

# Prakki Sai Rama Sridatta

Senior Bioinformatician

National Centre for Infectious Diseases

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## Education

- 2010–2012 **Nanyang Technological University (NTU), Singapore**, *Master of Science, Bioinformatics.*  
2006–2010 **Bharath University, Chennai, India**, *Bachelor of Technology, Bioinformatics.*

## Work Experience

Jul 2019 – Present **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.  
**Refer Poster Presentations: 1-3 at European ECCMID conference**

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

Jun 2012 – Dec 2016 **Bioinformatics Engineer**, TEMASEK LIFE SCIENCES LABORATORY, NUS, Singapore.

- 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: 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 Flexdashboard

## 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

2009 **Best performer award for poster presentation at National Symposium on "Insilico Drug Design against Leishmaniasis"**, SRM UNIVERSITY, Chennai, India.

- **Sai Rama Sridatta Prakki**, Oon Tek Ng et al. "Nosocomial infections within a tertiary-care hospital at Singapore caused by *Pseudomonas aeruginosa* *bla*<sub>NDM-1</sub> ST308 clone linked to Environmental isolates" Unpublished manuscript, National Centre for Infectious Diseases, Singapore.

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## International Peer Reviewed Publications

- 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 *bla*KPC2 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, Ngoh 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.

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## Poster Presentations at International Conferences

- 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 *bla*NDM 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 Ngoh, 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