

Surbhi Grewal

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

804
citations

687363

13
h-index

713466

21
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29
all docs

29
docs citations

29
times ranked

870
citing authors

#	ARTICLE	IF	CITATIONS
1	Adenosine Methylation in Arabidopsis mRNA is Associated with the 3' End and Reduced Levels Cause Developmental Defects. <i>Frontiers in Plant Science</i> , 2012, 3, 48.	3.6	213
2	A step change in the transfer of interspecific variation into wheat from <i>Amblyopyrum muticum</i> . <i>Plant Biotechnology Journal</i> , 2017, 15, 217-226.	8.3	124
3	Characterisation of <i>Thinopyrum bessarabicum</i> chromosomes through genome-wide introgressions into wheat. <i>Theoretical and Applied Genetics</i> , 2018, 131, 389-406.	3.6	74
4	Introgression of <i>Aegilops speltoides</i> segments in <i>Triticum aestivum</i> and the effect of the gametocidal genes. <i>Annals of Botany</i> , 2018, 121, 229-240.	2.9	57
5	Rapid identification of homozygosity and site of wild relative introgressions in wheat through chromosome-specific KASP genotyping assays. <i>Plant Biotechnology Journal</i> , 2020, 18, 743-755.	8.3	50
6	Development and validation of an exome-based SNP marker set for identification of the St, Jr and Jvs genomes of <i>Thinopyrum intermedium</i> in a wheat background. <i>Theoretical and Applied Genetics</i> , 2019, 132, 1555-1570.	3.6	37
7	Generation of amphidiploids from hybrids of wheat and related species from the genera <i>Aegilops</i> , <i>Secale</i> , <i>Thinopyrum</i> , and <i>Triticum</i> as a source of genetic variation for wheat improvement. <i>Genome</i> , 2015, 58, 71-79.	2.0	30
8	Detection of <i>T. urartu</i> Introgressions in Wheat and Development of a Panel of Interspecific Introgression Lines. <i>Frontiers in Plant Science</i> , 2018, 9, 1565.	3.6	27
9	Development and characterisation of interspecific hybrid lines with genome-wide introgressions from <i>Triticum timopheevii</i> in a hexaploid wheat background. <i>BMC Plant Biology</i> , 2019, 19, 183.	3.6	25
10	Exploiting the genome of <i>Thinopyrum elongatum</i> to expand the gene pool of hexaploid wheat. <i>Theoretical and Applied Genetics</i> , 2020, 133, 2213-2226.	3.6	25
11	Development of Stable Homozygous Wheat/ <i>Amblyopyrum muticum</i> (<i>Aegilops mutica</i>) Introgression Lines and Their Cytogenetic and Molecular Characterization. <i>Frontiers in Plant Science</i> , 2019, 10, 34.	3.6	21
12	Development of Wheat- <i>Aegilops caudata</i> Introgression Lines and Their Characterization Using Genome-Specific KASP Markers. <i>Frontiers in Plant Science</i> , 2020, 11, 606.	3.6	20
13	Mapping the "breaker" element of the gametocidal locus proximal to a block of sub-telomeric heterochromatin on the long arm of chromosome 4Ssh of <i>Aegilops sharonensis</i> . <i>Theoretical and Applied Genetics</i> , 2015, 128, 1049-1059.	3.6	15
14	Comparative Mapping and Targeted Capture Sequencing of the Gametocidal Loci in <i>Aegilops sharonensis</i> . <i>Plant Genome</i> , 2017, 10, plantgenome2016.09.0090.	2.8	13
15	The Use of Pentaploid Crosses for the Introgression of <i>Amblyopyrum muticum</i> and D-Genome Chromosome Segments Into Durum Wheat. <i>Frontiers in Plant Science</i> , 2019, 10, 1110.	3.6	13
16	Chromosome-specific KASP markers for detecting <i>Amblyopyrum muticum</i> segments in wheat introgression lines. <i>Plant Genome</i> , 2022, 15, e20193.	2.8	11
17	Whole-genome sequencing uncovers the structural and transcriptomic landscape of hexaploid wheat/ <i>Amblyopyrum muticum</i> introgression lines. <i>Plant Biotechnology Journal</i> , 2023, 21, 482-496.	8.3	10
18	Introgression of the <i>Triticum timopheevii</i> Genome Into Wheat Detected by Chromosome-Specific Kompetitive Allele Specific PCR Markers. <i>Frontiers in Plant Science</i> , 2022, 13, .	3.6	9

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19	Generation of Doubled Haploid Wheat-Triticum urartu Introgression Lines and Their Characterisation Using Chromosome-Specific KASP Markers. <i>Frontiers in Plant Science</i> , 2021, 12, 643636.	3.6	7
20	Resistance to wheat rusts identified in wheat/ <i>Amblyopyrum muticum</i> chromosome introgressions. <i>Crop Science</i> , 2020, 60, 1957-1964.	1.8	5
21	Development of a New A m “Genome” Specific Single Nucleotide Polymorphism Marker Set for the Molecular Characterization of Wheat “Triticum monococcum Introgression Lines. <i>Plant Genome</i> , 2019, 12, 180098.	2.8	4
22	Assessing the Potential of Using the Langdon 5D(5B) Substitution Line for the Introgression of <i>Aegilops tauschii</i> Into Durum Wheat. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	2
23	Exploring Untapped Wheat Genetic Resources to Boost Food Security. , 2022, , 319-340.		1