

Shuyu Liu

List of Publications by Year in descending order

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citations

293460

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83
docs citations

83
times ranked

2188
citing authors

#	ARTICLE	IF	CITATIONS
1	Population genomic analysis of <i>Aegilops tauschii</i> identifies targets for bread wheat improvement. <i>Nature Biotechnology</i> , 2022, 40, 422-431.	9.4	102
2	Development of the Wheat Practical Haplotype Graph database as a resource for genotyping data storage and genotype imputation. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	0.8	7
3	Assessment of floral characteristics for hybrid wheat (<i>Triticum aestivum</i> L.) production in Texas. , 2022, 5, .		3
4	Genomic variants affecting homoeologous gene expression dosage contribute to agronomic trait variation in allopolyploid wheat. <i>Nature Communications</i> , 2022, 13, 826.	5.8	31
5	A new strategy for using historical imbalanced yield data to conduct genome-wide association studies and develop genomic prediction models for wheat breeding. <i>Molecular Breeding</i> , 2022, 42, 1.	1.0	0
6	Genetic dissection of end-use quality traits in two widely adapted wheat cultivars TAM 111 and TAM 112. <i>Crop Science</i> , 2021, 61, 1944-1959.	0.8	9
7	Genomic selection of forage agronomic traits in winter wheat. <i>Crop Science</i> , 2021, 61, 410-421.	0.8	5
8	Thermal imaging to evaluate wheat genotypes under dryland conditions. , 2021, 4, e20152.		7
9	Characterization of wheat curl mite resistance gene <i>Cmc4</i> in OK05312. <i>Theoretical and Applied Genetics</i> , 2021, 134, 993-1005.	1.8	11
10	RNA-seq analysis reveals different drought tolerance mechanisms in two broadly adapted wheat cultivars TAM 111 and TAM 112. <i>Scientific Reports</i> , 2021, 11, 4301.	1.6	19
11	Population genomics and haplotype analysis in spelt and bread wheat identifies a gene regulating glume color. <i>Communications Biology</i> , 2021, 4, 375.	2.0	11
12	Function and evolution of allelic variations of <i>Sr13</i> conferring resistance to stem rust in tetraploid wheat (<i>Triticum turgidum</i> L.). <i>Plant Journal</i> , 2021, 106, 1674-1691.	2.8	15
13	Genome-wide QTL mapping of yield and agronomic traits in two widely adapted winter wheat cultivars from multiple mega-environments. <i>PeerJ</i> , 2021, 9, e12350.	0.9	6
14	Genetic Mapping of Quantitative Trait Loci for End-Use Quality and Grain Minerals in Hard Red Winter Wheat. <i>Agronomy</i> , 2021, 11, 2519.	1.3	8
15	Soil water extraction and use by winter wheat cultivars under limited irrigation in a semi-arid environment. <i>Journal of Arid Environments</i> , 2020, 174, 104046.	1.2	12
16	Genome wide identification of QTL associated with yield and yield components in two popular wheat cultivars TAM 111 and TAM 112. <i>PLoS ONE</i> , 2020, 15, e0237293.	1.1	17
17	RhizoVision Crown: An Integrated Hardware and Software Platform for Root Crown Phenotyping. <i>Plant Phenomics</i> , 2020, 2020, 3074916.	2.5	74
18	Use of NDVI for characterizing winter wheat response to water stress in a semi-arid environment. <i>Journal of Crop Improvement</i> , 2019, 33, 633-648.	0.9	29

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19	Drought-Stress Tolerance in Wheat Seedlings Conferred by Phenazine-Producing Rhizobacteria. <i>Frontiers in Microbiology</i> , 2019, 10, 1590.	1.5	39
20	Comparison of TaqMan, KASP and rhAmp SNP genotyping platforms in hexaploid wheat. <i>PLoS ONE</i> , 2019, 14, e0217222.	1.1	54
21	Using aerial imagery and digital photography to monitor growth and yield in winter wheat. <i>International Journal of Remote Sensing</i> , 2019, 40, 6905-6929.	1.3	5
22	Genomic Selection of Forage Quality Traits in Winter Wheat. <i>Crop Science</i> , 2019, 59, 2473-2483.	0.8	7
23	“TAM 204”™ Wheat, Adapted to Grazing, Grain, and Graze-out Production Systems in the Southern High Plains. <i>Journal of Plant Registrations</i> , 2019, 13, 377-382.	0.4	5
24	Genotype Imputation in Winter Wheat Using First-Generation Haplotype Map SNPs Improves Genome-Wide Association Mapping and Genomic Prediction of Traits. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 125-133.	0.8	22
25	Yield determination in winter wheat under different water regimes. <i>Field Crops Research</i> , 2019, 233, 80-87.	2.3	35
26	Developing KASP Markers on a Major Stripe Rust Resistance QTL in a Popular Wheat TAM 111 Using 90K Array and Genotyping-by-sequencing SNPs. <i>Crop Science</i> , 2019, 59, 165-175.	0.8	14
27	Physiological responses to water stress and yield of winter wheat cultivars differing in drought tolerance. <i>Journal of Agronomy and Crop Science</i> , 2018, 204, 347-358.	1.7	23
28	Canopy temperature depression at grain filling correlates to winter wheat yield in the U.S. Southern High Plains. <i>Field Crops Research</i> , 2018, 217, 11-19.	2.3	66
29	“TAM 114”™ Wheat, Excellent Bread-making Quality Hard Red Winter Wheat Cultivar Adapted to the Southern High Plains. <i>Journal of Plant Registrations</i> , 2018, 12, 367-372.	0.4	7
30	Mapping and KASP marker development for wheat curl mite resistance in “TAM 112” wheat using linkage and association analysis. <i>Molecular Breeding</i> , 2018, 38, 1.	1.0	30
31	Saturated Genetic Mapping of Wheat Streak Mosaic Virus Resistance Gene <i>Wsm2</i> in Wheat. <i>Crop Science</i> , 2017, 57, 332-339.	0.8	13
32	More Recent Wheat Cultivars Extract More Water from Greater Soil Profile Depths to Increase Yield in the Texas High Plains. <i>Agronomy Journal</i> , 2017, 109, 2771-2780.	0.9	17
33	Wheat Curl Mite Resistance in Hard Winter Wheat in the US Great Plains. <i>Crop Science</i> , 2017, 57, 53-61.	0.8	18
34	Development and Validation of KASP Markers for Wheat Streak Mosaic Virus Resistance Gene <i>Wsm2</i> . <i>Crop Science</i> , 2017, 57, 340-349.	0.8	25
35	Development and validation of KASP markers for the greenbug resistance gene Gb7 and the Hessian fly resistance gene H32 in wheat. <i>Theoretical and Applied Genetics</i> , 2017, 130, 1867-1884.	1.8	60
36	Mapping of quantitative trait loci for grain yield and its components in a US popular winter wheat TAM 111 using 90K SNPs. <i>PLoS ONE</i> , 2017, 12, e0189669.	1.1	55

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37	Phenotypic Plasticity of Winter Wheat Heading Date and Grain Yield across the US Great Plains. <i>Crop Science</i> , 2016, 56, 2223-2236.	0.8	75
38	Validation of Chromosomal Locations of 90K Array Single Nucleotide Polymorphisms in US Wheat. <i>Crop Science</i> , 2016, 56, 364-373.	0.8	26
39	Spectral Reflectance Models for Characterizing Winter Wheat Genotypes. <i>Journal of Crop Improvement</i> , 2016, 30, 176-195.	0.9	6
40	Cooler Canopy Contributes to Higher Yield and Drought Tolerance in New Wheat Cultivars. <i>Crop Science</i> , 2014, 54, 2275-2284.	0.8	22
41	Yield Determination and Water Use Efficiency of Wheat under Water Limited Conditions in the U.S. Southern High Plains. <i>Crop Science</i> , 2014, 54, 34-47.	0.8	74
42	Molecular Mapping of Stripe Rust Resistance in Hard Red Winter Wheat TAM 111 Adapted to the U.S. High Plains. <i>Crop Science</i> , 2014, 54, 1361-1373.	0.8	50
43	Mapping Net Blotch Resistance in 'Nomini'™ and Clho 2291 Barley. <i>Crop Science</i> , 2014, 54, 2596-2602.	0.8	23
44	Characterization of Fusarium Head Blight Resistance and Deoxynivalenol Accumulation in Hulled and Hulless Winter Barley. <i>Plant Disease</i> , 2014, 98, 599-606.	0.7	13
45	Physiology and transcriptomics of water-deficit stress responses in wheat cultivars TAM 111 and TAM 112. <i>Journal of Plant Physiology</i> , 2014, 171, 1289-1298.	1.6	52
46	Molecular Markers Linked to Important Genes in Hard Winter Wheat. <i>Crop Science</i> , 2014, 54, 1304-1321.	0.8	55
47	Molecular characterization of field resistance to Fusarium head blight in two US soft red winter wheat cultivars. <i>Theoretical and Applied Genetics</i> , 2013, 126, 2485-2498.	1.8	59
48	Marker-trait associations in Virginia Tech winter barley identified using genome-wide mapping. <i>Theoretical and Applied Genetics</i> , 2013, 126, 693-710.	1.8	78
49	Identification and mapping of adult plant stripe rust resistance in soft red winter wheat cultivar 'USG' 3555™. <i>Plant Breeding</i> , 2013, 132, 53-60.	1.0	33
50	Transcriptomics of induced defense responses to greenbug aphid feeding in near isogenic wheat lines. <i>Plant Science</i> , 2013, 212, 26-36.	1.7	32
51	Identification and Mapping of Adult Plant Stripe Rust Resistance in Soft Red Winter Wheat VA00W38. <i>Crop Science</i> , 2013, 53, 871-879.	0.8	16
52	Registration of 'Eve'™ Winter Hulless Barley. <i>Journal of Plant Registrations</i> , 2013, 7, 5-11.	0.4	1
53	Molecular Characterization of Resistance to Fusarium Head Blight in U.S. Soft Red Winter Wheat Breeding Line VA00W38. <i>Crop Science</i> , 2012, 52, 2283-2292.	0.8	35
54	Resistance to Fusarium Head Blight and Deoxynivalenol Accumulation in Virginia Barley. <i>Plant Disease</i> , 2012, 96, 279-284.	0.7	18

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55	Registration of â€˜Beckerâ€™/â€˜Masseyâ€™ Wheat Recombinant Inbred Line Mapping Population. Journal of Plant Registrations, 2012, 6, 358-362.	0.4	2
56	Registration of Fusarium Head Blightâ€‘Resistant Soft Red Winter Wheat Germplasm VA04Wâ€‘433 and VA04Wâ€‘474. Journal of Plant Registrations, 2012, 6, 111-116.	0.4	7
57	Registration of â€˜Merlâ€™ Wheat. Journal of Plant Registrations, 2011, 5, 68-74.	0.4	1
58	Registration of â€˜SW049029104â€™ Wheat. Journal of Plant Registrations, 2011, 5, 91-97.	0.4	1
59	Registration of â€˜Snowglennâ€™ Winter Durum Wheat. Journal of Plant Registrations, 2011, 5, 81-86.	0.4	2
60	Registration of â€˜Danâ€™ Winter Hulless Barley. Journal of Plant Registrations, 2011, 5, 1-4.	0.4	8
61	Registration of â€˜Vision 30â€™ Wheat. Journal of Plant Registrations, 2011, 5, 353-359.	0.4	6
62	Registration of â€˜Vision 40â€™ Wheat. Journal of Plant Registrations, 2011, 5, 360-366.	0.4	3
63	Construction of a BAC library and a physical map of a major QTL for CBB resistance of common bean (<i>Phaseolus vulgaris</i> L.). Genetica, 2010, 138, 709-716.	0.5	8
64	Registration of â€˜Jamestownâ€™ Wheat. Journal of Plant Registrations, 2010, 4, 28-33.	0.4	24
65	Registration of â€˜Shirleyâ€™ Wheat. Journal of Plant Registrations, 2010, 4, 38-43.	0.4	9
66	Registration of USG 3209/Jaypee Wheat Recombinant Inbred Line Mapping Population. Journal of Plant Registrations, 2010, 4, 159-162.	0.4	3
67	Registration of â€˜3434â€™ Wheat. Journal of Plant Registrations, 2010, 4, 44-49.	0.4	0
68	Registration of â€˜5205â€™ Wheat. Journal of Plant Registrations, 2009, 3, 283-288.	0.4	1
69	Metaâ€‘Analysis of QTL Associated with Fusarium Head Blight Resistance in Wheat. Crop Science, 2009, 49, 1955-1968.	0.8	234
70	Registration of â€˜USG 3555â€™ Wheat. Journal of Plant Registrations, 2009, 3, 273-278.	0.4	6
71	Quantitative Trait Loci Associated with Deoxynivalenol Content and Kernel Quality in the Soft Red Winter Wheat â€˜Ernieâ€™. Crop Science, 2008, 48, 1408-1418.	0.8	25
72	Development of STS markers and QTL validation for common bacterial blight resistance in common bean. Plant Breeding, 2007, 127, 070807025605005-???.	1.0	17

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73	QTL associated with Fusarium head blight resistance in the soft red winter wheat Ernie. Theoretical and Applied Genetics, 2007, 115, 417-427.	1.8	74
74	Inheritance of Fusarium head blight resistance in the soft red winter wheat Ernie. Theoretical and Applied Genetics, 2005, 110, 454-461.	1.8	27
75	Genetic Variation in PI 294994 Wheat for Resistance to Russian Wheat Aphid. Crop Science, 1998, 38, 527-530.	0.8	32
76	Middle portion of the wheat culm remobilizes more carbon reserve to grains under drought. Journal of Agronomy and Crop Science, 0, , .	1.7	6
77	Capturing Wheat Phenotypes at the Genome Level. Frontiers in Plant Science, 0, 13, .	1.7	8