



## DIVYANSHU SRIVASTAVA

Computational Biologist

### PHD STUDENT

Currently working in Computational Oncology, jointly affiliated with the University of Lausanne and EPFL at Lausanne, Switzerland. Focussed on understanding selection and adaptation in tumour evolution.

### ABOUT ME

I am a IT engineer turned computational biology researcher. I enjoy solving puzzles and problems. Interests include mathematics, electronics and sports.

### SKILLS AND INTERESTS

- ✓ ✓ ✓ Data Science
- ✓ ✓ ✓ Network Biology
- ✓ ✓ ✓ Statistical Computation
- ✓ ✓ ✓ Natural Language Processing
- ✓ ✓ ✓ Python, R, MATLAB, Bash
- ✓ ✓ Parallel Programming
- ✓ ✓ Graph Signal Processing
- ✓ Cuda C, Arduino, C, C++, JAVA

### REACH ME AT

- ☎ +41 78 223 43 34 | +91-9899522926
- 💻 [divyanshusrivastava.github.io](https://github.com/divyanshusrivastava)
- @ [divyanshu28jan@gmail.com](mailto:divyanshu28jan@gmail.com)
- 📍 232, Ch. de Mongevon 4, 1023 Crissier, Vaud, Switzerland

## WORK EXPERIENCE

Researcher | Life Sciences (Microbiology and Metagenomics)  
TATA Research, Development and Design Center (TRDDC) - Pune  
2018 - 2021

## ACADEMIC HISTORY

Indraprastha Institute of Information Technology, Delhi

M. Tech (Computational Biology) 2016-2018

- GPA - 9.26
- Thesis - Graph Signal processing based analysis of biological networks. (June 2017 - July 2018)
- Teaching Assistant - Introduction to Mathematical Biology, Linear Algebra

Cluster Innovation Center, University of Delhi

B. Tech (Information Technology and Mathematical Innovations) 2012-2016

- Percentage - 83.61
- Specialization - Electronics and Robotics
- Internships - Defence Terrain Research Laboratory (DRDO), Birla Institute of Technology, PHD Chamber of Commerce
- Purchase Manager - Robotics Society

## PUBLICATIONS

- Divyanshu Srivastava, Arvind Iyer, Vibhor Kumar, Debarka Sengupta; CellAtlasSearch: a scalable search engine for single cells, *Nucleic Acids Research*, Volume 46, Issue W1, 2 July 2018, Pages W141-W147.
- Divyanshu Srivastava, Krishanu Das Baksi, Kuntal K. Bhusan, Sharmila Mande; 'EviMass': A literature evidence based miner for human microbial associations, *Frontiers in Genetics*, August 2019.
- Sunil Nagpal, Divyanshu Srivastava, Sharmila S. Mande: What if we perceive SARS-CoV-2 genomes as documents? Topic modelling using Latent Dirichlet Allocation to identify mutation signatures and classify SARS-CoV-2 genomes, *bioRxiv* 2020.08.20.258772.

## AWARDS AND ACHIEVEMENTS

- IELTS Score (2019) - 8.5
- GATE Score (2016) - 532
- Awarded First Prize - "Best in Robo" @ Innovation Fair, 2015, National Science Centre, New Delhi for "Android Controlled Robot."