Jeremy Gresham

Phone Email LinkedIn Website (336) 972-6733 greshjs@gmail.com

edIn https://www.linkedin.com/in/JSGresham site https://ihautai.github.io

Technical Skills

- Experience teaching and training to diverse audiences, including undergraduates and surgeons
- Proficient in technical writing, such as software documentation and troubleshooting instructions
- Experience providing technical support, including hardware and software troubleshooting
- Understanding of basic electronics and soldering
- Experience with microcontrollers, including Arduino and BeagleBone Black
- Proficiency in LATEX, a technical typesetting language

Programming and Technologies

Python3 | Pandas | Django | R/Rcpp | C++11/14/17 | STL | Boost | JavaScript | Scala | Linux | Make | CMake | web development | HTML | CSS | Maple | Matlab | Docker and Singularity | AWS EC2/S3/EKS

Education

Master of Science (MS), Mathematics University of North Carolina Wilmington

2011

Bachelor of Science (BS), Mathematics Minor in Physics North Carolina State University

2007

Employment

Duke Cancer Institute, Bioinformatics Shared Resource *Bioinformatician I/II*

Durham, NC November 2015 - Present

- Developed reproducible pipelines for preprocessing and analysis of Next-Generation Sequencing data from a variety of platforms
- Adapted preprocessing and analysis pipelines for use on Amazon Web Services using container technology
- Developed command-line tools for creation of reproducible pipelines using container image metadata
- Developed pipelines for use on cloud computing platforms
- Collaboration with cancer researchers and statisticians
- Contributed to bcSeq software package for fast sequence alignment for high-throughput shRNA and CRISPR Screens
- Provided independent validation of analysis results generated by colleagues for various research groups
- Participated in grant writing, including design of a cloud-based data sharing and analysis platform using a Linked Data approach
- Programmed in a variety of environments and languages including Python, Scala, C/C++, R, Bash
- Supported Linux, pipeline, and AWS usage within the department
- · Created and maintained training materials for new hires

ShopBot Tools, Inc. Support Technician

Durham, NC November 2013 - May 2015

- Provided technical support for hardware, software, and mechanical aspects of CNC tools for ShopBot customers
- Collected information about customer issues with construction and installation, operation, configuration, customization, and usage of tools and software. Identify problems, investigate causes, generate tests, and analyze test results. Research technical issues to resolve complex issues. Provide specific instructions and guidance to customers to troubleshoot and resolve issues.
- Was responsible for supporting a range of technical issues, including machine repair, parts replacement, wiring instructions, general tool use, calibration, tool path generation (2D/3D CAD/CAM), coding, and integration into manufacturing processes
- Tracked new product development to prepare for new systems, tools and accessories and understand how they function and how they will be integrated with existing software.
- Assisted in general software testing and suggest improvements to user interface

Wake Forest Baptist Medical Center

WakeOne Credential Trainer

Winston-Salem, NC June-October, 2012

- Facilitated multiple specialized one-day training sessions for physicians and surgeons to orient them in the use of a new heavily customized Electronic Medical Record (EMR), focusing on accurate documentation of patient visits and surgical consultations
- Provided in-person follow-up support to medical teams upon system implementation
- Analyzed system roll-out and implementation and provided recommendations to programmers regarding design issues and bugs

Publications

As Co-Author:

- Lin J, Gresham J, Xie J and Owzar K (2017). bcSeq: Fast Sequence Alignment for High-Throughput shRNA and CRISPR Screens. R package version 1.0.1, https://github.com/jl354/bcSeq.
- Thomas Longo, Kathleen F. McGinley, Jennifer A. Freedman, Wiguins Etienne, Yuan Wu, Alexander Sibley, Kouros Owzar, Jeremy Gresham, Christopher Moy, Stephen Szabo, Joel Greshock, Hui Zhou, Yuchen Bai, Brant A. Inman, Targeted Exome Sequencing of the Cancer Genome in Patients with Very High-risk Bladder Cancer, European Urology, Volume 70, Issue 5, 2016, Pages 714-717, ISSN 0302-2838, https://doi.org/10.1016/j.eururo.2016.07.049.

(http://www.sciencedirect.com/science/article/pii/S0302283816304638)

Professional Training

Python Programming

The Iron Yard

Durham, NC May 2015 - July 2015

Final Project: 'Moovn On' webapp Description of Responsibilities:

Heroku management and deployment, creation of a django back-end with read-only RESTful API, data gathering, data cleaning, data visualization with D3.js, map making and bubble-charting *Technologies Used:*

Python3, Django, django-rest-framework, GIS data from Census.gov and Zillow, ogr2ogr, GeoJSON, TopoJSON, D3.js, JavaScript, jQuery, HTML, Heroku, PostgreSQL, SVG