Manuel Gloeckler

PhD student at the University of Tübingen

Date of Birth: May 29, 1998

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Google Scholar: M. Gloeckler

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Interests -

Bayesian inference

Simulation-based inference

Generative models

probabilistic and differentiable programming

Skills -

Programming:

Python

Bash scripting

Java

Cuda/C/C++

Tools:

JAX

PyTorch

Hydra/SLURM

TensorFlow/Keras

Languages

German (Mother Tongue)

English (Proficient)

French (B1, 2016)

Working Experience

Apr 2022 – PhD student

Working as a researcher at the University of Tübingen in the Cluster of Excellence - Machine Learning in Science, supervised by Prof. Dr. Jakob Macke. Member of the International Max Planck Research School for Intelligent Systems (IMPRS-IS). Research on simulation-based (or likelihood-free Bayesian) inference methods, focusing on robustness and efficiency.

Sep 2020 - Research Assistant

University of Tübingen, DE

OpenAI, Inc.

University of Tübingen, DE

Worked as a research assistant in the Computational Systems Biology Group, supervised by Dr. Reihaneh Mostolizadeh. The research focused on the computational analysis of microbiota interactions in the human nasal microbiome through genome-scale modeling.

Minijob

Feb 2022

ongoing

Nov 2024 – Independent Contractor

Jan 2025 Expert consultant for research related parts in PaperBench: Evaluating AI's Ability to

Replicate AI Research

Feb 2019 – Student Assistant University of Tübingen, DE

Oct 2019 Teaching assistant in the Theory of Machine Learning Group of Prof. Dr. Ulrike von

Luxburg.

Education

Mar 2022 – M.Sc. in Bioinformatics

University of Tübingen, DE

Oct 2019 Focused on machine learning methods and theory and their applications in life sciences. Graduation with distinction [certificate].

Thesis: Variational Methods for Simulation-Based Inference [pdf].

Supervisors: Dr. Jakob Macke, Dr. Manfred Classen

Average grade: 1.15 (3.9 GPA equivalent)

| Detailed List of Grades

University of Tübingen, DE

Sep 2019 - B.Sc. in Bioinformatics

Oct 2016 Graduation with distinction [certificate].

Thesis: The Landscapes of CD8+ T Cell Immunogenicity from a Self-

Tolerance-Based Perspective in Sequence Space [pdf]. **Supervisors**: Dr. Leon Kuchenbecker, Dr. Oliver Kohlbacher

Average grade: 1.31 (3.7 GPA equivalent)

| Detailed List of Grades

Summer'22.24

Other Training

Jul 2022 Cambridge ELLIS Machine Learning Summer School Cambridge, UK

Participation in the summer school and poster presentation [certifi-

cate].

Jul 2021 Machine Learning Summer School Taipei, Online (Cov19)

Participation in the summer school [certificate].

Teaching Experience

Teaching assistant

University of ML4202: Probabilistic Machine Learning

Tübingen Graduate course.

ML 4102: Data Literacy Winter'22

Graduate course about basic data science methods.

INFO4412: Algorithms and Complexity Winter'19
Undergraduate course about algorithms and complexity analysis.

Short Bio

Manuel Gloeckler is a PhD student in the lab of Prof. Jakob H. Macke at the University of Tübingen, with a background in Bioinformatics. His research focuses on (conditional) generative modeling for scientific data, particularly in the context of inverse problems. Since beginning his PhD in 2022, he has contributed to major international machine learning conferences. He enjoys applying his programming expertise to develop novel methods for solving scientific challenges.

Metrics



Profiles





Publications

Selected

- 2025 Compositional simulation-based inference for time series [arxiv]

 Manuel Gloeckler, Shoji Toyota, Kenji Fukumizu, Jakob H. Macke

 Accepted as poster for ICLR 2025
- 2024 All-in-one simulation-based inference [arxiv]

 Manuel Gloeckler, Michael Deistler, Christian Weilbach,
 Frank Wood, Jakob H. Macke
 Accepted as a oral for ICML 2024 (top 5%)
- 2023 Adversarial robustness of amortized Bayesian inference [arxiv].

 Manuel Gloeckler, Michael Deistler, Jakob H. Macke.

 Accepted as a poster for ICML 2023
- 2022 Variational methods for simulation-based inference [arxiv].

 Manuel Gloeckler, Michael Deistler, Jakob H. Macke.

 Accepted as a spotlight for ICLR 2022 (top 6 %).

Published

- 2025 **sbi reloaded: a toolkit for simulation-based inference workflows**Boelts et. al (role as core-contributor); JOSS
- A practical guide to statistical distances for evaluating generative models in science [arxiv]

 Bischoff et. al (alphabetically sorted, role as a co-organizer); TMLR
- 2024 Inferring stochastic low-rank recurrent neural networks from neural data [arxiv]

Matthijs Pals et. al; NeurIPS 2024

- 2023 Hierarchical modelling of microbial communities Manuel Gloeckler, A Dreager, R Mostolizadeh
- 2022 NCMW: a python package to analyze metabolic interactions in the nasal microbiome [frontiers].

Manuel Gloeckler, Andreas Draeger, Reihaneh Mostolizadeh

2022 **Towards the human nasal microbiome: Simulating D. pigrum**

2022 Towards the human nasal microbiome: Simulating D. pigrun and S. aureus [frontiers].
Reihaneh Mostolizadeh, Manuel Gloeckler, Andreas Draeger

Pre-prints

Differentiable simulation enables large-scale training of detailed biophysical models of neural dynamics [bioarxiv]
 Michael Deistler, Kyra L Kadhim, Matthijs Pals, Jonas Beck, Ziwei Huang, Manuel Gloeckler, Janne K Lappalainen, Cornelius Schröder, Philipp Berens, Pedro J Gonçalves, Jakob H Macke

Review Duties

Journal IOP Trusted reviewer [certificate]
Conferences ICLR 2024, ICLR 2025, ICML 2025

Open source contributions

sbi Core maintainer, Contributor 684 stars

References

Ref. 1	Prof. Dr. Jakob Macke	University of Tübingen, DE
	jakob.macke@uni-tuebingen.de	
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Ref. 2 **Dr. Reihaneh Mostolizadeh** University of Tübingen, DE reihaneh.mostolizadeh@uni-tuebingen.de