SEAN HACKETT

Summary of Qualifications:

- Actively plans and directs long-term strategy, near-term tactics, and day-to-day operations.
- Leverages statistics and programming to tackle challenging scientific problems. Google Scholar
- Excels at communicating with diverse audiences in both written and oral formats.

Skills: Analytics: statistics (GLMs, nonlinear regression, maximum likelihood, model comparison, Bayesian statistics), ML (LASSO, random forest, LDA, LLMs), optimization (linear/quadratic programming, gradient-based MAP estimation). Programming R (dplyr, purrr, ggplot2, devtools, Shiny, Connect), Python (pandas, Jupyter, igraph, package development) Tools Building and querying databases (SQL, graphQL, BigQuery), Docker, WDL, GCP, CI/CD with GitHub Actions.

| EDUCATION | Princeton University Ph.D., Quantitative and Computational Biology DOE Office of Science Graduate Fellowship (SCGF) | Princeton, NJ 2015 |
|----------------|---|-----------------------|
| | Cornell University B.S., Biological Sciences Magna Cum Laude with Distinction in Research | Ithaca, NY 2006 |
| RESEARCH | | |
| DIRECTOR | Calico Life Sciences LLC | S. San Francisco, CA |
| | Formed the Discovery Data Science team to balance collaborations on basic science programs with focal support and methods development. Established data systems to support the key outputs of the Discovery organization: mechanisms and targets. | Jan 2023 - Present |
| Assoc Director | • Collaborated with key stakeholders to develop long-term strategies for causal inference and systems biology. | Jan 2021 - Jan 2023 |
| MANAGER | Helped reorganize the Computing team to improve impact, collaboration, and accountability. Managed 4-6 data scientists, prioritizing high value projects in a problem-rich environment. Led initiatives around computational education, results sharing, and de-duplication of efforts. | Feb 2018 - Jan 2021 |
| Data Scientist | Created a genome-scale mechanistic network connected to known gene-disease associations to support indication prioritization. Improved approaches for finding causal connections in high-dimensional time series using a combination of parametric modeling and LASSO. Developed an automated metabolomics pipeline to streamline data normalization and compound identification. | Jan 2017 - Present |
| POSTDOCTORAL | Princeton University, Lewis-Sigler Institute | Princeton, NJ |
| ASSOCIATE | Supervisor: John Storey, Director of the Center for Statistics and ML Used Latent Dirichlet Allocation with Empirical Bayes priors to identify latent variables affecting sparse high-dimensional data. | 2015 - 2017 |
| GRADUATE | Princeton University, Quantitative and Computational Biology | Princeton, NJ |
| FELLOW | Adviser: Josh Rabinowitz, Professor of Chemistry and Genomics Supervised two systems biology graduate students. Developed a scalable algorithm for combining metabolomics, proteomics and fluxes to identify novel allosteric regulators and dissect how metabolite and enzyme concentrations jointly control metabolism. | 2010 - 2015 |