## Wenbo Li

(832) 409-4936 liwenbo4936@gmail.com

#### **EMPLOYMENT**

EMPLOYMEN I	
♦ Postdoctoral Research Associate, Division of Diabetes, Endocrinology and Metabolism, Baylor College of Medicine Supervisor: Dr. Zheng Sun.	Jul. 2017- ongoing
<ul> <li>Postdoctoral Research Associate, Department of Biology and Biochemistry, University of Houston.</li> <li>Supervisor: Dr. Li Chen.</li> </ul>	Jun. 2016- Jul. 2017
<ul> <li>Postdoctoral Research Associate, BioSciences Department, Rice University.</li> <li>Supervisor: Dr. Janet Braam</li> </ul>	Jan. 2014- Jul. 2015
EDUCATION	
Ph. D. <b>Developmental Physiology and Molecular Biology program</b> , Beijing Normal University, Beijing, China. Advisor: Dr. <u>Yingdian Wang</u> <i>Core courses</i> : Genomics, Immunology, Proteomics, Developmental Biology, Gene Engineering, and Biological Literature Study.	Sept. 2006- Jun. 2013
Botany and Plant Sciences Ph.D. joint training program, University of California, Riverside, USA.  Advisor: Dr. <u>Jian-Kang Zhu</u>	Sept. 2007- Jul. 2009
B. S. <b>Biological Sciences</b> , Beijing Normal University  Core courses: Molecular Biology, Genetics, Biochemistry, Cell Biology, Cell  Engineering, Cell Culture and Tissue Culture, Physiology, Zoology, Botany, and Microbiology.	Sept. 2002- Jul. 2006
SELECTED RESEARCH PROJECTS	
Hdac3 function during Heart failure in Type I Diabetes Mouse Model Supervisor: Dr. Zheng Sun I knocked out Hdac3 gene in Type I diabetes mouse model, to study how Hdac3 contribute to heart failure during diabetes.	Jul. 2017 - ongoing
♦ De facto Target of Histone Deacetylase Inhibitors Supervisor: Dr. Zheng Sun I am using lentivirus based CRISPR library to screen or genes that upon activation or silencing, confers resistance to HDI-mediated cell death in liver cancer cells.	Jul. 2017 - ongoing

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Aug.2016-

Jul. 2017

**♦** Numa function in lineage determination during embryo development.

Supervisor: Dr. Li Chen Group leader: Dr. Robert J. Schwartz

I used multiple markers to test Mesp1 lineage-specific Numa deletion cells in early embryo development stage and found out that cell differentiation in this group is dramatically reduced.

### **♦** Mesp1 lineage cells migration to endoderm.

Jul. 2016 -

Supervisor: Dr. Li Chen Group leader: Dr. Robert J. Schwartz

Jul. 2017

We discovered that one subgroup of Mesp1 lineage cells can migrate to endoderm. I found out that the cell number of Mesp1 lineage endoderm migration group increased in Mesp1 know-out embryos, and the endoderm cells also increased in non-Mesp1 lineage cells.

# ♦ How jasmonate and peroxisomes defend plant against fungus infection. Supervisor: Dr. Janet Braam

May. 2014 -Jul. 2015

I discovered that when infected by the fungus (*Botrytis cinerea*), jasmonate biosynthesis mutant *aos* died from infection while wild-type lived as the infection was blocked at the petiole and that the blockage of infection was jasmonate-dependent. I also found that the number of peroxisome organelles, inside which jasmonate acid was synthesized, increased after fungus infection.

#### **♦** Investigation of opr3 suppressors in *Arabidopsis*.

Jan. 2014 -

Supervisor: Dr. Janet Braam

Jul. 2015

I screened for *Arabidopsis* mutants that compromise infertile phenotype of *opr3* and found 3 qualified mutants. I used both genome sequencing method and map-based cloning and narrowed down one mutant to 3 possible genes.

# ♦ Investigation of the function of cytoskeleton organization in detoxification of ROS in *Arabidopsis* during salt stress tolerance.

Jan. 2008-Jan. 2012

Supervisor: Drs. Jian-Kang Zhu and Yingdian Wang

I screened for salt-sensitive *Arabidopsis* mutants, identified and cloned the mutant gene RSA3. I discovered that RSA3, along with other proteins, plays a critical role in salt stress tolerance, which is maintaining the proper organization of actin microfilaments and minimizing damage caused by excessive ROS.

### ♦ Investigation of the function of RDM12 in DNA methylation of Arabidopsis.

Sept.2007 -Jun. 2009

Supervisor: Dr. Jian-Kang Zhu. Collaborator: Zhi-min Zheng, Xinjian He We screened for second-site suppressors of ros1, and identified the RDM12 locus. We then investigated the function of the gene and discovered that RDM12 is a component of the RdDM pathway, and that RdDM may involve double-stranded RNAs with a 5' overhang and the partnering between RDM12 and RDR2.

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#### **PUBLICATIONS**

- Shiyang Song, Chih-Liang Tien, Hao Cui, Paul Basil, Ningxia Zhu, Yingyun Gong, Wenbo Li, Hui Li, Qiying Fan, Jong Min Choi, Weijia Luo, Yanfeng Xue, Rui Cao, Wenjun Zhou, Andrea R Ortiz, Brittany Stork, Vatsala Mundra, Nagireddy Putluri, Brian York, Maoping Chu, Jiang Chang, Sung Yun Jung, Liang Xie, Jiangping Song, Lilei Zhang, Zheng Sun. (2022) Myocardial Rev-erb-Mediated Diurnal Metabolic Rhythm and Obesity Paradox. Circulation.145(6), 448-464
- ❖ Guolian Ding, Xin Li, Xinguo Hou, Wenjun Zhou, Yingyun Gong, Fuqiang Liu, Yanlin He, Jia Song, Jing Wang, Paul Basil, Wenbo Li, Sichong Qian, Pradip Saha, Jinbang Wang, Chen Cui, Tingting Yang, Kexin Zou, Younghun Han, Christopher I Amos, Yong Xu, Li Chen, Zheng Sun. (2021). REV-ERB in GABAergic neurons controls diurnal hepatic insulin sensitivity. Nature. 592(7867), 763-767
- ♦ Wenbo Li and Zheng Sun. (2019). Mechanism of Action for HDAC Inhibitors— Insights from Omics Approaches. International journal of molecular sciences. 20(7), 1616.
- ♦ Shun Bai, Kaiqiang Fu, Huiqi Yin, Yiqiang Cui, Qiuling Yue, Wenbo Li, ..., Lan Ye. (2018). Sox30 initiates transcription of haploid genes during late meiosis and spermiogenesis in mouse testes. Development. 145(13): dev164855.
- ♦ Wenbo Li, Qingmei Guan, Zhen-Yu Wang, Yingdian Wang and Jianhua Zhu. (2013). A Bi-Functional Xyloglucan Galactosyltransferase Is an Indispensable Salt Stress Tolerance Determinant in *Arabidopsis*. Molecular Plant. 6 (4): 1344-1354.
- → Hai Liu, Defeng Shen, Shenghua Jia, Wenbo Li, Jin Liu, Shengcheng Han and Yingdian Wang. (2012). Microarray-based Screening of the MicroRNAs Associated with Caryopsis Development in *Oryza sativa*. Biologia Plantarum. 57(2): 255-261.

- ❖ Xiaojin Zhou, Jie Li, Wei Chen, Hai Liu, Mengmeng Li, Yuan Zhang, Wenbo Li, Shengcheng Han and Yingdian Wang. (2010). Gene Structure Analysis of Rice ADPribosylation Factors (OsARFs) and their mRNA Expression in Developing Rice Plants. Plant Molecular Biology Reporter. 28: 692-703.

#### **TECHNICAL EXPERTISE**

♦ Wetlab skills: CRISPR CAS9, DNA and RNA extraction, PCR, standard RT-PCR, semi-quantitative RT-PCR, map-based cloning, Northern Blotting, Western Blotting, DNA cloning, plasmid construction, Yeast Two-Hybrid, transformation, cell culture, cell transfection, stem cell culture, *in vitro* embryo culture, protein isolation and purification, UHPLC analysis, GC-MS analysis, spectrophotometry, fluorescence

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- microscopy, confocal microscopy, flow cytometry, mouse breeding and husbandry, and animal dissection.

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