

FLAVIA HODEL

Doctoral student in Human Genomics

✉ flavia.hodel@epfl.ch ☎ +41 (0)77 405 03 37 ⌂ Rue du Midi 10, 1003 Lausanne, Switzerland
🌐 https://flaviahodel.github.io ⌂ linkedin.com/in/flavia-hodel ⌂ @FlaviaHodel



Experiences

Visiting PhD student

Wellcome Trust Centre for Human Genetics, University of Oxford

📅 2019

I went to the WTCHG for a collaborative project to identify key genomic determinants of humoral immune responses against multiple human pathogens. This experience allowed the exchange of ideas and data, and provided me with new tools to bring back to my colleagues.

PhD student in Human genomics & Computational biology

Swiss Federal Institute of Technology (EPFL), Lausanne, Switzerland

📅 2018 - ongoing

Supervisor: Prof. Jacques Fellay

I use genome-wide genotyping data, bioinformatics tools and statistical methods to explore the interplay between human genetic variation, persistent infections and chronic inflammation, and understand their independent and combined effect on coronary artery disease. It is a multidisciplinary and collaborative project bringing together our in-house data with other biobanks.

Graduate teaching assistant

Swiss Federal Institute of Technology (EPFL), Lausanne, Switzerland

📅 2018 - ongoing

Subjects taught: Probabilities and statistics, Genetics and genomics, and Introduction to life sciences for School of Computer and Communication Sciences (IC).

Summer trainee

Institute of Human Virology, Nigeria

📅 2016

Worked with a team of scientists to address the HIV/AIDS crises in Nigeria through the development of infrastructure for treatment, care, prevention, and support.

Education

Ph.D. in Computational and Quantitative Biology (EDCB)

Swiss Federal Institute of Technology (EPFL), Lausanne, Switzerland

📅 2017 – ongoing

M.Sc. in Bioinformatics

University of Lausanne, Switzerland

📅 2016 – 2017

B.Sc. in Biology

University of Lausanne, Switzerland

📅 2012 – 2015

Volunteering

Nuit de la lecture

Caretaker of book-crossing boxes in Lausanne

📅 2022 - ongoing

EPFL – ETHZ Summer School

"Shaping the Future of (bio)-Medicine"

Co-organiser

📅 2019

Planned the event and location, made budget for the event, contacted sponsors and speakers, co-organised the health hackathon in collaboration with the MIT

Basic computer course for seniors

Instructor at ACLI Svizzera

📅 2016

Languages

French	Native
Swiss-German	Native
English	Full professional proficiency
Italian	Level C1
Spanish	Level C1

Strengths

Data analysis	Bioinformatics
Biostatistics	Data visualization
Genomics	Infectious diseases
GWAS	Linear/logistic regression
Univariate/multivariate analysis	
Survival analysis	Biobanks
Biomarkers	R
LaTeX	BASH
	Python
GitHub	

Team spirit	Leadership	Adaptability
Eye for detail	Emotional intelligence	
Conferences and events organisation		
Research presentations		

Publications

Hodel, F., Naret, O., Bonnet, C., Brenner, N., Bender, N., Waterboer, T., ... & Fellay, J. (2022). The combined impact of persistent infections and human genetic variation on C-reactive protein levels. *medRxiv*, doi 10.1101/2022.01.07.22268880.

Hodel, F., Chong, A. Y., Scepanovic, P., Xu, Z. M., Naret, O., Thorball, C. W., ... & Fellay, J. (2021). Human genomics of the humoral immune response against polyomaviruses. *Virus Evolution*, veab058.

Lawless, D., Allen, H. L., Thaventhiran, J., Hodel, F., Anwar, R., Fellay, J., ... & NIHR BioResource–Rare Diseases Consortium. (2019). Predicting the Occurrence of Variants in RAG1 and RAG2. *Journal of clinical immunology*, 39(7), 688-701.

Scepanovic, P., Hodel, F., Mondot, S., Partula, V., Byrd, A., Hammer, C., ... & Lantz, O. (2019). A comprehensive assessment of demographic, environmental, and host genetic associations with gut microbiome diversity in healthy individuals. *Microbiome*, 7(1), 130.

Scepanovic, P., Alanio, C., Hammer, C., Hodel, F., Bergstedt, J., Patin, E., ... & Quintana-Murci, L. (2018). Human genetic variants and age are the strongest predictors of humoral immune responses to common pathogens and vaccines. *Genome medicine*, 10(1), 59.

Hodel, F., Patxot, M., Snäkä, T., & Ciuffi, A. (2016). "HIV-1 latent reservoir: size matters". *Future Virology*, 11(12), 785-794.

Conferences

European Society of Human Genetics (ESHG)
Vienna, Austria - Berlin, Germany - Gothenburg, Sweden
2019, 2020, 2021
Early career poster award candidate in 2021

American Society of Human Genetics (ASHG)
San Diego, CA, USA
2018

Professional development

Machine Learning
Online course on Coursera
2020 - Ongoing Given by Stanford University, CA, USA

- Supervised learning (parametric/non-parametric algorithms, support vector machines, kernels, neural networks)
- Unsupervised learning (clustering, dimensionality reduction, recommender systems, deep learning)
- Best practices in machine learning (bias/variance theory; innovation process in machine learning and AI)

Creative thinking: techniques and tools for success

Online course on Coursera
Certificate earned in Oct. 2020 Given by Imperial College London, UK

- Creativity tools and Thinking styles
- Morphological analysis
- TRIZ: The theory of inventive problem solving
- The SCAMPER method

Summary

As a data analyst, I have expertise in computational biology. I analyze large-scale biological data and find solutions to better understand human diseases using genetics and environmental factors. In my PhD, I particularly enjoy the collaboration with my interdisciplinary colleagues and other institutes, the communication of data to enable future decisions, and the great freedom to create. I am enthusiastic and eager to learn new software and tools.