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Education

Tsinghua University, B.Sc. Computer Science and Engineering

Sep. 2021 - Present

- GPA: **3.93/4.0 (19/178)**; TOFEL: **119**; GRE: **336+5**
- Relevant Coursework: Fundamentals of Engineering Biology (A+), Fundamentals of Programming (A+), Theory of Computer Network (A+), Computational Biology (A), Introduction to Artificial Intelligence (A), Introduction to Machine Learning (A), Artificial Neural Network (A), Data Mining (A)

Research Experience

Carnegie Mellon University Research Assistant, Advisor: Prof. Jian Ma

Jun. 2024 - present

- Tissue Generator: Predicting Missing Tissue Regions in Spatial Transcriptomics
 - Developed a Variational Autoencoder (VAE) model to embed single cell gene expression profiles.
 - Developed a Vision Transformer model to generate spatial transcriptomics data for missing tissue regions.

BioMap Inc. Research Intern, Advisors: Prof. Le Song & Prof. Jie Tang

Aug. 2023 - Jun. 2024

- Protein Language Foundation Models
 - Discovered multiple forms of scaling law for different downstream tasks.
 - Enhanced model generative performance by 300% with supervised fine-tuning and reinforced self-training.
- Protein Multiple Sequence Alignment (MSA) Generation
 - Designed a simple yet effective MSA generation framework to aid protein structure prediction in low-MSA regime, achieving state-of-the-art performance.
 - Further enhanced model performance by leveraging feedback from AlphaFold2 for reinforcement learning.

Tsinghua Knowledge Engineering Group Advisors: Prof. Yuxiao Dong & Prof. Jie Tang

Nov. 2022 - May. 2023

- · Molecule Generation with Diffusion Model
 - Designed a mixed training strategy utilizing unlabeled protein and molecule for data augmentation.
 - Developed a Graph Neural Network (GNN) model in python for pocket-ligand pair affinity prediction.
 - Enhanced conditional generation performance by 13% using classifier guidance.

Tsinghua Machine Learning and Computational Biology Group Advisor: Prof. Jianyang Zeng Jun. 2022 - Dec. 2022

- Gene Selection and Drug Screening Based on Knowledge Map and Community Detection
 - Aligned embeddings from biological knowledge graphs with orthogonal gene and drug datasets for data integration.
 - Designed algorithms for community detection on heterogeneous knowledge graph.

Institute of Biophysics, Chinese Academy of Sciences Advisor: Prof. Pingsheng Liu

Jan. 2019 - Sep. 2020

- · Blocking the Binding of Bacterial Lipid Droplets to Phage DNA
 - Discovered an effective blocker for MLDS protein and viral DNA binding through in vitro experiments.
 - Acquired expertise in laboratory techniques, including confocal microscopy imaging, spectrophotometry, cell culturing and Western blotting.

Publication

- Bo Chen*, **Zhilei Bei***, Xingyi Cheng, Pan Li, Jie Tang, Le Song. "MSAGPT: Neural Prompting for Protein Structure Prediction via MSA Generative Pre-Training." In Proceedings of the 38th Annual Conference on Neural Information Processing Systems (NeurIPS), preprint at arXiv: 2406.05347, 2024.
- Bo Chen, Xingyi Cheng, Pan Li, Yangli-ao Geng, Jing Gong, Shen Li, **Zhilei Bei**, et al. "xTrimoPGLM: Unified 100B-Scale Pre-Trained Transformer for Deciphering the Language of Proteins." Acceptance in Principle at Nature Methods, preprint at arXiv: 2401.06199, 2024.

Honors & Awards

2023 **Comprehensive Scholarship**, Department of Computer Science and Technology, Tsinghua University *Beijing, China* 2022 **Comprehensive Scholarship**, Department of Computer Science and Technology, Tsinghua University *Beijing, China* 2020 **Finalist**, International Science and Engineering Fair (no specific awards announced due to COVID-19) *Virtual*

Skills

Programming Languages Python (Pytorch), C/C++, MATLAB, Rust, Javascript, HTML, CSS, SQL, x86 assembly **Tools & Frameworks** Git, Django, MySQL, React

November 7, 2024 Zhilei Bei