

Chao Hou, Ph.D.

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Academic Curriculum Vitae (Update: May 2025)

Research Interests:

- Large language model for biological sequences
 - Protein dynamics and multi-conformation
 - Functional analysis of genetic mutations in disease
 - Biomolecular interactions and subcellular compartmentalization
 - Drug designation and effect prediction
 - Transcription regulation and cell fate determination

Work Experience:

Education:

- Ph.D. in Biomedical Informatics Peking University Sep 2020 - Jul 2023
❖ Advisor: [Dr. Tingting Li](#)
 - Bachelor of Medicine and Economics Peking University Sep 2015 - Jul 2020

Main Publications:

(# indicates co-first author, * indicates co-corresponding author)

1. **Hou C***, Di L, Zafar A, Shen Y*. *Understanding Protein Language Model Scaling on Mutation Effect Prediction*. BioRxiv. Apr 2025.
 2. **Hou C**, Zhao H, Shen Y*. *Learning Biophysical Dynamics with Protein Language Models* (previous title: SeqDance: A Protein Language Model for Representing Protein Dynamic Properties). BioRxiv. Apr 2025.
 3. **Hou C#**, Wang X#, Xie H#, Chen T, Zhu P, Xu X, You K, Li T*. *PhaSepDB in 2022: annotating phase separation-related proteins with droplet states, co-phase separation partners and other experimental information*. Nucleic Acids Res. Jan 2023.
 4. **Hou C**, Li Y, Wang M, Wu H, Li T*. *Systematic prediction of degrons and E3 ubiquitin ligase binding via deep learning*. BMC Biology. Jul 2022.
 5. **Hou C#**, Xie H#, Fu Y#, Ma Y#, Li T*. *MloDisDB: a manually curated database of the relations between membraneless organelles and diseases*. Brief Bioinform. Jul 2021.
 6. Chen Z#, **Hou C#**, Wang L#, Yu C, Chen T, Shen B, Hou Y, Li P*, Li T*. *Screening membraneless organelle participants with machine-learning models that integrate multimodal features*. Proc Natl Acad Sci U S A. Jun 2022.
 7. Zhu P#, **Hou C#**, Liu M, Chen T, Li T*, Wang L*. *Investigating phase separation properties of chromatin-associated proteins using gradient elution of 1,6-hexanediol*. BMC Genomics. Aug 2023.

8. Han P, **Hou C**, Zheng X, Cao L, Shi X, Zhang X, Ye H, Pan H, Liu L, Li T*, Hu F*, Li Z*. *Serum Antigenome Profiling Reveals Diagnostic Models for Rheumatoid Arthritis*. Front Immunol. Apr 2022.

Other Publications:

9. Xu X, Li Y, Chen T, Hou C, Yang L, Zhu P, Zhang Y, Li T. *Investigating variant impact on phosphorylation events driving carcinogenesis*. Brief Bioinform. Dec 2023.
10. Yu C, Lang Y, Hou C, Yang E, Ren X, Li T. *Distinctive Network Topology of Phase-Separated Proteins in Human Interactome*. J Mol Biol. Jan 2022.
11. Shi M, You K, Chen T, Hou C, Liang Z, Liu M, Wang J, Wei T, Qin J, Chen Y, Zhang MQ, Li T. *Quantifying the phase separation property of chromatin-associated proteins under physiological conditions using an anti-1,6-hexanediol index*. Genome Biol. Aug 2021.
12. Chen T, Tang G, Li T, Yanghong Z, Hou C, Du Z, Ma L, Li T. *PhaSeDis: A Manually Curated Database of Phase Separation–Disease Associations and Corresponding Small Molecules*. Genomics Proteomics Bioinformatics. Mar 2025.

Oral Presentations

- 2024.09 *Learning Representation of Protein Dynamic Properties with a Language Model*. Retreat of Department of Systems Biology of Columbia University. PA, US.
- 2023.07 *The degradation regulation of phase separating proteins*. The student symposium in Fudan international summer school of life science, Shanghai, China.
- 2023.06 *The degradation regulation of phase separating proteins*. Excellent graduates symposium of Peking University School of Basic Medical Sciences, Beijing, China.
- 2023.04 *The degradation regulation of phase separating proteins*. Multidisciplinary Conference on New Ideas and New Technologies at Peking University, Beijing, China.
- 2022.12 *Targeting disordered degrons on phase separating proteins*. Silk Road International Symposium for Distinguished Young Scholars at Xi'an Jiaotong University, Virtual meeting.

Poster Presentations

- 2024.07 *Learning Representation of Protein Dynamic Properties with a Language Model*. ISMB 2024. Montreal, Canada.
- 2024.11 *Predicting missense mutation effects with biophysics-informed protein language model*. ASHG 2024. Denver, US.
- 2025.02 *Learning Biophysical Dynamics with Protein Language Models*. BPS 2025. Los Angeles, US.

Honors and Awards

- 2023 First prize in the student symposium in Fudan international summer school of life science
- 2023 Peking University Excellent Graduate
- 2022 Peking University President Scholarship
- 2021 Peking University Doctoral Innovation Scholarship
- 2020 Peking University Junior Scholar
- 2016 National Scholarship

2014 Silver medal in the final of Chinese Physics Olympiad (high school)

Journal Referee

Ebiomedicine, Plos Computational Biology, Protein Science

Teaching

2024.01 – 2024.09: Rotation student from Department of Biomedical Informatics. Mentor for Aziz Zafar. Columbia University.

2024.06 – 2024.09: Program for Mathematical Genomics (PMG) Undergrad Student Summer Program. Mentor for Jason Xie. Columbia University.

2025.01 – present: Rotation student from Department of Biomedical Informatics. Mentor for Di Liu. Columbia University.