Sreya Ghosh

List of Publications by Year in descending order

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933447 1199594 1,918 12 10 12 citations h-index g-index papers 16 16 16 2310 docs citations times ranked citing authors all docs

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Speed breeding is a powerful tool to accelerate crop research and breeding. Nature Plants, 2018, 4, 23-29. | 9.3 | 770 |
| 2 | Speed breeding in growth chambers and glasshouses for crop breeding and model plant research. Nature Protocols, 2018, 13, 2944-2963. | 12.0 | 286 |
| 3 | Resistance gene cloning from a wild crop relative by sequence capture and association genetics. Nature Biotechnology, 2019, 37, 139-143. | 17.5 | 280 |
| 4 | Rapid gene isolation in barley and wheat by mutant chromosome sequencing. Genome Biology, 2016, 17, 221. | 8.8 | 265 |
| 5 | Population genomic analysis of Aegilops tauschii identifies targets for bread wheat improvement. Nature Biotechnology, 2022, 40, 422-431. | 17.5 | 102 |
| 6 | A roadmap for gene functional characterisation in crops with large genomes: Lessons from polyploid wheat. ELife, 2020, 9, . | 6.0 | 78 |
| 7 | Creation and judicious application of a wheat resistance gene atlas. Molecular Plant, 2021, 14, 1053-1070. | 8.3 | 66 |
| 8 | The wheat <i>Sr22</i> , <i>Sr33</i> , <i>Sr35</i> and <i>Sr45</i> genes confer resistance against stem rust in barley. Plant Biotechnology Journal, 2021, 19, 273-284. | 8.3 | 14 |
| 9 | Mutagenesis ofÂPuccinia graminisÂf. sp.Âtritici and Selection of Gain-of-Virulence Mutants. Frontiers in Plant Science, 2020, 11, 570180. | 3.6 | 13 |
| 10 | Discovery and characterisation of a new leaf rust resistance gene introgressed in wheat from wild wheat Aegilops peregrina. Scientific Reports, 2020, 10, 7573. | 3.3 | 13 |
| 11 | Fine mapping of Aegilops peregrina co-segregating leaf and stripe rust resistance genes to distal-most end of 5DS. Theoretical and Applied Genetics, 2019, 132, 1473-1485. | 3.6 | 8 |
| 12 | Extensive Genetic Variation at the Sr22 Wheat Stem Rust Resistance Gene Locus in the Grasses Revealed Through Evolutionary Genomics and Functional Analyses. Molecular Plant-Microbe Interactions, 2020, 33, 1286-1298. | 2.6 | 6 |