Curriculum Vitae of Dr. Jia Wang

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Research interest

My research area includes **Machine learning** and **Bioinformatics**. I mainly pay my attention to construct diverse prediction models for different pathogenic microbes such as influenza A virus, West Nile virus, Pasteurella multocida and so on. These models could be applied in the predictions of interspecies transmission, protein interactions, antigenic variation, host tropism, etc.

Education & academic appointments

11/2018-present	Associate Professor in Computer Science, College of Informatics,
	Huazhong Agricultural University, Wuhan, China
07/2014-11/2018	Lecturer in Computer Science, College of Informatics, Huazhong
	Agricultural University, Wuhan, China
12/2012-06/2014	Lecturer in Computer Science, College of Science, Huazhong
	Agricultural University, Wuhan, China
09/2008-12/2012	PhD, School of Computer Science and Technology, Huazhong
	University of Science Technology, Wuhan, China
09/2006-06/2008	Master, School of Computer Science and Technology, Huazhong
	University of Science Technology, Wuhan, China
09/1998-06/2002	Bachelor, School of Computer Science and Information Engineering,
	Hubei University, Wuhan, China

Ongoing and finished third-party funding

01/2018 Research on breeding and disease control services for smart pig farm based on "Internet Plus" (Grant number: 2662018JC034), Fundamental Research Funds for the Central Universities, 2018-2020. (300,000 RMB Yuan for 3 years) 01/2015 Bioinformatics study on avian influenza virus transmission from avian to human (Grant number: 2662016PY035), Fundamental Research Funds for the Central Universities (80,000 RMB Yuan for 1 years)

Peer-review & editorial duties

I have served as a reviewer for Plos One and Journal of Biomolecular Structure & Dynamics.

Publications

(* for corresponding author, # for co-first author)

- Zhong Peng, Junyang Liu, Wan Liang, Fei Wang, Li Wang, Xueying Wang, Lin Hua, Huanchun Chen; Brenda A Wilson*; Jia Wang*; Bin Wu*. Development of an online tool for Pasteurella multocida genotyping and genotypes of Pasteurella multocida from different hosts, Frontiers in Veterinary Science, 2021, 8(771157)
- Jing Chen, Jun Sun, Xiangming Liu, Feng Liu, Rong Liu*, Jia Wang*. Structure-based prediction of West Nile virus-human protein-protein interactions. Journal of Biomolecular Structure & Dynamics. 2019, 37(9):2310-2321
- Muhammad Tahir ul Qamar, Amna Bari, Muhammad Muzammal Adeel, Arooma Maryam, Usman Ali Ashfaq*, Xiaoyong Du, Iqra Muneer, Hafiz Ishfaq Ahmad, Jia Wang*. Peptide vaccine against chikungunya virus: immuno-informatics combined with molecular docking approach. Journal of Translational Medicine. 2018, 16(1):298
- Xiaoxia Yang#, Jia Wang#, Jun Sun, Rong Liu*. SNBRFinder: A sequence-based hybrid algorithm for enhanced prediction of nucleic acid-binding residues. Plos One. 2015, 10(7): e0133260.
- Haibo Cui, Xiaomei Wei, Yu Huang, Bin Hu, Yaping Fang, Jia Wang*. Using multiple linear regression and physicochemical changes of amino acid mutations to predict antigenic variants of influenza A/H3N2 viruses. Bio-Medical Materials and Engineering. 2014, 24(6): 3729-3735.
- Jia Wang#, Dijun Chen#, Yang Lei#, JiWei Chang, BaoHai Hao, Feng Xing, Sen Li, Qiang Xu, XiuXin Deng, LingLing Chen*. Citrus sinensis Annotation Project (CAP): A comprehensive database for sweet orange genome. PLoS ONE. 2014, 9(1): e87723.
- Jia Wang, Zheng Kou, Mojie Duan, Chuang Ma*, Yanhong Zhou*. Using amino acid factor scores to predict avian-to-human transmission of avian influenza viruses: a machine learning study. Protein & Peptide Letters. 2013, 20(10): 1115-1121.

 Jia Wang, Chuang Ma, Zheng Kou, Yanhong Zhou, Huailan Liu*. Predicting transmission of avian influenza A viruses from avian to human by using informative physicochemical properties. International Journal of Data Mining and Bioinformatics. 2013, 7(2): 166-179.