

ADRIAN EDWARD THOMAS HENKEL

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30.10.1998 | Nationality: German

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EDUCATION

TECHNICAL UNIVERSITY MUNICH / LUDWIG-MAXIMILIANS-UNIVERSITY

Master of Science Bioinformatics

MUNICH, GERMANY

2021 - NOW

TECHNICAL UNIVERSITY MUNICH / LUDWIG-MAXIMILIANS-UNIVERSITY

Bachelor of science Bioinformatics

MUNICH, GERMANY

2017 - 2021

- Thesis topic: Communication-efficient approaches for federated deep neural networks (grade 1.0)

LUDWIG-MAXIMILIANS-UNIVERSITY

Bachelor of science Statistics (minor CS)

MUNICH, GERMANY

2016 - 2017

PROFESSIONAL EXPERIENCE

DISCOVERY LIFE SCIENCES BIOMARKER SERVICES GMBH

Working Student

MUNICH / KASSEL, GERMANY | ALABAMA, US

2020 - NOW

- Proposing, planning, managing, and implementing a video tutorial series (~20 videos) of the image analysis software Visiopharm to increase productivity and to avoid recurring time-consuming trainings.
- Data visualization of produced clinical data with Python.
- Development, testing and validation of multiple GCP compliant Application Protocols (APPs) for Visiopharm.

ASTRAZENECA COMPUTATIONAL PATHOLOGY

Working student

MUNICH, GERMANY

Mar. 2019 - MAR. 2020

- Planning and implementation of neural network with TensorFlow for segmentation of immunohistochemistry (IHC) tissue images.
- Analysis of different machine learning algorithms (NNs, RFs)
- Automation of training workflows with pytest and GitLab CI/CD tools.

DEFINIENS AG

Intern

MUNICH, GERMANY

MAR. 2017

- Implementation of a three-dimensional data-visualization based on JavaScript charting and python data processing.
- Annotation of glomeruli and epithelial cells in IHC images.

TARGOS MOLECULAR BIOLOGY GMBH

Intern

KASSEL, GERMANY

OCT. 2016

TARGOS MOLECULAR BIOLOGY GMBH

Administrative Assistant

KASSEL, GERMANY

JUN. 2015 - DEC. 2016

PROGRAMMING PROJECTS

- **ADVANCED MASTERS INTERNSHIP**

Shiny application for survival analysis based on deconvoluted bulk RNA sequencing data. (deconvsurvR)
(Ongoing and to be published)

- **PROTEIN PREDICTION I**
Detection of disordered post-translational modification with PyTorch
(grade bonus - exam 1.3)
- **PROTEIN PREDICTION II**
Prediction of transmembrane classes directly from protein sequence embedding using the class-attention mechanism.
<https://github.com/mainpyp/cls-protein-prediction>
(grade bonus - exam 1.3)
- **TABLE INSPECTOR**
Shiny application that allows a quick overview of the built-in data that comes with R.
https://github.com/mainpyp/builtin_inspectoR
- **MASTERS INTERNSHIP**
Determination of expression signatures in macrophages using machine learning feature importance.
(grade 1.7)
- **BACHELORS THESIS**
Communication-efficient approaches for Federated Deep Neural Networks. For the experiments I have used Tensorflow.
<https://github.com/mainpyp/ComEff>
(grade 1.0)
- **PROBLEM BASED LEARNING SEMINAR**
Implementation of a Python based web framework to receive and rank scientific papers with Flask.
<https://github.com/mainpyp/paper-miner>
(grade 1.0)
- **PROGRAMMING INTERNSHIP & GENOME-ORIENTED INTERNSHIP**
(grade 1.3 & 2.0)

EXTRACURRICULAR ACTIVITIES (OR OTHER EXPERIENCES & INTERESTS)

- Competitive basketball player at FC Bayern Basketball from 2013 - 2015. Played at national level in the Jugend-Basketball-Bundesliga (JBBL), won the German cup and ranked third in the Bundesliga in 2014.
- Coach at multiple youth basketball camps. (43 camp / Omari Knox Skill Camp)
- Volunteer as Basketball Instructor at DJK Würmtal from 2017 - 2021

LANGUAGES SKILLS

German (Native); **English** (Advanced);

COMPUTER SKILLS

LANGUAGES: **Python** (Advanced) • **R** (Intermediate) • **JavaScript - HTML - CSS** (Intermediate) • **Java** (Beginner)

FRAMEWORKS AND PACKAGES: **Pytorch (lightning)** • **TensorFlow - Keras** • **NumPy** • **Pandas** • **Flask** • **scikit-learn** • **Jupyter** • **Matplotlib - Seaborn** • **Plotly** • **pytest** • **Pillow**

ADDITIONAL SKILLS: **Office 365** • **iWork** • **Mac OS** • **Linux** • **JetBrain IDEs (DataSpell, PyCharm)** • **Conda** • **Git(Hub | Lab)** • **Bash**