

CONTACT INFORMATION	<p><i>Location:</i> Golden, CO 80401</p> <p><i>Email:</i> peter.stjohn@nrel.gov</p> <p><i>Phone:</i> (508) 494-2474</p>
EXPERIENCE	<p>National Renewable Energy Lab (2015 - Present)</p> <p><i>Postdoctoral Scholar</i></p> <ul style="list-style-type: none"> • 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	<p>University of California, Santa Barbara (2010-2015)</p> <p><i>Ph.D., Department of Chemical Engineering, Santa Barbara, California (GPA: 3.68)</i></p> <ul style="list-style-type: none"> • Thesis: Computational analysis of the mammalian circadian clock, with a focus on elucidating the functional design consequences of the underlying genetic regulatory network. <p>Tufts University (2006 - 2010)</p> <p><i>BS, Chemical and Biological Engineering, Medford, Massachusetts (GPA: 3.79)</i></p>
RESEARCH EXPERTISE	<ul style="list-style-type: none"> • <i>Machine Learning:</i> Neural networks, preprocessing methods, hyperparameter optimization • <i>Statistics:</i> uncertainty analysis, bayesian methods, model selection • <i>Optimization:</i> Linear programming, nonlinear programming, stochastic methods • <i>Nonlinear systems:</i> Ordinary differential equations, collocation methods, sensitivity analysis
SOFTWARE EXPERTISE	<ul style="list-style-type: none"> • <i>Python:</i> thorough familiarity with the PyData stack, including relational databases (pandas), machine learning methods (sklearn), and compiled extensions (cython, swig, numba) • <i>Development:</i> 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	<p>St. John, P. C., Crowley, M. F., & Bomble, Y. J. (2017). Efficient estimation of the maximum metabolic productivity of batch systems. <i>Biotechnology for Biofuels</i>, 10(1). doi:10.1186/s13068-017-0709-0</p> <p>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) <i>PNAS</i>, 113 (16) pp. 4512-4517</p> <p>St. John, P.C. and F.J. Doyle III. Quantifying stochastic noise in cultured circadian reporter cells (2015), <i>PLoS Computational Biology</i> 11(11): e1004451.</p> <p>St. John, P.C., Taylor, S.R., Abel, J.H., and F.J. Doyle III. Amplitude metrics for cellular circadian bioluminescence reporters (2014) <i>Biophysical Journal</i>, 107 (11) pp. 2712-2722</p> <p>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) <i>PNAS</i>, 111 (5) pp. 2040-2045.</p> <p>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) <i>Science</i>, 337 (6098) pp. 1094-1097.</p>
ADDITIONAL INFORMATION	<p><i>Website:</i> http://www.nrel.gov/bioenergy/bios/peter-stjohn.html</p> <p><i>Google Scholar:</i> https://scholar.google.com/citations?user=NdWzcVMAAAAJ</p> <p><i>Github:</i> https://github.com/pstjohn</p>