Randy Parrish

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EDUCATION

MSPH, Biostatistics May 2021

Rollins School of Public Health (RSPH), Emory University | Atlanta, GA

GPA: 3.75 | Thesis

BS, Biology (Cellular/Physiology Track)

May 2019 May 2019

BA, Mathematics

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University of Louisville (UofL) | Louisville, KY

Minor: Global Public Health | $Cum\ Laude$ | GPA: 3.71

RESEARCH OUTCOMES

Parrish, R., Gibson, G., Epstein, M. P., Yang, J. TIGAR-V2: Efficient TWAS Tool with Nonparametric Bayesian eQTL Weights of 49 Tissue Types from GTEx V8. (November, 2021). *HGG Advances*. https://doi.org/10.1016/j.xhgg.2021.100068

Yang, J., Parrish, R., Gibson, G., Epstein, M. P. (October, 2021). TIGAR-V2: Efficient TWAS Tool with Nonparametric Bayesian eQTL Weights of 49 Tissue Types from GTEx V8 [Poster]. ASHG Virtual Meeting 2021.

Tang, S., Parrish, R., Buchman, A. S., De Jager, P. L., Bennett, D. A., Epstein, M. P., Yang, J. (October, 2020). Powerful variance-component method for TWAS identifies novel and known risk genes for Alzheimer's dementia [Poster Presentation]. ASHG Virtual Meeting 2020.

Staples, A. K., King, R. A., Doughty, M. A., **Parrish, M. L.**, Klann, J. K., Rinehart, C. A., ... Hatfull, G. F. (2017). Mycobacterium phage TinaFeyge, complete genome [Direct Submission]. GenBank: KY471266.1.

Presentations

SR-TWAS: Leveraging multiple reference panels to improve TWAS power by ensemble machine learning [Platform Presentation]

October 2021

ASHG 2021 Virtual Meeting | Website

Correlation of geographical distribution of mycobacteriophages with genomic clusters and genome distance [Poster]

May 2013

 $12^{th}Annual\ UT\text{-}ORNL\text{-}KBRIN\ Bioinformatics\ Summit\ |\ Buchanan,\ TN\ |\ Abstract$

Mycobacteriophage sub-cluster distribution and correlation to hydrologic regions [Poster]

November 2012

BISC and HPCC Research Symposium, WKU | Bowling Green, KY

Research Experience

Transcriptome-Integrated Genetic Association Resource (TIGAR)

February 2020–Present

- Dr. Jingjing Yang | Emory University School of Medicine | Atlanta, GA
 preparing/analyzing large multi-omics data in HPC cluster environment
 - applying TIGAR tool to GTEx V8 data
 - optimizing code to reduce memory consumption and execution time in all TIGAR Python scripts
 - gene expression model training time for TIGAR example data on 1 core reduced from 17+ minutes to <4 for all models

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Modeling qPCR Curves

August 2018-May 2019

Dr. Gary Cobbs | *UofL* | Louisville, KY

• modified and developed R code to model qPCR curves in order to predict initial absolute concentration of template DNA from raw fluorescence values

- investigated kinetics of primer/probe annealing, processivity in literature
- generated visualizations from model output using ggplot2

Mycobacteriophage Sub-Cluster Distribution & Correlations

August 2012-May 2013

Dr. Claire Rinehart | Western Kentucky University (WKU) | Bowling Green, KY

- used Mathematica to analyze geographic distribution of phages and to generate visualizations
- assessed distribution of phage clusters and examined correlation between geographic distance and genetic distance within each cluster

Mycobacteriophage Isolation & Genome Annotation

August 2011–May 2012

Dr. Rodney King & Dr. Claire Rinehart | SEA-PHAGES & WKU | Bowling Green, KY

TECHNICAL SKILLS

- R, Python, Bash, AWK, SAS 9.4
- Linux, HPC Cluster, Sun Grid Engine, parallel computing in R/Python, Docker, AWS
- MS Office, Zotero reference manager, LATEX, BIBTEX

EMPLOYMENT

Research Specialist

June 2021–Present

Research Assistant February 2020–June 2021

YangLab, Center for Computational and Quantitative Genetics,

Emory University School of Medicine | Atlanta, GA

Small Group Tutor for Biostatistics & Microbiology REACH Learning Resource Center, UofL | Louisville, KY October 2016–April 2018

Professional Membership

American Society of Human Genetics American Statistical Association August 2020–Present July 2017–Present

COMMUNITY INVOLVEMENT

Humanitarian Emergencies Research Team (HERT)

October 2019–December 2020

GEB Data Analysis and Visualizations Co-Lead | RSPH, Emory University

Student Outbreak and Response Team (SORT)

October 2019–December 2020

Member | RSPH, Emory University

Summer Institute in Biostatistics (SIBS)

June 2017–July 2017

Student | RSPH, Emory University

Honors

• ASHG 2021 Charles J. Epstein Trainee Awards Semifinalist

2021

• UofL Dean's List Fall 2016, Spring 2017, Spring 2018

• UofL Dean's Scholar

Fall 2017

• Kentucky Educational Excellence Scholarship

2013, 2016, 2017

• NCWIT Aspirations in Computing Kentucky Affiliate

2013

• Carol Martin Gatton Academy of Mathematics and Science in Kentucky

2011-2013