Kunjulakshmi R

PhD Applicant

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• Bioinformatics

• Computational Genomics

• Aging Studies

• Database & Prediction Server

Impact: Developed databases and prediction tools in computational genomics and transcriptomics, driving breakthroughs in anti-aging research, cancer biology, and multi-omics integration. Contributions to projects resulted in scientific publications, fostering advancements in the field.

EDUCATION

Bachelor of Science - Master of Science (BS-MS), Major in Biology

Indian Institute of Science Education and Research (IISER) Berhampur, India

GPA: 7.7/10.00 2018 - 2023

Class XII, Science stream

Central Board of Secondary Education (CBSE), India

92.2%2015 - 2017

Experience

Intern (Computational Genomics & Transcriptomics Lab)

Department of Biotechnology, IIT Hyderabad

Apr 2022 — Present

- Led the development of AagingBase and AagingPEPred, advancing anti-aging research. Collaborated closely, contributing essential insights to ensure project success.
- Significantly contributed to MyeloDB and AMLdb, demonstrating proficiency in database development and multi-omics integration
- Co-authored influential book chapters, enriching scientific literature and emphasizing collaborative research dissemination

Expertise acquired: Database, Prediction server, Anti-aging studies, Multi-omics resource

PROJECTS

A supervised ML based classification model for anti-aging peptides

GitHub

Under Dr.Rahul Kumar, Assistant Professor, Department of Biotechnology, IIT Hyderabad

Jul 2023 — Present

- Engineered a supervised ML-based server for predicting anti-aging peptide efficacy, employing bioinformatics tools and curated datasets
- Achieved an 81% prediction accuracy through training various ML models
- Instigated the development of a user-friendly web interface, facilitating peptide design and identification of anti-aging peptide fragments, thus driving progress in anti-aging interventions

Skills acquired: Feature calculation, Feature ranking, ML algorithms, Bash scripting

AagingBase: A comprehensive database of anti-aging peptides

GitHub

Under Dr.Rahul Kumar, Assistant Professor, Department of Biotechnology, IIT Hyderabad

Jun 2022 - Jun 2023

- Proficiently utilized Linux command line operations in developing AagingBase, enabling efficient data processing, automation, and system management
- Spearheaded the development of "AagingBase," a comprehensive database housing 282 experimentally validated anti-aging peptides, leveraging expertise in database management
- Significantly contributed to anti-aging research by providing AgingBase as a valuable resource, advancing geriatric medicine

Skills acquired: Peptide sequence analysis, Protein structure prediction, Front-end, and back-end development

Exosomal gene expression in pancreatic cancer: Uncovering biomarker potential

GitHub

Under Dr.Rahul Kumar, Assistant Professor, Department of Biotechnology, IIT Hyderabad

Apr 2022 — May 2022

- Conducted robust exosomal gene expression analysis for pancreatic cancer using RNA-seq data, leveraging bioinformatics skills
- Led batch effects mitigation using the PyCombat library, ensuring data integrity
- Developed visualization techniques, revealing downregulation of HIST1H4A and HIST1H4B exosomal genes in pancreatic cancer

Skills acquired: Python & R programming

TECHNICAL SKILLS & TOOLS

Bioinformatics PyMol, MEME Suite, I-TASSER, DSSP, Biopython, BEDTools, SAMtools, AlphaFold

Genomics Sequence alignment, variant calling

Programming Python, R

Command Line Proficiency Unix/Linux, Bash scripting

Data Analysis and Visualization Pandas, matplotlib, seaborn, ggplot2

Machine learning Scikit-learn

Web development HTML5, PHP, CSS, Javascript

Database ManagementMySQLVersion ControlGitDocument TypesettingLaTeX

PUBLICATIONS

Articles

- 1. **R**, **K**., Kumar, A., Vinod Kumar, K. & et.al. AagingBase: A comprehensive database of anti-aging peptides. *Database*. DOI (2024).
- 2. Kumar, A., Vinod Kumar, K., Kundal, K., Sengupta, A., Sharma, S., R, K. & Kumar, R. MyeloDB: A multi-omics resource for Multiple Myeloma. Functional & Integrative Genomics. DOI (2024).
- 3. Vinod Kumar, K., Kumar, A., Kundal, K., Sengupta, A., R, K., Sharma, S., Nishana, M. & Kumar, R. AMLdb: A comprehensive multi-omics platform to understand the pathogenesis and discover biomarkers for acute myeloid leukemia. *Briefings in Functional Genomics*. Under_revision (2024).

Book Chapters (Co-first author)

- 4. R, K., Kumar, A., Vinod Kumar, K., Kundal, K., Sengupta, A. & Kumar, R. in Computational Biology for Stem Cell Research (eds Kumar Raghav, P., Kumar, R., Lathwal, A. & Sharma, N.) DOI, 3–16 (Academic Press, 2024).
- 5. R, K., Korra, B. T., Subashani, Kar, S. S., Kundal, K., Sengupta, A. & Kumar, R. in *Springer Handbook of Chem- and Bioinformatics* (ed Jerzy, L.) (Springer Handbooks, 2024). (In review).

Blogs

- 6. R, K. The Shedded Cells NGSF Intern's Article. (Under review).
- 7. R, K. The Brightly Colored Warning The Qrius Rhino (LINK).
- 8. R, K. In silico Platforms Syntillate: Blog of iGEM IISER Berhampur (LINK).
- 9. R, K. Tiny Plant Wanderers EPISTEME, Volume 2 (LINK).

AWARDS & HONORS

2023 GATE (Life Sciences)

2023 NGSF internship and dissertation program fellow (Next Generation Scientist's Foundation India)

RESPONSIBILITIES & VOLUNTEERING

2022 Advisory board memeber (LaVida-Biology Club: IISER BPR)

2022 Coordinator (Brain awareness week 2022: IISER BPR)

2021 Organizer (BiOlympics 2021 - The biology olympiad)

2019 Volunteer (STREAM-The annual science outreach (IISER BPR))

ENGLISH PROFICIENCY

TOFEL iBT Total (99/120); Speaking (25); Listening (25); Writing (22); Reading (27)

References

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Assistant Professor
Department of Biological Sciences
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