

# 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](mailto:yanshihu@zju.edu.cn)*

*Phone: (86) 188-8895-0980*

*Website: [yanshihu.github.io](http://yanshihu.github.io)*

## **EDUCATION**

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

## **EXPERIENCE**

- 
- |           |   |
|-----------|---|
| 2016-2017 | <b>Research Assistant</b><br>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 <i>Computational Biology and Chemistry</i>  |
| 2014  | The Sixth National Conference on Bioinformatics & Systems Biology of China,<br>Second-Class Prize for Excellent Poster |
| 2011  | Shandong First Medical University / Taishan Medical University,<br>Second-Class Scholarship                            |
| 2010  | Shandong First Medical University / Taishan Medical University,<br>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. W Wu, Y Wu, D Hu, Y Zhou, **Y Hu**, M Chen\* (*Accepted*). PncStress: a manually curated database of experimentally validated stress-responsive non-coding RNAs in plants. *Database*. (IF = **3.978**)

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. PubMed PMID: 30804456. DOI: 10.1038/s41388-019-0763-0. (IF = **6.933**)
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. PubMed PMID: 28446202. DOI: 10.1186/s13195-017-0252-z. (IF = **7.072**)
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. PubMed PMID: 27349437. DOI: 10.1007/s12035-016-9998-8. (IF = **6.190**)
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. PubMed PMID: 28790017. DOI: 10.1016/j.ymeth.2017.08.001. (IF = **3.998**)
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. PubMed PMID: 31725863. PubMed Central PMCID: PMC5750854. DOI: 10.1093/database/bax097. (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. *The Sixth National Conference on Bioinformatics & Systems Biology and International Workshop on Advanced Bioinformatics*, 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).

#### ACADEMIC TALKS

---

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

#### 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).