

Molecular and Phylogenetic Analyses of the Complete MADS-Box Gene Family in Arabidopsis

Plant Cell

15, 1538-1551

DOI: [10.1105/tpc.011544](https://doi.org/10.1105/tpc.011544)

Citation Report

#	ARTICLE	IF	CITATIONS
2	And then there were many: MADS goes genomic. <i>Trends in Plant Science</i> , 2003, 8, 475-483.	4.3	179
3	Expression and Maintenance of Embryogenic Potential Is Enhanced through Constitutive Expression of AGAMOUS-Like 15 Å. <i>Plant Physiology</i> , 2003, 133, 653-663.	2.3	225
4	Toward the Analysis of the Petunia MADS Box Gene Family by Reverse and Forward Transposon Insertion Mutagenesis Approaches: B, C, and D Floral Organ Identity Functions Require SEPALLATA-Like MADS Box Genes in Petunia. <i>Plant Cell</i> , 2003, 15, 2680-2693.	3.1	188
5	Systematic Reverse Genetic Screening of T-DNA Tagged Genes in Rice for Functional Genomic Analyses: MADS-box Genes as a Test Case. <i>Plant and Cell Physiology</i> , 2003, 44, 1403-1411.	1.5	95
6	Identification and Characterization of Four Chrysanthemum MADS-Box Genes, Belonging to the APETALA1/FRUITFULL and SEPALLATA3 Subfamilies. <i>Plant Physiology</i> , 2004, 134, 1632-1641.	2.3	111
7	Genome-Wide ORFeome Cloning and Analysis of Arabidopsis Transcription Factor Genes. <i>Plant Physiology</i> , 2004, 135, 773-782.	2.3	205
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19	Conservation and diversity in flower land. <i>Current Opinion in Plant Biology</i> , 2004, 7, 84-91.	3.5	143

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#	ARTICLE	IF	CITATIONS
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#	ARTICLE	IF	CITATIONS
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#	ARTICLE	IF	CITATIONS
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