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Ting-You Wang

Education

- 2013–present **PhD. Candidate, Bioinformatics and Genomics**, The University of Hong Kong, Hong Kong SAR, China.
 - 2008–2011 MSc. Bioinformatics, China Agricultural University, Beijing, China.
 - 2003–2007 **BEng. Mechanical Design, Manufacturing and Automation**, China Agricultural University, Beijing, China.

Publications

- 2016 "Regulatory hierarchy and signaling modules in systemic lupus erythematosus revealed by a systems biology approach", **Ting-You Wang**, Yong-Fei Wang, Yan Zhang, Jiangshan Jane Shen, Mengbiao Guo, Jing Yang, Yu-Lung Lau, Wanling Yang. Submitted in Arthritis & Rheumatology
- 2016 "Genome-wide association meta-analysis in Chinese and European individuals identifies ten new loci associated with systemic lupus erythematosus", David L Morris, Yujun Sheng, Yan Zhang, Yong-Fei Wang, Zhengwei Zhu, Philip Tombleson, Lingyan Chen, Deborah S Cunninghame Graham, James Bentham, Amy L Roberts, Ruoyan Chen, Xianbo Zuo, Tingyou Wang, Leilei Wen, Chao Yang, Lu Liu, Lulu Yang, Feng Li, Yuanbo Huang, Xianyong Yin, Sen Yang, Lars Rönnblom, Barbara G Fürnrohr, Reinhard E Voll, Georg Schett et al. Nature Genetics, 2016, 48, 940-946
- 2015 "Meta-analysis of GWAS on two Chinese populations followed by replication identifies novel genetic variants on the X chromosome associated with systemic lupus erythematosus", Yan Zhang, Jing Zhang, Jing Yang, Yongfei Wang, Lu Zhang, Xianbao Zuo, Liangdan Sun, Hai-Feng Pan, Nattiya Hirankarn, Tingyou Wang, Ruoyan Chen, Dingge Ying, et al. Human Molecular Genetics, 2015, 24(1), 274-284
- 2014 "Chromosome-8-Coded Proteome of Chinese Chromosome Proteome Data Set (CCPD) 2.0 with Partial Immunohistochemical Verifications", Yang Liu, Wantao Ying, Zhe Ren, Wei Gu, Yang Zhang, Guoquan Yan, Pengyuan Yang, Yinkun Liu, Xuefei Yin, Cheng Chang, Jing Jiang, Fengxu Fan, Chengpu Zhang, Ping Xu, Quanhui Wang, Bo Wen, Liang Lin, Tingyou Wang, Cahoqin Du et al., Journal of Proteome Research, 2014, 13(1), 126-136.
- 2013 "Qualitative and quantitative expression status of the human chromosome 20 genes in cancer tissues and the representative cell lines", Quanhui Wang[#], Bo Wen[#], Guangrong Yan[#], Junying Wei[#], Liqi Xie[#], Shaohang Xu, Dahai Jiang, **Tingyou Wang**, Ruo Zhou, Haiyi Zhao et al., *Journal of Proteome Research*, 2013, 12(1), 151-161. (*These authors contribute equally to this work.)

2011 "A predicted protein-protein interaction network of the filamentous fungus *Neurospora crassa*", **Ting-You Wang**, Fei He, Qi-Wen Hu, Ziding Zhang, *Molecular BioSystems*, 2011, 7, 2278-2285.

Presentations

2016 "Understanding gene expression changes in Systemic Lupus Erythematosus through a systems approach", Poster presentation at the 13th International Congress of Human Genetics, Kyoto, Japan

Industry and commercial experience

2011–2013 Bioinformatics engineer, BGI, Shenzhen, China.

This was a full-time position involving mass spectrometry based proteomics research , development of databases.

2006 Intern Engineer, China FAW Group Corporation, Changchun, China.

This was an internship job involving the improvement of machinery design and manufacturing process.

Teaching experience

2009 Teacher Assistant for undergraduate students in the course of Bioinformatics (full semester).

Awards and scholarships

2008-2009 Outstanding graduate student leader award (vice-president of the Graduate Students' Union)

2008-2009 Outstanding academic performance scholarship

2003-2005 Second-class scholarship winner for excellent academic performance (top 10%)

Conferences attended

- 2016 American Society of Human Genetics 2016 Annual Meeting, Vancouver, Canada
- 2016 The 13th International Congress of Human Genetics, Kyoto, Japan
- 2014 2014 East-West Alliance Global Symposia, Hong Kong SAR, China
- 2013 Frontiers in Biomedical Research HKU 2013, Hong Kong SAR, China
- 2012 ICG-7 & Bio-IT APAC 2012, Hong Kong SAR, China
- 2011 The 6th International Conference on Genomics (ICG-6), Shenzhen, China
- 2010 2010 International Summer School on Chips, Computers and Crops, Beijing, China

Skills

Bioinformatics

systems An extensive knowledge of systems biology

biology

genetics Expertise in genome-wide association study (GWAS)

genomics Proficient in exome sequencing and RNA sequencing data analysis

shotgun Familiar with methods and applications for identification and quantification in mass proteomics spectrometry proteomics. Participator of Chromosome-Centric Human Proteome Project(CHPP) in AOHUPO.

Computer skills

os Linux, Windows, Mac OS X

pc Python, Perl, PHP, R, Shell, Visual Basic, C/C++

applications Microsoft Office Suite, OpenOffice, Vim, Photoshop, LaTex, AutoCAD

web HTML5, CSS, JavaScript/jQuery, LAMP, Google Maps API, Django, TCP/IP stack

database MySQL

Personal

A fast learner and critical thinker

Willing to take the incredible effort on the area of interest and never get tired of it

Other skills

Data and information collection

Writing and presenting reports

Language

Mandarin, English, Cantonese (Basic), Spanish (Basic)

Other work-related interests if I had time

Online courses from edX and Coursera

Reading books involving data science and data visualization

Watching TED talks, since these videos facilitate me to synthesize new ideas.

References

name Assoc. Prof. Wanling Yang Ph. D.

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website http://paed.hku.hk/menu/staff/wanlingyang/wanlingyang.html

comments Dr. Wanling Yang is my Ph.D. primary supervisor.

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comments Prof. Yu-lung Lau is my Ph.D. co-supervisor.

name Prof. Ziding Zhang Ph. D.

address College of Biological Sciences, China Agricultural University, Beijing 100193, China

email zidingzhang@cau.edu.cn

website http://systbio.cau.edu.cn/zzdlab/faculty

comments Dr. Ziding Zhang is my M. Sc. thesis supervisor.