images. Images depicting bacteria treated with new compounds of unknown effect were classified into effect category. I developed a pipeline in R that has the potential to speed-up screening for drugs with a desired effect. Classical approaches such as support vector machine algorithms and t-SNE clustering provided good results, and a Genedata business poster I prepared for SLAS conference 2016 in San Diego set a new record for requested poster downloads at Screener Business Unit. Additionally I implemented a GUI for active learning in MATLAB, allowing for images classified with low probability to be presented to a human expert to be classified.
- 10/2015–03/2016 **Research assistant** at *Control Theory and Systems Biology group*, ETH Zurich, Basel, Switzerland.  
As a wetlab research assistant I performed standard tasks – such as Minipreps, PCRs, gel electrophoresis etc. – to assist projects lead by Post-Doctoral researchers.
- 09/2013–07/2014 **Academic research intern** at *Lawrence Berkeley National Laboratory*, Berkeley, USA.  
I worked at the SIBYLS beamline of the Physical Biosciences Division. I performed Protein Crystallography and Small Angle X-Ray Scattering experiments and subsequent data analysis. I worked on project, which eventually led to a publication in *Structure* while also providing guidance on data analysis to SIBYLS beamline users.  
Supervisor: Dr. Michal Hammel
- 04/2012–07/2012 **Summer research assistant** at *Chair of Crystallography and Structural Physics*, FAU Erlangen, Germany.  
In this summer intership I grew zinc oxide crystals, mounted them to a plate and aligned them using a goniometer.

## THESIS SUPERVISION

- 2020 Sebastian Balzer. *Improved MALDI-TOF based antimicrobial phenotype prediction through incorporating phylogenetic structure*  
Bachelor thesis, ETH Zurich
- 2019 Lucie Bourguignon. *Mortality prediction using self-reported health records and large scale genomic data*  
Master thesis, ETH Zurich

<sup>1</sup>ETH Zurich's Grading System: [https://ethz.ch/content/dam/ethz/special-interest/itet/department/Studies/Forms/20160112\\_Grading\\_System.pdf](https://ethz.ch/content/dam/ethz/special-interest/itet/department/Studies/Forms/20160112_Grading_System.pdf)

<sup>2</sup>German Grading System

<sup>3</sup>General qualification for university entrance

<sup>4</sup>Secondary school

## PUBLICATIONS

- In press **Caroline Weis**, Max Horn, Bastian Rieck, Aline Cuenod, Adrian Egli, Karsten Borgwardt. *Topological and kernel-based microbial phenotype prediction from MALDI-TOF mass spectra*. Bioinformatics, accepted at ISMB 2020.
- 2020 **Caroline Weis**, Catherine Jutzeler, Karsten Borgwardt. *Machine learning for microbial identification and antimicrobial susceptibility testing on MALDI-TOF mass spectra: a systematic review*. Clinical Microbiology and Infection, March 2020.  
<https://doi.org/10.1016/j.cmi.2020.03.014>
- 2019 **Caroline Weis**, Max Horn, Bastian Rieck, Karsten Borgwardt. *Sparse representations for MALDI-TOF based microbial classification*. Poster and peer-reviewed abstract. 14<sup>th</sup> Machine Learning in Computational Biology Meeting (MLCB).
- 2017 Jamie R. Wallen, Hao Zhang, **Caroline Weis**, Weidong Cui, Brittini M. Foster, Chris M. W. Ho, Michal Hammel, John A. Tainer, Michael L. Gross, Tom Ellenberger. *Hybrid Methods Reveal Multiple Flexibly Linked DNA Polymerases within the Bacteriophage T7 Replisome*. Structure, 25. 157–166., 2017.  
<https://doi.org/10.1016/j.str.2016.11.019>
- Oliver Ratmann, Emma B. Hodcroft, Michael Pickles, Anne Cori, Matthew Hall, Samantha Lycett, Caroline Colijn, Bethany Dearlove, Xavier Didelot, Simon Frost, A.S. Md Mukarram Hossain, Jeffrey B. Joy, Michelle Kendall, Denise Kühnert, Gabriel E. Leventhal, Richard Liang, Giacomo Plazzotta, Art F.Y. Poon, David A. Rasmussen, Tanja Stadler, Erik Volz, **Caroline Weis**, Andrew J. Leigh Brown, Christophe Fraser, on behalf of the PANGEA-HIV Consortium. *Phylogenetic Tools for Generalized HIV-1 Epidemics: Findings from the PANGEA-HIV Methods Comparison*. Molecular Biology and Evolution, Volume 34, Issue 1, January 2017, Pages 185–203, 2017.  
<https://doi.org/10.1093/molbev/msw217>

## TEACHING EXPERIENCE

- In each of the following courses, I have served as a teaching assistant. Duties are listed for each course individually.
- 2018 + 2020 Exercises *Data Mining II*, ETH Zurich  
The course duties consisted of programming exercises in Python, creation and grading of bi-weekly exercises, as well as as the development and correction of exam questions.
- 2012 – 2013 Exercises *Mathematical modeling and statistics for scientists*, Department Mathematics, FAU Erlangen  
This computer science tutorial consisted of modeling and statistical analyses programming exercises in R which I had to present and supervise. In addition I supervised and corrected the exam.
- 2012 Exercises *Structural physics*, Chair of Crystallography and Structural Physics, FAU Erlangen  
The course duties consisted of presenting written exercises in crystallography and structural physics, as well as as the development and correction of exam questions.
- 2011 – 2012 Microscopy course *Biology for physicians*, Animal Physiology, Department of Biology, FAU Erlangen  
In this microscopy course I facilitated laboratory exercises about plant biology and basic physiology for first-year physiology students.

## SKILLS

Strong knowledge of Python for data analysis (numpy, scipy, scikit-learn) and working knowledge of deep learning frameworks (TensorFlow and Keras).

Working knowledge of R and MATLAB.

Strong knowledge of data visualization tools matplotlib and seaborn in Python and ggplot2 and in R, plus Bokeh and Rshiny for interactive visualization.

Knowledge of digital typesetting languages (T<sub>E</sub>X, L<sup>A</sup>T<sub>E</sub>X) and of the Git revision control system.

Strong writing and public speaking skills.

## LANGUAGES

<i>German</i>	native speaker
<i>English</i>	professional proficiency
<i>French</i>	basic knowledge

## SERVICE TO THE COMMUNITY

- 02/2020–  
05/2020 *Academic Jury member*, St. Gallen Symposium, St. Gallen, Switzerland  
As Academic Jury member I evaluate essays on the topic of *Freedom revisited* and identify the top 100 contributions which receive an all-expenses-covered invitation to the 2020 St. Gallen Symposium. I will also participate in the three-day St. Gallen Symposium, to discuss the most urging problems threatening our freedom at this time and what actions should be taken.  
(symposium cancelled after evaluation round due to *SARS-CoV-2* pandemic)
- 06/2019–  
06/2020 *Founding member of peer mentoring group 'Women in Data Science'*  
Along with two colleagues I successfully applied for a grant of 5'000 CHF provided by ETH Zurich's Fix-the-Leaky-Pipeline program. Among my tasks were to approach industry partners who serve as mentors for our peer groups, as well as leading peer meetings with fellow Data Science PhD students and Post-Doctoral researchers.
- 2018 *ETH representative*, ETH Zurich pavillion, World Economic Forum, Davos, Switzerland  
I represented ETH Zurich at the universities pavillion in Davos during the 2018 World Economic Forum. Over the course of four days I made the research topic of 'significant pattern mining' accessible to visitors, which ranged from Switzerland's leading research representatives and honorary guests to several high-school groups.
- 02/2015–  
02/2016 *Student and Academic Affairs Commissioner*, Biotechnology Student Association, ETH Zurich, Switzerland  
Among my duties as Commissioner was preparing, holding and interpreting the lecture evaluations each semester. I was also the primary contact person for communication between student and professor concerning teaching matters. I served as the student body contact person to admitted students before their arrival, organized welcome events and helped them settle in at our department at ETH Zurich.
- 02/2013–  
08/2014 *Board Member*, Biotechnology Student Initiative Erlangen, Germany  
At the Biotechnology Student Initiative I organized talks of professors and company representatives, soft skill workshops and company tours, e.g. at Siemens Healthcare in Erlangen.

## REFERENCES

PROF. DR. KARSTEN BORGWARDT  
*Machine Learning and Computational Biology Lab*  
 ETH Zurich  
[karsten.borgwardt@bsse.ethz.ch](mailto:karsten.borgwardt@bsse.ethz.ch)

Further references and credentials are available on request.

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