Yeonji Ji, Ph.D.

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Summary

Data Scientist with a Ph.D. and experience in statistical modeling, large-scale data analysis, and machine learning (Python, SQL, Py-Torch). Skilled in building scalable pipelines and delivering insights that drive product decisions.

Technical Skills

Programming:

Python (pandas, scikit-learn, PyTorch, matplotlib, seaborn, etc.), Bash, Git, C++

Machine Learning:

Supervised/unsupervised learning, Feature Engineering, Hyperparameter Tuning, Model Evaluation, Ensembles, Deep Learning

Computing:

Linux, HPC (Slurm), Workflow Automation

Specialized Expertise:

Computational Drug Discovery, Molecular Simulations, Virtual Screening, Molecular Visualization

Education

Ph.D. in Biochemistry

The Graduate Center, CUNY 2018 – 2024 Dissertation Title:

"Incorporating Solvation Thermodynamic Mapping in Computer-Aided Drug Design"

B.S. in Chemistry

Kyung Hee University, Seoul 2012 – 2017

Certifications

Python for Machine Learning and Data **Science Masterclass (Udemy)**

Supervised/unsupervised ML, PCA, model evaluation

Deep Learning Specialization (Coursera)

Neural networks (CNN, RNN, LSTM, etc.), optimization, project structuring

Honors & Grants

CUNY DSRG (2023)

Penny J. Gilmer Grant, OpenEye (2023) CUNY Science Scholarship (2018-2024) Superiority Scholarship, KHU (2014–2015)

Machine Learning Projects

Amazon Review Sentiment Classification (GitHub)

2025

- Built a text classification model using TF-IDF and ML algorithms (Logistic Regression, XGBoost).
- · Conducted feature engineering, hyperparameter tuning, and evaluation on 3M+ reviews.

Teleco Customer Churn Prediction (GitHub)

2025

- Developed churn prediction models using Logistic Regression, Random Forest, and XGBoost.
- Applied SMOTE and feature engineering on 7k+ customer records to address class imbalance and improve prediction accuracy.

Research & Data Projects

Binding Site Prediction from Simulation Data (Publication)

- Processed large-scale molecular dynamics datasets, analyzing timeseries trajectories and 3D solvation data to identify cryptic pockets.
- Used statistical models to quantify solvation energy costs and assess pocket binding potential.

Data-driven Pharmacophore Modeling (Publication in process)

- Engineered automated Python workflows to extract hydration features and construct water-derived pharmacophore models.
- Evaluated performance on large compound libraries, enhancing screening reproducibility and scalability.

COVID-19 Solvation Mapping Repository (Publication)

 Contributed analysis tools and datasets to an open-source GitHub repository, leading to a peer-reviewed publication.

Professional Experience

Postdoctoral Researcher

Lehman College, CUNY 2024 – Present

- Leading independent research on binding site thermodynamics and molecular simulations.
- Developing Python workflows and preparing manuscripts; mentoring graduate students and collaborating across the lab.

Adjunct Lecturer

CUNY Research Foundation 2019 – 2024

 Taught General Chemistry Lab to 100+ undergraduates, simplifying complex concepts through hands-on instruction.

Cosmetic Chemist Intern

Englewood Lab, NJ 2015 – 2016

• Conducted R&D experiments to optimize cosmetic formulations, producing samples that met stability and performance standards.