

# Zujun Yang

## List of Publications by Year in descending order

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113  
papers

3,132  
citations

186265

28  
h-index

189892

50  
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115  
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115  
docs citations

115  
times ranked

2129  
citing authors

#	ARTICLE	IF	CITATIONS
1	Fine mapping and candidate gene analysis of dwarf gene Rht14 in durum wheat ( <i>Triticum durum</i> ). <i>Functional and Integrative Genomics</i> , 2022, 22, 141.	3.5	7
2	Molecular and Cytogenetic Identification of Stem Rust Resistant Wheatâ€™ <i>Thinopyrum intermedium</i> Introgression Lines. <i>Plant Disease</i> , 2022, 106, 2447-2454.	1.4	4
3	Oligoâ€™FISH Paints in Triticeae. <i>Current Protocols</i> , 2022, 2, e364.	2.9	7
4	Wheat- <i>Thinopyrum</i> Substitution Lines Imprint Compensation Both From Recipients and Donors. <i>Frontiers in Plant Science</i> , 2022, 13, 837410.	3.6	1
5	Genetic Incorporation of the Favorable Alleles for Three Genes Associated With Spikelet Development in Wheat. <i>Frontiers in Plant Science</i> , 2022, 13, 892642.	3.6	2
6	Genetic Diversity Assessment of the International Maize and Wheat Improvement Center and Chinese Wheat Core Germplasm by Non-Denaturing Fluorescence In Situ Hybridization. <i>Plants</i> , 2022, 11, 1403.	3.5	4
7	Characterization of a Wheat- <i>Dasypyrum breviaristatum</i> Chromosome Addition and Its Derived Progenies Carrying Novel <i>Dasypyrum</i> -Specific Gliadin Genes. <i>Agronomy</i> , 2022, 12, 1673.	3.0	1
8	An efficient Oligoâ€™FISH painting system for revealing chromosome rearrangements and polyploidization in Triticeae. <i>Plant Journal</i> , 2021, 105, 978-993.	5.7	58
9	Genetic Dissection of Adult Plant Resistance to Sharp Eyespot Using an Updated Genetic Map of Niavt14 Å– Xuzhou25 Winter Wheat Recombinant Inbred Line Population. <i>Plant Disease</i> , 2021, 105, 997-1005.	1.4	7
10	Physical mapping of chromosome 7J and a purple coleoptile gene from <i>Thinopyrum intermedium</i> in the common wheat background. <i>Planta</i> , 2021, 253, 22.	3.2	3
11	Molecular Cytogenetic and Agronomic Characterization of the Similarities and Differences Between Wheatâ€™ <i>Leymus mollis</i> Trin. and Wheatâ€™ <i>Psathyrostachys huashanica</i> Keng 3Ns (3D) Substitution Lines. <i>Frontiers in Plant Science</i> , 2021, 12, 644896.	3.6	3
12	A universal karyotypic system for hexaploid and diploid <i>Avena</i> species brings oat cytogenetics into the genomics era. <i>BMC Plant Biology</i> , 2021, 21, 213.	3.6	19
13	Development of Sequence-Tagged Site Marker Set for Identification of J, JS, and St Sub-genomes of <i>Thinopyrum intermedium</i> in Wheat Background. <i>Frontiers in Plant Science</i> , 2021, 12, 685216.	3.6	9
14	Characterization, Identification and Evaluation of Wheat- <i>Aegilops sharonensis</i> Chromosome Derivatives. <i>Frontiers in Plant Science</i> , 2021, 12, 708551.	3.6	0
15	Karyotyping <i>Dasypyrum breviaristatum</i> chromosomes with multiple oligonucleotide probes reveals the genomic divergence in <i>Dasypyrum</i> . <i>Genome</i> , 2021, 64, 789-800.	2.0	5
16	<i>Zanthoxylum</i> -specific whole genome duplication and recent activity of transposable elements in the highly repetitive paleotetraploid <i>Z. bungeanum</i> genome. <i>Horticulture Research</i> , 2021, 8, 205.	6.3	19
17	Molecular characteristics and inheritance of a chromosome segment from <i>Psathyrostachys huashanica</i> Keng in a wheat background. <i>Genetic Resources and Crop Evolution</i> , 2020, 67, 1245-1257.	1.6	5
18	Extraordinarily conserved chromosomal synteny of <i>Citrus</i> species revealed by chromosome-specific painting. <i>Plant Journal</i> , 2020, 103, 2225-2235.	5.7	33

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19	Sympatric speciation of wild emmer wheat driven by ecology and chromosomal rearrangements. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 5955-5963.	7.1	39
20	Identification and characterization of a new stripe rust resistance gene Yr83 on rye chromosome 6R in wheat. <i>Theoretical and Applied Genetics</i> , 2020, 133, 1095-1107.	3.6	136
21	Molecular cytogenetic characterization of a novel wheatâ€“ <i>Psathyrostachys huashanica</i> Keng T3DS-5NsLâ€“5NsS and T5DL-3DSâ€“3DL dual translocation line with powdery mildew resistance. <i>BMC Plant Biology</i> , 2020, 20, 163.	3.6	12
22	Molecular dissection of <i>Secale africanum</i> chromosome 6Rafr in wheat enabled localization of genes for resistance to powdery mildew and stripe rust. <i>BMC Plant Biology</i> , 2020, 20, 134.	3.6	14
23	Comparative FISH and molecular identification of new stripe rust resistant wheat- <i>Thinopyrum intermedium</i> ssp. <i>trichophorum</i> introgression lines. <i>Crop Journal</i> , 2019, 7, 819-829.	5.2	15
24	Identification of QTLs for Stripe Rust Resistance in a Recombinant Inbred Line Population. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3410.	4.1	32
25	Diversified Chromosome Rearrangements Detected in a Wheatâ€“ <i>Dasypyrum breviaristatum</i> Substitution Line Induced by Gamma-Ray Irradiation. <i>Plants</i> , 2019, 8, 175.	3.5	14
26	Molecular cytogenetic characterization of a novel wheatâ€“ <i>Psathyrostachys huashanica</i> Keng 5Ns (5D) disomic substitution line with stripe rust resistance. <i>Molecular Breeding</i> , 2019, 39, 1.	2.1	20
27	Genome-Wide Distribution of Novel Ta-3A1 Mini-Satellite Repeats and Its Use for Chromosome Identification in Wheat and Related Species. <i>Agronomy</i> , 2019, 9, 60.	3.0	8
28	New ND-FISH-Positive Oligo Probes for Identifying <i>Thinopyrum</i> Chromosomes in Wheat Backgrounds. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2031.	4.1	27
29	Characterization, identification and evaluation of a set of wheat- <i>Aegilops comosa</i> chromosome lines. <i>Scientific Reports</i> , 2019, 9, 4773.	3.3	24
30	A novel QTL QTrl.saw-2D.2 associated with the total root length identified by linkage and association analyses in wheat ( <i>Triticum aestivum</i> L.). <i>Planta</i> , 2019, 250, 129-143.	3.2	30
31	Genetic diversity in common wheat lines revealed by fluorescence in situ hybridization. <i>Plant Systematics and Evolution</i> , 2019, 305, 247-254.	0.9	9
32	Characterization of Chromosomal Rearrangement in New Wheatâ€“ <i>Thinopyrum intermedium</i> Addition Lines Carrying <i>Thinopyrum</i> â€“Specific Grain Hardness Genes. <i>Agronomy</i> , 2019, 9, 18.	3.0	28
33	Physical location of tandem repeats in the wheat genome and application for chromosome identification. <i>Planta</i> , 2019, 249, 663-675.	3.2	57
34	Precise identification of wheat â€“ <i>Thinopyrum intermedium</i> translocation chromosomes carrying resistance to wheat stripe rust in line Z4 and its derived progenies. <i>Genome</i> , 2018, 61, 177-185.	2.0	40
35	Characterization of New Wheat- <i>Dasypyrum breviaristatum</i> Introgression Lines with Superior Gene(s) for Spike Length and Stripe Rust Resistance. <i>Cytogenetic and Genome Research</i> , 2018, 156, 117-125.	1.1	6
36	Mutation Landscape of Base Substitutions, Duplications, and Deletions in the Representative Current Cholera Pandemic Strain. <i>Genome Biology and Evolution</i> , 2018, 10, 2072-2085.	2.5	7

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37	Molecular Cytogenetic Characterization of New Wheatâ€”Dasypyrum breviaristatum Introgression Lines for Improving Grain Quality of Wheat. <i>Frontiers in Plant Science</i> , 2018, 9, 365.	3.6	8
38	Developing New Oligo Probes to Distinguish Specific Chromosomal Segments and the A, B, D Genomes of Wheat ( <i>Triticum aestivum</i> L.) Using ND-FISH. <i>Frontiers in Plant Science</i> , 2018, 9, 1104.	3.6	82
39	QTL Mapping and Validation of Adult Plant Resistance to Stripe Rust in Chinese Wheat Landrace Humai 15. <i>Frontiers in Plant Science</i> , 2018, 9, 968.	3.6	54
40	Introduction of <i>Thinopyrum intermedium</i> ssp. <i>trichophorum</i> chromosomes to wheat by trigeneric hybridization involving <i>Triticum</i> , <i>Secale</i> and <i>Thinopyrum</i> genera. <i>Planta</i> , 2017, 245, 1121-1135.	3.2	30
41	Molecular cytogenetic characterization and phenotypic evaluation of new wheatâ€”rye lines derived from hexaploid triticales â€”Certaâ€™™—A common wheat hybrids. <i>Plant Breeding</i> , 2017, 136, 809-819.	1.9	6
42	The <i>Aegilops tauschii</i> genome reveals multiple impacts of transposons. <i>Nature Plants</i> , 2017, 3, 946-955.	9.3	164
43	Characterization of Stripe Rust Resistance Genes in the Wheat Cultivar Chuanmai45. <i>International Journal of Molecular Sciences</i> , 2016, 17, 601.	4.1	39
44	The Chinese bread wheat cultivar Xiaoyanmai 7 harbours genes encoding a pair of novel high-molecular-weight glutenin subunits inherited from cereal rye. <i>Crop and Pasture Science</i> , 2016, 67, 29.	1.5	1
45	Molecular Mapping of the Stripe Rust Resistance Gene <i>Yr69</i> on Wheat Chromosome 2AS. <i>Plant Disease</i> , 2016, 100, 1717-1724.	1.4	41
46	Molecular cytogenetic characterization of <i>Dasypyrum breviaristatum</i> chromosomes in wheat background revealing the genomic divergence between <i>Dasypyrum</i> species. <i>Molecular Cytogenetics</i> , 2016, 9, 6.	0.9	43
47	New molecular markers and cytogenetic probes enable chromosome identification of wheat- <i>Thinopyrum intermedium</i> introgression lines for improving protein and gluten contents. <i>Planta</i> , 2016, 244, 865-876.	3.2	45
48	Characterization of wheat- <i>Secale africanum</i> chromosome 5Ra derivatives carrying <i>Secale</i> specific genes for grain hardness. <i>Planta</i> , 2016, 243, 1203-1212.	3.2	15
49	Heat Shock Proteins in Wild Barley at â€œEvolution Canyonâ€, Mount Carmel, Israel. <i>Heat Shock Proteins</i> , 2016, , 79-102.	0.2	2
50	Oligonucleotide Probes for ND-FISH Analysis to Identify Rye and Wheat Chromosomes. <i>Scientific Reports</i> , 2015, 5, 10552.	3.3	138
51	Molecular and Cytogenetic Characterization of New Wheatâ€” <i>Dasypyrum breviaristatum</i> Derivatives with Post-Harvest Re-Growth Habit. <i>Genes</i> , 2015, 6, 1242-1255.	2.4	10
52	Identification of Novel Chromosomal Aberrations Induced by <sup>60</sup> Co-Î³-Irradiation in Wheat- <i>Dasypyrum villosum</i> Lines. <i>International Journal of Molecular Sciences</i> , 2015, 16, 29787-29796.	4.1	24
53	Transcriptome Analysis of Interspecific Hybrid between <i>Brassica napus</i> and <i>B. rapa</i> Reveals Heterosis for Oil Rape Improvement. <i>International Journal of Genomics</i> , 2015, 2015, 1-11.	1.6	20
54	Molecular and Cytogenetic Characterization of a Powdery Mildew-Resistant Wheat- <i>Aegilops mutica</i> Partial Amphiploid and Addition Line. <i>Cytogenetic and Genome Research</i> , 2015, 147, 186-194.	1.1	8

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55	Precise identification of two wheat–Thinopyrum intermedium substitutions reveals the compensation and rearrangement between wheat and Thinopyrum chromosomes. <i>Molecular Breeding</i> , 2015, 35, 1.	2.1	150
56	MDC-Analyzer-facilitated combinatorial strategy for improving the activity and stability of halohydrin dehalogenase from <i>Agrobacterium radiobacter</i> AD1. <i>Journal of Biotechnology</i> , 2015, 206, 1-7.	3.8	11
57	Molecular Characterization of Sec2 Loci in Wheat–Secale africanum Derivatives Demonstrates Genomic Divergence of Secale Species. <i>International Journal of Molecular Sciences</i> , 2015, 16, 8324-8336.	4.1	4
58	Molecular Characterization of a New Wheat–Thinopyrum intermedium Translocation Line with Resistance to Powdery Mildew and Stripe Rust. <i>International Journal of Molecular Sciences</i> , 2015, 16, 2162-2173.	4.1	25
59	Improvement of the thermostability and activity of halohydrin dehalogenase from <i>Agrobacterium radiobacter</i> AD1 by engineering C-terminal amino acids. <i>Journal of Biotechnology</i> , 2015, 212, 92-98.	3.8	10
60	Cytogenetic and molecular markers for detecting <i>Aegilops uniaristata</i> chromosomes in a wheat background. <i>Genome</i> , 2014, 57, 489-497.	2.0	17
61	A Novel Wheat–Dasypyrum breviaristatum Substitution Line with Stripe Rust Resistance. <i>Cytogenetic and Genome Research</i> , 2014, 143, 280-287.	1.1	18
62	Characterization and phylogenetic analysis of $\alpha$ -gliadin gene sequences reveals significant genomic divergence in Triticeae species. <i>Journal of Genetics</i> , 2014, 93, 725-731.	0.7	4
63	Adaptive evolution of duplicated hsp17 genes in wild barley from microclimatically divergent sites of Israel. <i>Genetics and Molecular Research</i> , 2014, 13, 1220-1232.	0.2	8
64	New St-chromosome-specific molecular markers for identifying wheat–Thinopyrum intermedium derivative lines. <i>Journal of Genetics</i> , 2014, 93, 69-74.	0.7	14
65	DC-Analyzer-facilitated combinatorial strategy for rapid directed evolution of functional enzymes with multiple mutagenesis sites. <i>Journal of Biotechnology</i> , 2014, 192, 102-107.	3.8	6
66	Oligonucleotides replacing the roles of repetitive sequences pAs1, pSc119.2, pTa-535, pTa71, CCS1, and pAWRC.1 for FISH analysis. <i>Journal of Applied Genetics</i> , 2014, 55, 313-318.	1.9	342
67	Chromosomal Location and Comparative Genomics Analysis of Powdery Mildew Resistance Gene Pm51 in a Putative Wheat–Thinopyrum ponticum Introgression Line. <i>PLoS ONE</i> , 2014, 9, e113455.	2.5	70
68	Development and utilization of new sequenced characterized amplified region markers specific for E genome of Thinopyrum. <i>Frontiers in Biology</i> , 2013, 8, 451-459.	0.7	1
69	Identification of wheat–Secale africanum chromosome 2Rafr introgression lines with novel disease resistance and agronomic characteristics. <i>Euphytica</i> , 2013, 194, 197-205.	1.2	22
70	Transcriptional abundance is not the single force driving the evolution of bacterial proteins. <i>BMC Evolutionary Biology</i> , 2013, 13, 162.	3.2	5
71	Evaluation of wheat seeds by terahertz imaging. , 2013, , .		3
72	Putative Thinopyrum intermedium-derived stripe rust resistance gene Yr50 maps on wheat chromosome arm 4BL. <i>Theoretical and Applied Genetics</i> , 2013, 126, 265-274.	3.6	104

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73	Introgression of a novel <i>Thinopyrum intermedium</i> St-chromosome-specific HMW-GS gene into wheat. <i>Molecular Breeding</i> , 2013, 31, 843-853.	2.1	29
74	Transcriptome Comparative Profiling of Barley <i>eibi1</i> Mutant Reveals Pleiotropic Effects of HvABCG31 Gene on Cuticle Biogenesis and Stress Responsive Pathways. <i>International Journal of Molecular Sciences</i> , 2013, 14, 20478-20491.	4.1	9
75	The <i>CentO</i> satellite confers translational and rotational phasing on cenH3 nucleosomes in rice centromeres. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E4875-83.	7.1	80
76	Molecular identification of a new wheat- <i>Thinopyrum intermedium</i> ssp. <i>trichophorum</i> addition line for resistance to stripe rust. <i>Cereal Research Communications</i> , 2013, 41, 211-220.	1.6	10
77	Isolation and phylogenetic analysis of novel $\beta$ -gliadin genes in genus <i>Dasyphyrum</i> . <i>Genetics and Molecular Research</i> , 2013, 12, 783-790.	0.2	0
78	Determination of Nine Microelements in <i>Nostoc commune</i> Vauch by ICP-AES. <i>Advanced Materials Research</i> , 2012, 518-523, 5020-5023.	0.3	2
79	Chromosomal distribution of a new centromeric Ty3-gypsy retrotransposon sequence in <i>Dasyphyrum</i> and related Triticeae species. <i>Journal of Genetics</i> , 2012, 91, 343-348.	0.7	8
80	Characterization of wheat <i>Secale africanum</i> introgression lines reveals evolutionary aspects of chromosome 1R in rye. <i>Genome</i> , 2012, 55, 765-774.	2.0	28
81	Characterization of a new T2DS.2DL-7R translocation triticale ZH-1 with multiple resistances to diseases. <i>Genetic Resources and Crop Evolution</i> , 2012, 59, 1161-1168.	1.6	18
82	Isolation and chromosomal localization of new MITE-like sequences from <i>Secale</i> . <i>Biologia (Poland)</i> , 2012, 67, 126-131.	1.5	3
83	Genomic rearrangement between wheat and <i>Thinopyrum elongatum</i> revealed by mapped functional molecular markers. <i>Genes and Genomics</i> , 2012, 34, 67-75.	1.4	31
84	New St-chromosome-specific molecular markers for identifying wheat- <i>Thinopyrum intermedium</i> derivative lines. <i>Journal of Genetics</i> , 2012, 91, e69-74.	0.7	10
85	Diversity and evolution of four dispersed repetitive DNA sequences in the genus <i>Secale</i> . <i>Genome</i> , 2011, 54, 285-300.	2.0	8
86	Characterization of wheat yellow rust resistance gene <i>Yr17</i> using EST-SSR and rice syntenic region. <i>Cereal Research Communications</i> , 2011, 39, 88-99.	1.6	22
87	Adaptive microclimatic evolution of the dehydrin 6 gene in wild barley at "Evolution Canyon", Israel. <i>Genetica</i> , 2011, 139, 1429-1438.	1.1	13
88	Molecular cytogenetic identification of a new wheat- <i>Thinopyrum</i> substitution line with stripe rust resistance. <i>Euphytica</i> , 2011, 177, 169-177.	1.2	42
89	The chromosome number, karyotype and genome size of the desert plant diploid <i>Reaumuria soongorica</i> (Pall.) Maxim. <i>Plant Cell Reports</i> , 2011, 30, 955-964.	5.6	20
90	Molecular cytogenetic characterization of a new wheat <i>Secale africanum</i> 2Ra(2D) substitution line for resistance to stripe rust. <i>Journal of Genetics</i> , 2011, 90, 283-287.	0.7	18

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91	A new long terminal repeat (LTR) sequence allows to identify J genome from JS and St genomes of <i>Thinopyrum intermedium</i> . <i>Journal of Applied Genetics</i> , 2011, 52, 31-33.	1.9	15
92	Molecular characterization of a wheat - <i>Thinopyrum ponticum</i> partial amphiploid and its derived substitution line for resistance to stripe rust. <i>Journal of Applied Genetics</i> , 2011, 52, 279-285.	1.9	28
93	Molecular cytogenetic characterization of a new leaf rolling triticale. <i>Genetics and Molecular Research</i> , 2011, 10, 2953-2961.	0.2	5
94	Molecular and cytogenetic identification of new wheat- <i>Dasypyrum breviaristatum</i> additions conferring resistance to stem rust and powdery mildew. <i>Breeding Science</i> , 2011, 61, 366-372.	1.9	15
95	Diversified chromosomal distribution of tandemly repeated sequences revealed evolutionary trends in <i>Secale</i> (Poaceae). <i>Plant Systematics and Evolution</i> , 2010, 287, 49-56.	0.9	26
96	Genome relationships in the genus <i>Dasypyrum</i> : evidence from molecular phylogenetic analysis and in situ hybridization. <i>Plant Systematics and Evolution</i> , 2010, 288, 149-156.	0.9	17
97	New PCR based markers allowed to identify <i>Secale</i> chromatin in wheat- <i>Secale africanum</i> introgression lines. <i>Frontiers in Biology</i> , 2010, 5, 187-192.	0.7	1
98	Sequence variations of PDHA1 gene in Triticeae species allow for identifying wheat-alien introgression lines. <i>Frontiers of Agriculture in China</i> , 2010, 4, 137-144.	0.2	4
99	Characterization of a partial wheat- <i>Thinopyrum intermedium</i> amphiploid and its reaction to fungal diseases of wheat. <i>Hereditas</i> , 2010, 147, 304-312.	1.4	22
100	Molecular Characterization and Evolutionary Analysis of Alpha-gliadin Genes from <i>Eremopyrum bonaepartis</i> (Triticeae). <i>Journal of Agricultural Science</i> , 2010, 2, .	0.2	0
101	Identification, Classification and Phylogenetic Analysis of SET Domain Gene in Barley. International Conference on Bioinformatics and Biomedical Engineering: [proceedings] International Conference on Bioinformatics and Biomedical Engineering, 2010, , .	0.0	0
102	Identification of $\beta$ -gliadin genes in <i>Dasypyrum</i> in relation to evolution and breeding. <i>Euphytica</i> , 2009, 165, 155.	1.2	15
103	Molecular cytogenetic characterization of wheat- <i>Secale africanum</i> amphiploids and derived introgression lines with stripe rust resistance. <i>Euphytica</i> , 2009, 167, 197-202.	1.2	21
104	Inheritance and mapping of powdery mildew resistance gene Pm43 introgressed from <i>Thinopyrum intermedium</i> into wheat. <i>Theoretical and Applied Genetics</i> , 2009, 118, 1173-1180.	3.6	150
105	Adaptive microclimatic structural and expressional dehydrin 1 evolution in wild barley, <i>Hordeum spontaneum</i> , at "Evolution Canyon", Mount Carmel, Israel. <i>Molecular Ecology</i> , 2009, 18, 2063-2075.	3.9	30
106	Isolation of a new repetitive DNA sequence from <i>Secale africanum</i> enables targeting of <i>Secale</i> chromatin in wheat background. <i>Euphytica</i> , 2008, 159, 249-258.	1.2	49
107	Analysis of DNA methylation variation in wheat genetic background after alien chromatin introduction based on methylation-sensitive amplification polymorphism. <i>Science Bulletin</i> , 2008, 53, 58-69.	1.7	22
108	Investigation on the electronically excited state properties of multiwalled carbon nanotube (MDDA) in solution. <i>Science in China Series B: Chemistry</i> , 2006, 49, 97-102.	0.8	2

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109	Molecular cytogenetic characterization and disease resistance observation of wheat-Dasypyrum breviaristatum partial amphiploid and its derivatives. <i>Hereditas</i> , 2006, 142, 80-85.	1.4	44
110	Molecular Characterization of a HMW Glutenin Subunit Allele Providing Evidence for Silencing of x-type Gene on Glu-B1. <i>Journal of Genetics and Genomics</i> , 2006, 33, 929-936.	0.3	9
111	Studies on genome relationship and species-specific PCR marker for <i>Dasypyrum breviaristatum</i> in Triticeae. <i>Hereditas</i> , 2006, 143, 47-54.	1.4	37
112	Molecular characterization of high molecular weight glutenin subunit allele 1Bx23 in common wheat introduced from hexaploid triticale. <i>Hereditas</i> , 2006, 143, 159-166.	1.4	7
113	The Physical Location of Stripe Rust Resistance Genes on Chromosome 6 of Rye ( <i>Secale cereale</i> L.) AR106BONE. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	4