

Surbhi Grewal

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2733433/publications.pdf>

Version: 2024-02-01

23
papers

804
citations

687363

13
h-index

713466

21
g-index

29
all docs

29
docs citations

29
times ranked

870
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | 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 |
| 2 | 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 |
| 3 | Exploring Untapped Wheat Genetic Resources to Boost Food Security. , 2022, , 319-340. | | 1 |
| 4 | Introgression of the <i>Triticum timopheevii</i> Genome Into Wheat Detected by Chromosome-Specific Competitive Allele Specific PCR Markers. <i>Frontiers in Plant Science</i> , 2022, 13, . | 3.6 | 9 |
| 5 | 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 |
| 6 | 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 |
| 7 | Development of Wheat-Aegilops caudata Introgression Lines and Their Characterization Using Genome-Specific KASP Markers. <i>Frontiers in Plant Science</i> , 2020, 11, 606. | 3.6 | 20 |
| 8 | Resistance to wheat rusts identified in wheat/ <i>Amblyopyrum muticum</i> chromosome introgressions. <i>Crop Science</i> , 2020, 60, 1957-1964. | 1.8 | 5 |
| 9 | 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 |
| 10 | 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 |
| 11 | 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 |
| 12 | 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 |
| 13 | 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 |
| 14 | 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 |
| 15 | 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 |
| 16 | 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 |
| 17 | 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 |
| 18 | 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 |

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Comparative Mapping and Targetedâ€ Capture Sequencing of the Gametocidal Loci in <i>Aegilops sharonensis</i> . <i>Plant Genome</i> , 2017, 10, plantgenome2016.09.0090. | 2.8 | 13 |
| 20 | Mapping the â€ breakerâ€™ 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 |
| 21 | 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 |
| 22 | Adenosine Methylation in Arabidopsis mRNA is Associated with the 3â€² End and Reduced Levels Cause Developmental Defects. <i>Frontiers in Plant Science</i> , 2012, 3, 48. | 3.6 | 213 |
| 23 | 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 |