

Awais Rasheed

List of Publications by Year in descending order

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Version: 2024-02-01

82
papers

4,383
citations

186265

28
h-index

114465

63
g-index

82
all docs

82
docs citations

82
times ranked

3819
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic gain and G \times E interaction in bread wheat cultivars representing 105 years of breeding in Pakistan. <i>Crop Science</i> , 2022, 62, 178-191.	1.8	8
2	Characterization of new COBRA like (COBL) genes in wheat (<i>Triticum aestivum</i>) and their expression analysis under drought stress. <i>Molecular Biology Reports</i> , 2022, 49, 1379-1387.	2.3	9
3	Root system architecture in cereals: progress, challenges and perspective. <i>Plant Journal</i> , 2022, 110, 23-42.	5.7	38
4	<i>Aegilops tauschii</i> presents a genetic roadmap for hexaploid wheat improvement. <i>Trends in Genetics</i> , 2022, 38, 307-309.	6.7	7
5	Genome-Wide Association and Genomic Prediction for Stripe Rust Resistance in Synthetic-Derived Wheats. <i>Frontiers in Plant Science</i> , 2022, 13, 788593.	3.6	7
6	Genetic Variability and Aggressiveness of <i>Tilletia indica</i> Isolates Causing Karnal Bunt in Wheat. <i>Journal of Fungi</i> (Basel, Switzerland), 2022, 8, 219.	3.5	2
7	High Resolution Genome Wide Association Studies Reveal Rich Genetic Architectures of Grain Zinc and Iron in Common Wheat (<i>Triticum aestivum</i> L.). <i>Frontiers in Plant Science</i> , 2022, 13, 840614.	3.6	15
8	Genome edited wheat- current advances for the second green revolution. <i>Biotechnology Advances</i> , 2022, 60, 108006.	11.7	19
9	Genome-wide analyses reveal footprints of divergent selection and popping-related traits in CIMMYT's maize inbred lines. <i>Journal of Experimental Botany</i> , 2021, 72, 1307-1320.	4.8	11
10	Quantifying senescence in bread wheat using multispectral imaging from an unmanned aerial vehicle and QTL mapping. <i>Plant Physiology</i> , 2021, 187, 2623-2636.	4.8	15
11	Genetic diversity and agronomic performance of wheat landraces currently grown in Tajikistan. <i>Crop Science</i> , 2021, 61, 2548-2564.	1.8	5
12	Genetic Gain for Grain Micronutrients and Their Association with Phenology in Historical Wheat Cultivars Released between 1911 and 2016 in Pakistan. <i>Agronomy</i> , 2021, 11, 1247.	3.0	12
13	Characterization of the genetic basis of local adaptation of wheat landraces from Iran and Pakistan using genome-wide association study. <i>Plant Genome</i> , 2021, 14, e20096.	2.8	8
14	Diversity and Adaptation of Currently Grown Wheat Landraces and Modern Germplasm in Afghanistan, Iran, and Turkey. <i>Crops</i> , 2021, 1, 54-67.	1.4	8
15	Harnessing Wheat <i>Fhb1</i> for Fusarium Resistance. <i>Trends in Plant Science</i> , 2020, 25, 1-3.	8.8	56
16	Appraisal of wheat genomics for gene discovery and breeding applications: a special emphasis on advances in Asia. <i>Theoretical and Applied Genetics</i> , 2020, 133, 1503-1520.	3.6	11
17	Genetic basis of spring wheat resistance to leaf rust (<i>Puccinia triticina</i>) in Kazakhstan and Russia. <i>Euphytica</i> , 2020, 216, 1.	1.2	13
18	Assessment of Water and Nitrogen Use Efficiencies Through UAV-Based Multispectral Phenotyping in Winter Wheat. <i>Frontiers in Plant Science</i> , 2020, 11, 927.	3.6	43

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19	Mobilizing Crop Biodiversity. <i>Molecular Plant</i> , 2020, 13, 1341-1344.	8.3	50
20	Dissection of Molecular Processes and Genetic Architecture Underlying Iron and Zinc Homeostasis for Biofortification: From Model Plants to Common Wheat. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9280.	4.1	27
21	Genome-Wide Association Mapping of Adult-Plant Resistance to Stripe Rust in Common Wheat (<i>Triticum aestivum</i>). <i>Plant Disease</i> , 2020, 104, 2174-2180.	1.4	6
22	Genome-Wide Association Analysis of Fusarium Head Blight Resistance in Chinese Elite Wheat Lines. <i>Frontiers in Plant Science</i> , 2020, 11, 206.	3.6	44
23	Genome-wide association analysis of stem water-soluble carbohydrate content in bread wheat. <i>Theoretical and Applied Genetics</i> , 2020, 133, 2897-2914.	3.6	20
24	Genomic Prediction for Grain Yield and Yield-Related Traits in Chinese Winter Wheat. <i>International Journal of Molecular Sciences</i> , 2020, 21, 1342.	4.1	27
25	Advanced Genomics and Breeding Tools to Accelerate the Development of Climate Resilient Wheat. , 2020, , 45-95.		1
26	Molecular Marker Development and Application for Improving Qualities in Bread Wheat. , 2020, , 323-345.		0
27	Identifying loci with breeding potential across temperate and tropical adaptation via EigenGWAS and EnvGWAS. <i>Molecular Ecology</i> , 2019, 28, 3544-3560.	3.9	32
28	From markers to genome-based breeding in wheat. <i>Theoretical and Applied Genetics</i> , 2019, 132, 767-784.	3.6	115
29	Genome-wide variation patterns between landraces and cultivars uncover divergent selection during modern wheat breeding. <i>Theoretical and Applied Genetics</i> , 2019, 132, 2509-2523.	3.6	56
30	Molecular Characterization of 87 Functional Genes in Wheat Diversity Panel and Their Association With Phenotypes Under Well-Watered and Water-Limited Conditions. <i>Frontiers in Plant Science</i> , 2019, 10, 717.	3.6	43
31	Genetic architecture of grain yield in bread wheat based on genome-wide association studies. <i>BMC Plant Biology</i> , 2019, 19, 168.	3.6	172
32	Genome-Wide Analyses Reveal Footprints of Divergent Selection and Drought Adaptive Traits in Synthetic-Derived Wheats. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 1957-1973.	1.8	53
33	Accuracy assessment of plant height using an unmanned aerial vehicle for quantitative genomic analysis in bread wheat. <i>Plant Methods</i> , 2019, 15, 37.	4.3	86
34	Breeding strategies for structuring salinity tolerance in wheat. <i>Advances in Agronomy</i> , 2019, 155, 121-187.	5.2	53
35	A rapid monitoring of NDVI across the wheat growth cycle for grain yield prediction using a multi-spectral UAV platform. <i>Plant Science</i> , 2019, 282, 95-103.	3.6	238
36	Allelic effects and variations for key bread-making quality genes in bread wheat using high-throughput molecular markers. <i>Journal of Cereal Science</i> , 2019, 85, 305-309.	3.7	26

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37	China-CIMMYT collaboration enhances wheat improvement in China. <i>Frontiers of Agricultural Science and Engineering</i> , 2019, 6, 233.	1.4	5
38	Fast-Forwarding Genetic Gain. <i>Trends in Plant Science</i> , 2018, 23, 184-186.	8.8	164
39	The goat grass genome's role in wheat improvement. <i>Nature Plants</i> , 2018, 4, 56-58.	9.3	21
40	Genotypic Variation and Genotype × Environment Interaction for Yield-Related Traits in Synthetic Hexaploid Wheats under a Range of Optimal and Heat-Stressed Environments. <i>Crop Science</i> , 2018, 58, 295-303.	1.8	23
41	QTL mapping for seedling morphology under drought stress in wheat cross synthetic (W7984)/Opata. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2018, 16, 359-366.	0.8	8
42	Identification of genome-wide single-nucleotide polymorphisms (SNPs) associated with tolerance to chromium toxicity in spring wheat (<i>Triticum aestivum</i> L.). <i>Plant and Soil</i> , 2018, 422, 371-384.	3.7	21
43	Wheat genetic resources in the post-genomics era: promise and challenges. <i>Annals of Botany</i> , 2018, 121, 603-616.	2.9	101
44	Time-Series Multispectral Indices from Unmanned Aerial Vehicle Imagery Reveal Senescence Rate in Bread Wheat. <i>Remote Sensing</i> , 2018, 10, 809.	4.0	98
45	Diversity in D-genome synthetic hexaploid wheat association panel for seedling emergence traits under salinity stress. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2017, 15, 488-495.	0.8	5
46	Physiological, biochemical and agronomic traits associated with drought tolerance in a synthetic-derived wheat diversity panel. <i>Crop and Pasture Science</i> , 2017, 68, 213.	1.5	17
47	Genome-wide association mapping of starch granule size distribution in common wheat. <i>Journal of Cereal Science</i> , 2017, 77, 211-218.	3.7	18
48	Genetic Progress in Grain Yield and Physiological Traits in Chinese Wheat Cultivars of Southern Yellow and Huai Valley since 1950. <i>Crop Science</i> , 2017, 57, 760-773.	1.8	94
49	Crop Breeding Chips and Genotyping Platforms: Progress, Challenges, and Perspectives. <i>Molecular Plant</i> , 2017, 10, 1047-1064.	8.3	380
50	Genome-wide association study for agronomic and physiological traits in spring wheat evaluated in a range of heat prone environments. <i>Theoretical and Applied Genetics</i> , 2017, 130, 1819-1835.	3.6	117
51	Genome-wide association mapping of black point reaction in common wheat (<i>Triticum aestivum</i> L.). <i>BMC Plant Biology</i> , 2017, 17, 220.	3.6	141
52	Comparison of Economically Important Loci in Landraces and Improved Wheat Cultivars from Pakistan. <i>Crop Science</i> , 2016, 56, 287-301.	1.8	18
53	Functional characterization of germin and germin-like protein genes in various plant species using transgenic approaches. <i>Biotechnology Letters</i> , 2016, 38, 1405-1421.	2.2	33
54	Development and validation of KASP assays for genes underpinning key economic traits in bread wheat. <i>Theoretical and Applied Genetics</i> , 2016, 129, 1843-1860.	3.6	357

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55	TaTGW6-A1, an ortholog of rice TGW6, is associated with grain weight and yield in bread wheat. <i>Molecular Breeding</i> , 2016, 36, 1.	2.1	163
56	Genome-Wide Association of Stem Water Soluble Carbohydrates in Bread Wheat. <i>PLoS ONE</i> , 2016, 11, e0164293.	2.5	50
57	Characterization of Synthetic Hexaploids Derived from Same <i>Aegilops tauschii</i> Accessions and Different Durum Cultivars. <i>Cytologia</i> , 2015, 80, 427-440.	0.6	10
58	Genome-wide association for grain yield under rainfed conditions in historical wheat cultivars from Pakistan. <i>Frontiers in Plant Science</i> , 2015, 6, 743.	3.6	169
59	Genome-Wide Linkage Mapping of QTL for Yield Components, Plant Height and Yield-Related Physiological Traits in the Chinese Wheat Cross Zhou 8425B/Chinese Spring. <i>Frontiers in Plant Science</i> , 2015, 6, 1099.	3.6	267
60	Cytological, Phenological and Molecular Characterization of B (S)-Genome Synthetic Hexaploids (2n =) Tj ETQq0 0 0 rgBT /Overlock 10 T	1.8	8
61	<i>Aegilops tauschii</i> Introgressions in <i>Wheat</i> . , 2015, , 245-271.		19
62	Characterization of D-genome diversity for tolerance to boron toxicity in synthetic hexaploid wheat and in silico analysis of candidate genes. <i>Acta Physiologiae Plantarum</i> , 2015, 37, 1.	2.1	11
63	Molecular characterization of the puroindoline-a and b alleles in synthetic hexaploid wheats and in silico functional and structural insights into Pina-D1. <i>Journal of Theoretical Biology</i> , 2015, 376, 1-7.	1.7	19
64	Comparative Assessment of Synthetic-derived and Conventional Bread Wheat Advanced Lines Under Osmotic Stress and Implications for Molecular Analysis. <i>Plant Molecular Biology Reporter</i> , 2015, 33, 1907-1917.	1.8	14
65	Genome-Wide Association Mapping for Seedling and Adult Plant Resistance to Stripe Rust in Synthetic Hexaploid Wheat. <i>PLoS ONE</i> , 2014, 9, e105593.	2.5	218
66	Association mapping identifies QTLs on wheat chromosome 3A for yield related traits. <i>Cereal Research Communications</i> , 2014, 42, 177-188.	1.6	14
67	Stripe rust resistance in <i>Triticum durum</i> " T. monococcum and <i>T. durum</i> " T. urartu amphiploids. <i>Australasian Plant Pathology</i> , 2014, 43, 109-113.	1.0	12
68	Wheat seed storage proteins: Advances in molecular genetics, diversity and breeding applications. <i>Journal of Cereal Science</i> , 2014, 60, 11-24.	3.7	139
69	Exploitation of synthetic-derived wheats through osmotic stress responses for drought tolerance improvement. <i>Acta Physiologiae Plantarum</i> , 2014, 36, 2453-2465.	2.1	13
70	Genome-wide association for grain morphology in synthetic hexaploid wheats using digital imaging analysis. <i>BMC Plant Biology</i> , 2014, 14, 128.	3.6	102
71	Analysis of Genetic Diversity in Synthetic Wheat Assemblage (<i>T. turgidum</i> ^ ^times; <i>Aegilops tauschii</i>); Tj ETQq1 1 0,784314 rgBT /Overlock 4	0.6	4
72	Genetic Diversity for Wheat Improvement as a Conduit to Food Security. <i>Advances in Agronomy</i> , 2013, , 179-257.	5.2	124

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73	Biotic Stress and Crop Improvement: A Wheat Focus Around Novel Strategies. , 2013, , 239-267.		2
74	An Overview of Omics for Wheat Grain Quality Improvement. , 2013, , 307-344.		2
75	Characterization of HMW-GS and evaluation of their diversity in morphologically elite synthetic hexaploid wheats. <i>Breeding Science</i> , 2012, 62, 365-370.	1.9	17
76	Stripe rust analysis of D-genome synthetic wheats (2n=6x=42, AABBDD) and their molecular diversity. <i>Archives of Phytopathology and Plant Protection</i> , 2012, 45, 1479-1487.	1.3	4
77	Powdery mildew resistance in some new wheat amphiploids (2n=6x=42) derived from A- and S-genome diploid progenitors. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2012, 10, 165-170.	0.8	7
78	High-molecular-weight (HMW) glutenin subunit composition of the Elite-II synthetic hexaploid wheat subset (<i>Triticum turgidum</i> – <i>Aegilops tauschii</i> ; 2n=6x=42; AABBDD). <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2012, 10, 1-4.	0.8	11
79	An overview of stripe rust of wheat (<i>Puccinia striiformis</i> f. sp. <i>tritici</i>) in Pakistan. <i>Archives of Phytopathology and Plant Protection</i> , 2012, 45, 2278-2289.	1.3	9
80	Molecular Basis of Disease Resistance in Cereal Crops: An Overview. , 2012, , 477-489.		2
81	Allelic variation and composition of HMW-GS in advanced lines derived from d-genome synthetic hexaploid / bread wheat (<i>Triticum aestivum</i> L.). <i>Journal of Crop Science and Biotechnology</i> , 2012, 15, 1-7.	1.5	14
82	Genetic Diversity and Selection Signatures in Synthetic-Derived Wheats and Modern Spring Wheat. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	2