

外国留学生研究生指导教师情况表 Resume of Supervisor (中英文版)

导师姓名: Name of supervisor:	刘登才 Deng-Cai Liu	导师类别: Supervisor Level:	博导 <input checked="" type="checkbox"/> 硕导 <input type="checkbox"/> Doctor <input checked="" type="checkbox"/> Master
院所 College/Institute:	小麦研究所 Triticeae Research Institute		
学科 Discipline:	作物学 Crop Science		
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拟接收留学生层次及人数 Levels and Numbers of International Students (此项不上网)	<input type="checkbox"/> 博士留学生__1__名; <input type="checkbox"/> 硕士留学生__1__名 Doctor Candidates __1__ persons; Master Candidates __1__ persons		
研究方向: Research Fields	小麦遗传育种 Wheat genetics and breeding		
教育背景: Educational Background:	09/1995-06/1998 博士, 四川农业大学, 作物遗传育种 09/1995-06/1998 Ph.D. in Crop genetics and breeding, Sichuan Agricultural University 09/1992-06/1995 硕士, 四川农业大学, 作物遗传育种 09/1988-06/1992 M.S. in Crop genetics and breeding, Sichuan Agricultural University 09/1988-06/1992 学士, 四川农业大学, 农学 09/1988-06/1992 B.S. in Agricultural, Sichuan Agricultural University		
工作经历: Professional Experience:	01/2004-现在 研究员, 四川农业大学小麦研究所 01/2004-present Professor, Triticeae Research Institute, Sichuan Agricultural University 10/2002-09/2003 访问学者, 南达科他州立大学 10/2002-09/2003 Visiting scholar, South Dakota State University in USA 11/2000-12/2003 副研究员, 四川农业大学小麦研究所 11/2000-12/2003 Associate Professor, Triticeae Research Institute, Sichuan		

	<p>Agricultural University</p> <p>11/1997-10/2000 助理研究员，四川农业大学小麦研究所</p> <p>12/2007-12/2010 Research Associate, Triticeae Research Institute, Sichuan Agricultural University</p> <p>07/1995-10/1997 研究实习员，四川农业大学小麦研究所</p> <p>07/1995-10/1997 Research Intern, Triticeae Research Institute, Sichuan Agricultural University</p>
<p>主要论著（10 篇代表论著） Publications</p>	<ol style="list-style-type: none"> 1. Making the bread: insights from newly synthesized allohexaploid Wheat. Molecular Plant, 2015, DOI: http://dx.doi.org/10.1016/j.molp.2015.02.016 2. Distant Hybridization: A tool for interspecific manipulation of chromosomes. In Alien Gene Transfer in Crop Plants, Volume 1 (pp. 25-42). Springer New York, 2014. 3. mRNA and small RNA transcriptomes reveal insights into dynamic homoeolog regulation of allopolyploid heterosis in nascent hexaploid wheat. Plant Cell (2014) 26: 1878–1900. 4. <i>QTug. sau-3B</i> Is a Major Quantitative Trait Locus for Wheat Hexaploidization. G3: Genes Genomes Genetics 2014 4(10): 1943-1953. 5. The detection of a de novo allele of the Glu-1Dx gene in wheat-rye hybrid offspring. Theor Appl Genet (2014) 127:2173–2182. 6. Production of hexaploid triticale by a synthetic hexaploid wheat-rye hybrid method. Euphytica, 193:347-357, 2013. 7. Stripe rust resistance in <i>Aegilops tauschii</i> germplasm. Crop Science, 53:2014-2020, 2013. 8. Microsatellite mutation rate during allohexaploidization of newly resynthesized wheat. International Journal of Molecular Sciences, 13:12533-12543, 2012. 9. Genetic map of <i>Triticum turgidum</i> based on a hexaploid wheat population without genetic recombination for D genome. BMC Genetics, 13:69, 2012. 10. Mitotic illegitimate recombination is a mechanism for novel changes in high-molecular-weight glutenin subunits in wheat-rye hybrids. PLoS ONE, 6(8): e23511, 2011.

<p>主要国际学术活动(5项以内): International Academic Activities:</p>	<ol style="list-style-type: none"> 1. Genetic variations during the extraction of bread wheat genome from wheat-rye hybrids. International Conference of Plant Chromosome Engineering and Functional Genomics for Breeding, 2015.06.29-07.01, Oral presentation (大会报告) 2. Chromosome translocation of wide hybrids in Triticeae is species-dependent. 7th International Triticeae Symposium, 2013.06.09-13, Abstract (会议摘要) 3. Amphitelic orientation of centromeres at metaphase I is an important feature for univalent-dependent meiotic nonreduction. 7th International Triticeae Symposium, 2013.06.09-13, Abstract (会议摘要) 4. Chromosomal Variation in Synthetic Hexaploid-Wheat. International Conference of Plant Chromosome Engineering and Functional Genomics for Breeding, 2015.06.29-07.01, Oral presentation (会议摘要) 5. Chromosome Variations Derived from the Hybridization between Common Wheat and <i>Aegilops variabilis</i>. International Conference of Plant Chromosome Engineering and Functional Genomics for Breeding, 2015.06.29-07.01, Abstract (会议摘要)
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