

# Gobikrishnan Subramaniam

Flat 1, 18 Rosemount Gardens, Belfast, BT15 5AG

Phone: 07436994977 | email: [g.subramaniam@qub.ac.uk](mailto:g.subramaniam@qub.ac.uk) | [LinkedIn](#) | [GitHub](#) | [ORCID](#) | [Website](#)

Aspiring to pursue a career in cancer biology to leverage system biology and multi-omics approaches in advancing precision oncology, with a focus on developing novel biomarkers and therapeutic targets for better diagnosis and treatment.

## Education

**M.Sc., Molecular Biology and Biotechnology, Queen's University Belfast, UK** **2022 - 2023**

- Achieved 2:1 degree with commendation
- **Relevant modules:** Advanced Molecular biology, Protein structure and function, Biotechnology, Foundation for research, and Bioinformatics and System Biology
- **Dissertation:** *Network Biology of Renal Cell Carcinoma Drug Resistance and Response*

**B.Sc., Biotechnology, PSG College of Arts & Science, Coimbatore-14, India** **2019 - 2022**

- CGPA: 7.62/10.00 (Equivalent to a UK 1<sup>st</sup>)
- **Relevant modules:** Biochemistry, Cell biology, Diagnostic biotechnology, Microbiology, Genetics, Molecular biology, Immunology, rDNA technology, Genomics and proteomics, Bioinformatics, animal biotechnology
- Mastered various molecular biology techniques through extensive laboratory practice
- Served as the voice of my peers in my capacity as class representative during the 2020-2021 academic year

## Research Experience

**Visiting scholar, Overton lab, Queen's University Belfast, Belfast, UK** **Dec 2023 – Present**

- Provided technical assistance to the group including managing user accounts, system updates and handling HPC queries
- Dockerized [NetNC](#) and updated its codebase, reducing deployment time by 70% and enabling scalable analysis of over 1000 samples
- Maintained and managed [TMA-Navigator](#), a web application for analysing tissue microarray data, ensuring maximum uptime and coordinated the rewriting of the website

**Research intern, Overton lab, Queen's University Belfast, Belfast, UK** **May 2023 – Sep 2023**

- Analysed microarray data from two different studies (SuMR clinical trial & SCOTRRCC study), comprising 192 samples from metastatic renal cell carcinoma (mRCC) patients administered Sunitinib
- Constructed a gene co-expression network using [NetNC](#) based on the ranked gene lists from DEA and discovered four putative modules involved in Sunitinib resistance in mRCC, interpreted in the context of current literature
- Presented and defended my findings to the research group and at a monthly journal club

**BioHackathon Europe 2024 (Perturb-bench)** **Nov 2024**

- Integrated the [GenKI](#) toolkit into a Nextflow pipeline and benchmarked it against other models with the help of teammates
- Implemented GPU acceleration for *GenKI* toolkit, enabling analysis of a 3000-cell scRNA-seq dataset in under 1 hour
- Presenting a tutorial on [scGen](#) at the ISMB/ECCB 2025 Tutorial track, having prepared all necessary materials

## **The Sixth Annual Structural Variant and Pangenomics Hackathon at BCM**

**Aug 2024**

- Developed the [MoVana](#) pipeline using Cromwell, processing 71,000 SVs from the ICGC dataset to predict functional outcomes of various mosaic variants based on allele frequency
- Act as tech lead to the development of the pipeline and assisted with maintaining the GitHub repository
- Presented project outcomes to other teams, culminating in a collaborative research publication in F1000Research highlighting potential changes for the following year

## **Collaborative Bioinformatics Hackathon**

**Mar 2025**

- Developed a Nextflow pipeline for GLIMPSE2, a tool for low-coverage whole-genome sequencing imputation for scalable and effective deployment
- Awarded “[the most collaborative team](#)” in the hackathon, highlighting our effective teamwork and communication and published the results in BioHackrXiv

## **Teaching Experience**

### **Teaching Assistant, Queen’s University Belfast**

**Jan 2025 – Apr 2025**

- Taught systems medicine module (SCM8152) for 20 students, covering network biology and patient stratification for multi-omics analysis

## **Publication**

Sabata, S., Kubica, J., Gupta, R., Ericson, L. W., Atanda, H. C., Subramaniam, G., ... Busby, B. (2025, June 4). Addressing Background Genomic and Environmental Effects on Health through Accelerated Computing and Machine Learning: Results from the 2025 Hackathon at Carnegie Mellon University. [https://doi.org/10.37044/osf.io/3a8cn\\_v1](https://doi.org/10.37044/osf.io/3a8cn_v1)

## **Invited talks & Conferences**

- July 20-24 2025 ISMB/ECCB 2025
- June 5-6 2025 Brunel Bioinformatics Symposium 2025
  - Invited for a
- 19 June 2024 BioFAIR Roadshow Queen's University Belfast
- 26 Feb 2024 - 1 Mar 2024 Building Transparent ML/AI Solutions to Advance Biological Research Codeathon

## **Funding & awards**

QUB-MDBS Travel Scholarship

May 2025

## **Volunteering and Extracurricular Experience**

### **Volunteer, Cancer Research UK, Botanic Avenue, Belfast**

**Apr 2023 – Sep 2024**

- Raised £2500+ through fundraising initiatives and managed inventory for a shop serving 200+ weekly customers

Member of Queen’s homework club programme, mentored 12 + students from disadvantaged backgrounds in STEM subjects 2022 – 2023

## **Skills**

**Laboratory skills:** DNA/RNA extraction, agarose gel electrophoresis, PCR, UV spectrophotometry, Western blot, ELISA, and mammalian/microbial cell culture

**Computing skills:** R, Python, Git, Docker, and Nextflow

**Other certifications:** Laboratory safety procedures, COSHH training, Bioinformatics, and Introduction to genomic technologies

**Professional society memberships**

- Associate member of the Royal Society of Biology
- Member of NC3Rs Oncology network
- Member of the Open Bioinformatics Foundation (OBF)

**Interests**

- Star gazer, and an advocate for open-source bioinformatics

**References**

Available upon request