

EDUCATION	Bachelor of Science, Computer Science Bachelor of Arts, Statistics University of Florida, GPA 3.84/4.0	Aug 2015 – May 2019
INDUSTRY & RESEARCH EXPERIENCE	Software Engineer Microsoft, Redmond, WA <ul style="list-style-type: none">Team: Microsoft Security Response Center (MSRC) Bioinformatics Research Assistant Lab of Dr. Lei Zhou, University of Florida <ul style="list-style-type: none">Identified significant mutation patterns in p53 pathway using dataset of 70m+ genome-wide mutationsCreated a Python package to parse and extract taxonomy data from hundreds of protein sequences in an object-oriented fashionApplied machine learning models and visualizations on thousands of genomic data entire Teaching Assistant EML 6934 (Python Programming), University of Florida <ul style="list-style-type: none">Worked with instructor in structuring a new course with 30+ graduate studentsCovered Python basics, NumPy, pandas, Matplotlib, SciPy, scikit-learn Research Intern U.S. Army Research Laboratory, Adelphi, MD <ul style="list-style-type: none">Developed Python script to parse data files and determine circuit design efficiencyFacilitated development of efficient wideband power amplifiers for on-field transmittersNominated by branch chief as outstanding intern for ARL fellowship Software Development/IT Intern Acceleration.net, Gainesville, FL <ul style="list-style-type: none">Designed Flask web app for device backup to facilitate management of 30+ offsite devicesManaged 10+ client websites using WordPress and Linux command lineUtilized Git and Trac to coordinate efforts with co-workers	August 2019 – Present May 2016 – Aug 2019 Sep 2017 – Dec 2017 Jun 2017 – Aug 2017 Jul 2015 – Jan 2017
PROJECTS	serratus.io – Front-facing interface for novel coronavirus homology search effort uniprot-taxonomy – Python library for extracting taxonomy information from UniProt database p53-chip-seq-data – Machine learning and visualizations of lab-generated genomic data	
PUBLICATIONS	Varsha Sundaresan, Victor T. Lin , et al. <i>Significantly Mutated Genes and Regulatory Pathways in SCLC – A Meta-analysis</i> . Cancer Genetics. 2017.	