



MATT RALSTON

Bioinformatics Research Scientist

@ mralston.development@gmail.com

+1-302-547-2437

Philadelphia PA, USA

matthewralston

MatthewRalston

matthewralston.github.io

STRENGTHS

Quantitative modeling

Statistical reasoning Logic

Technical Expertise

Programming Python

Pandas Numpy Scipy

AWS

PROGRAMMING

Python • R • NodeJS • Emacs
Linux • AWS • Docker
D3.js • Perl • bash • grep/sed
L^AT_EX • Matlab • Mongo •
PostgreSQL

LABORATORY

Cell Biology Illumina NGS
library prep • aseptic tech-
nique • Northern & Western
blot • Spectrometry • Mi-
croscopy • qRT-PCR
IF • IHC • FACS Human cell
culture • Anaerobic fermenta-
tion

Analytical Chemistry GC-
MS • HPLC • FTIR

REFERENCES

Kyle Lambert

kyle-lambert-9879848b

Mark Russo

mark-russo-24950a

Malcolm Davis

malcolm-davis-4b37586

ABOUT ME

Combines a strong understanding of the physicochemical basis of biological phenomenon with intermediate computer science and data science abilities. Expert in data visualization, database applications, pipeline development, and analytical problem solving.

EXPERIENCE

Genomics Pipeline Developer | Bayer Crop Sciences

06 2020 - 12 2020

St. Louis MO, USA

- Used AWS and AWFL to get to production (24 steps) within 6 months.
- Learned new methodologies: biosynthetic gene discovery, phylogenetics, completeness, and other functional genomics annotations.
- Developed parsers, data structures, and scripts in Python.

Research Scientist II | Bristol Myers Squibb

05 2015 - 05 2019

Lawrenceville NJ, USA

- Galaxy bioinformatics server administrator, sequencer automation, AWS (Comp. Bio)
- LevTech Innovation award for a dataset version control system git-lfs clone (Comp. Bio)
- Expanded a multi-platform theoretical chemistry framework to enumerate all protomers, tautomers, and rotamers/conformers (2D->3D enumeration) (CADD)
- Parameter optimization and scalability investigation of the framework across multiple chemotypes, SMARTS filters for implausible substructures. (CADD)

EDUCATION

M.Sc. in Bioinformatics and Computational Biology | Univ. of Delaware

08 2012 - 02 2015

Newark DE, USA

- GPA: 3.97

B.Sc. in Biochemistry | Univ. of Delaware

08 2008 - 06 2012

Newark DE, USA

- Minor in Biochemical Engineering

PROJECTS

Kmerdb | |

10 2019 - 01 2021

- A new file format based on the bgzf spec for genomic investigations.
- Normalization, PCA, t-SNE, distance matrices, Markov probabilities, and more!