

Skills

Programming:

Others: R, Java, C, C++, SQL,

Machine Learning, Deep Learning Neural Networks, Computational

Biology, Artificial Intelligence,

Genomics,

Git, Unit Testing, Gurobi, UML,

Transversal:

Languages

Native

Fluent

TOEFL iBT (04.2020): 99/120

Personnality

POO, PowerBI, Cluster Usage

Discovery,

Modelling, Problem Solving

Windows, GNU/Linux

Project Management

Some words to describe me:

Involved, serious, versatile, curious, with

a strong critical mind. Perfectionist,

Highly interested in sciences since

childhood, nurtured by self motivation

via documentaries, lectures, scientific articles, online courses, newsgroups,

popular science movies and online

Hobbies: Photography, Travels, Sport

(Cycling, Hiking, Skiing, Climbing,

Badminton), DJing, Cooking, Reading,

pedagogue, rigorous and perseverant.

Microsoft Office

French:

English:

PHP,

HTML/CSS,

Precision

Systems

Languages: Main: Julia, Python

Mathlab,

Drug

Medicine,

Tools.

Mathematica

Fields of Interest:

Simon CROUZET

PHD CANDIDATE IN COMPUTATIONAL BIOLOGY





Qualifications

Ecole Polytechnique Fédérale de Lausanne (EPFL) – Lausanne (Switzerland) Ph.D. in Computational and Quantitative Biology

2021 - 2022 Ecole Normale Supérieure (ENS-PSL, part of Paris Science et Lettres) – Paris (France) M.Sc. in Systems biology, Genomics and Bioinformatics - with First Class Honours

2020 - 2021 Sorbonne University - Paris (France)

M.Sc. in Bioinformatics and Modelling - with High Honours

Linnaeaus University - Växjö (Sweden) 2019

Semester Abroad during B.Sc. in Computer Science

Semester 5 of my Bachelor of Computer Science through the mobility program ISEP Study Abroad

2017 - 2020 Aix-Marseille University - Marseille Luminy (France)

B.Sc. in Computer Sciences - with First Class Honours

2016 - 2017 Interruption of studies due to health issues

Grenoble Alpes University - Saint-Martin-d'Hères (France) 2013 - 2016



Work experiences

from 2023 (Ongoing)

Doctoral Assistant at EPFL School of Life Sciences

Ecole Polytechnique Fédérale de Lausanne (Switzerland)

Development of AI approaches and computational methods in structural biology, with diverse applications in protein and drug design.

Summer and Autumn 2022

Computational Biologist Intern (IP2TIS Fellow) at Roche pRED-PMDA

Roche pRED Pharma Research and Early Development (Switzerland)

Analysis of data generated by phenotypic screens with a focus on causal inference, network-based machine learning and interpretable graph neural network, to identify putative high-potential targets for novel drug discovery programmes.

Winter and

Research Intern at ETH Zürich D-BSSE

Spring 2022 ETH Zurich (Switzerland)

Design, implementation, and evaluation of computational methods for predicting HIV evolutionary pathways and for designing optimal drug combinations.

Summer 2021

Research Assistant at Wellcome Sanger Institute

University of Cambridge (England)

Structural variation of genomes and pangenomics exploration of Anopheles Funestus (mosquito, major vector

Spring and Summer 2020

Research Intern at the Institute of Biodiversity, Animal Health and Comparative Medicine

University of Glasgow (Scotland)

Development and improvement of a viral host prediction model in Julia through genomic feature, phylogenetic neighbourhoods (using BLAST search) and protein characteristics, following the first version available on https://bioinformatics.cvr.ac.uk/software/viral-host-predictor/.

Summer 2018

Research Intern at Laboratoire d'Informatique & Systèmes

Aix-Marseille University - Campus of Marseille Luminy (France)

Development in C of an automaton network solver using data-flow algorithms to model genetic regulatory networks as the control of floral morphogenesis in *Arabidopsis Thaliana*.

ORCID: 0000-0002-5012-4621



Publications & Projects

Publications & Conferences:

- Yasser Mohseni Behbahani, Simon Crouzet, Élodie Laine, Alessandra Carbone, Deep Local Analysis evaluates protein docking conformations with Locally oriented Cubes (Bioinformatics 2022, doi: 10.1093/bioinformatics/btac551)
- Simon Crouzet, Matthew Arnold, Nardus Mollentze, Richard J. Orton, Daniel G. Streicker, Simon A. Babayan, Highperformance and modular identification of virus hosts of origin utilising machine learning of genomic signatures in Julia (work in progress, talk presented on eSCAMPS 2021)

Internet & Video Games...

searches.

Contact

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Phone: +33 6 88 71 65 38

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Volunteering

- International Ecovolunteering during the month of June 2019 with the NGO Archelon on a sea turtle conservation project in Kyparissia Bay (Greece), and during the month of November 2022 with the NGO WFFT on a elephant refuge project in Phetchaburi (Thailand)
- Volunteering in several student associations, including Asso'M (event-driven student association) as well as FAMI (representation student association), TACT (large-scale sport event, as a Financial Officer), BDE Luminy (student association, as a Treasurer) or Multicultural Växjö (charitable student association).
- Representative student at CVEC (Contribution à la Vie Étudiante du Campus) and at Department "Informatique et Interaction" council of Aix-Marseille University.
- Ambassador for COP2 Étudiante, a national summit to negotiate a global agreement on ecological and social transition within French higher education.