

Juliana Benevenuto

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

618
citations

932766

10
h-index

940134

16
g-index

19
all docs

19
docs citations

19
times ranked

720
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic prediction for canopy height and dry matter yield in alfalfa using family bulks. <i>Plant Genome</i> , 2022, 15, .	1.6	10
2	Complete Chromosome-Scale Genome Sequence Resource for <i>Sporisorium panici-leucophaei</i> , the Causal Agent of Sourgrass Smut Disease. <i>Molecular Plant-Microbe Interactions</i> , 2021, 34, 448-452.	1.4	3
3	Genomic Selection in an Outcrossing Autotetraploid Fruit Crop: Lessons From Blueberry Breeding. <i>Frontiers in Plant Science</i> , 2021, 12, 676326.	1.7	26
4	Genome-wide association of volatiles reveals candidate loci for blueberry flavor. <i>New Phytologist</i> , 2020, 226, 1725-1737.	3.5	84
5	High-Resolution Linkage Map and QTL Analyses of Fruit Firmness in Autotetraploid Blueberry. <i>Frontiers in Plant Science</i> , 2020, 11, 562171.	1.7	19
6	Impact of dominance effects on autotetraploid genomic prediction. <i>Crop Science</i> , 2020, 60, 656-665.	0.8	28
7	How can a high-quality genome assembly help plant breeders?. <i>GigaScience</i> , 2019, 8, .	3.3	67
8	Cost-effective detection of genome-wide signatures for 2,4-D herbicide resistance adaptation in red clover. <i>Scientific Reports</i> , 2019, 9, 20037.	1.6	6
9	Molecular and Genetic Bases of Fruit Firmness Variation in Blueberry—A Review. <i>Agronomy</i> , 2018, 8, 174.	1.3	35
10	Insights Into the Genetic Basis of Blueberry Fruit-Related Traits Using Diploid and Polyploid Models in a GWAS Context. <i>Frontiers in Ecology and Evolution</i> , 2018, 6, .	1.1	60
11	Comparative Genomics of Smut Pathogens: Insights From Orphans and Positively Selected Genes Into Host Specialization. <i>Frontiers in Microbiology</i> , 2018, 9, 660.	1.5	33
12	Progress in understanding fungal diseases affecting sugarcane: smut. <i>Burleigh Dodds Series in Agricultural Science</i> , 2018, , 221-243.	0.1	2
13	Using Population and Comparative Genomics to Understand the Genetic Basis of Effector-Driven Fungal Pathogen Evolution. <i>Frontiers in Plant Science</i> , 2017, 8, 119.	1.7	135
14	Molecular variability and genetic relationship among Brazilian strains of the sugarcane smut fungus. <i>FEMS Microbiology Letters</i> , 2016, 363, fnw277.	0.7	6
15	Complete Genome Sequence of <i>Sporisorium scitamineum</i> and Biotrophic Interaction Transcriptome with Sugarcane. <i>PLoS ONE</i> , 2015, 10, e0129318.	1.1	93
16	Conservation study of an endangered stingless bee (<i>Melipona capixaba</i>) (Hymenoptera: Apidae) with restricted distribution in Brazil. <i>Journal of Insect Conservation</i> , 2014, 18, 317-326.	0.8	6