

# Sanjiv Kumar

PH.D. BIOTECHNOLOGY

+46 076 057 62 08 | [sanjiv@kth.se](mailto:sanjiv@kth.se) | 0000-0003-3514-8999 | [sanjiv856](#) | [sanjiv-kumar-b246b44a](#)

Molecular Biology, Biotechnology, Protein Biochemistry, Bacteriology, Bioinformatics

## Research Interests

Highly efficient molecular biologist, with focus on Protein Biochemistry and Infection Biology • Adept at *in-silico* characterization of proteins (phylogenetic analysis, homology modeling, drug discovery, molecular docking and simulations etc.) • Knowledge and experience of Genomics, metagenomics and RNA-Seq data analyses tools and software • Efficient at the techniques in the context of drug discovery against infectious pathogens

## Technical Skills

Molecular biology techniques, Microbiology techniques, Protein biochemistry, Protein crystallography, Drug discovery, Bioinformatics (Pharmaceutical Bioinformatics, Genomic and Metagenomics data analysis), R[Intermediate], Python[Intermediate], bash. Adept user of UNIX.

## Research Experience

**Division of Glycoscience, School of Biotechnology, KTH Royal Institute of Technology** Stockholm, Sweden  
Researcher, Oomycetes Biology August 2017 - Present

As a researcher with the Vaibhav Srivastava's Lab, I am responsible for the development and implementation of research strategy towards novel anti-infectives against Oomycete pathogens. Including following roles:

- Expression, purification and biochemical characterization of proteins involved in cell-wall biogenesis from Oomycete pathogens, Horizon 2020 project
- Cell wall linkage analysis of various Oomycete pathogens, Horizon 2020 project
- Identification and characterization of protein targets and inhibitors important for disease control strategies against *Saprolegnia parasitica* and their *in-vitro* assessment.

**Justin Radolf Lab at University of Connecticut Health Center** Connecticut, USA  
Post-Doctoral Fellow, Infectious Biology April 2015 - August 2017

As a Post-Doctoral Fellow at UConn Health, I was involved with the characterization of rare Outer Membrane Proteins (OMPs) in *Treponema pallidum*. Including following roles:

- Consensus computational framework to identify rare OMPs in *T. pallidum*
- Biophysical and experimental characterization of BAM – complex in *T. pallidum*
- Identification and analysis of sequence variations in *T. pallidum* from clinical samples

**Vineet Sharma Lab at Indian Institute of Science Education and Research, Bhopal (IISER-B)** Bhopal, India  
Post-Doctoral Fellow, Metagenomics and Systems Biology December 2013 - February 2015

As a Post-Doctoral Fellow at MetaBioSys lab, I was responsible for the Metagenomic analysis of human gut and polluted river microbiomes. Some of the roles included:

- Reconstruction of bacterial and viral genomes from multiple human gut metagenomes
- Development of prediction tool for peptidoglycan hydrolases from bacterial genomes
- De novo assembly of genomes from human gut metagenomic data

**S. Ramachandran Lab, Institute of Genomics and Integrative Biology** New Delhi, India  
Junior Research Fellow, Genomics and Integrative Biology December 2007 - February 2015

As a Junior Research Fellow, I was responsible for the identification of novel adhesins from *Mycobacterium tuberculosis*, including • Identification of novel potential adhesins of *M. tuberculosis* H37RV and their experimental validation • Characterization of a novel N-acetylmuramoyl-L-alanine amidase Rv3717 from *M. tuberculosis* H37Rv

## Patent Filed

- Vaibhav Srivastava, **Sanjiv Kumar**, and Vincent Bulone (2020). "Treatment of saprolegniasis." International Patent Application No.PCT/SE2020/050468, Filing date: May 7, 2020. [Link]

## Awards

---

KTH Innovation Challenge Tech for the Global Goals award of 50000 SEK, jointly shared with Vaibhav Srivastava, awarded by KTH Royal Institute of Technology, Stockholm, Sweden (May 2018) • Senior Research Fellowship from CSIR, New Delhi, India (March 2009 – February 2012) • Junior Research Fellowship from CSIR, New Delhi, India, (March 2007 – February 2009) • Defense Scholarship during graduation (B. V. Sc. & A. H.) (1998-2003)

## Education

---

### Institute of Genomics and Integrative Biology (CSIR-IGIB)

Ph.D. in Biotechnology

Delhi, India

2014

### Lala Lajpat Rai University of Veterinary and Animal Sciences

Masters in Veterinary Sciences (M.V.Sc.), Veterinary Microbiology

Haryana, India

2007

### Veterinary College, KVAFS University

Bachelor of Veterinary Sciences & Animal Husbandry (B.V.Sc. & A.H.)

Bangalore, Karnataka, India

2004

## Training

---

**Pharmaceutical Bioinformatics**, Grade: **VG**, Uppsala University, *Uppsala, Sweden*, Aug. 31 - Nov. 01, 2020 [Credential] • **Applied Pharmaceutical Structural Bioinformatics**, Grade: **VG**, Uppsala University, *Uppsala, Sweden*, Aug. 31 - Oct. 15, 2020 [Credential] • **RNA-Seq Data Analysis**, Workshop by National Bioinformatics Infrastructure Sweden (NBIS) and **SciLifeLab**, *Uppsala, Sweden*, May 13 - 15, 2019 [Credential].

## Independent Coursework

---

**Intro to Machine Learning**, Kaggle, Mar. 2021 [Credential] • **Python**, Kaggle, Mar. 2021 [Credential] • **Automate the Boring Stuff with Python Programming**, Udemy, Jun. 2019 [Credential] • **Python Programming Bootcamp**, Udemy, Jun. 2019 [Credential] • **Python for Absolute Beginners**, Udemy, Jun. 2019 [Credential] • **Introduction to Python**, DataCamp, April 2019 [Credential] • **Introduction to R Course**, DataCamp, Nov. 2017 [Credential].

## Publications (Since 2018)

---

For a complete list of publications please see *My Google Scholar Profile*.

First Author Peer-Reviewed Publications – **6** • Total Peer-Reviewed Publications – **19** • Popular Science Articles – **1**.

1. Srivastava, A., Biswas, S., Yadav, S., **Kumar, S.**, Srivastava, V., & Mishra, Y. (2021). Acute cadmium toxicity and post-stress recovery: Insights into coordinated and integrated response/ recovery strategies of *Anabaena sp.* PCC 7120. **Journal of Hazardous Materials**. [ScienceDirect]
2. **Kumar, S.**, Mandal, R.S., Bulone, V. & Srivastava, V. (2020). Identification of growth inhibitors of the fish pathogen *Saprolegnia parasitica* using in silico subtractive proteomics, computational modelling, and biochemical validation. **Frontiers in Microbiology** 11, 2533. [PubMed]
3. Murugan, N.A., **Kumar, S.**, Jeyakanthan, J. & Srivastava, V. (2020). Searching for target-specific and multi-targeting organics for COVID-19 in the Drugbank database with a double scoring approach. **Scientific Reports** 10(1), 19125. [Scientific Reports]
4. Yadav, S., Srivastava, A., Biswas, S., Chaurasia, N., Singh, S.K., **Kumar, S.**, Srivastava, V. & Mishra, Y. (2020) Comparison and optimization of protein extraction and two-dimensional gel electrophoresis protocols for liverworts. **BMC Research Notes** 13, 60. [PubMed]
5. Mittal, P., PK, V.P., Dhakan, D.B., **Kumar, S.** & Sharma, V.K. (2019) Metagenome of a polluted river reveals a reservoir of metabolic and antibiotic resistance genes. **Environmental Microbiome**, 14: 5. [BioMed Central]
6. Kaur, S., Srivastava, A., **Kumar, S.**, Srivastava, V., Ahluwalia, A.S. and Mishra, Y., (2019). Biochemical and proteomic analysis reveals oxidative stress tolerance strategies of *Scenedesmus abundans* against allelochemicals released by *Microcystis aeruginosa*. **Algal Research**, 41, p.101525. [ScienceDirect]
7. **Kumar, S.**, Caimano, M. J., Anand, A., Dey, A., Hawley, K. L., LeDoyt, M., La Vake, C., Cruz, A. R., Ramirez, L. G., Pastekova, L., Bezsonova I, Smajs D, Salazar J. C., & Radolf J. D. (2018) Sequence variation of rare outer membrane protein  $\beta$ -barrel domains in clinical strains provides insights into the evolution of *Treponema pallidum* subsp. *pallidum*, the syphilis spirochete. **mBio** 9(3): e01006-18.[PubMed]

## References

---

References will be provided upon request.