Joshua J. Hamilton

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Employment History

_	Federation Bio, Inc.	South San Francisco, CA
•	Data Scientist	$\it 2019-present$
•	University of Wisconsin-Madison (UW-Madison) Postdoctoral Associate, Department of Biochemistry	Madison, WI 2018-2019
•	UW-Madison Postdoctoral Associate, Department of Bacteriology	Madison, WI 2014-2017

Education and Academic Positions

_	UW-Madison	Madison, WI
•	Ph.D., Chemical Engineering	2009-2014
_	Case Western Reserve University (CWRU)	Cleveland, OH
•	B.S., Chemical Engineering	2005-2009

Patents

1. Swem L, Ricci D, Brumbaugh AR, Cremin J, **Hamilton JJ**, Tripathi S, Wong L, Romasko H, Bracken R, Drabant Conley E. (2021) *Microbial consortia for the treatment of disease*. US Patent Application No. PCT/US2021/021,790.

Publications

- 14. Clark RL, Connors B, Stevenson DM, Hromada SE, **Hamilton JJ**, Amador-Noguez D, and OS Venturelli. (2021) Design of synthetic human gut microbiome assembly and function. Nature Communications. 12: 3254. doi:10.1038/s41467-021-22938-y.
- 13. Scarborough MJ, **Hamilton JJ**, Erb EA, Donohue TJ, and DR Noguera. (2020)

 Diagnosing and Predicting Mixed-Culture Fermentations with Unicellular and Guild-Based

 Metabolic Models. mSystems. 5(5):e00755-20 doi:10.1128/mSystems.00755-20.
- 12. Cao X*, **Hamilton JJ***, and OS Venturelli. (2019) *Understanding and Engineering Distributed Biochemical Pathways in Microbial Communities*. Biochemistry. 58(2): 94-107. doi:10.1021/acs.biochem.8b01006.
- 11. Dwulit-Smith JR, **Hamilton JJ**, Stevenson DM, He S, Oyserman BO, Moya-Flores F, Amador-Noguez D, McMahon KD, and KT Forest. (2018) acI Actinobacteria Assemble a Functional Actinorhodopsin with Natively-synthesized Retinal. Applied and Environmental Microbiology. 84(24): e01678-18. doi:10.1128/AEM.01678-18.

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- Scarborough MJ, Lawson CE, Hamilton JJ, Donohue TJ, and DR Noguera. (2018)
 Metatranscriptomic and Thermodynamic Insights into Medium-Chain Fatty Acid Production Using an Anaerobic Microbiome. mSystems. 3(6): e00221-18.
 doi:10.1128/mSystems.00221-18.
- 9. Rohwer RR, **Hamilton JJ**, Newton, RJ, and KD McMahon. (2018) *TaxAss: Leveraging a Custom Freshwater Database Achieves Fine-Scale Taxonomic Resolution*. mSphere. 3(5): e00327-18. doi:10.1128/mSphere.00327-18.
- 8. Garcia SL, Buck M, **Hamilton JJ**, Wurzbacher C, Grossart HP, McMahon KD, and A Eiler. (2018) *Model Communities Hint at Promiscuous Metabolic Linkages between Ubiquitous Free-Living Freshwater Bacteria*. mSphere. 3(3): e00202-18. doi:10.1128/mSphere.00202-18.
- 7. Hamilton JJ, Garcia SL, Brown BS[†], Oyserman BO, Moya F, Bertilsson S, Malmstrom RR, Forest KT, and KD McMahon. (2017) Metabolic Network Analysis and Metatranscriptomics Reveals Auxotrophies and Nutrient Sources of the Cosmopolitan Freshwater Microbial Lineage acI. mSystems. 2(4): e00091-17. doi:10.1128/mSystems.00091-17.
- 6. Lawson CE, Wu S, Bhattacharjee AS, **Hamilton JJ**, McMahon KD, Goel R, and DR Noguera. (2017) *Metabolic network analysis reveals microbial community interactions in anammox granules*. Nature Communications. 8: 15416. doi:10.1038/ncomms15416.
- 5. **Hamilton JJ**, Calixto Contreras M[†], and JL Reed. (2015) Thermodynamics and H₂ Transfer in a Methanogenic, Syntrophic Community. PLoS Computational Biology. 11(7): e1004364. doi:10.1371/journal.pcbi.1004364.
- 4. Vinay-Lara E, **Hamilton JJ**, Stahl B, Broadbent JR, Reed JL, and JL Steele. (2014) Genome-Scale Reconstruction of Metabolic Networks of Lactobacillus casei ATCC 334 and 12A. PLoS ONE. 9(11): e110785. doi:10.1371/journal.pone.0110785.
- 3. Hamilton JJ and JL Reed. (2014) Software platforms to facilitate reconstructing genome-scale metabolic networks. Environmental Microbiology. 16(1): 49-59. doi:10.1111/1462-2920.12312.
- 2. **Hamilton JJ**, Dwivedi V[†], and JL Reed. (2013) Quantitative Assessment of Thermodynamic Constraints on the Solution Space of Genome-Scale Metabolic Models. Biophysical Journal. 105(2): 512-522. doi:10.1016/j.bpj.2013.06.011.
- 1. **Hamilton JJ** and JL Reed. (2012) *Identification of Functional Differences in Metabolic Networks Using Comparative Genomics and Constraint-Based Models.* PLoS ONE. 7(4): e34670. doi:10.1371/journal.pone.0034670.

† indicates an undergraduate student author

Oral Presentations

- 7. **Hamilton JJ**, Garcia SL, Brown BS, Oyserman BO, Moya F, Bertilsson S, Malmstrom RR, Forest KT, and KD McMahon. *Microbial Ecosystems Biology: A Framework for Scaling from Genomes to Ecosystems*. Association of Environmental Engineering and Science Professors Research and Education Conference. Ann Arbor, MI, June 2017.
- 6. **Hamilton JJ**. Diving Into Freshwater Microbial Genomes to Infer Bacterial Traits. Wisconsin Ecology 22nd Annual Fall Symposium. Madison, WI, October 2016.

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^{*} indicates equal contribution

- 5. **Hamilton JJ** and JL Reed. Thermodynamics and H₂ Transfer in a Methanogenic, Syntrophic Community. COBRA 2014: 3rd Conference on Constraint-Based Reconstruction and Analysis, Charlottesville, VA, May 2014.
- 4. Hamilton JJ and JL Reed. *Identification of Functional Differences in Cyanobacterial Metabolic Networks Using Constraint-Based Models*. American Society of Microbiology (ASM) General Meeting, Denver, CO, May 2013.
- 3. **Hamilton JJ**, Dwivedi V, and JL Reed. Thermodynamics-Based Flux-Balance Analysis: Incorporation of Thermodynamic and Metabolomic Data Into Genome-Scale Constraint-Based Models. American Institute of Chemical Engineers Annual Meeting, Pittsburgh, PA, October 2012.
- Hamilton JJ and JL Reed. Comparison of Microbial Metabolic Networks to Guide Background Strain Selection. American Chemical Society National Meeting, San Diego, CA, March 2012.
- 1. **Hamilton JJ** and JL Reed. Genome Alignment and Pair-Wise Comparison of Microbial Metabolic Networks Identifies Novel Potential Drug Targets. American Institute of Chemical Engineers Annual Meeting, Minneapolis, MN, October 2011.

Poster Presentations

- 5. **Hamilton JJ**, Clark RL, Groblewski ES, and OS Venturelli. A chemically-defined growth medium for a model human gut microbiome. Madison Microbiome Meeting (M3), Madison, WI. April 2018.
- 4. Hamilton JJ, Garcia SL, Bendall ML, Brown BS, Stevens SLR, Bertilsson S, Forest KT, Malmstrom RR, Stepanauskas R, Tringe SG, Woyke T, and KD McMahon. Metabolic Network Analysis and Metatranscriptomics of a Cosmopolitan and Streamlined Freshwater Lineage. 16th International Symposium on Microbial Ecology, Montreal, Canada. August 2016.
- 3. Hamilton JJ, Bendall M, Glavina del Rio T, Bertilsson S, Malmstrom R, Stepanauskas R, Tringe S, Woyke T, and KD McMahon. *Ecological Niches of Freshwater Actinobacteria Revealed through Comparative Genomics and Reverse Ecology*. Gordon Research Conference on Applied and Environmental Microbiology, Mount Holyoke College, South Hadley, MA. July 2015.
- 2. Hamilton JJ, He S, Bendall M, Glavina del Rio T, Bertilsson S, Malmstrom R, Stepanauskas R, Tringe S, Woyke T, and KD McMahon. *Ecological Niches of Uncultivated Freshwater Phyla Revealed Through Comparative Genomics*. UMX: 10th Annual Department of Energy Joint Genome Institute User Meeting, Walnut Creek, CA. March 2015.
- 1. **Hamilton JJ** and JL Reed. *Thermodynamics and H*₂ *Transfer in a Methanogenic, Syntrophic Community.* COBRA 2014: 3rd Conference on Constraint-Based Reconstruction and Analysis, Charlottesville, VA. May 2014.

Teaching Experience

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Critical Analyses in Microbiology **UW-Madison** InstructorSpring 2016 **Environmental Microbiology UW-Madison** Guest Lecturer Spring 2015, Spring 2017 Computational Tools for Analyzing Microbial Metabolism ASM General Meeting InstructorMay 2013 **Biochemical Engineering UW-Madison** Guest Lecturer Fall 2011, Fall 2012, Fall 2015 **Biochemical Engineering UW-Madison** Teaching Assistant Fall 2011 **Mass Transfer Operations UW-Madison** Teaching Assistant Spring 2011 **Organic Chemistry CWRU** $Supplemental\ Instructor$ Fall 2008 Chemistry of Materials **CWRU** Supplemental Instructor Fall 2007, Spring 2008, Spring 2009

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