Junli Zhang

POSTDOCTORAL SCHOLAR · UC DAVIS

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Education

Department of Plant Sciences, University of California, Davis

POSTDOCTORAL SCHOLAR IN DUBCOVSKY'S LAB

Davis, California, USA

Nov 2013 - Present

Department of PSES, University of Idaho

PH.D. IN PLANT SCIENCE

Moscow, Idaho, USA

Aug 2008 - Dec 2013

College of Agronomy, Northwest A&F University

B.S. IN AGRONOMY

Yangling, Shaanxi, China Sep. 2004 - Jul. 2008

Skills

Basic Programming Python, Perl, Markdown, LaTeX, R, SAS, Shell Script

Quantitative Genetics Linkage mapping, association mapping, fine mapping

Lab Skills Common lab procedures used in molecular genetics **Statistics** Good at experiment design and statistical analysis **Bioinformatics** Shell Script, Slurm usage, blast, exome capture

Software Microsoft Office, R, SAS, MapMaker 3.0b, QTL Cartographer, TASSEL, GAPIT, R/qtl

Languages English, Chinese

Experience _____

Dubcovsky's lab, Department of Plant Sciences, University of California, Davis

Davis, CA, USA

POSTDOCTORAL SCHOLAR

Nov. 2013 - Present

- Genome-wide Association Study on Water Use Efficiency and Yield Components of Wheat Using an Association Mapping Panel and Nested Association Mapping Panel
- Genome-wide Association Mapping of Stripe Rust Resistance in Spring Hexaploid Wheat from National Small Grains Collection (NSGC)

Chen's Lab, Department of PSES, University of Idaho

Moscow, ID, USA

RESEARCH ASSISTANT

Aug 2008 - Oct 2013

- Association Mapping of Hagberg Falling Number in Hard White Spring Wheat
- Water and Nitrogen Use Efficiency Study of Triticeae CAP (TCAP) Wheat and Barley Collections (Canopy spectrum reflectance and genome-wide association mapping)
- Quantitative Trait Locus (QTL) Mapping of Grain Yield and Physiological Traits Using a Population of Recombinant Inbred Lines of Common Wheat
- Study on Barley Yellow Dwarf Virus (BYDV) Resistance of Bread Wheat by RNAi Technique Using Both Virus-Induced Gene Silencing (VIGS) and Transgenic Methods

Honors & Awards

2011-2012 Easton, Gene and Marlene PSES Scholarship

2010-2011 Pure Line Scholarship & Toevs, John L. & Lois K. Scholarship

University of Idaho University of Idaho

Professional Affiliation

Since 2011 American Society of Agronomy (ASA)
Since 2011 Crop Science Society of America (CSSA)

USA

USA

Publications

8 Google Scholar Citation Metrics

R⁶ ResearchGate Profile

2016

- Bulli, P., **J. Zhang**, S. Chao, X. Chen, and M. Pumphrey. 2016. Genetic architecture of resistance to stripe rust in a global winter wheat germplasm collection. G3 6(8): 2237–2253.
- Chen, J., M.J. Guttieri, **J. Zhang**, D. Hole, E. Souza, and B. Goates. 2016a. A novel QTL associated with dwarf bunt resistance in Idaho 444 winter wheat. Theor Appl Genet: 1–10.
- Chen, J., J. Wheeler, K. O'Brien, W. Zhao, N. Klassen, **J. Zhang**, B. Bowman, Y. Wang, C. Jackson, J.M. Marshall, and X.M. Chen. 2016b. Registration of "UI Platinum" Hard White Spring Wheat. Journal of Plant Registrations 10(1): 36–40.

2015

- Bonman, J.M., E.M. Babiker, A. Cuesta-Marcos, K. Esvelt-Klos, G. Brown-Guedira, S. Chao, D. See, J. Chen, E. Akhunov, **J. Zhang**, H.E. Bockelman, and T.C. Gordon. 2015. Genetic diversity among wheat accessions from the USDA National Small Grains Collection. Crop Science 55(3): 1243–1253.
- Bowman, B.C., J. Chen, **J. Zhang**, J. Wheeler, Y. Wang, W. Zhao, S. Nayak, N. Heslot, H. Bockelman, and J.M. Bonman. 2015. Evaluating grain yield in spring wheat with canopy spectral reflectance. Crop Science 55(5): 1881–1890.
- Maccaferri, M., J. Zhang, P. Bulli, Z. Abate, S. Chao, D. Cantu, E. Bossolini, X. Chen, M. Pumphrey, and J. Dubcovsky. 2015. A genome-wide association study of resistance to stripe rust (*Puccinia striiformis* f. sp. *tritici*) in a worldwide collection of hexaploid spring wheat (*Triticum aestivum* L.). G3 5(3): 449–465. (First 3 authors contributed equally)

2014

- **Zhang, J.**, J. Chen, B.C. Bowman, K. O'Brien, J.M. Marshall, and J.M. Bonman. 2014a. Association mapping of Hagberg falling number in hard white spring wheat. Crop Science 54(3): 1243–1252.
- **Zhang, J.**, J. Chen, C. Chu, W. Zhao, J. Wheeler, E.J. Souza, and R.S. Zemetra. 2014b. Genetic dissection of QTL associated with grain yield in diverse environments. Agronomy 4(4): 556–578.
- Chen, J., G. Hu, **J. Zhang**, C. Chu, and Y. Wu. 2014. Mapping of STS markers developed from drought tolerance candidate genes and preliminary analysis of their association with yield-related traits in common wheat (*Triticum aestivum*). Cereal Research Communications 42(2): 199–208.

2013

• Chen, J., J. Wheeler, J. Clayton, W. Zhao, K. O'Brien, **J. Zhang**, C. Jackson, J.M. Marshall, B.D. Brown, K. Campbell, X.M. Chen, R. Zemetra, and E.J. Souza. 2013. Registration of "UI Stone" soft white spring wheat. Journal of Plant Registrations 7(3): 321–326.

2011

• Li, P., J. Chen, P. Wu, **J. Zhang**, C. Chu, D. See, G. Brown-Guedira, R. Zemetra, and E. Souza. 2011. Quantitative trait loci analysis for the effect of *Rht-B1* dwarfing gene on coleoptile length and seedling root length and number of bread wheat. Crop Sci. 51(6): 2561–2568.