

Harry Petch-Smith

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Msc Bioinformatics

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EDUCATION

- MSc Bioinformatics - Distinction**, *University of Birmingham* 09 2022
Modules: *Essentials of Mathematics and Statistics, Genomics and Next Generation Sequencing, Data Analytics and Statistical Machine Learning, Metabolomics and Advanced omics Technologies, Computational Biology for Complex Systems*
- BSc Biomedical Science With a Year In Industry - 2:1**, *The University of Sheffield* 07 2021
- A levels**, *South Hunsley School and Sixth Form College* 07 2017
Psychology (A*), Chemistry (B), Biology (B)

EXPERIENCE

- Volunteer Instructor/Introduction to Python Programming** Oct 2021 — Dec 2021
Code First Girls Remote
- Taught beginners python course with the aim of introducing more women to the field of tech.
 - Collaborated in a group of three instructors to deliver weekly lessons.
 - Managed lesson planning alongside masters studies.
 - Led interactive sessions in an engaging manner, with a mix of activities and lecture format.
 - Clear communication of programming principles to women from many backgrounds.
- Student Biomedical Scientist** 06 2019 — 07 2020
Hammersmith Medicines Research Wembley, London
- Year in Industry Placement at UK leading clinical trials company.
 - Responsibilities: Sample processing, Monitoring patient data, Validation of new assays, Quality control of documentation, Stock/supplies management
 - Gained experience working in a fast paced professional environment that valued both team work and independence

PROJECTS

- A Bayesian non Parametric Approach to Time Series Modelling**
Machine Learning
- **Research Question:** Can multiple output Gaussian processes accurately model and predict time series data.
 - **Objective:** Test multiple output GPs on two datasets; Simulated dependant time series data, and blood glucose monitoring data.
 - Implemented single output gaussian process using **R**.
 - **Simulated** time series datasets in **R and Python**.
 - Developed a Multiple Output Gaussian Process from scratch in **R**.
 - Results reported in **nature article format** using **Latex**
- Identification of Cyp21a2-dependent, glucocorticoid regulated target genes in zebrafish larvae**
Data Science/Bioinformatics
- **Research question:** What are the effects of glucocorticoid deficiency on gene expression in Zebrafish larvae?
 - **Objective:** Given a **large dataset** of zebrafish gene expression counts; **combine and analyse** the data to identify differentially expressed genes.
 - constructed a Bioinformatics pipeline to conduct **data preprocessing** and **analysis and visualisation** in **R**.
 - Used **R** to combine multiple datasets for easy comparison
 - Performed **quality filtering and normalisation**
 - Implemented **principle component analysis, MA plotting, Volcano plots** and **downstream gene ontology analysis**
 - Maintained a lab book recording each step generated as an R markdown file.
 - Work was awarded a **first class grade**

SKILLS

Languages Bash, Python, R, HTML/CSS, Javascript
Tools Linux command line, Rstudio, Jupyter Notebook, Spyder, LaTeX, Markdown, Visual Studio Code.