

# CURRICULUM VITAE



## **YANSHI HU, M.S., Ph.D. Candidate**

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## **EDUCATION**

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- 2017-      **Doctor of Philosophy in Bioinformatics**  
Zhejiang University, Hangzhou, China
- 2013-2016   **Master of Science in Biomedical Engineering**  
Tianjin Medical University, Tianjin, China
- 2009-2013   **Bachelor of Science in Biomedical Engineering**  
Shandong First Medical University / Taishan Medical University, Taian, China

## **EXPERIENCE**

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- 2016-2017   **Research Assistant**  
School of Biomedical Engineering, Tianjin Medical University, Tianjin, China

## **ACADEMIC HONORS & AWARDS**

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- 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**

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- Bioinformatics (IF = **8.470**)
- Molecular Neurobiology (IF = **5.590**)
- Science China-Life Sciences (IF = **4.754**)
- IEEE/ACM Transactions on Computational Biology and Bioinformatics (IF = **3.395**)
- Computational Biology and Chemistry (IF = **2.411**)
- Current Bioinformatics (IF = **2.048**)

## JOURNAL PUBLICATIONS

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1. W Wu, Y Wu, D Hu, Y Zhou, **Y Hu**, Y Chen, M Chen\*. PncStress: a manually curated database of experimentally validated stress-responsive non-coding RNAs in plants. *Database*, 2020, Volume 2020, baaa001. (IF = **4.159**)  
DOI: [10.1093/database/baaa001](https://doi.org/10.1093/database/baaa001)
2. T Wang, P Song, T Zhong, X Wang, X Xiang, Q Liu, H Chen, T Xia, H Liu, Y Niu, **Y Hu**, L Xu, Y Shao, L Zhu, H Qi, J Shen, T Hou, R Fodde\*, J Shao\*. The inflammatory cytokine IL-6 induces FRA1 deacetylation promoting colorectal cancer stemness and malignancy. *Oncogene*, 2019, 38:4932-4947. (IF = **9.867**)  
DOI: [10.1038/s41388-019-0763-0](https://doi.org/10.1038/s41388-019-0763-0)
3. **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 = **8.076**)  
DOI: [10.1186/s13195-017-0252-z](https://doi.org/10.1186/s13195-017-0252-z)
4. **Y 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 = **5.590**)  
DOI: [10.1007/s12035-016-9998-8](https://doi.org/10.1007/s12035-016-9998-8)
5. 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 = **4.669**)  
DOI: [10.1016/j.ymeth.2017.08.001](https://doi.org/10.1016/j.ymeth.2017.08.001)
6. Z Fang, Y Yang, **Y Hu**, MD Li\*, J Wang\*. GRONS: a comprehensive genetic resource of nicotine and smoking. *Database*, 2017, Volume 2017, bax097. (IF = **4.159**)  
DOI: [10.1093/database/bax097](https://doi.org/10.1093/database/bax097)
7. T Ling, Y Zhou, C Xu, X Shao, **Y Hu**, K Ding\*, M Chen\* (*Submitted*). Colorectal cancer computer-aided image analysis: the teenager in the new era of deep learning. *Briefings in Bioinformatics*, 2022 (IF = **11.622**)
8. Y Chen, **Y Hu**, X Hu, C Feng, M Chen\* (*Submitted*). CoGO: a contrastive learning framework to predict disease similarity based on gene network and ontology structure. *Bioinformatics*, 2022 (IF = **8.470**)
9. Y Zhou, **Y Hu**, D Hu, C Feng, MA Ahsan, Y Liu, T Ling, S Li, X Yang, R Hofestädt, M Chen\* (*Submitted*). DaTo: a repertoire dedicated to biological online resources. *Bioinformatics*, 2021 (IF = **8.470**)
10. Y Zhou, J Xue, MA Ahsan, D Hu, **Y Hu**, Y Liu, Y Jiang, W Ni, M Chen\* (*In Preparation*). CytoSEE: a web-based toolkit for automatic computation and evaluation of cytometry data.
11. **YS Hu\***, Z Fang, Y Niu, J Wang\*, M Chen\* (*In Preparation*). A systems biology framework identifies latent molecular relationships between Alzheimer's and Parkinson's disease.

## CONFERENCE PAPERS OR ABSTRACTS

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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
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
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*

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*
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*
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*
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*
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*

**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)