

Portland, OR

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Summary_

Currently an Associate Data Scientist at Vacasa in Portland, OR.

Skills_

Programming Python, PostgreSQL, Git, Bash

Data Science Pandas, Numpy, Jupyter, Data Visuzaliation, Feature Engineering

Other Sphinx Documentation, Statistics

Experience _____

Vacasa Portland, OR

ASSOCIATE DATA SCIENTIST

Jul 2018 - PRESENT

• Working with software engineers to create machine learning models to help with business needs.

JUNIOR ANALYST Sep 2017 - Jul 2018

• Continuing work from Intern projects

ANALYST INTERN Jun 2017 - Sep 2017

• Building training set for machine learning model to predict owner churn, identifying gaps in owner data, and identifying indicators to help determine at risk owners.

University of Oregon - Institute of Molecular Biology

Eugene, OR

BIOINFORMATICIST - SELKER LAB

Oct 2016 - May 2017

- Updated and managed scripts for analysis of ChIP-seq and RNA-seq data.
- Compared ChIP-seq datasets looking at differences in RNA pol II binding between different strains of n. crassa. Done using various data analysis programs and automated using R and bash scripting.

SCIENTIFIC PROGRAMMER - HARMS LAB

Sep 2015 - May 2017

- Developed a GUI using PyQt5 for python API that analyses isothermal calorimetry (ITC) data. Along with this, wrote documentation for GUI using sphinx on Read the Docs as well as a C extension to calculate the binding polynomial in the API.

 Repo: https://github.com/harmslab/pytc-gui and https://github.com/harmslab/pytc
- Developed an extension of a phage display analysis pipeline. Researched and implemented different methods of data clustering. Repo: https://github.com/harmslab/phagedisplay

Publications

HIRANMAYI DUVVURI, LUCAS C. WHEELER, AND MICHAEL J. HARMS

Biochemistry

PYTC: OPEN-SOURCE PYTHON SOFTWARE FOR GLOBAL ANALYSES OF ISOTHERMAL TITRATION CALORIMETRY DATA

2018

Education

University of Oregon Eugene, OR

B.S. IN BIOCHEMISTRY

Sep 2011 - Sep 2016

· focus on biology, chemistry, and computer science