
CONTACT
INFORMATION

Location: Golden, CO 80401
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Phone: (508) 494-2474

EXPERIENCE

National Renewable Energy Lab (2015 - Present)

Postdoctoral Scholar

- Used machine learning techniques to develop a model to predict the tendency of a molecule to form soot from directly from its structure.
- Developed models of microbial metabolism to determine means of improving product yields through genetic engineering. Helped to develop and maintain python packages for such purposes, including *cobrapy* and *d3flux*.

EDUCATION

University of California, Santa Barbara (2010-2015)

Ph.D., Department of Chemical Engineering, Santa Barbara, California (GPA: 3.68)

- Thesis: Computational analysis of the mammalian circadian clock, with a focus on elucidating the functional design consequences of the underlying genetic regulatory network.

Tufts University (2006 - 2010)

BS, Chemical and Biological Engineering, Medford, Massachusetts (GPA: 3.79)

RESEARCH EXPERTISE

- *Machine Learning:* Neural networks, preprocessing methods, hyperparameter optimization
- *Statistics:* uncertainty analysis, bayesian methods, model selection
- *Optimization:* Linear programming, nonlinear programming, stochastic methods
- *Nonlinear systems:* Ordinary differential equations, collocation methods, sensitivity analysis

SOFTWARE EXPERTISE

- *Python:* thorough familiarity with the PyData stack, including relational databases (*pandas*), machine learning methods (*sklearn*), and compiled extensions (*cython*, *swig*, *numba*)
- *Development:* *unittests*, continuous integration, and helped to develop software for large open-source projects.
- Comfortable with unix environments, HPC, and front-end stack languages

SELECTED
PEER-REVIEWED
PUBLICATIONS

St. John, P. C., Crowley, M. F., & Bomble, Y. J. (2017). Efficient estimation of the maximum metabolic productivity of batch systems. *Biotechnology for Biofuels*, 10(1). doi:10.1186/s13068-017-0709-0

Abel, J.H., Meeker, K., Granados-Fuentes, D., **St. John, P.C.**, Wang, T.J., Bales, B.B., Doyle F.J. III, Herzog, E.D., and L.R. Petzold. Functional network inference of the suprachiasmatic nucleus (2016) *PNAS*, 113 (16) pp. 4512-4517

St. John, P.C. and F.J. Doyle III. Quantifying stochastic noise in cultured circadian reporter cells (2015), *PLoS Computational Biology* 11(11): e1004451.

St. John, P.C., Taylor, S.R., Abel, J.H., and F.J. Doyle III. Amplitude metrics for cellular circadian bioluminescence reporters (2014) *Biophysical Journal*, 107 (11) pp. 2712-2722

St. John, P.C., Hirota, T., Kay, S.A. and F.J. Doyle III. Spatiotemporal separation of PER and CRY posttranslational regulation in the mammalian circadian clock (2014) *PNAS*, 111 (5) pp. 2040-2045.

Hirota, T., Lee, J.W., **St. John, P.C.**, Sawa, M., Iwaisako, K., Noguchi, T., Pongsawakul, P.Y., Sonntag, T., Welsh, D.K., Brenner, D.A., Doyle, F.J. III, Schultz, P.G., Kay, S.A., Identification of small molecule activators of cryptochrome (2012) *Science*, 337 (6098) pp. 1094-1097.

ADDITIONAL
INFORMATION

Website: <http://www.nrel.gov/bioenergy/bios/peter-stjohn.html>
Google Scholar: <https://scholar.google.com/citations?user=NdWzcVMAAA>
Github: <https://github.com/pstjohn>