Lydia Gramzow

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8100146/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The Norway spruce genome sequence and conifer genome evolution. Nature, 2013, 497, 579-584.	27.8	1,303
2	The Selaginella Genome Identifies Genetic Changes Associated with the Evolution of Vascular Plants. Science, 2011, 332, 960-963.	12.6	794
3	A hitchhiker's guide to the MADS world of plants. Genome Biology, 2010, 11, 214.	9.6	252
4	On the origin of MADS-domain transcription factors. Trends in Genetics, 2010, 26, 149-153.	6.7	123
5	Horizontal gene transfer and functional diversification of plant cell wall degrading polygalacturonases: Key events in the evolution of herbivory in beetles. Insect Biochemistry and Molecular Biology, 2014, 52, 33-50.	2.7	116
6	MADS goes genomic in conifers: towards determining the ancestral set of MADS-box genes in seed plants. Annals of Botany, 2014, 114, 1407-1429.	2.9	101
7	Live and Let Die - The Bsister MADS-Box Gene OsMADS29 Controls the Degeneration of Cells in Maternal Tissues during Seed Development of Rice (Oryza sativa). PLoS ONE, 2012, 7, e51435.	2.5	73
8	The Molecular Evolution of Cytochrome P450 Genes within and between Drosophila Species. Genome Biology and Evolution, 2014, 6, 1118-1134.	2.5	72
9	Phylogenomics of MADS-Box Genes in Plants — Two Opposing Life Styles in One Gene Family. Biology, 2013, 2, 1150-1164.	2.8	70
10	Phylogenomics reveals surprising sets of essential and dispensable clades of MIKC ^c â€group MADSâ€box genes in flowering plants. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2015, 324, 353-362.	1.3	69
11	GORDITA (AGL63) is a young paralog of the Arabidopsis thaliana Bsister MADS box gene ABS (TT16) that has undergone neofunctionalization. Plant Journal, 2010, 63, 914-924.	5.7	49
12	SR1—a small RNA with two remarkably conserved functions. Nucleic Acids Research, 2012, 40, 11659-11672.	14.5	42
13	Two independent duplications forming the Cyp307a genes in Drosophila. Insect Biochemistry and Molecular Biology, 2007, 37, 1044-1053.	2.7	37
14	Array of MADS-Box Genes: Facilitator for Rapid Adaptation?. Trends in Plant Science, 2018, 23, 563-576.	8.8	35
15	Selaginella Genome Analysis – Entering the "Homoplasy Heaven―of the MADS World. Frontiers in Plant Science, 2012, 3, 214.	3.6	31
16	Structure and Evolution of Plant MADS Domain Transcription Factors. , 2016, , 127-138.		30
17	Did Convergent Protein Evolution Enable Phytoplasmas to Generate â€~Zombie Plants'?. Trends in Plant Science, 2015, 20, 798-806	8.8	28
18	<i>Aethionema arabicum</i> genome annotation using PacBio fullâ€length transcripts provides a valuable resource for seed dormancy and Brassicaceae evolution research. Plant Journal, 2021, 106, 275-293.	5.7	20

Lydia Gramzow

#	Article	IF	CITATIONS
19	Nonâ€canonical structure, function and phylogeny of the B sister MADS â€box gene O s MADS 30 of rice () Tj ET	Qq1_1 0.7	784314 rgBT 16
20	Plant miRNA Conservation and Evolution. Methods in Molecular Biology, 2019, 1932, 41-50.	0.9	14
21	A Dead Gene Walking: Convergent Degeneration of a Clade of MADS-Box Genes in Crucifers. Molecular Biology and Evolution, 2018, 35, 2618-2638.	8.9	10
22	Independent origin of <i>MIRNA</i> genes controlling homologous target genes by partial inverted duplication of antisenseâ€transcribed sequences. Plant Journal, 2020, 101, 401-419.	5.7	7
23	Comparative transcriptomics identifies candidate genes involved in the evolutionary transition from dehiscent to indehiscent fruits in Lepidium (Brassicaceae). BMC Plant Biology, 2022, 22, .	3.6	3
24	Stranger than Fiction: Loss of MADS-Box Genes During Evolutionary Miniaturization of the Duckweed Body Plan. Compendium of Plant Genomes, 2020, , 91-101.	0.5	1