Muhammad Iqbal

List of Publications by Year in descending order

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567281 526287 34 839 15 27 citations h-index g-index papers 34 34 34 747 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Genomic Predictions for Common Bunt, FHB, Stripe Rust, Leaf Rust, and Leaf Spotting Resistance in Spring Wheat. Genes, 2022, 13, 565.	2.4	13
2	Comparison of single-trait and multi-trait genomic predictions on agronomic and disease resistance traits in spring wheat. Theoretical and Applied Genetics, 2022, 135, 2747-2767.	3.6	4
3	Genomic Prediction Accuracy of Stripe Rust in Six Spring Wheat Populations by Modeling Genotype by Environment Interaction. Plants, 2022, 11, 1736.	3.5	3
4	Physical Mapping of QTL in Four Spring Wheat Populations under Conventional and Organic Management Systems. I. Earliness. Plants, 2021, 10, 853.	3.5	13
5	Phenotypic performance and associated QTL of  Peace' ×  CDC Stanley' mapping population under conventional and organic management systems. Crop Science, 2021, 61, 3469-3483.	1.8	8
6	Physical mapping of QTL associated with agronomic and end-use quality traits in spring wheat under conventional and organic management systems. Theoretical and Applied Genetics, 2021, 134, 3699-3719.	3.6	23
7	Genome-based prediction of agronomic traits in spring wheat under conventional and organic management systems. Theoretical and Applied Genetics, 2021, 135, 537.	3.6	10
8	Genetic diversity and selective sweeps in historical and modern Canadian spring wheat cultivars using the 90K SNP array. Scientific Reports, 2021, 11, 23773.	3.3	10
9	The performance of spring wheat cultivar mixtures under conventional and organic management in Western Canada., 2020, 3, e20003.		4
10	Genetic analyses of native Fusarium head blight resistance in two spring wheat populations identifies QTL near the B1, Ppd-D1, Rht-1, Vrn-1, Fhb1, Fhb2, and Fhb5 loci. Theoretical and Applied Genetics, 2020, 133, 2775-2796.	3 . 6	9
11	Mapping genomic regions controlling agronomic traits in spring wheat under conventional and organic managements. Crop Science, 2020, 60, 2038-2052.	1.8	16
12	Mapping QTL Associated with Stripe Rust, Leaf Rust, and Leaf Spotting in a Canadian Spring Wheat Population. Crop Science, 2019, 59, 650-658.	1.8	15
13	There are Different Pathways to Stable Spring Wheat Grain Yield and Nitrogen Utilization Efficiency in Conventional and Organicallyâ€Managed Systems. Agronomy Journal, 2019, 111, 2370-2377.	1.8	3
14	Agronomic and physiological aspects of nitrogen use efficiency in conventional and organic cereal-based production systems. Renewable Agriculture and Food Systems, 2018, 33, 443-466.	1.8	26
15	Investigating Genetic Progress and Variation for Nitrogen Use Efficiency in Spring Wheat. Crop Science, 2018, 58, 1542-1557.	1.8	15
16	Mapping QTLs Controlling Agronomic Traits in the †Attila†× †CDC Go†Spring Wheat Population unc Organic Management using 90K SNP Array. Crop Science, 2017, 57, 365-377.	der 1.8	30
17	Allelic variation and effects of 16 candidate genes on disease resistance in western Canadian spring wheat cultivars. Molecular Breeding, 2017, 37, 1.	2.1	11
18	Genome-wide association mapping of genomic regions associated with phenotypic traits in Canadian western spring wheat. Molecular Breeding, 2017, 37, 1.	2.1	30

#	Article	IF	CITATIONS
19	Mapping of QTLs associated with resistance to common bunt, tan spot, leaf rust, and stripe rust in a spring wheat population. Molecular Breeding, 2017, 37, 1.	2.1	21
20	Population Structure and Genomewide Association Analysis of Resistance to Disease and Insensitivity to Ptr Toxins in Canadian Spring Wheat Using 90K SNP Array. Crop Science, 2017, 57, 1522-1539.	1.8	24
21	QTLs associated with agronomic traits in the Attila $ ilde{A}-$ CDC Go spring wheat population evaluated under conventional management. PLoS ONE, 2017, 12, e0171528.	2.5	68
22	QTLs Associated with Agronomic Traits in the Cutler $\tilde{A}-$ AC Barrie Spring Wheat Mapping Population Using Single Nucleotide Polymorphic Markers. PLoS ONE, 2016, 11, e0160623.	2.5	36
23	Genetic Improvement in Grain Yield and other Traits of Wheat Grown in Western Canada. Crop Science, 2016, 56, 613-624.	1.8	45
24	Effect of Lr34/Yr18 on agronomic and quality traits in a spring wheat mapping population and implications for breeding. Molecular Breeding, 2016, 36, 1.	2.1	18
25	Genetic variation for flowering time and height reducing genes and important traits in western Canadian spring wheat. Euphytica, 2016, 208, 377-390.	1.2	35
26	Earliness per se quantitative trait loci and their interaction with Vrn-B1 locus in a spring wheat population. Molecular Breeding, 2015, 35, 1.	2.1	19
27	Flowering time in wheat (Triticum aestivum L.): a key factor for global adaptability. Euphytica, 2014, 197, 1-26.	1.2	168
28	Genetic Analysis of Yield and Yield Contributing Quantitative Traits in Bread Wheat Under Sodium Chloride Salinity. Journal of Agricultural Science, 2013, 5, .	0.2	2
29	Determination of Rust Resistance Gene Complex <i><scp>L</scp>r34/<scp>Y</scp>r18</i> in Spring Wheat and its Effect on Components of Partial Resistance. Journal of Phytopathology, 2012, 160, 628-636.	1.0	5
30	Molecular characterization of vernalization response genes in Canadian spring wheat. Genome, 2007, 50, 511-516.	2.0	57
31	The Effect of Vernalization Genes on Earliness and Related Agronomic Traits of Spring Wheat in Northern Growing Regions. Crop Science, 2007, 47, 1031-1039.	1.8	22
32	Genetic analysis of flowering and maturity time in high latitude spring wheat. Euphytica, 2007, 154, 207-218.	1.2	55
33	Allelic variation at the Vrn-A1, Vrn-B1, Vrn-D1, Vrn-B3 and Ppd-D1a loci of Pakistani spring wheat cultivars. Electronic Journal of Biotechnology, 2001, 14, .	2.2	16
34	Genomeâ€wide association mapping of agronomic traits and grain characteristics in spring wheat under conventional and organic management systems. Crop Science, 0, , .	1.8	5