

Sai R Sridatta Prakki

Doctoral Candidate

Helmholtz Center Munich

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Education

- 2023–present **Helmholtz Center Munich, Germany**, *Doctoral Candidate, Bioinformatics.*
- 2010–2012 **Nanyang Technological University (NTU), Singapore**, *Master of Science, Bioinformatics.*
- 2006–2010 **Bharath University, Chennai, India**, *Bachelor of Technology, Bioinformatics.*

Work Experience

- May 2023 – present **Doctoral Candidate**, RESEARCH UNIT FOR PRECISION REGENERATIVE MEDICINE/INSTITUTE OF COMPUTATIONAL BIOLOGY, Helmholtz Munich, Germany.
- Working on sc/snRNAseq data analysis generated from lung tissues (human/mouse)
 - Generating Applications useful for lung research community (Please refer below: Bleomycin Aging App)
 - Training colleagues on improving their Bioinformatics skills

Supervisor: Prof. Dr. Herbert Schiller, (Director, Research Unit for Precision Regenerative Medicine)
Co-Supervisor: Prof. Dr. Fabian Theis, (Director, Institute of Computational Biology)

- Jul 2019 – May 2023 **Senior Bioinformatician**, NATIONAL CENTRE FOR INFECTIOUS DISEASES, Tan Tock Seng Hospital, Singapore.
- Preparation of bacterial genome sequence data, preprocess the datasets, data analysis, reporting to infection control at Tan Tock Seng Hospital, Singapore
 - Writing programming scripts (using R/Bash/PERL) to analyse sequence data
 - Preparation of appropriate documentation and generating analysis reports and summary dashboards (using Github and R Markdown)
 - Leverage on the HPC servers for data storage and managing data transfers with collaborators
 - Developing light-weight tools (developed and published an R package "**CPgeneProfiler**")
 - Automate and optimize clinical bioinformatics processes using workflow management systems

Supervisor: Dr. Ng Oon Tek, (Senior Consultant, National Center for Infectious Diseases)

- Aug 2017 – Jun 2019 **Research Associate - External Staff at NCID**, SAW SWEE HOCK SCHOOL OF PUBLIC HEALTH, NUS, Singapore.
- Development of standard operating procedures for the bioinformatics data analysis
 - Bacterial Genome Data and Metadata management and data transfer
 - Training peer researchers in the work group to achieve insightful results
 - Analyse High Performance Computing (HPC) system computational performance, investigate and resolve performance issues
 - Participate in the preparation and presentation of peer-reviewed articles, posters, reports and grants.

Supervisor: Dr. Ng Oon Tek, (Senior Consultant, National Center for Infectious Diseases)

- Jun 2012 – **Bioinformatics Engineer**, TEMASEK LIFE SCIENCES LABORATORY, NUS, Singapore.
Dec 2016
- Published "**The Asian Seabass Genome Project**", spearheaded by the Reproductive Genomics Group initiated in 2012
 - Published Asian Seabass Transcriptome
 - Experience working on genomics data of other popular aquaculture fishes Tilapia, Arowana

Supervisor: Professor Dr. Laszlo Orban (Senior Principal Investigator)

- Sep 2011 – **Research Intern**, GENOME INSTITUTE OF SINGAPORE, A*STAR, Singapore.
Apr 2012
- Computational analysis of algal organism **Botryococcus** genomes and transcriptomes

Supervisor: Dr. Jianhua Liu (Senior Principal Investigator)

Technical Skills & Abilities

- Programming: Python, PERL, UNIX/Linux, Shell Scripting, AWK, SED
Server Skills: High Performance Computing (HPC), Data management
Statistical Computing: R programming (comfortable writing light-weight packages, using tidyverse, ggplot2 etc)
Applications: MS Word, MS Excel, MS Powerpoint, Vi, LaTeX, LibreOffice
Operating systems: Windows, Ubuntu, CentOS
Version Control: Github (<https://github.com/ramadatta>)
Data Visualizations: R Shiny Bleomycin Aging App (<https://hschillerlabshiny.shinyapps.io/BleomycinAging>)

Certifications

- Data Analysis and Statistical Inference**, *authorized by Duke University.*
The Data Scientist's Toolbox, *authorized by John Hopkins University.*
Data Wrangling in R, *from Linkedin Learning.*
Data Visualization in R with ggplot2, *from Linkedin Learning.*
Big Data Systems Management, *Temasek Polytechnic, Singapore.*
Tools for Data Science, *authorized by IBM.*
Mastering Data Analysis in Excel, *authorized by Duke University.*
R Programming, *authorized by John Hopkins University.*
Bioinformatics: Life Sciences on Your Computer, *authorized by John Hopkins University.*
Hands-on SAS Base, Advanced, Graph and Statistics, *SCRI, A*star, Singapore.*

Other Contributions

- Biostar member**, (*user profile: Prakki Rama*).
Web Blogging, (*A search for solutions*).
Writing computational tools and maintaining Github repositories, (*user profile: ramadatta*).
Publish onscreen tutorials, (*Beginner Bioinformatics commands, Genome alignment using BRIG tool*).
Data wrangling and visualization from Tidyuesday project, (*Analysis of Nurses dataset*).

Awards

- 2025 **EHLRS Grant Award (2,000 EUR) for the project "Plasticity Unforeseen: Loss of PTK7/SOX4 in human fibrotic lung slices drives Pulmonary Epithelial AT2 Reprogramming (PULP-in-PEAR)"**, HELMHOLTZ MUNICH, COMPREHENSIVE PNEUMOLOGY CENTER, Munich, Germany.
- 2009 **Best performer award for poster presentation at National Symposium on "Insilico Drug Design against Leishmaniasis"**, SRM UNIVERSITY, Chennai, India.

International Peer Reviewed Publications

- Janine Gote-Schniering, M. Camila Melo-Narváez, Gowtham Boosarpu, David Lauer, **Sai Rama Sridatta Prakki**, et al. "Single cell decomposition of multicellular aging programs associated with impaired lung regeneration." bioRxiv. 2025. doi: 10.1101/2025.07.24.666371.
- Koh, V., Cabrera, R., **Sai Rama Sridatta Prakki**, et al. "Plasmid dynamics driving carbapenemase gene dissemination in healthcare environments: a nationwide analysis of closed Enterobacterales genomes." Nat Commun. 2025;16:9522. doi: 10.1038/s41467-025-64515-7.
- Sai Rama Sridatta Prakki**, Oon Tek Ng et al. "Dissemination of Pseudomonas aeruginosa bla_{NDM-1} positive ST308 clone in Singapore" Microbiol Spectr. 2023 Jun 15;11(3):e0403322. doi: 10.1128/spectrum.04033-22.
- Kalisvar Marimuthu, Indumathi Venkatachalam, **Sai Rama Sridatta Prakki**, Oon Tek Ng et al. "Whole Genome Sequencing Reveals Hidden Transmission of Carbapenemase-producing Enterobacterales: A retrospective cohort study" Nature communications (2022), in press.
- Sridatta, Prakki Sai Rama**, et al. "CPgeneProfiler: A lightweight R package to profile the Carbapenemase genes from genome assemblies." Journal of Open Source Software 5.54 (2020): 2473.
- Chen, Yahua, Kalisvar Marimuthu, Jeanette Teo, Indumathi Venkatachalam, Benjamin Pei Zhi Chong, Liang De Wang, **Sai Rama Sridatta Prakki** et al. "Acquisition of Plasmid with Carbapenem-Resistance Gene blaKPC2 in Hypervirulent Klebsiella pneumoniae, Singapore." Emerging infectious diseases 26, no. 3 (2020): 549.
- Octavia, Sophie, Marimuthu Kalisvar, Indumathi Venkatachalam, Oon Tek Ng, Weizhen Xu, **Prakki Sai Rama Sridatta**, Yi Fa Ong et al. "Klebsiella pneumoniae and Klebsiella quasipneumoniae define the population structure of bla KPC-2 Klebsiella: a 5 year retrospective genomic study in Singapore." Journal of Antimicrobial Chemotherapy 74, no. 11 (2019): 3205-3210.
- Komissarov A, Vij S, Yurchenko A, Trifonov V, Thevasagayam N, Saju J, **Sridatta P**, Purushothaman K, Graphodatsky A, Orban L, Kuznetsova I. **B chromosomes of the Asian seabass (Lates calcarifer) contribute to genome variations at the level of individuals and populations.** Genes. 2018 Sep 20;9(10):464.
- Shen X, Ngoh SY, Thevasagayam NM, **Prakki SR**, Bhandare P, Tan AW, Tan GQ, Singh S, Phua NC, Vij S, Orban L. **BAC-pool sequencing and analysis confirms growth-associated QTLs in the Asian seabass genome.** Scientific reports. 2016 Nov 8;6:36647.
- Vij S, Kuhl H, Kuznetsova IS, Komissarov A, Yurchenko AA, Van Heusden P, Singh S, Thevasagayam NM, **Prakki SR**, Purushothaman K, Saju JM. **Chromosomal-level assembly of the Asian seabass genome using long sequence reads and multi-layered scaffolding.** PLoS genetics. 2016 Apr 15;12(4):e1005954.
- Ngoh SY, Tan D, Shen X, Kathiresan P, Jiang J, Liew WC, Thevasagayam NM, Kwan HY, Saju JM, **Prakki SR**, Goh CH. **Nutrigenomic and nutritional analyses reveal the effects of pelleted feeds on Asian seabass (Lates calcarifer).** PloS one. 2015 Dec 22;10(12):e0145456.
- Thevasagayam NM, **Sridatta PS**, Jiang J, Tong A, Saju JM, Kathiresan P, Kwan HY, Ngoh SY, Liew WC, Kuznetsova IS, Shen X. **Transcriptome survey of a marine food fish: Asian seabass (Lates calcarifer).** Journal of Marine Science and Engineering. 2015 Jun 2;3(2):382-400.
- Kuznetsova IS, Thevasagayam NM, **Sridatta PS**, Komissarov AS, Saju JM, Ngoh SY, Jiang J, Shen X,

Orban L. **Primary analysis of repeat elements of the Asian seabass (*Lates calcarifer*) transcriptome and genome.** *Frontiers in genetics*. 2014 Jul 25;5:223.

- Shen XY, Kwan HY, Thevasagayam NM, **Prakki SR**, Kuznetsova IS, Ngho SY, Lim Z, Feng F, Chang A, Orban L. **The first transcriptome and genetic linkage map for Asian arowana.** *Molecular ecology resources*. 2014 May;14(3):622-35.

Poster Presentations at International Conferences

- Sai Rama Sridatta Prakki**, Hon Pei Yun, Oon Tek Ng et al., **Dissemination of *Pseudomonas aeruginosa* blaNDM-1 positive ST308 clone in Singapore** at Singapore Health and Biomedical Conference (SHBC), Singapore, 2022
- Weizhen Xu, **Sai Rama Sridatta Prakki**, Natascha May Thevasagayam, Liang De Wang, Wei Xin Khong, Jia Jun Lee, Victor Yi Fa Ong, Kalisvar Marimuthu, Jeanette Teo, Indumathi Venkatachalam, Oon Tek Ng **Tracking plasmid-mediated horizontal gene transmission of blaNDM across Enterobacteriaceae: whole-genome analysis of an international multi-centre dataset** at European Congress of Clinical Microbiology and Infectious Diseases (ECCMID), Amsterdam, Netherlands, 2019
- Glorijoy Shi En Tan, Brenda Ang, Eli Perencevich, Jia Qi Kum, Bee Fong Poh, **Sai Rama Sridatta Prakki**, Liang De Wang, Victor Yi Fa Ong, Natascha May Thevasagayam, Weizhen Xu, Indumathi Venkatachalam, Jeanette Teo, Oon Tek Ng, Kalisvar Marimuthu, **Patient-level risk factors for transmission of carbapenemase-producing Enterobacteriaceae (CPE) in an endemic setting: Findings from contact tracing and whole genome sequencing (WGS) over 8 years** at European Congress of Clinical Microbiology and Infectious Diseases (ECCMID), Amsterdam, Netherlands, 2019
- Kalisvar Marimuthu, Partha Pratim De, Bee Fong Poh, **Sai Rama Sridatta Prakki**, Weizhen Xu, Natascha May Thevasagayam, Victor Yi Fa Ong, Indumathi Venkatachalam, Jeanette Teo, Prabha U. Krishnan, Brenda Ang, Oon Tek Ng. **Population whole-genome sequencing uncovers plasmid-mediated NDM persistence despite high compliance to World Health Organisation (WHO)'s carbapenem-resistant Enterobacteriaceae (CRE) guideline** at European Congress of Clinical Microbiology and Infectious Diseases (ECCMID), Madrid, Spain, 2018
- Sai Rama Sridatta Prakki**, Natascha May Thevasagayam, Junhui Jiang, Jolly M. Saju, Hsiao Yuen Kwan, Woei Chang Liew, Purushothaman Kathiresan, Jun Hong Xia, Gen Hua Yue, Si Lok, Laszlo Orban, **Improved Assembly of the Asian Seabass Transcriptome towards obtaining Full-Length Coding Sequences** at Plant and Animal Genome (PAG) Asia Conference, Singapore, 2014
- Woei Chang Liew, Natascha May Thevasagayam, **Sai Rama Sridatta Prakki**, Jolly M. Saju, Xueyan Shen, Si Yan Ngho, Junhui Jiang, Laszlo Orban, **Sequencing and de novo Assembly of the Mozambique Tilapia Transcriptome** at Plant and Animal Genome (PAG) Asia Conference, Singapore, 2014
- Xueyan Shen, Hsiao Yuen Kwan, Natascha May Thevasagayam, **Sai Rama Sridatta Prakki**, Inna Kuznetcova, Zijie Lim, Felicia Feng, Alex Chang, Laszlo Orban, **Transcriptome Characterization and Construction of a First Generation Genetic Linkage Map for Asian Arowana (*Scleropages formosus*)** at Plant and Animal Genome (PAG) Asia Conference, Singapore, 2014
- Natascha May Thevasagayam, **Sai Rama Sridatta Prakki**, Junhui Jiang, Jolly M. Saju, Hsiao Yuen Kwan, Woei Chang Liew, Purushothaman Kathiresan, Jun Hong Xia, Gen Hua Yue, Laszlo Orban, **Sequencing and Assembly of the Asian seabass Transcriptome** in at Plant and Animal Genome (PAG) Conference, San Diego, California, 2013