### Akanksha Yadav

#### Education

2015-2019 Bachelor of Science in Chemistry (4 Year BS)

Along with a Minor degree in Biosciences and Bioengineering Indian Institute of Technology (IIT) Bombay, Mumbai

CGPA: 8.86/10

2019-2021 Master of Science in Molecular Biology

(Ongoing) International Max Planck Research School (IMPRS) Molecular Biology, Göttingen

### Research Experience

Oct 2020 Master's project - Developing reverse regression method for detecting trans-eQTLs

- Present Guide: Dr. Johannes Söding, Quantitative and Computational Biology group, MPI-BPC

- Created simulated-gene expression data to include confounding, cis and trans effects
- Calculated likelihood ratios under the null by fitting a reverse logistic lasso on scrambled data
- Fitted exponential tails to ECDF of test statistic obtained under the null
- Performed ROC curve analysis to evaluate predictive performance of the method

May-June Rotation III - Investigating Ubiquitin dynamics using NMR-based relaxation dispersion 2020 Guide: Prof. Christian Griesinger, NMR-based Structural Biology group, MPI-BPC

- Analyzed  ${}^1H^{-15}N$  **TROSY** experiments of 15N-labeled ubiquitin in water/d8-glycerol, 900MHz
- Calculated  $au_c$  for ubiquitin samples from  $R_1$  and  $R_2$  relaxation data using Ccp-Nmr Analysis
- Analyzed  $^1H^{-15}N$  CPMG relaxation dispersion data and wrote scripts to fit into two/three-state fast-exchange Luz-Meiboom model

Mar-Apr Rotation II - Analyzing MD simulations of ArfB in solution v/s in ribosome complex 2020 Guide: Prof. Helmut Grubmüller, Theoretical and Computational Biophysics group, MPI-BPC

- Performed 600ns of molecular dynamics simulation of free protein ArfB in explicit solvent starting from a cryo-EM derived structure in ribosome complex, using **GROMACS**
- Analyzed secondary structure composition along trajectories and performed principal component analysis (**PCA**) with respect to different domains to identify dominant modes of motion
- Investigated mechanism of C-terminal of ArfB helping to position N-terminal, in the ribosome, for catalyzing **peptidyl-tRNA hydrolysis** using RMSF, RMSD and distance analyses

Jan-Feb Rotation I - Prediction of gene expression measurements using cis and trans-eQTLs 2020 Guide: Dr. Johannes Söding, Quantitative and Computational Biology group, MPI-BPC

- Modified the PredictDB pipeline from PrediXcan to build **gene expression prediction** models using cis- and trans-eQTLs (discovered using **Tejaas**) from GTEx data
- Compared models' (different methods for **confounder correction** used on gene expression input data) performance using summary statistics
- Compared models' predictive performance across different races using nested cross-validation

2016-2019 Molecular Dynamics Studies of Small Molecule Ligands with G-Quadruplex DNA Guide: Prof. Pradeepkumar P.I., Nucleic Acids Chemical Biology lab, IIT Bombay

- Modelled and optimized potential G-quadruplex binding ligands using Gaussian 09
- Performed rigid docking of ligand with receptor G4 DNA in silico using Autodock 4.2
- Performed molecular dynamics simulations of ligand-DNA complex using AMBER 16
- Assisted performing biophysical assays to study interactions of ligands with G-quadruplexes
- Kumari, B.; Yadav, A.; Pany, S.P.; Pradeepkumar P.I; Kanvah, S.; Cationic red emitting fluorophore: A light up NIR fluorescent probe for G4-DNA. *J. Photochem. Photobiol., B*, 2019, 190, 128-136

#### Summer GUI development for a data-based chemical modelling software suite, CANDIY

2018 Guide: Prof. Gaurav Chopra, Dept. of Chemistry, Purdue University

- Studied features of CANDOCK, part of CANDIY, designed for **docking** of small molecule ligands with biomolecules, coded in C++ for adding in the GUI
- Integrated 3D molecule renderer from **Avogadro** (open source) into **Qt**-based GUI of CANDOCK
- Parsed atomic and bond information from CANDOCK to Avogadro for visualization of input molecule integrated with the functions to run the program from the GUI

#### **Scholastic Achievements**

2019-2021	Recipient of stipend by the International Max Planck Research School (IMPRS)
2019	Ranked <b>2nd</b> in the 2015 batch, Department of Chemistry
2015-2019	Granted INSPIRE Scholar Award by the Dept. of Science and Technology, Govt. of India
2018	Recipient of summer stipend under Purdue Undergraduate Research Experience program
2016,2017	Awarded the Institute Academic Prize for two consecutive years
2011-2015	Qualified the National Talent Search Examination and received a scholarship
2013	Selected for the Indian National Mathematical Olympiad from Mumbai zone conducted by
	HBCSE under the National Board of Higher Mathematics
2009	Among top 50 scholarship holders in state-level two-tier Mathematics Prodigy Competition

### **Academic Projects**

## Apr 2019 Macro-molecular Crystallography course project - raw data processing Guide: Prof. Ruchi Anand, Dept. of Chemistry

- Processed diffraction data starting from raw images on iMOSFLM including peak picking, indexing, refining, integration, merging and scaling
- Determined space group, unit cell parameters, mosaicity and processing parameters
- Used COOT to visualize a given electron density and predict structure of putative ligand

# Oct 2018 Bioinformatics course project - Understanding de novo DNA synthesis Guide: Prof. Prasenjit Bhaumik, Dept. of Biosciences and Bioengineering

- Literature review regarding DNA polymerase showing de novo synthesis activity
- Recognizing **primase** and **polymerase** active sites in available PDB structures, understanding the mechanism of action and comparing key residues involved
- Utilizing multiple sequence alignment, secondary and tertiary structure prediction and fold recognition tools to gain insight into structure of **PrimPol**

# Sep 2016 **Molecular biology course project - Modelling a Genetic Switch** *Guide: Prof. Swati Patankar, Dept. of Biosciences and Bioengineering*

- Designed a hypothetical switch to drive a targeted system for checking cancer cell growth
- Incorporated an **optoelectronic gate**, based on a **series of porphyrins**, stimulated using an input photon of green light produced by an intracellular GFP construct
- Applied small molecule targeted therapy strategy by utilizing the switch to activate a repressor ligand photochemically, which then binds to promoter region and arrests transcription

# Nov 2015 Computer Programming and Utilization course project - Text Processor Guide: Prof. Varsha Apte, Dept. of Computer Science and engineering

- Coded an editing software that took raw electronic text as input and used simple commands fed by the user to invoke desired utilities
- Utilized basic structures in C++ to create editing tools like Find and Replace, Encode, Decode

### Skill set

**Languages** C/C++, R, Python, Bash, LATEX

Software PyMol, Chimera, Coot, ClustalX, GROMACS, CCP4i, CcpNmr, Autodock, Gaussian,

AMBER, Rosetta, 3DNA, Adobe Illustrator, AutoCAD, Solidworks

Wet lab UV/Visible and IR spectroscopy, agarose gel electrophoresis, Gibson assembly, PCR and

cloning, DNA quantification, protein purification, SDS-PAGE, cell culture, transfection, mRNA microinjection, fast kinetics using stopped flow, single-molecule FRET using

TIRF microscopy, light microscopy, NMR and Mass Spectroscopy of proteins

### **Key Courses Undertaken**

Major Organic: Reactions of carbonyl compounds, Thermal and photochemical reactions

(Chemistry) Inorganic: Magnetochemistry, Organometallic Chemistry

**Physical:** Chemical Thermodynamics, Data Analysis, Introduction to computational quantum mechanics, Molecular energetics and dynamics, Molecular spectroscopy, Computational Chemistry

Others Calculus, Linear Algebra, Differential equations, Quantum Physics, Basics of electricity/magnetism Minor Molecular Biology, Genetic Engineering, Introduction to Computational Biology, Metabolism

and Bioenergetics, Molecular cell biology and Genetics, Bioinformatics, Molecular biophysics

Master **DNA and Gene Expression:** DNA Structure, Replication and Repair, Genomics, Epigenetics, (2 lectures Transcription, Translation, RNA Splicing, RNA-based regulation, Protein Structures and Folding

Metabolic and Genetic Networks: Enzyme Mechanisms and Regulation, Biological Membranes,

Photosynthesis, Metabolic Networks, Signal Transduction, Microbiomes

**Cell Biology, Immunology, Neuroscience:** Protein Sorting and Processing, Ubiquitin, Nucleocytoplasmic Transport, Cytoskeleton, Cell Cycle, Meiosis, Apoptosis, Cancer, Infectious Diseases, Immunology: Innate Immunity, T Cell Development and Function, B Cells, Sensory Systems, Synapses and Synaptic Transmission, Neurons: Structure, Function, Electrical Properties, Nervous Systems: Early Development, Development of Networks, Glial Cells and Brain Vasculature.

#### **Extracurricular Activities**

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Misc. - Member of **Hertha Sponer College** of Multiscale Bioimaging (MBExC) platform, Göttingen targeting select students to provide interdisciplinary training in natural sciences/biomedicine

- Attended advanced statistical methods course from the Institute for Mathematical Statistics
- Attended UK Biobank Scientific Conference 2020, Lindau Online Science Days 2020
- Joined **The Physics of Life** online summer school 2020 by Princeton University
- Successfully completed the workshop Statistics for Life Scientists conducted by the SIB, Basel
- Qualified for finals of Bioinformatics Contest 2018 organized by Stepik and Rosalind
- Completed courses on **Coursera**: Introduction to genomic technologies JHU, Bioinformatics-I (with honors) UC San Diego, Machine Learning Stanford
- Cleared level A1.1 of **German language** course
- Volunteered to organize Asia-Pacific Conference of Theoretical and Computational Chemistry, 8

Sports - Secured **gold medal** in the 200m event at the annual athletic championship conducted by Bombay City District Amateur Athletic Association (BCDAAA)

- Bagged bronze medal in 4x100m relay at Bombay YMCA annual athletic meet
- Won bronze medals in the events 100,200m at the institute Athletic General Championship
- Part of the IIT Bombay Athletics team selected to participate in Inter IIT competition