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
	Taishan Medical University / Shandong First 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
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	Taishan Medical University / Shandong First Medical University,
	Second-Class Scholarship
2010	Taishan Medical University / Shandong First Medical University,
	Third-Class Scholarship

JOURNAL PEER REVIEW

IEEE/ACM Transactions on Computational Biology and Bioinformatics (IF = 2.428)

Computational Biology and Chemistry (IF = 1.412)

Current Bioinformatics (IF = 0.627)

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 = **6.206**)
- 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, 1 January 2017, bax097. PubMed Central PMCID: PMC5750854. DOI: 10.1093/database/bax097. (IF = 3.978)
- 6. Y Zhou, J Xue, MA Ahsan, D Hu, Y Hu, Y Liu, Y Jiang, W Ni, M Chen* (*In Peer Review*). CytoSEE: a web-based toolkit for automatic computation and evaluation of cytometry data. *Nucleic Acids Research* Webserver Issue (IF = 11.561)
- 7. **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).
- R Fan, M Liu, X Liu, Y Hu, Y Hu, X Yi, T Zhang, J Wang*. The functional divergence analysis of neuronal nicotinic acetylcholine receptor subunits. <u>The Sixth National Conference on Bioinformatics & Systems Biology and International Workshop on Advanced Bioinformatics</u>, 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).
- 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. <u>The Sixth National Conference</u>
 on Bioinformatics and Systems Biology and International Workshop on Advanced Bioinformatics, Nanjing,
 China, (2014).