

Rachael M. Cox

2500 Speedway, MBB 3.128B, Austin, TX 78705

email: rachaelcox@utexas.edu

SUMMARY

Highly interdisciplinary and well-rounded computational biochemist with a passion for chemistry, systems biology, machine learning and large-scale data analysis; aspires to leverage comparative evolutionary frameworks to inform technology development, with current projects having applications to agricultural engineering and human health.

EDUCATION

Doctor of Philosophy, Biochemistry

The University of Texas at Austin, Austin, Texas

Jun. 2018 – present

Bachelor of Science, Chemistry

Texas A&M University, College Station, Texas

Aug. 2009 – Dec. 2013

RESEARCH EXPERIENCE

Graduate Research Assistant

Prof. Edward Marcotte, The University of Texas at Austin

Jun. 2018 – present

- Validated machine learning predictions for plant protein complexes using gene knock-out and “guilt-by-association” phenotype predictions
- Use machine learning and statistical models for comparative evolution of protein organization across multicellular organisms
- Lead bioinformatician for engineering non-model plants (e.g., dandelions) to elicit a specific response, given a specific input, by integrating comparative proteomic and transcriptomic approaches
- Developed a novel de Bruijn graph-based algorithm for multiple sequence alignment; the algorithm scales linearly with input size and detects evolutionarily conserved sequence motifs and clusters

Research Technician

Eastman Chemical Company

Mar. 2015 – Jun. 2018

- Evaluated new inorganic catalysts to make a process more efficient or environmentally friendly using reaction kinetics and yield as metrics
- Researched new chemistry to anticipate market demands (e.g., designing a less volatile solvent that functions as well as existing products), while working in conjunction with plant engineers to tune reaction conditions to ensure the new product would be realistic given existing manufacturing infrastructure
- Designed and constructed continuous reaction systems at various scales (i.e., 1g/min to 1kg/min process rates) that model existing industrial-scale processes to generate reliable data for commercialization of new processes

RESEARCH EXPERIENCE (cont.)

Research Technician

Sept. 2014 – Mar. 2015

Prof. Deepak Nijhawan, University of Texas Southwestern Medical Center

- Pursued drug discovery and characterization using mammalian cell tissue culture techniques in conjunction with bio-orthogonal click chemistry and immunohistochemical experiments
- Identified drug targets using streptavidin-biotin affinity purification, immunoprecipitation, silver staining and LC-MS
- Evaluated synthetic lethality with drug titrations and luminescent cell viability assays to generate dose-response curves

Research Technician (Undergraduate)

May 2013 - Jun. 2014

Prof. David Barondeau, Texas A&M University

- Evaluated mechanistically significant residues on the surface of the human ISC protein frataxin
- Helped design, optimize and perform fluorescence anisotropy experiments for characterizing binding kinetics associated with FXN and the NFS1-ISD11-ISCU2 complex in the human ISC system
- Processed kinetic data and computed binding constants
- Trained in biosynthetic protocols such as recombinant protein expression (DNA transfection, PCR cloning and sequencing) and purification (FPLC affinity chromatography, aerobic and anaerobic)

SKILLS

- **Computational:** Python for scripting and machine learning; R for statistical analysis and data visualization; Bash for scripting, string manipulation and large-scale data management; proficient in a Linux environment
- **Organic and biochemistry:** Organic synthesis and inorganic catalysis; small and large molecule chromatography (i.e., gas and liquid chromatography); recombinant protein expression and purification; gel electrophoresis
- **Molecular and synthetic biology:** Mammalian, yeast and microbial tissue culture; plasmid engineering, Sanger and Illumina sequencing and analysis
- **Instrumentation:** Mass spectrometry, NMR ($^1\text{H}/^{13}\text{C}/^{32}\text{P}$), FTIR spectrometry, UV-Vis and fluorescence spectroscopy

PUBLICATIONS

- Lee, C.; Cox, R. M.; Papoulas, O.; Horani, A.; Drew, K.; Devitt, C. C.; Brody, S. L.; Marcotte, E. M.; Wallingford, J. B. "Functional partitioning of a liquid-like organelle during assembly of axonemal dyneins." <http://biorxiv.org/lookup/doi/10.1101/2020.04.21.052837> (2020)

- Drew, K.; Lee, C.; **Cox, R. M.**; Dang, V.; Devitt, C. C.; Papoulas, O.; Huizar, R. L.; Marcotte, E. M.; Wallingford, J. B. “A systematic label-free method for identifying RNA-associated proteins *in vivo* provides insights into vertebrate ciliary beating.” <http://biorxiv.org/lookup/doi/10.1101/2020.02.26.966754> (2020)
- McWhite, C. D.; Papoulas, O.; Drew, K.; **Cox, R. M.**; Oldfield-June, V.; Dong, O. X.; Kwon, T.; Wan, C.; Salmi, M. L.; Emili, A.; Roux, S. J.; Chen, J. Z.; Browning, K. S.; Ronald, P. C.; Marcotte, E. M. “Deep conservation and interaction-to-phenotype in a pan-plant protein complex map” *Cell* **181**, 460-474.e14 (2020).
- Vranish, J.; Russell, W.; Yu, L. E.; **Cox, R. M.**; Russell, D. H.; Barondeau, D. P. “Fluorescent probes for Tracking metal the Transfer of Iron-Sulfur Cluster and other Metal Cofactors in Biosynthetic Reaction Pathways” *J. Am. Chem. Soc.* **137**, 390–398 (2015).

TALKS & PRESENTATIONS

- **Cox, R.M.** “Engineering endogenous promoters for non-model plant species.” Biology Talks About Computers, Organisms and Systems (BioTACOS) (**February 2020**)
- **Cox, R.M.** “Predicting Legendary Pokemon Using a Random Forest Algorithm (a Machine Learning Demo)” UT Austin Open Coding Hour, Special Edition (**December 2019**, https://github.com/rachaelcox/pokemon_machine_learning_demo)
- **Cox, R.M.** “Falling Forward: Lessons in Failure” UT Austin Chips and Dip Seminar Series (**December 2019**)
- **Cox, R.M.** “Leveraging comparative proteomics for health and engineering” UT Austin Chips and Dip Seminar Series (**November 2019**)
- **Cox, R. M.** “The role of systems biology in health and engineering” They Blinded Me With Science Radio Show 91.7HD2 (**October 2019**)
- **Cox, R. M.** “Protein lightning: multi-species sequence alignments visualized via de Bruijn graphs” UT Austin College of Natural Science’s Art in Science Event (**October 2019 – Poster**)
- **Cox, R. M.**; McWhite, C. D.; Dinkeloo, K. D.; Tran, J.; Lloyd, A.; Qiao, H.; Ellington, A.; Marcotte, E. M. “Leveraging orthology and systems biology for engineering in non-model organisms” UT Austin Institute for Cell and Molecular Biology Annual Retreat (**September 2019 – Poster**)
- **Cox, R. M.**; McWhite, C. D.; Dinkeloo, K. D.; Tran, J.; Lloyd, A.; Qiao, H.; Ellington, A.; Marcotte, E. M. “Leveraging orthology and systems biology for engineering in non-model organisms” UT Austin Synthetic Biology Symposium (**August 2019 – Poster**)

TEACHING

- Teaching Assistant
 - **Computational Biology and Bioinformatics**; UT Austin (Spring 2020)
- Lead instructor
 - **Practical Approaches to Analyzing Biological Data in R**; Big Data in Biology Summer School, Center for Biomedical Research Support (Summer 2020)

ORGANIZATIONS & OUTREACH

Open Coding Hour

Jan. 2019 – present

- In charge of organizing and scheduling a weekly help session designed to help students with data science and programming-related problems
- Spearheaded a new monthly program where computational scientists demo a useful programming skill (e.g., programmatic figure making, machine learning models)

Graduate Student Association

Jun. 2019 – present

- First Year Student Liaison (2019-2020)
- Vice President (2020-present)

Junior League of Longview

Dec. 2016 – Jun. 2018

- Project coordinator for the program “Buy a Box, Feed a Child”
- This project raised more than \$10,000 towards filling “School Break” boxes with food for children who usually depend on school-provided meals; my group converted that money to food supplies and delivered the organized boxes to East Texas schools
- Awarded “Provisional of the Year” for this effort

REFERENCES

- Dr. Edward Marcotte (**email:** edward.marcotte@gmail.com)
- Dr. Andrew Ellington (**email:** ellingtonlab@gmail.com)
- Dr. Claus Wilke (**email:** wilke@austin.utexas.edu)
- Dr. Damon Billodeaux (**email:** dbillodeaux@eastman.com)
- Dr. David Barondeau (**email:** barondeau@chem.tamu.edu)