

# BOBBY RANJAN

+65 9647 4046 | [bbbranjan@gmail.com](mailto:bbbranjan@gmail.com) | <http://www.bobbyranjan.com>

---

## Education

---

### Nanyang Technological University (NTU)

Singapore

Ministry of Education Tuition Grant Recipient

Bachelor of Engineering (Computer Engineering)

Graduated in June 2018

Minor in Entrepreneurship

Minor in Life Sciences

GPA: 4.60/5.00

GRE General Test: 162 Verbal, 167 Quantitative, 4.0 Writing

March 2019

### Vikhe Patil Memorial School

Pune, India

Grade 12 | CBSE Board Examinations (A-level equivalent)

Graduated in May 2014

School Rank 1 with 96.8%

---

## Selected Publications

---

- **Ranjan, B.**, Sun W., Park J., Xie R., Alipour F., Singhal V., Prabhakar S. (2020). DUBStepR: correlation-based feature selection for clustering single-cell RNA sequencing data. *bioRxiv*.
  - DUBStepR (Determining the Underlying Basis using Stepwise Regression) uses information contained in pairwise correlations of mRNA levels of genes to select features for clustering.
  - **Received ISMB Fellowship award** for presenting DUBStepR at ISMB (virtual) in June 2020.
- Lee, H. O.\*, Hong, Y.\*, Etlioglu, H. E.\*, Cho, Y. B.\*, Pomella, V., ..., **Ranjan, B.**, ..., Park, W. Y. (2020). Lineage-dependent gene expression programs influence the immune landscape of colorectal cancer. *Nature Genetics*, 1-10.
  - Analysis of 91,103 unsorted single cell transcriptomes from 23 Korean and 6 Belgian CRC patients.
  - Cancer cells displayed normal differentiation programs & genetic alterations that fostered immunosuppressive microenvironments directed by T-reg cells, myofibroblasts and myeloid cells.
- **Ranjan, B.\***, Schmidt, F.\*, Sun, W., Park, J., Honardoost, M. A., Tan, J., ... & Prabhakar, S. (2020). scConsensus: combining supervised and unsupervised clustering for cell type identification in single-cell RNA sequencing data. *bioRxiv*.
  - Hybrid approach to obtain cell type labels in scRNA-seq data using a consensus of supervised and unsupervised clustering.
  - **Selected for flash talk** at Single Cell Analyses Meeting, CSHL in November 2019.
- **Ranjan, B.**, Chong, K. H., & Zheng, J. (2018). Composite mathematical modeling of calcium signaling behind neuronal cell death in Alzheimer's disease. *BMC systems biology*, 12(1), 10.
  - An integrated mathematical model depicting the relationship among amyloid depositions, calcium signaling and mitochondrial permeability transition pore (PTP) related cell apoptosis in Alzheimer's disease.
  - **Selected for talk** at Asia-Pacific Bioinformatics Conference in April 2018
- **Ranjan, B.** (2018). Detection of functional and topological modules in protein interaction networks. *Final Year Project, Nanyang Technological University*.
  - Applied module detection and clustering algorithms to identify functionally relevant protein modules in protein-protein interaction networks.

---

## Achievements & Awards

---

**ISMB 2020 Fellowship Award** | International Society for Computational Biology | June 2020  
**Top 8 & Most Socially Useful Award** | NUS Hack & Roll 2018 | January 2018  
**People's Choice Award** | NUS Hack & Roll 2017 | January 2017  
**President's Research Scholar** | NTU | August 2015  
**Best Freshmen Award** | NUS Hack & Roll 2015 | January 2015  
**All India Rank 1820** | IIT JEE Advanced (India) | May 2014  
**All India Rank 3 scoring 98.2%** | ICSE Board (O-level equivalent) | May 2012

---

## Experience

---

**Bioinformatics Specialist | Genome Institute of Singapore** Singapore  
August 2018 – Present

Group Leader: Dr. Shyam Prabhakar

Ongoing work:

- Robust clustering and interpretation of scRNA-seq data using reference component analysis (manuscript in preparation)
- Analysis of single-cell data as part of an integrative multi-omics study on the mechanism of anti-depressant drug action in the rodent brain (manuscript in preparation)

**Software Design Engineer Intern | BitTitan** Singapore  
May 2017 – August 2017

- Built customer-facing license consumption report for all BitTitan products
- Conducted tech feasibility analysis to improve BitTitan's reporting capacity
- Built code analysis tool to clean up database references across codebase

**Technology Analyst Intern | Bank of America, Merrill Lynch** Singapore  
August 2016 – December 2016

- Used message queues to improve efficiency of payment processing system
  - Built a proof-of-concept (POC) to help onboard new testers onto platform
- 

## Skills

---

- **Genomic data analysis:** scRNA-seq • Bulk RNA-seq • scATAC-seq
  - **Programming languages:** R • Java • Python • C# • C
  - **Wet lab:** 10X scRNA-seq Chemistry • RNAscope
- 

## Hobbies & Interests

---

Hackathons • Science Communication • Football • Table Tennis • Music

---

## Other Activities

---

**Project Officer | Stronghold Diagnostics Labs (SDL) [COVID-19 Testing Facility]**

July 2020 – October 2020 | Singapore

- Assisted in submission of daily clinical reports for SDL
- Performed software integration testing for automated clinical reporting
- Developed visualization tools for staff members to communicate and receive updates

**Website Manager | PGMedOnline.com**

January 2018 – Present | India

- PGMedOnline is a home-based coaching website for all medical examinations
- My responsibilities involve regular website maintenance and liaison with website developers