Shailya Verma

Reseach Scholar (Int. Ph.D)



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National Centre for Biological Sciences Tata Institute of Fundamental Research UAS-GKVK Campus, Bellary Road Bangalore 560065, INDIA



Work Experience / Internship



Key Research Areas

- Understanding the sequence and structure of various Toll/interleukin-1 (IL-1) receptor (TIR) domains containing proteins.
- Studying the structural effect of mutations with the help of modeling, molecular dynamics simulation, protein residue network analysis.
- Structure based drug designing through virtual screening pipeline and in vitro experimental validation.

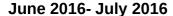
Bhabha Atomic Research Centre | (SRFP) INSA-IASc-NASI | June 2017- July



Key Research Areas

 Structure based drug design using Ribosome as a target.

ANDC, Delhi University | Summer Elite Project |





Key Research Areas

Studied Interaction of Plasmodium vivax
 Duffy Binding Protein (PvDBP) with its host
 receptors the Duffy Antigen (Fy) on RBC
 Surface using Bioinformatics Tools



Education

- Integrated PhD, Biological Sciences NCBS - TIFR |2018 - Present
- B.Sc. (Hons.), Biomedical Sciences ANDC, Delhi University 2015 2018
- Sunbeam School CBSE| 2009-2014



Awards and Achievements

- Selected under top 6 in the Pharmaceutical Science category for Sunpharma Science Scholar Award 2023
- Organized National Post doc symposium (NPDF) 2023 as a lead organizer at Ashoka University, in collaboration with NCBS
- Selected as speaker for NCBS annual talk 2023 (Patterns of Biology)
- Selected for European Conference on Computational Biology held in Barcelona, Spain 2022 with the prestigious DBT-CTEP travel grant.
- Best poster award in NCBS annual talk 2022 (Facets of Biology).
- Volunteered for COVID-19 testing at inStem -NCBS testing facility (2020).
- Secured AIR 82 in CSIR-UGC-NET, 2019.
- Granted DAE TIFR GS fellowship for pursuing Integrated PhD (2018-2024).
- Biomedical science **department topper** for three consecutive years (2016, 2017, 2018)
- Selected for Ideathon, Nobel Prize Series in Ahmedabad, Gujarat - 2017



· English & Hindi



- **Verma S**, Sowdhamini R. A genome-wide search of Toll/Interleukin-1 receptor (TIR) domain-containing adapter molecule (TICAM) and their evolutionary divergence from other TIR domain containing proteins. Biol Direct. 2022 Sep 2;17(1):24. doi: 10.1186/s13062-022-00335-9. PMID: 36056415; PMCID: PMC9440496.
- Bhattacharyya T, Ghosh A, Verma S, Raghu P, Sowdhamini R. Structural rationale to understand the effect of disease-associated mutations on Myotubularin. Curr Res Struct Biol. 2023 Mar 22;5:100100. doi: 10.1016/j.crstbi.2023.100100. PMID: 37101954; PMCID: PMC10123148.
- **Verma S**, Reddy P, Sowdhamini R. Integrated approaches for the recognition of small molecule inhibitors for Toll-like receptor 4. Comput Struct Biotechnol J. 2023 Jul 22;21:3680-3689. doi: 10.1016/j.csbj.2023.07.026. PMID: 37576745; PMCID: PMC10412839.
- **Verma S,** Menon R, Sowdhamini R. Structural insights into the role of deleterious mutations at the dimeric interface of Toll-like receptor interferon-β related adaptor protein. Proteins. Published online May 30, 2024. doi:10.1002/prot.26707
- Verma S, Sharma A, Pathak A, Menon R, and Sowdhamini R. Evolution of protein families, Encyclopedia of Bioinformatics and Computational Biology, 2nd Edition (Ed. Shoba Ranganathan), Elsevier Press (accepted) - Book Chapter
- **Verma, S**, Sowdhamini, R. Sequence and structural analysis of adaptors of Toll-like receptor 4 sheds light on the evolutionary trajectory and functional emergence, bioRxiv, 2024, doi: 10.1101/2024.07.30.605793
- Sharma A, CR Chandrashekar, **Verma S**, Krishna S, Sowdhamini R, Development of a Machine learning model to forecast potential variable residues within pathogenically and therapeutically important proteins in RNA pandemics (Manuscript under preparation)
- Mam B, **Verma S**, Sowdhamini R, Virtual Screening in OBPs to identify natural repellents and metabolite partners (Manuscript under preparation)



References

•	Prof. R. Sowdhamini	(NCBS)	mini@ncbs.res.in	+91-8023666250
•	Dr. Vinothkumar Kutti Ragunath	(NCBS)	vkumar@ncbs.res.in	+91-8061948060
•	Dr. Praveen Kumar Vemula	(inStem)	praveenv@instem.res.in	+91-8023666395
•	Dr. Arvind Ramanathan	(inStem)	arvind@instem.res.in	+91-8061948101



Skills and Competency

Bioinformatics

- Phylogenetics, Sequence based evolutionary analysis, Mutational studies, Protein Modeling, Molecular Docking, Virtual screening, Molecular Dynamics simulations, Protein residue network analysis

Biochemistry

- Cloning, Protein expression and purification, Western blotting, Chromatography, Site directed Mutagenesis, Isothermal Calorimetry

Cell Biology

-Tissue culture handling, Cell based reported assay, Cell viability assay, Polymerase chain reaction