

Pablo G. Romano

DATA SCIENTIST

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Summary

Data Scientist, PhD Physical Chemistry, Texan

Skills

Programming	Python, Fortran, R, Matlab, Cython SQL, Bash
Data Analysis Tools	NumPy, Scikit-Learn, SciPy, Pandas, spaCy, NLTK, Matplotlib, ggplot2
Statistical Methods	Markov Chains, Mixture Models, Regression (Linear and Logistic), kNN
Unsupervised Learning	Clustering (K-Means, K-Medoids, Fuzzy C-Means, and Ward), PCA, Time-lagged ICA
Supervised Learning	Random Forest, Naive Bayes, and SVM Classification
Natural Language Processing	Topic Modeling (NMF, LDA), Text Processing, tf-idf, word2vec, doc2vec

Research Experience

Insight Data Science

Remote

FELLOW

Jan. 2018 - Present

- Implemented a rating system to score dogs based on the similarity between searched personalities and pet description using NMF topic modeling and word2vec.
- Trained and validated Naïve Bayes classifier to predict whether sentences described pets for adoption.
- Queried, processed, and stored pet-descriptions from Petfinder's RESTful API using NLTK, spaCy, pandas, and PostgreSQL.
- Built interactive front-end web-app to select features in search queries using Bootstrap and Flask.

University of Oregon

Eugene, Oregon

GRADUATE RESEARCH FELLOW

Sep. 2012 - Sep. 2017

- Predicted macroscopic conformational states from 100+ GB DNA simulations using k-means and Ward clustering.
- Evaluated kinetic mechanisms in DNA un-stacking by modeling inter-state transitions as a Markov chain and analyzing through transition path theory.
- Estimated relaxation timescales for DNA un-stacking by spectral decomposition of transition likelihoods.
- Coarse-grained atomistic representation from 1000+ atoms to <10 features using PCA and time-lagged ICA.
- Designed and implemented "Gradient Adaptive Decomposition" algorithm, improving predictive accuracy of kinetic classification by over 20% using NumPy, SciPy, and Scikit-learn.
- Facilitated training of incoming undergraduate and graduate students in Bash shell, coding with Python and Fortran, and molecular dynamics simulations.

St. Edward's University

Austin, Texas

RESEARCH ASSISTANT

Jan. 2009 - Jun. 2012

- Collaborated between synthetic and computational groups in the design of novel inhibitors targeting precursors to Rheumatoid Arthritis.
- Validated the efficacy of inhibitors through docking studies to reduce the number of synthesized compounds from 1000+ to 100.

Education

Sep. 2012 - Sep. 2017 **Physical Chemistry, PhD**

University Of Oregon

Aug. 2008 - May. 2012 **Chemistry, BS**

St. Edward's University

Projects

LaTeX CV 📄

Modification of the posquit0 Awesome CV template. Includes more classes, and modified design. They made a great template!

Publications

The Gradient Adaptive Decomposition (GRAD) Method: Optimized Refinement Along Macrostate Borders in Markov State Models

PG ROMANO, MG GUENZA

J. Chem. Info. & Mod.

Acetylenic inhibitors of ADAM10 and ADAM17: In silico analysis of potency and selectivity

EF HEALY, PG ROMANO, M MEJIA, G LINDFORS III

J. Mol. Graph. & Model.

Presentation & Workshops

Chiral Pathways in DNA dinucleotides using gradient optimized refinement along metastable borders

New Orleans, Louisiana

NATIONAL AMERICAN PHYSICAL SOCIETY MEETING

Mar. 2017

Presented novel method for improving accuracy by greater than 20% in Markov models of macromolecular simulation data.

CECAM Macromolecular Simulation Software Workshop

Jülich, Germany

CENTRE EUROPÉEN DE CALCUL ATOMIQUE ET MOLÉCULAIRE

Oct. 2015

Attended workshop series designed to improve computational methods and software design aimed at open and reproducible science.

Analyzing Small DNA Constructs via a Chromophore Model within the Point Dipole Approximation.

Baltimore, Maryland

BIOPHYSICAL SOCIETY: NATIONAL MEETING

Mar. 2015

Presented structural model of organic chromophores to reproduce, with 95% confidence, experimental results of Adenine dinucleotides.

Acetylenic inhibitors of ADAM10 and ADAM17: In silico analysis of potency and selectivity

New Orleans, Louisiana

AMERICAN CHEMICAL SOCIETY, SOUTHWEST REGIONAL MEETING

Sep. 2012

Presented talk on the analysis of novel acetylenic TACE inhibitors.