

# Liang-Bo Wang

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## EDUCATION

**Washington University in St. Louis**, United States  
**Ph.D. candidate** in Computational and Systems Biology Aug 2016 – (present)

**National Taiwan University**, Taipei, Taiwan  
**M.S.** in Biomedical Electronics and Bioinformatics Jan 2014 – Jul 2016  
**B.S.** in Electrical Engineering Sep 2009 – Jan 2014  
Minor in Economics

## AWARDS & SCHOLARSHIPS

**McDonnell International Scholars Academy** 2016 – Present  
Washington University in St. Louis

**Outstanding Summer Research Award** 2012  
Center of Genomic Medicine, National Taiwan University

## RESEARCH EXPERIENCE

**PhD Thesis**, Washington University in St. Louis May 2017 – (present)

- Advisor: Dr. Li Ding
- Topic: Proteogenomic and Spatial Characterization of Glioblastoma Tumors at Bulk and Single Cell Level

**Microsoft Research Asia**, Beijing, China Jul 2014 – Feb 2015  
Internship

- Advisor: Dr. Eric Chang
- Topic: Cancer Image and Pathway Analysis
- Applied convolutional neural network to histopathology image analysis

**Biomedical Engineering and Bioinformatics**, National Taiwan University  
Graduate Research Jan 2014 – Jun 2016

- Advisor: Professor Eric Y. Chuang
- Topic: Integrative analysis of copy number variation, gene expression, and miRNA regulation on cancer; drug treatment prediction on lung cancer cell line

Undergraduate Research Sep 2012 – Jan 2014

- Advisor: Professor Eric Y. Chuang
- Topic: DNA-seq, RNA-seq and small RNA-seq data processing
- Project: Nextbiopsy, a web-based NGS pipeline report generator

Undergraduate Research Sep 2011 – Nov 2012

- Advisor: Associate Professor Wei-Cheng Tian
- Project: CMOS-based tactile sensor system for blood pressure measurement

## WORK EXPERIENCE

**Pinkoi**, Taipei, Taiwan Jun 2015 – Sep 2015  
Internship

- Pinkoi is an online marketplace for unique and original design goods in Asia
- Topic: Product search improvement
- Improved Chinese text segmentation on ElasticSearch and added preliminary semantic search built on Word2Vec

## PUBLICATIONS

## JOURNALS

- [1] **LB Wang\***, A Karpova\*, MA Gritsenko\*, JE Kyle\*, S Cao\*, Y Li\*, *et al.* “Proteogenomic and Metabolomic Characterization of Human Glioblastoma.” *Cancer Cell* 39 (4). Feb 2021
- [2] F Petralia, *et al.* “Integrated Proteogenomic Characterization across Major Histological Types of Pediatric Brain Cancer.” *Cell* 183 (7), 1962-1985. Dec 2020
- [3] MH Bailey, *et al.* “Retrospective evaluation of whole exome and genome mutation calls in 746 cancer samples.” *Nature Communications* 11 (1), 1-27. Sep 2020
- [4] MA Gillette, *et al.* “Proteogenomic characterization reveals therapeutic vulnerabilities in lung adenocarcinoma.” *Cell* 182 (1), 200-225. Jul 2020
- [5] O Rozenblatt-Rosen, *et al.* “The Human Tumor Atlas Network: charting tumor transitions across space and time at single-cell resolution.” *Cell* 181 (2), 236-249. Apr 2020
- [6] JE McDermott, *et al.* “Proteogenomic Characterization of Ovarian HGSC Implicates Mitotic Kinases, Replication Stress in Observed Chromosomal Instability.” *Cell Reports Medicine*, 1 (1) 100004. Apr 2020
- [7] DJ Clark, *et al.* “Integrated proteogenomic characterization of clear cell renal cell carcinoma.” *Cell* 179 (4), 964-983. Oct 2019
- [8] GF Gao\*, JS Parker\*, SM Reynolds\*, TC Silva\*, **LB Wang\***, W Zhou\*, *et al.* “Before and After: Comparison of Legacy and Harmonized TCGA Genomic Data Commons’ Data.” *Cell Systems* 9 (1), 24-34. Jul 2019
- [9] SQ Sun, RJ Mashl, S Sengupta, AD Scott, W Wang, P Batra, **LB Wang**, M A Wyczalkowski, L Ding. “Database of evidence for precision oncology portal.” *Bioinformatics* 34 (24), 4315-4317. Jul 2018.
- [10] KL Huang, *et al.* “Pathogenic germline variants in 10,389 adult cancers.” *Cell*, 173 (2), 355-370. e14. Apr 2018.
- [11] Y Xu, Z Jia, **LB Wang**, Y Ai, F Zhang, M Lai, I Eric, C Chang. “Large scale tissue histopathology image classification, segmentation, and visualization via deep convolutional activation features.” *BMC Bioinformatics*, 18 (1), 281. Dec 2017.
- [12] YP La\*, **LB Wang\***, WA Wang\*, LC Lai, MH Tsai, TP Lu, E Y Chuang. “iGC—an Integrated Analysis Package of Gene Expression and Copy Number Alteration.” *BMC Bioinformatics*, 18 (1), 35. Jan 2017.
- [13] **LB Wang**, E Y Chuang, TP Lu. “Identification of predictive biomarkers for ZD-6474 in lung cancer.” *Translational Cancer Research*, 4 (4), 324-331. Aug 2015.
- [14] CY Lee\*, YC Chiu\*, **LB Wang\***, YL Kuo\*, EY Chuang, LC Lai, MH Tsai. “Common applications of next-generation sequencing technologies in genomic research.” *Translational Cancer Research*, 2 (1), 33-45. Feb 2013.

## CONFERENCE PAPERS

- [1] **LB Wang**, SY Dai, TP Lu, LC Lai, MH Tsai, E Y Chuang. “Nextbiopy: Utilities for next-generation sequencing data processing”, *9th Asian Pacific Conference on Medical and Biological Engineering (APCMBE)*, Tainan, Taiwan, Oct 2014.
- [2] CY Lee\*, **LB Wang\***, MH Tsai, LC Lai, E Y Chuang. “Identification of novel miRNAs in breast data of the next generation sequencing using miRDeep2 and Galaxy”, *AACR 104th Annual Meeting*, Washington, DC, U.S.A., Apr 2013.
- [3] YC Lin, CJ Hsieh, **LB Wang** (presenter), JC Liou, WC Tian. “CMOS MEMS metal-based tactile sensors development”, *AVS 59th International Symposium & Exhibition*, Florida, U.S.A., Oct 2012.
- [4] CT Sun, YC Lin, CJ Hsieh, JC Liou, **LB Wang**, WC Tian. “A linear-response CMOS-MEMS capacitive tactile sensor”, *IEEE Sensors 2012*, Taipei, Taiwan, Oct 2012.

## PROGRAMMING SKILLS

### LANGUAGES

Python (both scientific and web stack), R, Web frontend (HTML/CSS/Javascript), SQL, C

## COMMUNITY ENGAGEMENT

**PyCon Taiwan** Organizing Committee, Taipei, Taiwan                      Jul 2015 – Jul 2016  
Program Committee Chair

**PyCon APAC** Organizing Committee, Taipei, Taiwan                      2014, 2015  
Public Relations Manager

**Open Source Communities**, Taipei, Taiwan                      2013 – 2016  
Regular Speaker

- Including Taiwan PyCon, Taipei.py, Taiwan R User Group, and PyCon China
- Gave over 10 talks, topics included convolutional neural network, text search, HDF5, parallel computing, and R ggplot2 plotting.

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