Heng Yang

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OBJECTIVE

Seeking a research position in AI and NLP, leveraging expertise in biological sequence modeling, sentiment analysis, and adversarial attacks to drive innovation and contribute to impactful research projects.

EDUCATION

• University of Exeter
PhD in NLP/Genomics Modelling

• South China Normal University

Master's in NLP

• Yangtze University
Bachelor's in Computer Science

Sept. 2021 - Sept. 2025 (Expected) Exeter, UK

> Sept. 2018 - June 2021 Guangzhou, China Sept. 2014 - June 2018

> > Jingzhou, China

University of Exeter, 2021-2025

AWARDS

• PhD Studentship with Research Grant

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• National (Academic) Graduate Scholarship

South China Normal University, 2020

• Outstanding Graduate

Yangtze University, 2018

PUBLICATIONS

- OmniGenBench: Automating Large-scale Benchmarking for Genomic Foundation Models Heng Yang, Jack Cole, Ke Li (Arxiv 2024)
- OmniGenome: Aligning RNA Sequences with Secondary Structures in Genomic Foundation Models Heng Yang, Ke Li (AAAI 2025)
- PlantRNA-FM: An Interpretable RNA Foundation Model for Exploration Functional RNA Motifs in Plants Haopeng Yu#, Heng Yang#, et al. (Co-first Author, Nature Machine Intelligence, 2024)
- Modeling Aspect Sentiment Coherency via Local Sentiment Aggregation Heng Yang, Ke Li (EACL 2024)
- The Best Defense is Attack: Repairing Semantics in Textual Adversarial Examples Heng Yang, Ke Li (EMNLP 2024)
- PyABSA: A Modularized Framework for Reproducible Aspect-based Sentiment Analysis Heng Yang, Cheng Zhang (CIKM 2023)
- Evolutionary Multi-objective Instruction Optimization via Large Language Model-based Instruction Operators Heng Yang, Ke Li (EMNLP 2023)
- DaNuoYi: Evolutionary Multi-Task Injection Testing on Web Application Firewalls Ke Li, Heng Yang (IEEE TSE, 2023)
- Boosting Text Augmentation via Hybrid Instance Filtering Framework Heng Yang, Ke Li (ACL 2023)
- A Multi-task Learning Model for Chinese-oriented Aspect Polarity Classification and Aspect Term Extraction Heng Yang, Biqing Zeng (Neurocomputing 2021)

Personal Open-source Projects

• OmniGenBench – 4.7k Installations
First large-scale in-silico benchmarking framework for genomic foundation models

Sept. 2024 - Present GitHub

June 2020 - Present GitHub

• PyABSA – 340k Installations, 1k GitHub stars
Aspect-based sentiment analysis framework, serving commercials, developers and scholars

COMMUNITY CONTRIBUTION STATISTICS

- GitHub: 1.5k stars, ≈ 160 commits and 60 PRs/year. I am a open-source lover since the beginning of my research. I am grateful to the developers whose projects helped me a lot. Therefore, I am committed to share all my open-source projects on GitHub with friendly MIT lenience.
- Huggingface: 8 Models and 10 Spaces with $\approx 400 \text{k}$ downloads and 10k access, respectively. Thanks to the Hungingface platform, I am glad to share all my pretrained and fine-tuned state-of-the-art sentiment analysis and genomic foundation models. e.g., deberta-v3-base-absa-v1.1 and OmniGenome-186M. Moreover, I have been to releasing demos for low-resource research topics, like RNA secondary structure prediction and RNA Design.
- PyPi: 8 python wheels with 800k downloads. I have made efforts to simplify the workflows and pipelines by packing and releasing open-access Python wheels. My wheels have been widely used in by the community. These wheels can be easily distributed via PyPi and installed via the pip tool.