

Aiswarya Prasad

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PhD candidate in Quantitative Biology with expertise in microbiome research, bioinformatics, and hands-on lab experience. Strong focus on microbiome ecology, metagenomics, and diverse next-generation sequencing (NGS) approaches. Proven track record in leading multi-disciplinary research projects and international collaboration. Seeking opportunities to leverage my skills in a challenging innovation-focused role.

Education

PhD in Quantitative Biology

University of Lausanne, Switzerland | AUG 2020 - AUG 2025 (planned)

- Awarded the Faculty of Biology and Medicine (FBM) PhD Fellowship and Mathilde Agassiz Scholarship (awarded to top-placed FBM fellow).
- Presented research as posters at 3 prominent international conferences; invited as a speaker at 3, awarded best selected talk. Co-authored 4 publications and performed the peer-review of 2 papers.
- 4 years of Teaching experience as TA for the Master-level “bioinformatics through research” course; developed an R markdown template to improve and standardize course material as interactive HTML documents. Supervised semester-long student projects presented at the course.
- Engaged as a student representative on the department faculty hiring committee and the organizing committee for the annual department scientific retreat.

B.S. and M.S. in Biology

Indian Institute of Science, Bengaluru | AUG 2015 - JUL 2020

- KVPY National Science Fellowship (All-India rank: 335)

Research Experience

PhD candidate

University of Lausanne | AUG 2020 - PRESENT

- Led honeybee gut microbiome evolution research using next-generation sequencing and custom-made Snakemake pipelines (Eg., <https://github.com/Aiswarya-prasad/honeybee-cross-species-metagenomics/>).
- Designed and conducted experiments on microbiota-free bees with synthetic bacterial communities to study processes influencing community assembly using PacBio long-read sequencing.
- Collaborated with multi-disciplinary teams in Malaysia and Kenya and initiated a new collaboration in India, expanding the study's scope with cross-country data and mitigating delays due to pandemic restrictions in Malaysia by over one year.

Master thesis research

Indian Institute of Science, Bengaluru | MAY 2019 - MAY 2020

- Established Nanopore sequencing system in a low-resource setting at 10% of the cost of other commonly available systems.
- Developed an end-to-end pipeline from sample collection to bioinformatic analysis for metagenomics; tested various approaches for sample preparation and metaproteomic analysis of fecal samples using LC-MS/MS.
- Analyzed the human gut microbiome in chronic pancreatitis patients, studying disease-related changes in collaboration with doctors.

iGEM undergraduate research

Indian Institute of Science, Bengaluru | MAY 2016 - OCT 2016

- Co-founded the first team from the institute and secured a bronze medal at iGEM in Boston, a competition for innovative biotech solutions using genetically engineered machines.
- Contributed to the designing and cloning of a quorum sensing-based auto-inducible system for protein overexpression in bacteria.
- Engaged in teamwork to secure funding for research and travel via three grants.

Publications

Mazel, F., [Prasad, A.](#), Engel, P. (2024) Host specificity of gut microbiota associated with social bees: patterns and processes. *Microbiology and Molecular Biology Reviews* (In Review)

[Prasad, A.](#), Pallujam, AD., Siddaganga, R., Suryanarayanan, A., Mazel, F., Brockmann, A., Yek SH., Engel, P. (2024) Evolution of the honeybee gut microbiome. *Nature communications*. (In Review)

Baud, G. L., [Prasad, A.](#), Ellegaard, K. M., & Engel, P. (2023). Turnover of strain-level diversity modulates functional traits in the honeybee gut microbiome between nurses and foragers. *Genome Biology*, 24(1), 283.

Sarton-Lohéac, G., Nunes da Silva, C. G., Mazel, F., Baud, G., de Bakker, V., Das, S., El Chazli, Y., Ellegaard, K., Carcia-Carcera, M., Glover, N., Liberti, J., Nacif Marçal, L., [Prasad, A.](#), Sommerville, V., SAGE class 2019-2020 and 2020-2021, Bonilla-Rosso, G., & Engel, P. (2023). Deep divergence and genomic diversification of gut symbionts of neotropical stingless bees. *Mbio*, 14(2), e03538-22.

Extracurricular Engagement

Director, Strategy and Expert Sourcing
Nucleate Switzerland | OCT 2024 (Onwards)

The Consulting Society – Consultant and Learning Unit Manager
École Polytechnique Fédérale de Lausanne (EPFL) | MAY 2024 (Onwards)

Workshop on Decolonizing Global North-South Research Collaboration
Centre for Development and Environment, University of Bern (Online) | AUG 2022

Moderator, Health & Medical Data Privacy Committee
iGEM India BIOSUMMIT (Remote) | JUL 2020

iGEM Delegate, United Nations Convention on Biological Diversity
Sharm El-Sheikh, Egypt | NOV 2018