

Arnav Kumar Upadhyay

LinkedIn: <https://www.linkedin.com/in/arnavu/> | arnav.ku.upadhyay@email.com | Mob- +91 9650237371

EDUCATION

Institute of Bioinformatics and Applied Biotechnology (IBAB) , Bangalore, India Post Graduate Diploma Big data Biology	Jan 2021 - Jan2022
Tel Aviv University , Tel-Aviv, Israel Research Scholar Phylogeny, Transcriptomics	Sep 2018 - Sep2020
Gautam Buddha University (GBU) , Noida, India B. tech + M. tech Integrated Biotechnology	July 2013 – July 2018

WORK EXPERIENCE

Medgenome , Bangalore, India Bioinformatics Analyst	Jan 2022 – Current
<ul style="list-style-type: none">Developed and Integrated pipeline using nextflow and GNU makeSupported clinical bioinformatics team (Resolving JIRA, Redmine issues)Created scripts for data processing using python and MySQL	
Publication/Projects	

IBAB, IIIT- Bangalore, India	Dec 2020 – Dec 2021
<ul style="list-style-type: none">Defining Digital RNA signature through Circulating Tumour Cells for scoring stemness and therapeutic resistance in Breast Cancer Patients GitHub - https://github.com/hmalavia/Genomics-Project-IBAB-2021	

Tel Aviv University , Tel-Aviv, Israel	Sep 2018 - Sep2020
<ul style="list-style-type: none">And then there were three...: Extreme regeneration ability of the solitary chordate <i>Polycarpa mytiligera</i> Gordon et al (Front. Cell Dev. Biol.), 2021, 9, 793 DOI: https://doi.org/10.3389/fcell.2021.652466Phylogenetic relationships of Aplousobranchia (Tunicata) based on mitochondrial genomes Arnav Kumar Upadhyay, Noa Shenkar, Dorothée Huchon, Dec 2019 (ISEB2019-Tel-Aviv) Poster presentation - https://www.iseb.org.il/previous-conference/2019-conference/	

Gautam Buddha University (GBU) , Noida, India	Jan 2021 - Jan2022
<ul style="list-style-type: none">Interspersed 5'cis-regulatory elements ascertain the spatio-temporal transcription of cytoskeletal profilin gene family in Arabidopsis, Upadhyay et al (Comput Biol Chem), 2019, 80, 177-186 DOI: https://doi.org/10.1016/j.combiolchem.2019.03.023CytoMegoVirus Infection Database: A Public Omics Database for Systematic and Comparable Information of CMV Kaushik et al, (Interdiscip Sci: Computational Life Sciences), 2020, 12, 2, 169-177 DOI: https://doi.org/10.1007/s12539-019-00350-x	

Skills & Interest

- Technical:** Python (scikit learn, pandas, NumPy), Bash, R (Statistical analysis, Data Visualization)
- Molecular biology:** DNA and RNA isolation, Plasmid isolation, SDS-PAGE, Cloning, PCR, spectrometry, NGS library preparation
- Microbiology:** Media preparation, Maintaining bacterial cultures
- Biological sequence analysis:** Single-cell RNA sequencing, Genomics, Transcriptomics, Phylogenetic analysis
- Version control and Ticket management system:** Git, SVN, JIRA, Redmine, OTRS, and LIMS
- Interest:** Evolutionary biology, Playing Football & Guitar, Pencil Sketching