Saul Pierotti

Computational genetic ist — PhD fellow at EMBL-EBI

Born in 1995 — Italian citizen

saulpierotti@gmail.comEmail: Website: GitHub: ORCiD:

Research experience

European Bioinformatics Institute (EMBL-EBI)

University of Cambridge

Predoctoral fellow; Supervisor: Prof. Ewan Birney

QTL mapping and exploration of GxE, GxG, and indirect (social) genetic

effects in medaka fish.

Science for Life Laboratory (SciLifeLab)

Master thesis; Supervisor: Prof. Arne Eloffson

Prediction of protein contact maps from deep mutational scanning data

using machine learning

Max Planck Institute for Evolutionary Biology

Intern; Supervisors: Prof. Arne Traulsen, Dr. Jenna Gallie

Experimental evolution of Escherichia coli and Pseudomonas fluorescens in

long-term batch cultures

University of Perugia

Bachelor thesis; Supervisor: Dr. Manlio di Cristina

Characterization of vacuolar proteins in Toxoplasma gondii

EDUCATION

European Bioinformatics Institute (EMBL-EBI)

University of Cambridge

PhD candidate at the University of Cambridge as part of the the EMBL

International PhD Program (EIPP) — Applicant acceptance rate 2.7%

University of Bologna International Master Degree in Bioinformatics (MSc)

Final mark: 110/110 cum laude

University of Perugia

Bachelor's degree in Biotechnology (BSc)

Final mark: 110/110 cum laude and honorable mention

Published works

S. Pierotti, B. Welz, M. Osuna-López, T. Fitzgerald, J. Wittbrodt, E. Birney, Genotype imputation in F2 crosses of inbred lines, Bioinformatics Advances, 2024, 10.1093/bioadv/vbae107

Planned publications

- Main PhD work on the detection of 16 QTLs for heart rate in medaka, their CRISPR validation, and the characterization of GxE and GxG effects — collaboration with Bettina Welz (Wittbrodt group, Heidelberg University, Germany).
- Pilot study on the effect of direct and indirect (social) genetic effects on medaka behaviour collaboration with Ian Brettel (Birney group, EMBL-EBI) and the group of Felix Loosli (KIT, Karlsruhe, Germany).
- Application note on a mixed linear model genetic association pipeline.

saulpierotti.github.io saulpierotti $0000\hbox{-}0002\hbox{-}59\bar{4}2\hbox{-}0793$ Linkedin: saulpierotti

Cambridge, UK

Jan 2021 — Jun 2021

Stockholm, Sweden

Jan 2021 — Jun 2021

Plön, Germany Jun 2019 — Sep 2019

Perugia, Italy

 ${\rm Oct}\ 2018 - {\rm Oct}\ 2019$

Cambridge, UK Expected Oct 2025

> Bologna, Italy Jul 2021

Perugia, Italy Oct 2019

- Application note on a genetic variant prioritization pipeline.
- GWAS for behavioural phenotypes in medaka collaboration with the group of Felix Loosli (KIT, Karlsruhe, Germany).

Conference presentations

The Biology of Genomes, Cold Spring Harbor Laboratory (Cold Spring Harbor, USA), May 7th to 11th, 2024 — poster

NRBP Medaka International Workshop, National Institute for Basic Biology (Okazaki, Japan), April 26th to 27th, 2024 — talk

Training delivered

EMBL-EBI course "Exploring Gene and Environmental Exposure Interactions to Understand Human Health and Disease", February 19th to 23rd, 2024

Nextflow practical for the EBI predoc course (online), November 21st to 25th, 2022

LANGUAGES

English: proficient, CEFR level C2. IELTS score 8.0. CAE with grade A awarded by Cambridge ESOL.

Italian: native speaker. Others: Currently learning Turkish, German and Dutch (elementary).

REFERENCES

Ewan Birney, current PhD supervisor at EMBL-EBI (Cambridge, UK): birney@ebi.ac.uk Arne Elofsson, master thesis supervisor at SciLifeLab (Stockholm, Sweden): arne@bioinfo.se