

## List of Publications by Year in descending order

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Version: 2024-02-01

22  
papers

1,041  
citations

566801

15  
h-index

676716

22  
g-index

22  
all docs

22  
docs citations

22  
times ranked

1036  
citing authors

#	ARTICLE	IF	CITATIONS
1	Wheat Fhb1 encodes a chimeric lectin with agglutinin domains and a pore-forming toxin-like domain conferring resistance to Fusarium head blight. <i>Nature Genetics</i> , 2016, 48, 1576-1580.	9.4	299
2	A Pipeline Strategy for Grain Crop Domestication. <i>Crop Science</i> , 2016, 56, 917-930.	0.8	101
3	Establishment and Optimization of Genomic Selection to Accelerate the Domestication and Improvement of Intermediate Wheatgrass. <i>Plant Genome</i> , 2016, 9, plantgenome2015.07.0059.	1.6	86
4	New Insights into the Organization, Recombination, Expression and Functional Mechanism of Low Molecular Weight Glutenin Subunit Genes in Bread Wheat. <i>PLoS ONE</i> , 2010, 5, e13548.	1.1	74
5	Composition and functional analysis of low-molecular-weight glutenin alleles with Aroona near-isogenic lines of bread wheat. <i>BMC Plant Biology</i> , 2012, 12, 243.	1.6	68
6	â€œMNâ€œClearwaterâ€œ™, the first foodâ€œgrade intermediate wheatgrass (Kernza perennial grain) cultivar. <i>Journal of Plant Registrations</i> , 2020, 14, 288-297.	0.4	58
7	Development of the first consensus genetic map of intermediate wheatgrass ( <i>Thinopyrum</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tr 5	1.8	43
8	Genome-Wide Association Study of Yield Component Traits in Intermediate Wheatgrass and Implications in Genomic Selection and Breeding. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 2429-2439.	0.8	34
9	Development and verification of wheat germplasm containing both Sr2 and Fhb1. <i>Molecular Breeding</i> , 2016, 36, 1.	1.0	32
10	Genome mapping of quantitative trait loci (QTL) controlling domestication traits of intermediate wheatgrass ( <i>Thinopyrum intermedium</i> ). <i>Theoretical and Applied Genetics</i> , 2019, 132, 2325-2351.	1.8	30
11	Enhancing Crop Domestication Through Genomic Selection, a Case Study of Intermediate Wheatgrass. <i>Frontiers in Plant Science</i> , 2020, 11, 319.	1.7	28
12	Uncovering the Genetic Architecture of Seed Weight and Size in Intermediate Wheatgrass through Linkage and Association Mapping. <i>Plant Genome</i> , 2017, 10, plantgenome2017.03.0022.	1.6	26
13	Development of genotyping by sequencing (GBS)- and array-derived SNP markers for stem rust resistance gene Sr42. <i>Molecular Breeding</i> , 2015, 35, 1.	1.0	24
14	New insights into high-molecular-weight glutenin subunits and sub-genomes of the perennial crop <i>Thinopyrum intermedium</i> (Triticeae). <i>Journal of Cereal Science</i> , 2014, 59, 203-210.	1.8	22
15	Towards the understanding of end-use quality in intermediate wheatgrass ( <i>Thinopyrum intermedium</i> ): High-molecular-weight glutenin subunits, protein polymerization, and mixing characteristics. <i>Journal of Cereal Science</i> , 2015, 66, 81-88.	1.8	20
16	Dominance and GÃ—E interaction effects improve genomic prediction and genetic gain in intermediate wheatgrass ( <i>Thinopyrum intermedium</i> ). <i>Plant Genome</i> , 2020, 13, e20012.	1.6	19
17	Floret site utilization and reproductive tiller number are primary components of grain yield in intermediate wheatgrass spaced plants. <i>Crop Science</i> , 2021, 61, 1073-1088.	0.8	19
18	Fifty years of a public cassava breeding program: evolution of breeding objectives, methods, and decision-making processes. <i>Theoretical and Applied Genetics</i> , 2021, 134, 2335-2353.	1.8	18

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19	Characterization of Genetic Resistance to Fusarium Head Blight and Bacterial Leaf Streak in Intermediate Wheatgrass ( <i>Thinopyrum intermedium</i> ). <i>Agronomy</i> , 2019, 9, 429.	1.3	14
20	Correlation of cooking time with water absorption and changes in relative density during boiling of cassava roots. <i>International Journal of Food Science and Technology</i> , 2021, 56, 1193-1205.	1.3	11
21	Identifying New Resistance to Cassava Mosaic Disease and Validating Markers for the CMD2 Locus. <i>Agriculture (Switzerland)</i> , 2021, 11, 829.	1.4	8
22	High-Throughput Virus Screening in Crosses of South American and African Cassava Germplasm Reveals Broad-Spectrum Resistance against Viruses Causing Cassava Brown Streak Disease and Cassava Mosaic Virus Disease. <i>Agronomy</i> , 2022, 12, 1055.	1.3	7