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EDUCATION

- 2015-present Ph.D. **Plant Genetics and Breeding**, China Agricultural University, Beijing, CHN Supervised by Dr. Patrick Schnable
- 2010-2013 Master Plant Genetics and Breeding, China Agricultural University, Beijing,
 CHN Supervised by Dr. Patrick Schnable
- 2006-2010 B.A. Biology Science, Inner Mongolia Agricultural University, Hohhot, CHN

Publications

- Li L, Li D, Liu S, Ma X, Dietrich CR, Hu H-C, et al. (2013) The Maize glossy13 Gene, Cloned via BSR-Seq and Seq-Walking Encodes a Putative ABC Transporter Required for the Normal Accumulation of Epicuticular Waxes. PLoS ONE 8(12): e82333. doi:10.1371/journal.pone.0082333
- 2. Nestler J, Liu S, Wen TJ, Paschold A, *et al.* (2014)Roothairless5, which functions in maize (*Zea mays L.*) root hair initiation and elongation encodes a monocot-specific NADPH oxidase. The Plant Journal.79(5):729-40. doi: 10.1111/tpj.12578.
- 3. Wang Z, Cui Y, Chen Y, Zhang D, et al. (2014) Comparative genetic mapping and genomic region collinearity analysis of the powdery mildew resistance gene *Pm41*. Theor Appl. Genet. 127:1741-1751
- Liu Z, Lu P, Liang Y, Li D et al. (2015) Fine genetic mapping of spot blotch resistance gene Sb2 caused by Bipolaris sorokiniana in wheat (*Triticum aestivum*). Theor Appl. Genet. In Press
- Wu Q, Chen Y, Zhou S, Fu L, Chen J, et al. (2015) High-Density Genetic Linkage Map Construction and QTL Mapping of Grain Shape and Size in the Wheat Population Yanda1817 × Beinong6. PLoS ONE 10(2): e0118144. doi: 10.1371/journal.pone.0118144

Experience

- 2015-Present Senior Bioinformatics Engineer. <u>Data2Bio</u>
 - GWAS
 - Fast Mapping
- 2013-2015 Bioinformatics Engineer. Data2Bio

- GWAS
- Fast Mapping

Technical

- 1. Perl
- 2. R
- 3. Linux