aryakaul.github.io • github.com/aryakaul • a1kaul@ucsd.edu

Education

University of California, San Diego

B.S. Bioengineering: Bioinformatics (GPA: 3.75)

Sep '15 – Present

Graduate Coursework: Algorithms for Scalable Biological Data Analysis, Big Data in Computational Molecular Biology, Network Biology & Biomedicine, Medical and Population Genetics

Undergrad Coursework: Biology Meets Computing, Mathematical Analysis, Data Structures, Advanced Algorithms, Probability and Statistics in Bioinformatics, Biological Databases

Publications & Presentations

Identifying statistically significant chromatin contacts from Hi-C data with Fit-Hi-C *Nature Protocols.* First author. Submitted after invitation and revision October 2018.

Measuring the quality and reproducibility of Hi-C data *Genome Biology*. Sole undergraduate author. Submitted September 2017

Small Molecule Inhibition of cAMP Response Element Binding Protein in Human Acute Myeloid Leukemia Cells *Molecular Cancer Therapeutics.* 2nd undergraduate author. Published June 2016

Research Experience

Sunyaev Lab

DEPARTMENT OF BIOMEDICAL INFORMATICS, HARVARD MEDICAL SCHOOL

Computational Research Assistant

Jun '18 – Present

Developed a computational tool, NovaSplice to predict novel splice sites due to single nucleotide polymorphisms that fall within non-coding DNA regions. Additionally, developed Simdigree, a computational tool to simulate pedigrees to study the overlap between monogenic and polygenic rare diseases.

Ay Lab

VACCINE DISCOVERY, LA JOLLA INSTITUTE FOR ALLERGY AND IMMUNOLOGY

Computational Research Assistant

Jun '16 – Present

Implemented a Python based fast matrix balancing algorithm (HiCKRy) for contemporary chromatin conformation capture techniques. Coded in Numpy and Scipy. Streamlined the installation for Ay Lab software packages using the Anaconda Python distribution system, PyPi, and Github.

Gleeson Lab

DEPARTMENT OF NEUROSCIENCES, UC SAN DIEGO

Biological Research Assistant

Oct '15 - Apr '17

Performed Whole Exome Sequencing (WES) and Whole Genome Sequencing (WGS) to investigate the molecular mechanism between WNT secretion levels in early embryonic development and microcephaly through in vitro and in vivo models. Developed knowledge of a variety of Wet Lab techniques: DNA extraction and sequencing, CRISPR, and DNA Gels.

Professional Activities

Undergraduate Bioinformatics Club

UC San Diego

A community of undergraduate bioinformaticians at UC San Diego serving to strengthen, expand, and support UCSD's bioinformatics program.

President Apr 17 – Present

Created a community service collaboration with Illumina, initiated planning for a future Bioinformatics Department within the Jacobs School of Engineering , and grew mentorship program participation 4-fold.

Vice President of External Affairs

Jun '16 – Apr '17

Organized a year-long speaker series where faculty members and industry leaders within bioinformatics could advise our member base. These included talks from leaders in academia, such as, Dr. Trey Ideker and Dr. Pavel Pevzner. As well as, speakers in industry, ranging from large companies like Illumina to promising startups like Animantis.

Biomedical Engineering Society

UC San Diego

Lab Expo Co-Chair

Mar 16 – Jun 18

Oversaw the planning and hosting of UCSD's largest on-campus research fair with a goal to increase appreciation of science and stress the importance of an interdisciplinary mindset in science. Planned and implemented attendee recruiting efforts that resulted in event registration increasing from under 200 to 500+ attendees. Presenters came from 7 diverse departments and presented research ranging from Bioengineering to pure Mathematics.

Health Hack 17

UC San Diego

Finalist

Mar '17

Annual interdisciplinary two-day hackathon seeking solutions for health care delivery and refugee health. 35 proposals submitted. Our team devised a Raspberry Pi based, RFID solution to address family separation during the refugee experience.

Skills

Computer Science: Python, C++, Java, R, LaTeX, UNIX, HTML, CSS, Javascript, Github, Anaconda/PyPi

Wet Lab: Western Blots, DNA gels, Tissue Culture, DNA Extraction/Sequencing, CRISPR