Ryan M. Moore

Graduate Research Assistant, Center for Bioinformatics & Computational Biology College of Agriculture & Natural Resources, University of Delaware

Delaware Biotechnology Institute 15 Innovation Way, Suite 205 Newark, DE 19711-5449 moorer@udel.edu

Education

- Ph.D. (in progress), Bioinformatics & Systems Biology. **University of Delaware**, Newark, DE. K. Eric Wommack, advisor.
- B.S., Mathematics (Hons.); Minors in Biology, Music. (2011). **Chestnut Hill College**, Philadelphia, PA.

Study abroad

- Université Catholique de Lyon, Lyon, France. (2012).
- University of Canterbury, Christchurch, New Zealand. (2008).

Awards & Distinctions

- University of Delaware Doctoral Fellowship. (2018 2019).
- Magna Cum Laude, Mathematics, Chestnut Hill College. (2011).
- Departmental honors in mathematics, Chestnut Hill College. (2011).
- Kappa Gamma Pi honors society, Chestnut Hill College. (2011).
- Sigma Zeta honors society, Chestnut Hill College. (2011).
- Delta Epsilon Sigma honors society (President), Chestnut Hill College. (2010).
- Alpha Lambda Delta honors society (Senior Advisor), Chestnut Hill College. (2007).

Professional Society Membership

• American Society for Microbiology. (2015).

Research & Scholarly Activity

Manuscripts & preprints

- 1. DJ Nasko, BD Ferrell, **RM Moore**, JD Bhavsar, SW Polson, KE Wommack. (Mar 2019). *CRISPR Spacers Indicate Preferential Matching of Specific Virioplankton Genes.* mBio, 10 (2) e02651-18; doi: 10.1128/mBio.02651-18.
- 2. AO Harrison, **RM Moore**, SW Polson, KE Wommack. (Feb 2019). *Reannotation of the Ribonucleotide Reductase in a Cyanophage Reveals Life History Strategies Within the Virioplankton*. Frontiers in Microbiology, 10, 134; doi: 10.3389/fmicb.2019.00134.

- 3. SM McAllister, **RM Moore**, A Gartman, GW Luther, D Emerson, CS Chan. (Jan 2019). *The Fe(II)-oxidizing Zetaproteobacteria: Historical, ecological, and genomic perspectives.* FEMS Microbiology Ecology, fizo15, doi; 10.1093/femsec/fizo15.
- 4. SM McAllister*, **RM Moore***, CS Chan. (Aug 2018). ZetaHunter, a Reproducible Taxonomic Classification Tool for Tracking the Ecology of the Zetaproteobacteria and Other Poorly Resolved Taxa. Microbiology Resource Announcements, 7 (7) e00932-18; doi: 10.1128/MRA.00932-18. (*Contributed equally)
- 5. **RM Moore**, AO Harrison, SM McAllister, KE Wommack. (Aug 2018). *Iroki: automatic customization and visualization of phylogenetic trees.* bioRxiv 106138; doi: 10.1101/106138.
- 6. JM Chopyk, **RM Moore**, ZR Stromberg, GL Lewis, DG Renter, N Cernicchiaro, RA Moxley, KE Wommack. (Mar 2016). *Presence of pathogenic* Escherichia coli *is correlated with bacterial community diversity and composition on pre-harvest cattle hides*. Microbiome 4:9; doi: 10.1186/s40168-016-0155-4

Theses

1. **RM Moore.** (2010). A Survey of Applications of Linear Algebra to Other Branches of Mathematics. Honors thesis in Mathematics. Chestnut Hill College.

Posters & Presentations

- 1. KE Wommack, SW Polson, BD Ferrell, DJ Nasko, JM Chopyk, EG Sakowski, RM Marine, AO Harrison, and **RM Moore**. (Jun 2019). *Genes Demonstrating Important Phenomic Connections Can Improve Ecological Understanding Of Unknown Viruses Of Microbes*. ASM Microbe 2019, San Francisco, CA. (Poster)
- 2. **RM Moore**, AO Harrison, SW Polson, and KE Wommack. (Mar 2019). *InteinFinder:* automated intein detection from large protein datasets. The 4th DENIN Symposium for Environmental Research, University of Delaware, Newark, DE. (Poster)
- 3. AO Harrison, **RM Moore**, BD Ferrell, SW Polson, and KE Wommack. (Mar 2019). Reannotation of the ribonucleotide reductase in a cyanophage reveals life history strategies. The 4th DENIN Symposium for Environmental Research, University of Delaware, Newark, DE. (Poster)
- 4. K Haramoto, **RM Moore**, P Joglekar, V Richards, SW Polson, KE Wommack, and J Fuhrmann. (Feb 2019). *Host Range Comparison of Lytic Rhizobiophages.* The 4th DENIN Symposium for Environmental Research, University of Delaware, Newark, DE. (Poster)
- 5. **RM Moore**. (Feb 2019). *InteinFinder: automated intein detection from large protein datasets*. The 7th Annual UD Microbial Systems Symposium, University of Delaware, Newark, DE. (Presentation)
- 6. K Haramoto, **RM Moore**, P Joglekar, V Richards, SW Polson, KE Wommack, and J Fuhrmann. (Feb 2019). *Host Range Comparison of Lytic Rhizobiophages*. The 7th Annual UD Microbial Systems Symposium, University of Delaware, Newark, DE. (Poster)
- 7. KE Wommack, SW Polson, BD Ferrell, DJ Nasko, JM Chopyk, EG Sakowski, RM Marine, AO Harrison, JF Dums, and **RM Moore**. (Jan 2019). *Squeezing out understanding from sequences: Genome to Phenome Connections in Viruses of Microbes.* The 4th Xiamen Symposium on Marine Environmental Sciences, Xiamen, China. (Poster)

- 8. AO Harrison, **RM Moore**, BD Ferrell, SW Polson, KE Wommack. (Jun 2018). *Ribonucleotide reductase as a marker for marine virioplankton communities.* Aquatic Virus Workshop 9, University of Nebraska-Lincoln, Lincoln, NE. (Poster)
- 9. AO Harrison, **RM Moore**, BD Ferrell, SW Polson, KE Wommack. (May 2018). *Ribonucleotide reductase as a marker for marine virioplankton communities.* College of Earth, Ocean, and Environment Graduate Student Symposium, University of Delaware, Newark, DE.
- 10. AO Harrison, **RM Moore**, BD Ferrell, SW Polson, KE Wommack. (Mar 2018). *Ribonucleotide reductase as a marker for marine virioplankton communities.* Viral EcoGenomics and Applications Symposium (VEGA), San Francisco, CA. (Poster)
- 11. AO Harrison, **RM Moore**, KE Wommack. (Feb 2018). *Ribonucleotide reductase as a marker for marine virioplankton populations.* Sixth Annual Microbial Systems Symposium, University of Delaware, Newark, DE. (Poster)
- 12. **RM Moore**, JM Chopyk, AO Harrison, SM McAllister, D Renter, N Cernicchiaro, R Moxley, and KE Wommack. (Jun 2017). *Analyzing connections between microbial community structure and pathogenic* E. coli *in cattle hide and feces.* 2017 Annual STEC-CAP Conference, Lincoln, NE. (Poster)
- 13. **RM Moore**, AO Harrison, SM McAllister, RL Marine, CS Chan, and KE Wommack. (Jun 2017). *Iroki: automatic customization for phylogenetic trees.* 2017 CANR Research Symposium, University of Delaware, Newark, DE. (Poster)
- 14. AO Harrison, **RM Moore**, JM Chopyk, D Renter, N Cernicchiaro, R Moxley, KE Wommack. (Feb 2017). *Examining connections between viral community diversity and pathogenic* E. Coli *in pre-harvest cattle feces using DNA Polymerase A.* Fifth Annual Microbial Systems Symposium, University of Delaware, Newark, DE. (Poster)
- 15. **RM Moore**, JM Chopyk, PR Joglekar, AO Harrison, N Cernicchiaro, D Renter, R Moxley, KE Wommack. (Jun 2016). *Exploring connections between pathogenic* E. coli and shifts in viral and bacterial community structure and metabolic potential in pre-harvest cattle feces. 2016 Annual STEC-CAP Conference, Lincoln, NE. (Poster)
- 16. AO Harrison, **RM Moore**, JM Chopyk, D Renter, N Cernicchiaro, R Moxley, KE Wommack. (Jun 2016). *Examining connections between viral community diversity and pathogenic* E. Coli *in pre-harvest cattle feces using DNA Polymerase A.* 2016 Annual STEC-CAP Conference, Lincoln, NE. (Poster)
- 17. AO Harrison, **RM Moore**, RM Marine, SW Polson, KE Wommack. (May 2016). *Fine scale temporal dynamics of estuarine bacterioplankton and virioplankton*. 2016 DENIN Scholars Symposium, University of Delaware, Newark, DE. (Poster)
- 18. CE Shatley, **RM Moore**, and SW Polson. (Aug 2015). *Examining the accuracy of recruitment based abundance estimations in metagenomes.* 2015 Summer Scholars poster presentation, University of Delaware, Newark, DE. (Poster)
- 19. JM Chopyk, **RM Moore**, AMC Ferelli, D Renter, N Cernicchiaro, R Moxley, KE Wommack. (Jun 2015). *Microbiome influences on STEC prevalence in pre-harvest cattle.* 2015 Annual STEC-CAP Conference, Manhattan, KS. (Presentation)
- 20. **RM Moore**, J Chopyk, N Cernicchiaro, D Renter, R Moxley, KE Wommack. (Jun 2015). *Shiga toxigenic* Escherichia coli *is associated with shifts in the bacterial community structure of pre-harvest cattle hides.* 2015 Annual STEC-CAP Conference, Manhattan, KS. (Poster)
- 21. JM Chopyk, DJ Nasko, EG Sakowski, **RM Moore**, SW Polson, KE Wommack. (Jun 2015). *Using Polymerase A to explore evolutionary realtionships, replicase organization and diversity in viral assemblages.* 115th General Meeting of the American Society for Microbiology, New Orleans, LA. (Poster)
- 22. **RM Moore**, J Chopyk, N Cernicchiaro, D Renter, R Moxley, KE Wommack. (Jun 2015). *Shiga toxigenic* Escherichia coli *is associated with shifts in the bacterial*

- *community structure of pre-harvest cattle hides.* 115th General Meeting of the American Society for Microbiology, New Orleans, LA. (Poster)
- 23. JM Chopyk, ZF DiSpirito, AMC Ferelli, **RM Moore**, D Renter, N Cernicchiaro, R Moxley, KE Wommack. (March 2015). *The Influence of viral and bacterial community diversity on pathogenic E. coli prevalence in pre-harvest cattle.* Polony Workshop, Haifa, Israel. (Poster)
- 24. JM Chopyk, ZF DiSpirito, **RM Moore**, D Renter, N Cernicchiaro, R Moxley, KE Wommack. (May 2014). *The correlation between microbial community diversity and STEC serotypes in beef processing samples.* 2014 Annual STEC-CAP Conference, Lincoln, NE. (Poster)
- 25. ZF DiSpirito, JM Chopyk, AMC Ferelli, **RM Moore**, D Renter, N Cernicchiaro, R Moxley, KE Wommack. (Aug 2014). *Viral community diversity correlation with STEC presence in cattle fecal samples.* Undergraduate Research and Service Celebratory Symposium, University of Delaware, Newark, DE. (Poster)
- 26. SA Smith, **RM Moore**, KE Wommack, S Polson. (Aug 2014). *Assembly-based metrics of abundance in shotgun metagenomes may provide inaccurate estimations of community structure.* Undergraduate Research and Service Celebratory Symposium, University of Delaware, Newark, DE. (Poster)
- 27. **RM Moore.** (Feb 2010). *Two Approaches to Finding Maxima & Minima of Multivariate Functions*. The 24th Annual Student Mathematics Conference, Moravian College, Bethlehem, PA. (Presentation)
- 28. **RM Moore.** (2009). *A New Spiritual Sublime: The Spiritual Revolution in the Works of Kasimir Malevich.* Southeastern Pennsylvania Consortium for Higher Education Honors Conference, Arcadia University, Glenside, PA. (Presentation).

Professional meetings attended

- 1. 2017 Annual STEC-CAP Conference. (Jun 2017). Lincoln, NE.
- 2. 2016 Annual STEC-CAP Conference. (Jun 2016). Lincoln, NE.
- 3. 2015 Annual STEC-CAP Conference. (Jun 2015). Manhattan, KS.
- 4. 115th General Meeting of the American Society for Microbiology. (May 2015). New Orleans, LA.
- 5. Environmental Virology Workshop. (Jan 2013). Biosphere 2, Tucson, AZ.

Relevant Skills

Programming languages, frameworks, and statistical packages

• Ruby, Ruby on Rails, C, R, Bash, Python, JavaScript

Professional Service & Outreach

Publicly available tools & software

- AAI. Multi-genome amino acid similarity pipeline. https://github.com/mooreryan/aai.
- BigSimon. An ensemble strategy for predicting virus-host interactions. https://github.com/moorervan/big_simon.
- IdentiBin. Dereplicating genomic bins from metagenomes. https://github.com/mooreryan/IdentiBin.

- InteinFinder. Automated intein detection from large protein datasets. https://github.com/mooreryan/InteinFinder
- Iroki. Automatic customization for phylogenetic trees. https://www.iroki.net.
- LSAForGenomes. Partitioning genomes and metagenomes with latent semantic analysis. https://github.com/mooreryan/lsa_for_genomes.
- markdown-dnd-images. Drag and drop images directly into a markdown-mode Emacs buffer. https://github.com/mooreryan/markdown-dnd-images.
- ParseFasta. Ruby library providing programmatic access to FASTA and FASTQ files. https://github.com/mooreryan/parse_fasta.
- PASV (Protein Active Site Validation). Protein partitioning and filtering using multiple sequence alignments and conserved residues. https://github.com/moorervan/pasv.
- RubyLikeC. C library of functions that behave like their Ruby counterparts. https://github.com/mooreryan/ruby like c.
- ZetaHunter. Automatic, stable OTU calling using user curated databases. https://github.com/mooreryan/ZetaHunter.

Open source contributions (pull requests, issues opened, etc.)

- awesome-c. (2018). https://github.com/aleksandar-todorovic/awesome-c
- swarm. (2018). https://github.com/torognes/swarm
- WISH. (2018). https://github.com/soedinglab/WISH
- VirHostMatcher. (2018). https://github.com/jessieren/VirHostMatcher
- jsPhyloSVG. (2016). https://github.com/guyleonard/jsPhyloSVG
- ListCompare. (2016). https://github.com/seanmcallister/list_compare
- Metagenomics Workshop. (2016).
 https://github.com/EnvGen/metagenomics-workshop
- BEAR. (2015). https://github.com/sej917/BEAR
- SeqComplex. (2015). https://github.com/caballero/SeqComplex
- Philly Open Health. (2015). https://github.com/PhillyHealthDataRepo/philly-open-health

Undergraduate research mentorship

- Amelia Harrison. (2016).
- Courtney Shatley. (2015).
- Steve Smith. (2013 2014).

Peer-review of manuscripts

Microbiome

Outreach

- Outreach with Havre de Grace High School Biomed class. (2016 2017). Havre de Grace, MD.
- Outreach with Cub Scouts of Delaware. (2015 2017). Granogue, DE.
- Outreach at the Delaware Biotechnology Institute. (2014 2015). Newark, DE.
- Marvelous Microbes 4-H camp. (2015). Newark, DE.

- Lab Assistant. (2011). St. Genevieve School, Flourtown, PA.
- Math tutor. (2006 2007, 2011). Chestnut Hill College, Philadelphia, PA.