

# CURRICULUM VITAE



## **YANSHI HU, PhD, MPhil**

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## **EDUCATION**

- 2017-2023 **Doctor of Philosophy in Bioinformatics**  
Zhejiang University, Hangzhou, China
- 2013-2016 **Master of Philosophy in Biomedical Engineering**  
Tianjin Medical University, Tianjin, China
- 2009-2013 **Bachelor of Science in Biomedical Engineering**  
Shandong First Medical University (Shandong Academy of Medical Sciences), Taian, China

## **EXPERIENCE**

- 2023- **Research Assistant**  
College of Life Sciences, Zhejiang University, Hangzhou, China
- 2016-2017 **Research Assistant**  
School of Biomedical Engineering, Tianjin Medical University, Tianjin, China

## **ACADEMIC HONORS & AWARDS**

- 2017- Bioinformatics Society of Zhejiang Province of China, membership
- 2019 Zhejiang University Award of Honor for Graduate
- 2018 Zhejiang University First-Class Scholarship for Doctoral Mid-term Examination (¥20,000)
- 2018 Zhejiang University Award of Honor for Graduate
- 2017 Zhejiang University Excellent Doctoral Freshman Scholarship (¥10,000)
- 2017 Certificate of Reviewing for *Computational Biology and Chemistry*
- 2014 The Sixth National Conference on Bioinformatics & Systems Biology of China,  
Second-Class Prize for Excellent Poster
- 2011 Shandong First Medical University & Shandong Academy of Medical Sciences,  
Second-Class Scholarship
- 2010 Shandong First Medical University & Shandong Academy of Medical Sciences,  
Third-Class Scholarship

## **JOURNAL PEER REVIEW**

Science China-Life Sciences (IF = 9.1), Bioinformatics (IF = 8.3), International Journal of Intelligent Systems (IF = 7.2), Frontiers in Neuroscience (IF = 5.2), Molecular Neurobiology (IF = 5.1), Scientific Reports (IF = 4.9), IEEE/ACM Transactions on Computational Biology and Bioinformatics (IF = 4.5), Current Bioinformatics (IF = 4.0), Frontiers in Neurology (IF = 3.9), BMC Medical Genomics (IF = 3.2), Journal of Molecular Neuroscience (IF = 3.1), Computational Biology and Chemistry (IF = 3.1)

## PUBLICATIONS AND PREPRINTS

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1. Z Wu, C Feng, **YS Hu**, Y Zhou, S Li, S Zhang, Y Hu, Y Chen, H Chao, Q Ni, M Chen\* (*Accepted in Principle*). HALD, a human aging and longevity knowledge graph for precision gerontology and geroscience analyses. *Scientific Data*, 2023 (IF = 10.8)
2. C Feng<sup>#</sup>, RX Tie<sup>#</sup>, SG Xin<sup>#</sup>, Y Chen, S Li, X Hu, Y Zhou, Y Liu, Y Hu, **YS Hu**, H Pan, Z Wu, H Chao, S Zhang, Q Ni, J Huang, W Luo\*, H Huang\*, M Chen\*. Systematic single-cell analysis reveals dynamic control of transposable element activity orchestrating the endothelial-to-hematopoietic transition. *bioRxiv*, 2023.06.19.545461. (*Under Review in BMC Biology*, IF = 7.1)  
DOI: [10.1101/2023.06.19.545461](https://doi.org/10.1101/2023.06.19.545461)
3. Y Chen<sup>#</sup>, **YS Hu**<sup>#</sup>, X Hu, C Feng, M Chen\*. CoGO: a contrastive learning framework to predict disease similarity based on gene network and ontology structure. *Bioinformatics*, 2022, 38(18):4380-4386. (IF = 8.3)  
DOI: [10.1093/bioinformatics/btac520](https://doi.org/10.1093/bioinformatics/btac520)
4. H Chen, X Hu, **YS Hu**, J Zhou, M Chen\*. CoVM<sup>2</sup>: Molecular Biological Data Integration of SARS-CoV-2 Proteins in a Macro-to-Micro Method. *Biomolecules*, 2022, 12(8):1067. (IF = 5.8)  
DOI: [10.3390/biom12081067](https://doi.org/10.3390/biom12081067)
5. B Tan, S Xin, **YS Hu**, C Feng\*, M Chen\*. LBD: a manually curated database of experimentally validated lymphoma biomarkers. *Database*, 2022, Volume 2022, baac051. (IF = 5.8)  
DOI: [10.1093/database/baac051](https://doi.org/10.1093/database/baac051)
6. WY Wu, Y Wu, DH Hu, YC Zhou, **YS Hu**, YJ Chen, M Chen\*. PncStress: a manually curated database of experimentally validated stress-responsive non-coding RNAs in plants. *Database*, 2020, Volume 2020, baaa001. (IF = 5.8)  
DOI: [10.1093/database/baaa001](https://doi.org/10.1093/database/baaa001)
7. TY Wang, P Song, TT Zhong, XJ Wang, XP Xiang, Q Liu, HY Chen, T Xia, H Liu, YM Niu, **YS Hu**, L Xu, YK Shao, LJ Zhu, HY Qi, J Shen, TJ Hou, R Fodde\*, JM Shao\*. The inflammatory cytokine IL-6 induces FRA1 deacetylation promoting colorectal cancer stemness and malignancy. *Oncogene*, 2019, 38:4932-4947. (IF = 8.8)  
DOI: [10.1038/s41388-019-0763-0](https://doi.org/10.1038/s41388-019-0763-0)
8. **YS Hu**, J Xin, Y Hu, L Zhang\*, J Wang\*. Analyzing the genes related to Alzheimer's disease via a network and pathway-based approach. *Alzheimer's Research & Therapy*, 2017, 9(1):29. (IF = 9.2)  
DOI: [10.1186/s13195-017-0252-z](https://doi.org/10.1186/s13195-017-0252-z)
9. **YS Hu**, Z Pan, Y Hu, L Zhang\*, J Wang\*. Network and Pathway-Based Analyses of Genes Associated with Parkinson's Disease. *Molecular Neurobiology*, 2017, 54(6):4452-4465. (IF = 5.1)  
DOI: [10.1007/s12035-016-9998-8](https://doi.org/10.1007/s12035-016-9998-8)
10. Y Hu, Y Yang, Z Fang, **YS Hu**, L Zhang\*, J Wang\*. Detecting pathway relationship in the context of human protein-protein interaction network and its application to Parkinson's disease. *Methods*, 2017, 131:93-103. (IF = 4.8)  
DOI: [10.1016/j.ymeth.2017.08.001](https://doi.org/10.1016/j.ymeth.2017.08.001)
11. ZH Fang, YC Yang, **YS Hu**, MD Li\*, J Wang\*. GRONS: a comprehensive genetic resource of nicotine and smoking. *Database*, 2017, Volume 2017, bax097. (IF = 5.8)  
DOI: [10.1093/database/bax097](https://doi.org/10.1093/database/bax097)
12. TY Ling, YC Zhou, CF Xu, XT Shao, **YS Hu**, KF Ding\*, M Chen\* (*In Preparation*). Colorectal cancer computer-aided image analysis: the teenager in the new era of deep learning.
13. YC Zhou, **YS Hu**, DH Hu, C Feng, MA Ahsan, YJ Liu, TY Ling, SD Li, XC Yang, R Hofestädt, M Chen\* (*In Preparation*). DaTo: a repertoire dedicated to biological online resources.
14. **YS Hu**<sup>#</sup>, YH Chen<sup>#</sup>, M Chen\* (*In Preparation*). DeePAX: Decrypting human biochemical pathway

crosstalk landscape in a deep learning architecture.

15. **YS Hu\***, YM Hu, M Chen\* (*In Preparation*). AWmeta powers adaptively-weighted meta-analysis for gene expression data.
16. **YS Hu\***, BT She, ZN Yin, XJ Yu, W Wu, M Chen\* (*In Preparation*). Systems biology framework unravels molecular substrates underlying comorbidity between Parkinson's and Crohn's disease.
17. YC Zhou, JT Xue, MA Ahsan, DH Hu, **YS Hu**, Y Liu, Y Jiang, W Ni, M Chen\* (*In Preparation*). CytoSEE: a web-based toolkit for automatic computation and evaluation of cytometry data.
18. **YS Hu\***, ZH Fang, J Wang\*, M Chen\* (*In Preparation*). A systems biology framework identifies latent molecular relationships between Alzheimer's and Parkinson's disease.

## CONFERENCE PROCEEDINGS

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1. **Y Hu\***, J Wang\*, M Chen. Analyzing the genes related to Alzheimer's disease via a network and pathway-based approach. *First Sino-Russian Workshop on Integrative Bioinformatics and Systems Biology (WIBSB-2018)* @ Novosibirsk, Russia  
DOI: [10.18699/WIBSB-2018-28](https://doi.org/10.18699/WIBSB-2018-28)
2. **Y Hu**, Z Pan, Y Hu, J Wang\*. Network and pathway based analyses of genes associated with Parkinson's disease. *The Seventh National Conference on Bioinformatics and Systems Biology* @ Chengdu, China  
Availability: [ResearchGate Link](#)
3. Y Hu, **Y Hu**, Y Yang, Z Fang, J Wang\*. Uncovering the common pathogenesis in neurodegenerative and psychiatric disorder via network approaches. *The Seventh National Conference on Bioinformatics and Systems Biology* @ Chengdu, China  
Availability: [ResearchGate Link](#)
4. Z Fang, Y Yang, Y Hu, **Y Hu**, J Wang\*. Identifying the enriched biological pathways in genes related to nicotine dependence via a network-based gene-weighting algorithm. *The Seventh National Conference on Bioinformatics and Systems Biology* @ Chengdu, China  
Availability: [ResearchGate Link](#)
5. **Y Hu**, R Fan, X Li, M Liu, X Liu, X Yi, T Zhang, J Wang\*. Common characteristics of Alzheimer's disease and Parkinson's disease based on AlzGene and PDGene databases. *The Sixth National Conference on Bioinformatics and Systems Biology* @ Nanjing, China  
Availability: [ResearchGate Link](#)
6. X Liu, X Li, M Liu, R Fan, **Y Hu**, Y Hu, X Yi, T Zhang, J Wang\*. Computing the phenotype similarity based on OMIM database and MESH vocabulary. *The Sixth National Conference on Bioinformatics and Systems Biology* @ Nanjing, China  
Availability: [ResearchGate Link](#)
7. X Li, M Liu, X Liu, R Fan, **Y Hu**, Y Hu, X Yi, T Zhang, J Wang\*. TarPriGO: a new method to prioritize miRNA targets based on Gene Ontology. *The Sixth National Conference on Bioinformatics and Systems Biology* @ Nanjing, China  
Availability: [ResearchGate Link](#)
8. M Liu, X Liu, X Li, R Fan, **Y Hu**, Y Hu, X Yi, T Zhang, J Wang\*. A comprehensive pathway and network analysis of candidate genes associated with nicotine addiction. *The Sixth National Conference on Bioinformatics and Systems Biology* @ Nanjing, China  
Availability: [ResearchGate Link](#)
9. R Fan, M Liu, X Li, X Liu, **Y Hu**, Y Hu, X Yi, T Zhang, J Wang\*. The functional divergence analysis of neuronal nicotinic acetylcholine receptor subunits. *The Sixth National Conference on Bioinformatics and Systems Biology* @ Nanjing, China  
Availability: [ResearchGate Link](#)

## ACADEMIC TALKS

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- 09/2019    Zhejiang University-Bielefeld University Joint Symposium @ Bielefeld, Germany
- 08/2018    First Sino-Russian Workshop on Integrative Bioinformatics and Systems Biology (WIBSB-2018)  
              @ Novosibirsk, Russia

## POSTERS

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1.    **Y Hu**, Z Pan, Y Hu, and J Wang. Network and pathway based analyses of genes associated with Parkinson's disease. *The Seventh National Conference on Bioinformatics and Systems Biology* @ Chengdu, China  
DOI: [10.13140/RG.2.2.20162.27844](https://doi.org/10.13140/RG.2.2.20162.27844)
2.    **Y Hu**, R Fan, X Li, M Liu, X Liu, X Yi, T Zhang and J Wang. Common characteristics of Alzheimer's disease and Parkinson's disease based on AlzGene and PDGene databases. *The Sixth National Conference on Bioinformatics and Systems Biology* @ Nanjing, China  
DOI: [10.13140/RG.2.2.14971.82725](https://doi.org/10.13140/RG.2.2.14971.82725)