Thomas M Davis

List of Publications by Year in descending order

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1040056 1125743 1,731 14 9 13 citations h-index g-index papers 15 15 15 2346 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	The genome of woodland strawberry (Fragaria vesca). Nature Genetics, 2011, 43, 109-116.	21.4	1,091
2	Multiple Models for Rosaceae Genomics. Plant Physiology, 2008, 147, 985-1003.	4.8	291
3	Development and preliminary evaluation of a 90ÂK Axiom® SNP array for the allo-octoploid cultivated strawberry Fragaria × ananassa. BMC Genomics, 2015, 16, 155.	2.8	179
4	RosBREED: bridging the chasm between discovery and application to enable DNA-informed breeding in rosaceous crops. Horticulture Research, 2020, 7, 177.	6.3	34
5	A New Perspective on Polyploid Fragaria (Strawberry) Genome Composition Based on Large-Scale, Multi-Locus Phylogenetic Analysis. Genome Biology and Evolution, 2017, 9, 3433-3448.	2.5	29
6	Simple Sequence Repeat Marker Development and Mapping Targeted to Previously Unmapped Regions of the Strawberry Genome Sequence. Plant Genome, 2011, 4, 165-177.	2.8	28
7	Editorial: Wild Plants as Source of New Crops. Frontiers in Plant Science, 2020, 11, 591554.	3.6	27
8	A Highâ€Density Linkage Map of the Ancestral Diploid Strawberry, Fragaria iinumae , Constructed with Single Nucleotide Polymorphism Markers from the IStraw90 Array and Genotyping by Sequencing. Plant Genome, 2016, 9, plantgenome2015.08.0071.	2.8	23
9	Insight into octoploid strawberry (<i>Fragaria</i>) subgenome composition revealed by GISH analysis of pentaploid hybrids. Genome, 2016, 59, 79-86.	2.0	11
10	Developing Chenopodium ficifolium as a potential B genome diploid model system for genetic characterization and improvement of allotetraploid quinoa (Chenopodium quinoa). BMC Plant Biology, 2021, 21, 490.	3.6	8
11	A pentaploid-based linkage map of the ancestral octoploid strawberry Fragaria virginiana reveals instances of sporadic hyper-recombination. Horticulture Research, 2020, 7, 77.	6.3	4
12	Defining the mutation sites in chickpea nodulation mutants PM233 and PM405. BMC Plant Biology, 2022, 22, 66.	3.6	3
13	Evaluation of Disease Severity and Molecular Relationships Between <i>Peronospora variabilis</i> Isolates on <i>Chenopodium</i> Species in New Hampshire. Plant Disease, 2022, 106, 564-571.	1.4	2
14	Enhanced Documentation of <i>Chenopodium foggii < /i> (Amaranthaceae) in Northern New England. Rhodora, 2018, 120, 257-259.</i>	0.1	0