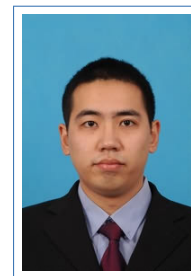


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Education

- 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 "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 Benthams, 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**, Wanling Yang 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 design and development of databases, MS-proteomics research and technical documentation.
- 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

- 2003-2005 Second-class scholarship winner for excellent academic performance (top 10%)
- 2008-2009 Outstanding graduate student leader award (vice-president of the Graduate Students' Union)
- 2008-2009 Outstanding academic performance scholarship

Conferences and courses attended

- 2009 Google Developer Day 2009, Beijing, China
- 2010 Dragon-Star Program 2010, Wuhan, China
- 2010 The 8th International Bioinformatics Workshop (IBW2010), Wuhan, China
- 2010 2010 International Summer School on Chips, Computers and Crops, Beijing, China
- 2011 Google Developer Day 2011, Guangzhou, China
- 2011 The 6th International Conference on Genomics (ICG-6), Shenzhen, China
- 2012 ICG-7 & Bio-IT APAC 2012, Hong Kong SAR, China
- 2013 Frontiers in Biomedical Research HKU 2013, Hong Kong SAR, China
- 2014 2014 East-West Alliance Global Symposia, Hong Kong SAR, China
- 2016 The 13th International Congress of Human Genetics, Kyoto, Japan

Skills

Bioinformatics

- systems biology An extensive knowledge of systems biology as they pertain to PPI network in my thesis.
- shotgun proteomics Familiar with methods and applications for identification and quantification in MS-proteomics. Participator of Chromosome-Centric Human Proteome Project(CHPP) in AOHUPO.
- genomics Proficient in exome sequencing and genome-wide association study(GWAS).

Computer skills

os Linux, Windows 7, Windows XP, 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
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

Other work-related interests if I had time

Online bioinformatics course, such as bioinformatics.ca

Reading books published by O'Reilly media, especially the books involving data science and data visualization

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

References

- name **Assoc. Prof. Wanling Yang Ph. D.**
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- name **Prof. Yu-lung Lau**
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comments Dr. Ziding Zhang is my M. Sc. thesis supervisor.