

Shuyu Liu

List of Publications by Year in descending order

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77
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257429
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docs citations

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times ranked

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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. | 17.5 | 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, . | 1.8 | 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. | 12.8 | 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. | 2.1 | 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. | 1.8 | 9 |
| 7 | Genomic selection of forage agronomic traits in winter wheat. <i>Crop Science</i> , 2021, 61, 410-421. | 1.8 | 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. | 3.6 | 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. | 3.3 | 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. | 4.4 | 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. | 5.7 | 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. | 2.0 | 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. | 3.0 | 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. | 2.4 | 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. | 2.5 | 17 |
| 17 | RhizoVision Crown: An Integrated Hardware and Software Platform for Root Crown Phenotyping. <i>Plant Phenomics</i> , 2020, 2020, 3074916. | 5.9 | 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. | 1.7 | 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. | 3.5 | 39 |
| 20 | Comparison of TaqMan, KASP and rhAmp SNP genotyping platforms in hexaploid wheat. <i>PLoS ONE</i> , 2019, 14, e0217222. | 2.5 | 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. | 2.9 | 5 |
| 22 | Genomic Selection of Forage Quality Traits in Winter Wheat. <i>Crop Science</i> , 2019, 59, 2473-2483. | 1.8 | 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. | 0.5 | 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. | 1.8 | 22 |
| 25 | Yield determination in winter wheat under different water regimes. <i>Field Crops Research</i> , 2019, 233, 80-87. | 5.1 | 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. | 1.8 | 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. | 3.5 | 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. | 5.1 | 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. | 0.5 | 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. | 2.1 | 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. | 1.8 | 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. | 1.8 | 17 |
| 33 | Wheat Curl Mite Resistance in Hard Winter Wheat in the US Great Plains. <i>Crop Science</i> , 2017, 57, 53-61. | 1.8 | 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. | 1.8 | 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. | 3.6 | 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. | 2.5 | 55 |

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|----|---|-----|-----------|
| 37 | Phenotypic Plasticity of Winter Wheat Heading Date and Grain Yield across the US Great Plains. Crop Science, 2016, 56, 2223-2236. | 1.8 | 75 |
| 38 | Validation of Chromosomal Locations of 90K Array Single Nucleotide Polymorphisms in US Wheat. Crop Science, 2016, 56, 364-373. | 1.8 | 26 |
| 39 | Spectral Reflectance Models for Characterizing Winter Wheat Genotypes. Journal of Crop Improvement, 2016, 30, 176-195. | 1.7 | 6 |
| 40 | Cooler Canopy Contributes to Higher Yield and Drought Tolerance in New Wheat Cultivars. Crop Science, 2014, 54, 2275-2284. | 1.8 | 22 |
| 41 | Yield Determination and Water Use Efficiency of Wheat under Water Limited Conditions in the U.S. Southern High Plains. Crop Science, 2014, 54, 34-47. | 1.8 | 74 |
| 42 | Molecular Mapping of Stripe Rust Resistance in Hard Red Winter Wheat TAM 111 Adapted to the U.S. High Plains. Crop Science, 2014, 54, 1361-1373. | 1.8 | 50 |
| 43 | Mapping Net Blotch Resistance in "Nomini"™ and Clho 2291 Barley. Crop Science, 2014, 54, 2596-2602. | 1.8 | 23 |
| 44 | Characterization of Fusarium Head Blight Resistance and Deoxynivalenol Accumulation in Hulled and Hulless Winter Barley. Plant Disease, 2014, 98, 599-606. | 1.4 | 13 |
| 45 | Physiology and transcriptomics of water-deficit stress responses in wheat cultivars TAM 111 and TAM 112. Journal of Plant Physiology, 2014, 171, 1289-1298. | 3.5 | 52 |
| 46 | Molecular Markers Linked to Important Genes in Hard Winter Wheat. Crop Science, 2014, 54, 1304-1321. | 1.8 | 55 |
| 47 | Molecular characterization of field resistance to Fusarium head blight in two US soft red winter wheat cultivars. Theoretical and Applied Genetics, 2013, 126, 2485-2498. | 3.6 | 59 |
| 48 | Marker-trait associations in Virginia Tech winter barley identified using genome-wide mapping. Theoretical and Applied Genetics, 2013, 126, 693-710. | 3.6 | 78 |
| 49 | Identification and mapping of adult plant stripe rust resistance in soft red winter wheat cultivar "USG 3555"™. Plant Breeding, 2013, 132, 53-60. | 1.9 | 33 |
| 50 | Transcriptomics of induced defense responses to greenbug aphid feeding in near isogenic wheat lines. Plant Science, 2013, 212, 26-36. | 3.6 | 32 |
| 51 | Identification and Mapping of Adult Plant Stripe Rust Resistance in Soft Red Winter Wheat VA00W-38. Crop Science, 2013, 53, 871-879. | 1.8 | 16 |
| 52 | Registration of "Eve"™ Winter Hulless Barley. Journal of Plant Registrations, 2013, 7, 5-11. | 0.5 | 1 |
| 53 | Molecular Characterization of Resistance to Fusarium Head Blight in U.S. Soft Red Winter Wheat Breeding Line VA00W-38. Crop Science, 2012, 52, 2283-2292. | 1.8 | 35 |
| 54 | Resistance to Fusarium Head Blight and Deoxynivalenol Accumulation in Virginia Barley. Plant Disease, 2012, 96, 279-284. | 1.4 | 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. | 0.5 | 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. | 0.5 | 7 |
| 57 | Registration of â€˜Merlâ€™ Wheat. Journal of Plant Registrations, 2011, 5, 68-74. | 0.5 | 1 |
| 58 | Registration of â€˜SW049029104â€™ Wheat. Journal of Plant Registrations, 2011, 5, 91-97. | 0.5 | 1 |
| 59 | Registration of â€˜Snowglennâ€™ Winter Durum Wheat. Journal of Plant Registrations, 2011, 5, 81-86. | 0.5 | 2 |
| 60 | Registration of â€˜Danâ€™ Winter Hulless Barley. Journal of Plant Registrations, 2011, 5, 1-4. | 0.5 | 8 |
| 61 | Registration of â€˜Vision 30â€™ Wheat. Journal of Plant Registrations, 2011, 5, 353-359. | 0.5 | 6 |
| 62 | Registration of â€˜Vision 40â€™ Wheat. Journal of Plant Registrations, 2011, 5, 360-366. | 0.5 | 3 |
| 63 | Construction of a BAC library and a physical map of a major QTL for CBB resistance of common bean (Phaseolus vulgaris L.). Genetica, 2010, 138, 709-716. | 1.1 | 8 |
| 64 | Registration of â€˜Jamestownâ€™ Wheat. Journal of Plant Registrations, 2010, 4, 28-33. | 0.5 | 24 |
| 65 | Registration of â€˜Shirleyâ€™ Wheat. Journal of Plant Registrations, 2010, 4, 38-43. | 0.5 | 9 |
| 66 | Registration of USC 3209/Jaypee Wheat Recombinant Inbred Line Mapping Population. Journal of Plant Registrations, 2010, 4, 159-162. | 0.5 | 3 |
| 67 | Registration of â€˜3434â€™ Wheat. Journal of Plant Registrations, 2010, 4, 44-49. | 0.5 | 0 |
| 68 | Registration of â€˜5205â€™ Wheat. Journal of Plant Registrations, 2009, 3, 283-288. | 0.5 | 1 |
| 69 | Metaâ€‘Analysis of QTL Associated with Fusarium Head Blight Resistance in Wheat. Crop Science, 2009, 49, 1955-1968. | 1.8 | 234 |
| 70 | Registration of â€˜USC 3555â€™ Wheat. Journal of Plant Registrations, 2009, 3, 273-278. | 0.5 | 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. | 1.8 | 25 |
| 72 | Development of STS markers and QTL validation for common bacterial blight resistance in common bean. Plant Breeding, 2007, 127, 070807025605005-??? | 1.9 | 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. | 3.6 | 74 |
| 74 | Inheritance of Fusarium head blight resistance in the soft red winter wheat Ernie. Theoretical and Applied Genetics, 2005, 110, 454-461. | 3.6 | 27 |
| 75 | Genetic Variation in PI 294994 Wheat for Resistance to Russian Wheat Aphid. Crop Science, 1998, 38, 527-530. | 1.8 | 32 |
| 76 | Middle portion of the wheat culm remobilizes more carbon reserve to grains under drought. Journal of Agronomy and Crop Science, 0, , . | 3.5 | 6 |
| 77 | Capturing Wheat Phenotypes at the Genome Level. Frontiers in Plant Science, 0, 13, . | 3.6 | 8 |