Ryan P Smith

http://www.linkedin.com/in/rpseq

EDUCATION

Washington University

PhD, Molecular Genetics and Genomics

St Louis, MO

Aug. 2014 - Present

University of Iowa

BS, Microbiology and Informatics; GPA: 3.8

Iowa City, IA

Aug. 2009 – May. 2014

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EXPERIENCE

The McDonnell Genome Institute, Washington University

St Louis, MO

Graduate Research Scientist, Ira Hall Lab, Computational Genetics

Feb 2015 - Present

- **Distributed Computing**: Leverage a high-performance distributed computing cluster to process population-scale genome sequencing data, improving our understanding of the causes and consequences of structural variation in human genomes. (10,000 CPU cores across 550 nodes. 1TB RAM in total.)
- Single-cell Sequencing: Develop novel computational and molecular biology methods to sequence the genomes of single mammalian neurons in collaboration with The Scripps Research Institute at University of California, San Diego. (Often processing up to 10 TB of raw Illumina sequencing data within a 36 hour period.)
- Teaching Assistant: Advised students in characterizing the genome sequences of novel bacteriophages. Hands-on computer lab course using a series of bioinformatics tools to identify genes and predict their function as well as classifying phages into existing phylogenetic groups.

University of Iowa

Iowa City, IA

Undergraduate Research Fellow, Adam Dupuy Lab, Cancer Genetics Aug 2010 - May 2014

- **Genome Editing**: Designed methods for viral genetic engineering in mouse models of human cancers and a sequencing method for detecting resulting transgene insertions.
- **Bioinformatics**: Processed high-throughput genome sequencing data using Linux bioinformatics tools followed by ad-hoc statistical analyses and data visualization in Python and R.

Programming Skills

- Languages: Bash, Python, R (ggplot), SQL, C++, awk
- **Technologies**: Linux/Unix, OSX, Windows, sed, git, Docker, Distributed Computing (IBM Platform LSF, Oracle Grid Engine)