Aroon Chande

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Summary

Experience Over 5 years of industry experience delivering turn-key bioinformatics and software solutions

Client focus Customized existing software tools and analysis platforms to keep costs down

Productivity Authored or contributed to 18 peer-reviewed scientific publications and 2 patents in 8 years

COVID-19 Georgia Tech asymptomatic COVID-19 testing system

Experience

COVID-19 pandemic response projects, Georgia Institute of Technology.

- Lead developer and social media manager for the nationally recognized COVID-19 Event Risk Assessment Planning Tool: https://covid19risk.biosci.gatech.edu/
 - Served over 3 million users and 15 million COVID-19 risk predictions
 - Received national press-coverage in Scientific American, Politico, Bloomberg, and elsewhere
- o Developer of the user-portal for asymptomatic COVID-19 testing system: https://mytest.gatech.edu/
 - Tested over 30,000 samples from faculty, students, and staff

Seattle Genetics, *Seattle*, *WA*, Software Developer II. 2020

- o Created a R Shiny-based analysis and visualization platform for cancer RNA-seq data to empower bench scientists to generate and test hypotheses
 - Facilitated analysis of >20,000 harmonized patient-derived tumor RNA-seg samples
 - Supports machine learning-based clustering approaches and differential gene expression analysis
- o Developed and applied single-sample gene set enrichment techniques

Applied Bioinformatics Laboratory, Atlanta, GA, Bioinformatics Scientist. 2016-2020

- o Partnered with government and industry leaders to deliver client-focused solutions
- o Project lead and software developer for global pathogen surveillance software in collaboration with the Centers for Disease Control and Prevention, shortening turn around time from ≥ 1 day to ≤ 60 seconds and decreasing computational costs by >1000x
- o Developed algorithms and platforms to simplify analysis of next-generation sequencing data

Georgia Institute of Technology, Atlanta, GA, Graduate student. Jordan Lab

- o Developed models for disease burden in Colombia and South America, focused on complex diseases such as type 2 diabetes and coronary artery disease.
- o Redesigned and taught two of the capstone classes for the Bioinformatics Master's Program
- o Trained and mentored multiple Master's and PhD students

Skills

Languages Bash, Git, Python, R, Shell, Perl, PHP, SQL, Unix systems administration

- o Full-stack development: Python with Flask/Django and R with Shiny
- o Data ETL and analysis pipelines in Python and R for use in machine learning and scientific research
- o Server administration and orchestration with Docker/Kubernetes, Ansible and Terraform

NGS Whole Genome Sequencing (WGS), Whole Exome Sequences (WES), RNA-seq, microarray

Wet-lab Tissue culture, confocal and electron microscopy, select pathogens/BSL3, protein purification and crystallography, ELISA and other immunoassays

Education

Georgia Institute of Technology, Atlanta, GA.

Ph.D. Bioinformatics (2020), M.S. Bioinformatics (2016)

Awards: Visiting Scientist Fellowship, São Paulo Research Foundation. Graduate Research Award (2016-2020)

University of Iowa, Iowa City, IA.

B.S. Microbiology (2015). B.S. Biology (2013)