# **CURRICULUM VITAE**



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

Department of Bioinformatics, State Key Laboratory of Plant Physiology and Biochemistry College of Life Sciences, Zhejiang University

866 Yuhangtang Road, Hangzhou 310058, P. R. China

Email: yanshihu@zju.edu.cn Phone: (86) 188-8895-0980 Website: yanshihu.github.io

### **EDUCATION**

| 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**

2016-2017 Research Assistant

School of Biomedical Engineering, Tianjin Medical University, Tianjin, China

## ACADEMIC HONORS AND 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)                   |
| 2014  | The Sixth National Conference on Bioinformatics & Systems Biology of China,             |
|       | Second-Class Prize for Excellent Poster   |
| 2011  | Shandong First Medical University / Taishan Medical University,                         |
|       | Second-Class Scholarship  |
| 2010  | Shandong First Medical University / Taishan Medical University,                         |
|       | Third-Class Scholarship   |

#### **JOURNAL PEER REVIEW**

IEEE/ACM Transactions on Computational Biology and Bioinformatics (IF = 2.896)

Computational Biology and Chemistry (IF = 1.581)

Current Bioinformatics (IF = 1.189)

## **JOURNAL PUBLICATIONS**

1. 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. PubMed PMID: 30804456. DOI:

- 10.1038/s41388-019-0763-0. (IF = **6.933**)
- 2. **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. PubMed PMID: 28446202. DOI: 10.1186/s13195-017-0252-z. (IF = 7.072)
- 3. 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. PubMed PMID: 27349437. DOI: 10.1007/s12035-016-9998-8. (IF = 6.190)
- 4. 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. PubMed PMID: 28790017. DOI: 10.1016/j.ymeth.2017.08.001. (IF = **3.998**)
- 5. Z Fang, Y Yang, Y Hu, MD Li\*, J Wang\*. GRONS: a comprehensive genetic resource of nicotine and smoking. *Database*, 2017, Volume 2017, bax097. PubMed PMID: 31725863. PubMed Central PMCID: PMC5750854. DOI: 10.1093/database/bax097. (IF = 3.978)
- 6. W Wu, Y Wu, D Hu, Y Zhou, Y Hu, M Chen\* (*In Minor Revision*). PncStress: a manually curated database of experimentally validated stress-responsive non-coding RNAs in plants. *Database*. (IF = 3.978)
- 7. Y Zhou, J Xue, MA Ahsan, D Hu, Y Hu, Y Liu, Y Jiang, W Ni, M Chen\* (*To Be Submitted*). CytoSEE: a web-based toolkit for automatic computation and evaluation of cytometry data.
- 8. **YS Hu\***, Z Fang, Y Niu, J Wang\*, M Chen\* (In preparation). A systems biology framework identifies latent molecular relationships between Alzheimer's disease and Parkinson's disease.

## CONFERENCE PAPERS OR ABSTRACTS

- 1. Y Hu\*, J Wang\*, M Chen. Analyzing the genes related to Alzheimer's disease via a network and pathway-based approach. DOI: 10.18699/WIBSB-2018-28. *Integrative Bioinformatics and Systems Biology (WIBSB-2018), First Sino-Russian Workshop*, Novosibirsk, Russia, (2018).
- 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 & Systems Biology and International Workshop on Advanced Bioinformatics & Precision Medicine*, Chengdu, China, (2016).
- 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 & Systems Biology and International Workshop on Advanced Bioinformatics & Precision Medicine*, Chengdu, China, (2016).
- 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 & Systems Biology and International Workshop on Advanced Bioinformatics & Precision Medicine*, Chengdu, China, (2016).
- 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 & Systems Biology and International Workshop on Advanced Bioinformatics*, Nanjing, China, (2014).
- 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 & Systems Biology and International Workshop on Advanced Bioinformatics*, Nanjing, China, (2014).
- 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 & Systems Biology and International Workshop on Advanced Bioinformatics*, Nanjing, China, (2014).
- 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. <u>The Sixth National Conference on Bioinformatics & Systems Biology and International Workshop on Advanced Bioinformatics</u>, Nanjing, China, (2014).
- 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 & Systems Biology and International Workshop on Advanced Bioinformatics*, Nanjing, China, (2014).

#### **POSTERS**

- 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 & Systems Biology and International Workshop on Advanced Bioinformatics & Precision Medicine*, Chengdu, China, (2016).
- 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 and International Workshop on Advanced Bioinformatics*, Nanjing, China, (2014).