## RAJEE GANESAN

302-932-4861 || rigan9@gmail.com || @rajeeganesan || Pittsburgh, PA

#### **EDUCATION**

**CARNEGIE MELLON UNIVERSITY** 

Second Year Ph.D. Student in Department of Biology

Pittsburgh, PA Exp. 2026/2027

#### UNIVERSITY OF NORTH CAROLINA, CHAPEL HILL

**Bachelor of Science in Quantitative Biology, Minors in Data Science and Statistics** [Cum Laude, Dean's List]

Chapel Hill, NC May 2022

Relevant Coursework: Italics indicate graduate level coursework

- Bioinformatics: Bioinformatics Data Practicum Genomics and Epigenetics of the Brain, Bioalgorithms, Biostatistics
- Computer Science: Practical Computing and Data Analysis, Object-Oriented Design, Data Structures
- Biology: Advanced Genetics, Cell Biology, Molecular Biology, Evolutionary Genetics
- Mathematics & Physics: Vector Calculus, Decision Sciences, Statistics II, Mechanics, Electricity and Magnetism

## **SKILLS**

Computational - Languages: Python, R, Bash, Java, SAS; Tools: JMP, Jupyter Notebooks, Vim, RMarkdown, ImageJ, Imaris, Slidebooks, ZEN, Job Managers (SLURM), Microsoft Office Products, Package management and installation Experimental - Research Project Design, Scientific and Technical Writing, Conference Presentation, Scientific Literature Research Antibody Staining, Fluorescence and Confocal Microscopy, Cell and RNA isolation, Sequencing preparation

## SELECTED EXPERIENCE

Carnegie Mellon University, Departments of Biology, Computational Biology

Pittsburgh, PA July 2022 - Present

Ph.D. Student in Pfenning Lab, Rotation Student in Pfenning, Ettensohn, and McManus Labs

- Executing projects that use computational approaches to study vocal learning and convergent evolution at a single cell level in order to uncover variants affected in pathways related to speech for experimental validation.
- Evaluated gaps in ongoing research and designed feasible projects to contribute meaningfully to the field.
- Developed an internal pipeline using Bash scripting and public tool integration that determined 3'/5' bias in RNA sequencing was not indicative of mRNA decay, computationally disproving the common hypothesis.
- Executed experimental analysis on urchin developmental stages and prepared a dozen samples for external sequencing.
- Regularly compiled, organized and presented research updates and findings at departmental and large group conferences.

## Furey Lab, UNC Department of Medicine

Senior Research Intern

Chapel Hill, NC Aug 2019 – Apr 2022

- Implemented a bioinformatics pipeline using Perl, Bash and Python scripting in a supercomputing environment to identify sites of allelic imbalance in Crohn's disease patients to better understand potential variants to treat using gene therapies.
- Designed, implemented and troubleshot pipeline testing guanine quadruplex formation in stimulated IL 10 KO macrophages.
- Completed and presented major Honors Thesis with results revealing regional correlation to disease-related genes.
- Mentored and managed two undergraduate students through reimplementation of the pipeline for alternative datasets.

# Watts Lab at National Institutes of Environmental Health Sciences, National Institutes of Health Scholars Connect Program Research Fellow

Durham, NC June 2021 - April 2022

- Developed computational methods using R, Bash and Python scripting to accurately identify regions forming quadruplexes, and integrated experimental results confirming a correlation between these regions and polymerase pausing/gene expression.
- Delivered communicative research presentations biweekly and quarterly at internal and external meetings.

#### **UNC Division One Baseball**

Chapel Hill, NC Dec 2019 – Dec 2020

**Data Analyst** 

- Collaborated with 18 analysts to develop game insights using R, assessing likelihoods of certain pitches in various situations.
- Created easy-to-read scouting reports for upcoming opponents by compiling and organizing data for coaches and players.

## **ADDITIONAL EXPERIENCE**

Residential Computing (IT) Consultant | UNC Housing [AUG 2020 - MAY 2021, CHAPEL HILL,NC]

Summer Research Intern | Meyer Lab at Cold Spring Harbor Laboratory [JUNE 2021 - AUG 2021, COLD SPRING, NY]

Summer Research Intern | Colbert Lab at NIAMS, National Institutes of Health [JUNE 2019 - AUG 2019, BETHESDA, MD]

Research Intern | Ley Lab at La Jolla Institute for Allergy and Immunology [JAN 2019 - JUNE 2019, LA JOLLA, CA]

STEM Head Instructor, Administrative Assistant | Zaniac Parkside [JUN 2017 - MAY 2021, CARY, NC]

## **PUBLICATIONS**

- Rajee Ganesan, Terrence S. Furey. (2022) Impact of guanine quadruplex formation on transcription and open chromatin regions in stimulated IL 10 KO macrophages. [doi: 10.17615/4vg8-aj33]
- Alex Marki, ..., **Rajee Ganesan**, ... and Klaus Ley. (2020) Elongated Neutrophil-Derived Structures (ENDS) form from tethers and are highly elevated in sepsis. Journal of Experimental Medicine. [doi: 10.1084/jem.20200551]