Bomin (David) Wei

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EDUCATION

University of California, Los Angeles, Los Angeles, CA

Sep. 2023 – Jun. 2027

Freshmen, Linguistic and Computer Science

Core Courses Taken:

AP Calculus BC (5), AP Computer Science A (5), Linear Algebra, Multivariable Calculus, Math 32A, CS 31

RESEARCH & INTERSHIP EXPERIENCE

Deep Learning-based Model for Drug Repurposing

Mar. 2021 - Oct. 2022

University of Utah

- Designed an embedding method based on Word2Vec to obtain complete drugs and proteins representation
- Designed CNN + LSTM Deep Neural Network to extract non-local features for binding interactions
- Developed **novel testing methods** with special data splitting to evaluate models' performance in real-world applications; and automated data cleaning and pre-processing of 2 million text string data; optimized the **classification criteria** for better model selection by **statistical analysis** on data distributions.
- Presented in professional conferences Intelligent Systems for Molecular Biology (ISMB, July. 2022) and the IDWeek (Oct. 2022); and published (first author) on Scientific Reports.
- https://github.com/David-BominWei/DeepLPI

SARS-COV-2 Genetic Mutation Modeling Prediction

Oct. 2020 - Mar. 2021

Hong Kong University

- · Modeled and visualized the SARS-CoV-2 mutation based on global geographical regions and time
- Used ARIMA model to analyze SARS-CoV-2 RNA sequence mutations as time series and predicted most likely
 mutation sites for different regions.

Machine Learning Algorithm (NLP Group) Intern

Jul. 2023 - Sep. 2023

4Paradigm, Beijing

- Compared and analyzed existing passage retrieval methods, such as BM25, using an efficient self-designed dataset; compiled query similarities datasets with a focus on banking question answering and integrated difficult negative samples; designed and finetuned a new query similarity model now used by the group.
- Finetuned and evaluated the basic model and found a problem that cannot distinguish proper nouns and extract important words. Finetuned the embedding module to reached better model result.

INTERESTS PROJECTS

Volunteer Management & Information Platform

Sep. 2020 - Dec. 2022

- Conducted assessment for Little Oaks Charity Center, leading to a customized volunteer management system.
- Achieved 1,614 users and 320 project registrations within a year, significantly improving operational efficiency.

Q2Q Dataset (https://github.com/David-BominWei/Q2QDataset)

Jul. 2023 - Aug. 2023

- Created an open-source Chinese query similarity dataset to train question-answering models.
- Utilized the BM25 model to construct negative samples for training.

Personal Blog Designer and Developer

Sep. 2020 – Sep. 2023

• Developed a blog platform using Hexo and deployed it on GitHub Pages.

Machine Learning for Movie Recommendation

Mar. 2023 – Jun. 2023

• Built a movie recommendation engine using inception networks, compared with a bag-of-words model.

Computational Graphics

Mar. 2023 – Jun. 2023

Developed a 3D GUI for matrix projection using Pygame, applying object-oriented programming.

SKILLS

- Programming Languages: Python (PyTorch, Scikit-learn, Pandas), C++, Java, R
- Frameworks: LSTM, ResNet, Transformer, CNN, MLP; Hexo, WordPress
- Software: Fusion360, OnShape, AutoCAD, Premiere Pro

HONORS AND AWARDS

•	First Place in the Computer Science category and Air Force Research Laboratory Award,	
	Mercer Science and Engineering Fair (ISEF affiliated)	2023
•	Silver Medal, ST. Yau High School Science Award, USA Regional; ranked 2nd in CS category	2022
•	1st Place & Best Poster (in Biology and Medicine) at IEEE-ISEC 2021 Conference	2021
•	Gold Division, United States of America Computing Olympiad (USACO)	2022
•	1st Place Kaggle Science Olympiad National Invitational 2022 Machine Learning Event	2022
•	6th place in total & 3rd place in AI round in the hackathon CMU Info & Math Competition (CMIMC)	2022

PUBLICATIONS

- 1. Wei, B., et al. "DeepLPI: A Deep Learning Model for Drug Repurposing." Sci Rep 12, 18200 (2022) https://www.nature.com/articles/s41598-022-23014-1
- 2. Wei, B., et al. "Modeling SARS-CoV-2 Mutations Based on Geography and Time." bioRxiv (2021) https://www.biorxiv.org/content/10.1101/2021.08.11.455941