

Ofere Francis Emeriewen

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

245
citations

1163117

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1199594

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times ranked

159
citing authors

#	ARTICLE	IF	CITATIONS
1	Malus Hostsâ€™Erwinia amylovora Interactions: Strain Pathogenicity and Resistance Mechanisms. <i>Frontiers in Plant Science</i> , 2019, 10, 551.	3.6	38
2	Status of fire blight resistance breeding in Malus. <i>Journal of Plant Pathology</i> , 2021, 103, 3-12.	1.2	33
3	Fire blight resistance of Malus Ã—arnoldiana is controlled by a quantitative trait locus located at the distal end of linkage group 12. <i>European Journal of Plant Pathology</i> , 2017, 148, 1011-1018.	1.7	32
4	Towards map-based cloning of FB_Mfu10: identification of a receptor-like kinase candidate gene underlying the Malus fusca fire blight resistance locus on linkage group 10. <i>Molecular Breeding</i> , 2018, 38, 106.	2.1	28
5	Mapping of fire blight resistance in Malus Ã—robusta 5 flowers following artificial inoculation. <i>BMC Plant Biology</i> , 2019, 19, 532.	3.6	24
6	Apple blotch disease (<i>Marssonina coronaria</i> (Ellis & Davis) Davis) â€™ review and research prospects. <i>European Journal of Plant Pathology</i> , 2019, 153, 657-669.	1.7	22
7	Construction of a dense genetic map of the Malus fusca fire blight resistant accession MAL0045 using tunable genotyping-by-sequencing SNPs and microsatellites. <i>Scientific Reports</i> , 2020, 10, 16358.	3.3	17
8	The fire blight resistance QTL of Malus fusca (Mfu10) is affected but not broken down by the highly virulent Canadian Erwinia amylovora strain E2002A. <i>European Journal of Plant Pathology</i> , 2015, 141, 631-635.	1.7	16
9	Genetic Analysis and Fine Mapping of the Fire Blight Resistance Locus of Malus Ã—arnoldiana on Linkage Group 12 Reveal First Candidate Genes. <i>Frontiers in Plant Science</i> , 2021, 12, 667133.	3.6	12
10	Recent Developments and Strategies for the Application of Agrobacterium-Mediated Transformation of Apple Malus Ã— domestica Borkh. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	7
11	Mapping of the Waxy Bloom Gene in â€™Black Jewelâ€™™ in a Parental Linkage Map of â€™Black Jewelâ€™™ Ã— â€™Glen Ampleâ€™™ (Rubus) Interspecific Population. <i>Agronomy</i> , 2020, 10, 1579.	3.0	5
12	Characterization of genomic DNA sequence of the candidate gene for FB_Mfu10 associated with fire blight resistance in Malus species. <i>BMC Research Notes</i> , 2021, 14, 291.	1.4	5
13	Evidence of apple blotch resistance in wild apple germplasm (Malus spp.) accessions. <i>European Journal of Plant Pathology</i> , 2021, 159, 441-448.	1.7	1