GILLIAN REYNOLDS

Worcestershire, United Kingdom gillianlfreynolds@gmail.com

PERSONAL NOTES

Formerly known as Gillian Woolard

EDUCATION

Individual Interdisciplinary PhD- Computer Science and Plant Genetics

August 2019 - present

Co-Supervisors: Dr. Brendan Mumey, Dr. Jennifer Lachowiec

The Gianforte School of Computing and The Department of Plant Sciences and Plant Pathology

Montana State University, USA

PhD in Plant Genetics - changed programs

January 2018 - August 2019

Interim Supervisor: Dr. Tracy Dougher August 2018- August 2019 Former Supervisor: Dr. Hikmet Budak January 2018 - August 2018

Department of Plant Sciences and Plant Pathology

Montana State University, USA

MRes Biology: Bioinformatics and Genomics

September 2015 - August 2018

The University of Worcester, UK

Biology Bsc(Hons)

September 2011- July 2014

The University of Worcester, UK

SKILLS

Biological

As a result of the varied projects of which I have been a part of I have developed a number of skills in computational biology. These include gene and genome alignments, phylogenetic tree construction, RNA-Seq analysis, GBS data analysis, sequence read mapping, genome assembly, genome annotation, comparative genomics and developing machine learning models for the analysis of short-read sequencing data and biological scientific texts. I also have a sound understanding of a range of biological topics including genomics, genetics and evolutionary biology. I have undertaken a number of self-directed courses in topics including genomic technologies and systems biology.

Computational

I am proficient the python coding language and have extensive experience work on the the linux command line. I am also competent in the R programming language. I have extensive experience with working with high performance computers, including those that use the SLURM workload manager. I am also experienced with working with SciKit learn tools for both supervised and unsupervised machine learning tasks including the analysis of short-read sequencing data and biological natural lanuage processing (BioNLP) tasks. I also have experience in algorithm design and implementation, including CPU parallelization.

Teaching and Supervision

I have enjoyed several undergraduate teaching and supervision opportunities at the University of Worcester. In addition, I have also been involved in the development of educational tools in Bioinformatics for both undergraduates and pre-university students. At Montana State University I have also mentored an undergraduate student on the analysis of large scale RNA-Seq data.

Interdisciplinary

As a PhD student who works across two departments at MSU I have been able to hone my skills in interdisciplinary communication through presentations, seminars, meetings, collaborations and initiating and developing a joint PhD proposal.

Outreach

I have participated in a number of outreach projects to engage students in STEM with audiences from primary schools, high schools and sixth form colleges.

RESEARCH PUBLICATIONS AND CONFERENCE PRESENTATIONS

Reynolds,G., Strnadova-Neeley,V., Lachowiec,J. MinHash k-mer sketching highlights allopolyploid subgenome sequence differentiation. *ISCB-Africa ASBCB (Accepted)*. 2021

Reynolds,G., Lachowiec,J., Strnadova-Neeley,V. Identifying features for subgenomic sequence identification in a De Bruijn Graph (DBG). Tech talks at the CRA-WP Grad Cohort Workshop for Women Conference. 2021

Williams, L., Reynolds, G., Mumey, B. (2019). RNA Transcript Assembly Using Inexact Flows. In 2019 IEEE International Conference on Bioinformatics and Biomedicine (BIBM). IEEE, 2019

Manuweera, B., **Reynolds**, **G**., Kahanda, I. (2019). Computational Methods for the *Ab initio* Identification of Novel microRNA in Plants: A Systematic Review. *PeerJ Computer Science* 5:e233 2019.

Pourreza Shahri, Morteza, Gillian Reynolds, Mandi Marie Roe, and Indika Kahanda. "PPPred: Classifying Protein-phenotype Co-mentions Extracted from Biomedical Literature." In Proceedings of the 10th ACM International Conference on Bioinformatics, Computational Biology and Health Informatics, pp. 414-422. ACM, 2019.

Shahri, Morteza Pourreza, Madhusudan Srinivasan, **Gillian Reynolds**, Diane Bimczok, Indika Kahanda, and Upulee Kanewala."Metamorphic Testing for Quality Assurance of Protein Function Prediction Tools." In 2019 IEEE International Conference On Artificial Intelligence Testing (AITest), pp. 140-148. IEEE, 2019.

Rajasekar, Karthik V., Shuangxi Ji, Rachel J. Coulthard, Jon P. Ride, **Gillian L. Reynolds**, Peter J. Winn, Michael J. Wheeler, Eva I. Hyde, and Lorna J. Smith. "Structure of SPH (self-incompatibility protein homologue) proteins: a widespread family of small, highly stable, secreted proteins." Biochemical Journal 476, no. 5 (2019): 809-826.

Sherrad, G., Woolard, G., Wheeler, M. Investigating the function of a small secreted protein family in physcomitrella patens. *MOSS 2016 Conference*. 2nd-5th September 2016, University of Leeds, U.K.

TEACHING & UNDERGRADUATE SUPERVISION

2020
2017-2018
2017
2014-2016
2014-2015

EMPLOYMENT

Graduate Research Assistantship, Montana State University	2018 - present
Associate Lecturer in Biology, University of Worcester	2016 - 2018
Independent Study Co-Supervisor, University of Worcester	2016 - 2017
Data Quality Improvement Officer, National Health Service (NHS)	2015 - 2017
Guest Lecturer in Biology, University of Worcester	2014 - 2016
Research Assistant, University of Worcester	2014 - 2015

PROFESSIONAL MEMBERSHIPS

REFERENCES

Dr. Jennifer Lachowiec Plant Biosciences Building 305 jennifer.lachowiec@montana.edu (406) 994-2903

Dr. Brendan Mumey Norm Asbjornson Hall 253B brendan.mumey@montana.edu (406) 994-7811