Charles Pepe-Ranney

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Professional Preparation

B.S. Engineering (high honors) - Environmental Science Specialty, Colorado School of Mines 2006.

M.S. Environmental Engineering - Biotechnology and Environmental Microbiology Emphasis, Colorado School of Mines 2009.

PhD Environmental Science and Engineering Division, Colorado School of Mines 2012.

Relevant Experience

Proficient with Python, R, Latex, Bash scripting and Linux system administration. Experience with Perl, javascript, Post-greSQL, MySQL

Thorough understanding and fluent with many data science/bioinformatics libraries including IPython notebooks, GG-Plot2 (R), Matplotlib (Python), Pandas (Python), phyloseq (R), plyR/dplyR/tidyR (R), QIIME, Mothur, Khmer, and BioPython. Experience with Bokeh (Python), D3.js (my bl.ocks: bl.ocks.org/chuckpr), ggvis (R) and lattice (R).

Currently taking online courses for the **Data Science Signature Track** with Coursera – Passed and received verified certificate with distinction for The Data Scientist's Toolbox, R Programming, Exploratory Data Analysis, Statistical Inference, and Regression Models courses.

Experience developing amplicon sequencing protocols from the ground up for next-generation-sequencing technologies (454 and Illumina) with SSU rRNA genes and Fungal ITS amplicons.

Appointments

Research Assistant, Environmental Science and Engineering Division, Colorado School of Mines (2006-2012)

Postdoctoral Researcher, Laboratory of Daniel H Buckley, Department of Crop and Soil Sciences, Cornell University (2013-present)

Teaching Fellow, Marine Biology Laboratory (Microbial Diversity Course) (2010-2014)

Awards and Fellowships

2010, **2011**, **2012** and **2013** Teaching fellow for the Microbial Diversity Course at the Marine Biological Laboratory, Woods Hole. Course Directors: Daniel Buckley and Steve Zinder.

2006 Outstanding Graduating Senior Award, Colorado School of Mines - Environmental Science and Engineering Division

2005 and 2006 Department of Energy Science Undergraduate Laboratory Internship (SULI) at Idaho National Lab

2006 INL Undergraduate Scholarship

Publications in Refereed Journals

Pepe-Ranney C, Berelson WM, Corsetti FA, Treants M, Spear JR. Cyanobacterial construction of hot spring siliceous stromatolites in Yellowstone National Park, Wyoming, 2012, Environmental Microbiology 14(5), 1182-1197. link

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Berelson WM, Corsetti FA, Pepe-Ranney C, Hammond DE, Beaumont W, Spear JR. **Hot spring siliceous stromatolites in Yellowstone National Park: assessing growth rates and laminae formation**, 2011, *Geobiology* 9(5), 411-424. link

Osburn MR, Sessions AL, Pepe-Ranney C, Spear JR. Hydrogen-isotopic variability in fatty acids from Yellowstone National Park hot spring microbial communities, 2011, Geochimica et Cosmochimica Acta 75(17), 4830-4845. link

Bräuer S, Vuono D, Carmichael M, Pepe-Ranney C, Strom A, Rabinowitz E, Buckley DH, Zinder S. **Microbial sequencing analyses suggest the presence of a fecal veneer on indoor climbing wall holds.**, 2014, Current Microbiology link

Pepe-Ranney C, Koechli C, Potrafka R, Garcia-Pichel F, Andam C, Eggleston E, Buckley DH. **Non-cyanobacterial** diazotrophs mediate dinitrogen fixation in biological soil crusts during early crust formation.

Accepted by at ISMEJ, May 2015. Preprint:

http://dx.doi.org/10.1101/013813

Code for sequence analysis and manuscript figures can be found here:

github.com/chuckpr/NSIP_data_analysis

Submitted Journal Articles

Pepe-Ranney C and Hall EK. The effect of carbon subsidies on planktonic niche partitioning and recruitment during biofilm assembly.

In review at Frontiers of Aquatic Microbiology. Preprint:

http://dx.doi.org/10.1101/013938

Manuscript figures and corresponding code can be found here:

github.com/chuckpr/BvP_manuscript_figures/

Journal Articles in Preparation

Pepe-Ranney C* and Campbell A*, Koechli C, Berthrong S, Buckley DH. Charting the flow of carbon through a soil microbial community with high resolution DNA stable isotope probing.

*co-first authors

Code for manuscript figures can be found here:

nbviewer.ipython.org/github/chuckpr/CSIP_succession_data_analysis

Wallace B, Roberts A, Pollet R, Venkatesh M, Guthrie L, OâĂŹNeal S, Ingle J, Robinson S, Dollinger M, Figueroa E, McShane S, Jin J, Frye S, Zamboni W, Pepe-Ranney C, Mani S, Kelly L, and Redinbo M. **Structure and Inhibition of Firmicutes Bacterial b-Glucuronidases to Alleviate Drug-Induced GI Toxicity**Contributed figure: link

Pepe-Ranney C, Campbell A, Buckley DH. Community genomics of soil cellulose degraders discovered by nucleic acid stable isotope probing

Code for manuscript figures can be found here:

nbviewer.ipython.org/github/chuckpr/CG-SIP/tree/master/

Koechli C, Pepe-Ranney C, Campbell A, Buckley DH. Mapping carbon flow through both 16S rRNA and 16S rRNA genes from an agricultural soil using stable isotope probing provides insights into bacterial metabolism.

Hahn C, Hall EK, Pepe-Ranney C, Oyler-McCance S. Evaluating the gut and cloacal bacterial community of cowbirds: a potential mechanism for enhanced immunity.

Code for sequence analysis and manuscript figures can be found here:

nbviewer.ipython.org/github/chuckpr/cowbird

Invited Talks

¹⁴ C and microbial diversity study of Yellowstone siliceous stromatolites: searching for the depositional community. 2009. Microbiology Supergroup, University of Colorado - Boulder.

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Cyanobacterial construction of finely laminated siliceous stromatolites in a Yellowstone National Park hot spring. 2012. Astrobiology Science Conference - Microbes in Lithifying Systems.

Last updated: May 28, 2015