CURRICULUM VITAE



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

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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 & 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

Bioinformatics (IF = 9.853)

Molecular Neurobiology (IF = 4.519)

Science China-Life Sciences (IF = 4.611)

IEEE/ACM Transactions on Computational Biology and Bioinformatics (IF = 3.015)

Computational Biology and Chemistry (IF = 1.850)

Current Bioinformatics (IF = 2.068)

JOURNAL PUBLICATIONS

- 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. PubMed Central PMCID: PMC7078067. DOI: 10.1093/database/baaa001. (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 = 7.971)
- 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 = 7.114)
- 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 = **4.061**)
- 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, 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 = **9.853**)
- 8. 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.
- 9. **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. First Sino-Russian Workshop on Integrative Bioinformatics and Systems Biology (WIBSB-2018) @ Novosibirsk, Russia
- 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

- 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

- 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
- 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