

Heng Yang



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[Homepage](#) | [GitHub](#) | [Huggingface](#) | [Google Scholar](#)
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Latest
Version

♥ OBJECTIVE

Seeking an opportunity to apply my expertise in genomics modeling and language modeling to contribute to impactful research and push the boundaries of AI4Science.

🎓 EDUCATION

- University of Exeter**
PhD in NLP/Genomic and Language Modeling, Adversarials
Sept. 2021 - Sept. 2025
Exeter, United Kingdom
- South China Normal University**
Master's in NLP/Sentiment Analysis
Sept. 2018 - June 2021
Guangzhou, China
- Yangtze University**
Bachelor's in Computer Science
Sept. 2014 - June 2018
Jingzhou, China

📖 MAIN PUBLICATIONS

- OmniGenBench: Automating Large-scale Benchmarking for Genomic Foundation Models**
Heng Yang, Jack Cole, Ke Li
ArXiv Preprint
- OmniGenome: Aligning RNA Sequences with Secondary Structures in Genomic Foundation Models**
Heng Yang, Ke Li
AAAI 2025
- MPRNA: Unleashing Multi-species RNA Foundation Model via Calibrated Secondary Structure Prediction**
Heng Yang, Ke Li
EMNLP 2024
- The Best Defense is Attack: Repairing Semantics in Textual Adversarial Examples**
Heng Yang, Ke Li
EMNLP 2024
- PlantRNAFM: 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
- PyABSA: A Modularized Framework for Reproducible Aspect-based Sentiment Analysis**
Heng Yang, Chen Zhang, Ke Li
CIKM 2023
- InstOptima: Evolutionary Multi-objective Instruction Optimization via LLM-based Instruction Operators**
Heng Yang, Ke Li
EMNLP 2023
- BoostAug: Boosting Text Augmentation via Hybrid Instance Filtering Framework**
Heng Yang, Ke Li
ACL 2023
- DaNuoYi: Evolutionary Multi-Task Injection Testing on Web Application Firewalls**
Ke Li, Heng Yang, Willem Visser
IEEE Trans. on Software Engineering 2023

🔗 PERSONAL OPEN-SOURCE PROJECTS

- OmniGenBench** – 5k Installations
First large-scale in-silico benchmarking framework for genomic foundation models
Sept. 2024 - Present
[GitHub](#)
- PyABSA** – 350k Installations, 1k GitHub stars
Aspect-based sentiment analysis framework, serving commercials, developers and scholars
June 2020 - Present
[GitHub](#)

👤 COMMUNITY CONTRIBUTION STATISTICS

- GitHub:** 1.5k stars, \approx 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 1,000k downloads and 10k access, respectively. Thanks to the Huggingface 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 \approx 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.

🏆 AWARDS

- PhD Scholarship, Research Grant**
University of Exeter, 2021-2025
- Chinese National Scholarship, First-class Academic Scholarships**
South China Normal University, 2018-2020
- Outstanding Bachelor Graduate**
Yangtze University, 2018