

Martin Meinel

PHD CANDIDATE · MACHINE LEARNING

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Profile

Highly motivated machine learning researcher with a passion for translating machine learning research to healthcare, primarily focused on inflammatory skin diseases. Developed SAFARI a framework for sparse biomarker detection for rare diseases. At the moment leveraging genetics in combination with single-cell expression data to identify causal risk factors.

Education

PhD Candidate in Computational Biology

HELMHOLTZ CENTER MUNICH | TECHNICAL UNIVERSITY MUNICH | LMU MUNICH

- Supervised by Dr. Michael Menden & Dr. Natalie Garzorz-Stark.

Munich, Germany

Oct 2021 - Present

M. Sc. in Data Engineering & Analytics

TECHNICAL UNIVERSITY OF MUNICH

- GPA: 1.7
- Relevant courses: Machine Learning, Machine Learning for Crowd Modeling & Simulation, Information Retrieval, Applied Regression, Foundations of Data Analysis, Foundations of Data Engineering.

Munich, Germany

Oct 2018 - Nov 2020

B. Sc. in Applied Computer Science

UNIVERSITY OF BAYREUTH

- GPA: 2.1

Bayreuth, Germany

Oct 2014 - Oct 2017

Research Experience

SAFARI - Sparse biomarkers For rARe diseases.

PHD STUDENT

- Developed the model agnostic framework SAFARI to identify sparse & robust biomarkers for rare diseases.
- SAFARI works directly on the gene space, enabling clinical translation.
- SAFARI ensures high expression levels of genes, simplifying their detection in clinical applications.
- Applied to the essential clinical problem of distinguishing cutaneous T Cell lymphoma from eczema, I extracted three potential biomarkers out of 20,000 genes from bulk RNA-seq data, empowering early diagnosis and proper treatment.
- Leveraging the three discovered biomarkers, I achieve a sensitivity and specificity score of 85 % and 93 %, respectively, using a logistic regression model on an imbalanced data set.

Helmholtz Center Munich

Feb 2021 - Present

Identification of diagnostic markers enabling precision medicine for psoriasis

PHD STUDENT

- Developed a linear imputation solution for missing therapy response data of patients, enabling the analysis of complete patient datasets.
- Created an objective metric to assess therapy response, allowing for more accurate in-between patient comparisons and categorization of patients.
- Implemented statistical tests identifying best drug target per endotype.

Helmholtz Center Munich

Apr 2021 - Present

Leveraging single-cell expression data with genetics to discover risk factors in inflammatory skin diseases

PHD STUDENT

- Leveraging Mendelian Randomization to infer causal associations of cell-type specific gene expression levels and inflammatory skin diseases.
- Implemented pipeline to detect cis eQTLs of single-cell expression data from Peripheral Blood Mononuclear Cells.

Helmholtz Center Munich

October 2023- Present

Efficient implementation of deep convolutional Gaussian Processes

RESEARCHER

- Implemented deep convolutional Gaussian processes to mimic convolutional neural networks (CNNs) to obtain uncertainty estimations for predictions of the MNIST dataset with the goal of faster computation (see here).
- Conducted a comprehensive evaluation of various imputation algorithms to determine their ability to efficiently approximate the kernel matrix and deliver optimal performance on test data.
- Accomplished a substantial 40% reduction in training time by using only 20% of the original training data while obtaining a high classification accuracy of 86% on the MNIST dataset (98% accuracy using the entire training set).

TU Munich

Jan 2020 - Nov 2020

- Implemented variational autoencoder generating MNIST digits.
- Trained Word2Vec embeddings on a review dataset to showcase embeddings store language characteristics.
- Implemented spectral clustering to users on the social network Yelp. Clustered users based on food preference similarity. A comprehensive evaluation was conducted to minimize the ratio cut and the normalized cut of the graph regarding how this affects spectral clustering.
- Implemented a graph convolutional neural network performing semi-supervised node classification on the CORA ML citation network. Analyzed how oversmoothing affects the network. Implemented approximate personalized propagation of neural predictions to overcome this problem.

Publications

- Garger, D., **Meinel, M.**, Dietl, T., Hillig, C., Garzorz-Stark, N., Eyerich, K., de Angelis, M., Eyerich, S., Menden M.P. (2023). The impact of the cardiovascular component and somatic mutations on ageing. *Aging Cell*. <https://doi.org/10.1111/ace.13957>
- Fischer, F., Doll, A., Uereyener, D., Roenneberg, S., Hillig, C., Weber, L., Hackert, V., **Meinel, M.**, Farnoud, A., Seiringer, P., et al. (2023). Gene expression based molecular test as diagnostic aid for the differential diagnosis of psoriasis and eczema in formalin fixed and paraffin embedded tissue, microbiopsies and tape strips. *The Journal of Investigative Dermatology*, S0022-202X(23)00156-2. <https://doi.org/10.1016/j.jid.2023.02.015>
- Farnoud, A., Ohnmacht, A. J., **Meinel, M.**, & Menden, M. P. (2022). Can artificial intelligence accelerate preclinical drug discovery and precision medicine?. *Expert opinion on drug discovery*, 17(7), 661–665. <https://www.tandfonline.com/doi/epdf/10.1080/17460441.2022.2090540>

Working experience

Opitz Consulting GmbH

Munich, Germany

SOFTWARE ENGINEER WORKING STUDENT

Oct 2019 - Oct 2020

- Built a docker skeleton for Apache Zeppelin, enabling users to start a docker container running Apache Zeppelin on openShift with one click.
- Created a web application for Allianz IDS GmbH, allowing employees to track, assign, and rate investment orders within the department.

REHAU Industries SE & Co. KG

Bayreuth, Germany

SOFTWARE ENGINEER WORKING STUDENT

Feb 2018 - Mar 2019

- Developed ALEXA skills for the 2018 FRONTALE window exhibition, demonstrating several features of the smart window.
- Implemented a web application for Medi Bayreuth to assess players' fatigue after training sessions and their performance during regular fitness checks to visualize and track their development and prevent injuries.

REHAU Industries SE & Co. KG

Rehau, Germany

SOFTWARE ENGINEER INTERN

Nov 2017 - Jan 2018

- Developed a plugin for the open-source smart home system openHAB, enabling users to connect their automated sensor-based garden watering system with other IoT devices.

Skills

Programming	Python, Java, R, SQL, shell
Libraries / Frameworks	NumPy, Pandas, PyTorch, scikit-learn, Spring, scanpy, DESeq2, edgeR
Tools / Platforms	AWS, Docker, Git, LaTeX, slurm
Languages	German, English

Honors & Awards

2023	Poster at ML4H , Accepted Findings Paper at “Machine Learning for Health” workshop	New Orleans, US
2023	Inventor , Registered patent “Differential diagnosis of mycosis fungoides, method and system” for discovered biomarkers to distinguish Eczema from Cutaneous T Cell lymphoma using SAFARI	Munich, Germany
2023	Speaker , Selected Speaker about “Biocomputational analysis of big data” at EADV-ESDR Summer Research Workshop: Big Data and Translation in Dermatology.	Freiburg, Germany
2023	Participant , Selected as participant for summer school OxML - MLxHealth.	Oxford, UK
2021	PhD , Selected as a member of Munich School for Data Science (MuDS)	Munich, Germany

Extracurricular Activity

TUM MINGA Mentoring Program

Munich, Germany

MENTOR

Oct 2019 - Oct 2020

- Assisted incoming master students from abroad with navigating TU Munich and life in Germany by offering guidance on university procedures, exam preparations, and insights on local resources and cultural differences.

Lern-Fair

Munich, Germany

MATH TEACHER

Feb 2021 - Jul 2021

- Delivered personalized online math lessons to 9th-grade students during the COVID-19 home-schooling period, promoting understanding and growth in mathematical concepts.
- Developed and curated exercise sheets to reinforce knowledge, deepen understanding, and facilitate independent practice.