Yi 7hou

Education

- 2021 2024 **Sichuan University**, M.S. in Computer Science and Technology, *GPA*: 3.76/4. Advisor: Prof. Min Zhu, Lab: Vision Computing Lab
- 2017 2021 Sichuan University, B.E. in Computer Science and Technology, GPA: 3.77/4. Under the Wu Yuzhang Honors program

Research Interests

Al4Science, Graph Machine Learning & Data Mining, Trustworthy Al

Publications & Preprints

- J Journal, C Conference, P Preprint; UR Under Review
- J1 Yi Zhou, Xinyi Wang, Lin Yao, Min Zhu. "LDAformer: Predicting LncRNA-Disease Associations based on Topological Feature Extraction and Transformer Encoder". Briefings in Bioinformatics (**BIB**), 2022. (JCR-Q1, IF: 13.994)
- J2 Wenwen Gao, Shangsong Liu, Yi Zhou, Fengjie Wang, Feng Zhou, Min Zhu. "GBDT4CTRVis: Visual Analytics of Gradient Boosting Decision Tree for Advertisement Click-Through Rate Prediction". Journal of Visualization (JoV), 2024.
- J3 Jiamin Zhu, Meixuan Wu, Yi Zhou, Nan Cao, Haotian Zhu, Min Zhu. "Dowsing: A Task-Driven Approach for Multiple-View Visualizations Dynamic Recommendation". Journal of Visualization (**JoV**), 2024.
- P1 Yi Zhou, Meixuan Wu, Chengzhou Ouyang, Min Zhu. "Generalizable and Explainable Prediction of Potential MiRNA-Disease Association based on Heterogeneous Graph Learning".

Research Experience

Jan 2021 - Research in Link Prediction on Biomedical Interaction Graph, Project "Visual Analysis Jun 2024 of Heterogeneous Graph for Disease-Regulatory Factor", supported by the General Program of National Natural Science Foundation of China (Grant No.62172289).

- Wrote the content about link prediction methods for disease-regulator factor in the proposal.
- Proposed a Link Prediction method for IncRNA-disease that outperformes sota baseline methods. It extracts multi-hop pathways between node pairs from similarity and association information and utilizes a Transformer encoder to learn the interdependencies among pathways. [J1, <u>D01</u>, PT2 (USE)]
- Developed a visual analytic system with collaborators, which displays the heterogeneous graph and explains heuristic similarities and logistic regression-based predictions. [Online Demo]
- Proposed a Link Prediction method for miRNA-disease that focuses more on Generalizability and Explainability. Constructed a miRNA-PCG-disease graph, proposed a new benchmark for generalizability, developed a prediction framework that combines pre-trained language models and heterogeneous graph neural networks, and designed a specialized scheme of explanation. [P1, arXiv, GitHub]
- Jan 2022 Research in Visualization Recommendation.
- May 2022 Implemented an LSTM-based model to perceive users' analytical tasks from historical actions.
 - Developed a recommendation system for multiple-view visualizations with collaborators, which exposes and utilizes users' potential analysis tasks. [J3, PT4 (USE), PT5 (USE), Webpage, Online Demo]

- Jan 2023 Research in Visualizational Explanation of GBDT.
 - Apr 2023 Implemented a demo Gradient Boosting Decision Tree (GBDT) model for click-through rate prediction.
 - Developed a visual analytic system with collaborators, which can assist advertising analysts in understanding the working mechanism of GBDT and facilitating the tuning process. [J2, <u>Video</u>]
- Dec. 2020 Research in Visualization of Chromatin Structure, Project "Platform for Visual Analysis of
 - Sep. 2021 Chromatin Multi-Level Structures and Gene Regulation Relationships", supported by Chengdu Science and Technology Program (Grant No.2021-YF05-02071-SN).
 - Wrote the project proposal. Developed a visual analytic platform with a function for topological association domain prediction. [PT1, PT3 (USE)]
- Sep 2022 Sichuan University Huawei MindSpore Application Case Implementation Project.
- Nov 2022 Reproduced Swin Transformer with MindSpore and illustrated a demo of image classification in a Jupyter Notebook file. It is selected as an application case on the official webpage of MindSpore. [GitHub]

Patents

- PT- Patent: USE Under Substantial Examination
- PT1 Min Zhu, Fuqiu Chen, Chunlin Long, **Yi Zhou**, Xinyi Wang. "A Visualization Method for Chromatin Hierarchy Analysis Based on Genetic Data". CN113946730A.
- PT5 (USE) Min Zhu, Meixuan Wu, Jiamin Zhu, **Yi Zhou**, Haotian Zhu. "An Analytical Task Perception Method that Integrates Deep Learning Models and Rules". CN116303737A.
- PT4 (USE) Min Zhu, Jiamin Zhu, Meixuan Wu, **Yi Zhou**, Haotian Zhu. "A Dynamic Visualization Recommendation Method Based on User Tasks". CN116204704A.
- PT3 (USE) Min Zhu, Xiyao Li, Chunlin Long, **Yi Zhou**, Xinyi Wang. "Chromatin Topologically Associating Domains Boundary Prediction Method Based on Multimodal Fusion". CN115831217A.
- PT2 (USE) Min Zhu, **Yi Zhou**, Xinyi Wang, Lin Yao. "Method and System for Long Non-coding RNA-Disease Association Prediction Based on Self-Attention Mechanism". CN115171780A.

Teaching & Services

- Fall 2023 **Teaching Assistant**, Sichuan University.
 - Data Visualization (311301030)

Journal Reviewer.

- Expert Systems With Applications (ESWA)
- Jul 2022 Mentor of Bioinformatics Group, Vision Computing Lab.
- Jun 2024 Mentored Meixuan Wu, Xiyao Li, Wanjing Zhang, Lin Gan and Xian Guan.
- Sep 2021 Living Manager, Vision Computing Lab.
 - Aug 2023 Responsible for all non-working tasks in the lab.

Honors & Awards

- 2024 Outstanding Graduate of Sichuan University
- 2023 Outstanding Graduate Student of Sichuan University
- 2023 First-class of Excellent Graduate Scholarship by Sichuan University
- 2023 Tencent Scholarship of Sichuan University
- 2021 Certificate of Honor from Wu Yuzhang Honors College

Skills

Programming Python, PyTorch, PyTorch Geometric, SQL, Linux command, Git

Languages Chinese, English (IELTS: 7.0), Fuzhou dialect