

Michael G Muszynski

List of Publications by Year in descending order

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32
papers

2,722
citations

236925

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454955

30
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36
all docs

36
docs citations

36
times ranked

3325
citing authors

#	ARTICLE	IF	CITATIONS
1	The Control of Spikelet Meristem Identity by the branched silkless1 Gene in Maize. <i>Science</i> , 2002, 298, 1238-1241.	12.6	270
2	The <i>FT</i> -Like <i>ZCN8</i> Gene Functions as a Floral Activator and Is Involved in Photoperiod Sensitivity in Maize. <i>Plant Cell</i> , 2011, 23, 942-960.	6.6	265
3	Conserved plant genes with similarity to mammalian de novo DNA methyltransferases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 4979-4984.	7.1	222
4	Duplicated <i>ft</i> Genes in Maize. <i>Plant Cell</i> , 2003, 15, 425-438.	6.6	173
5	delayed flowering1 Encodes a Basic Leucine Zipper Protein That Mediates Floral Inductive Signals at the Shoot Apex in Maize. <i>Plant Physiology</i> , 2006, 142, 1523-1536.	4.8	161
6	Dynamic Changes in <i>ANGUSTIFOLIA3</i> Complex Composition Reveal a Growth Regulatory Mechanism in the Maize Leaf. <i>Plant Cell</i> , 2015, 27, 1605-1619.	6.6	154
7	The Maize Homologue of the Cell Cycle Checkpoint Protein <i>MAD2</i> Reveals Kinetochore Substructure and Contrasting Mitotic and Meiotic Localization Patterns. <i>Journal of Cell Biology</i> , 1999, 145, 425-435.	5.2	125
8	Genome-Wide Distribution of Transposed <i>Dissociation</i> Elements in Maize. <i>Plant Cell</i> , 2010, 22, 1667-1685.	6.6	123
9	A Maize Homolog of Mammalian <i>CENPC</i> Is a Constitutive Component of the Inner Kinetochore. <i>Plant Cell</i> , 1999, 11, 1227-1238.	6.6	122
10	Maize Chromomethylase <i>Zea methyltransferase2</i> Is Required for CpNpG Methylation. <i>Plant Cell</i> , 2001, 13, 1919-1928.	6.6	120
11	Involvement of the <i>MADS</i> -Box Gene <i>ZMM4</i> in Floral Induction and Inflorescence Development in Maize. <i>Plant Physiology</i> , 2008, 147, 2054-2069.	4.8	117
12	The Boron Efflux Transporter <i>ROTTEN EAR</i> Is Required for Maize Inflorescence Development and Fertility. <i>Plant Cell</i> , 2014, 26, 2962-2977.	6.6	91
13	Altered expression of maize <i>PLASTOCHRON1</i> enhances biomass and seed yield by extending cell division duration. <i>Nature Communications</i> , 2017, 8, 14752.	12.8	89
14	Maize Chromomethylase <i>Zea methyltransferase2</i> Is Required for CpNpG Methylation. <i>Plant Cell</i> , 2001, 13, 1919-1928.	6.6	86
15	Characterization of Grain Filling Patterns in Diverse Maize Germplasm. <i>Crop Science</i> , 2009, 49, 999-1009.	1.8	74
16	Maximum Likelihood Methods Reveal Conservation of Function Among Closely Related Kinesin Families. <i>Journal of Molecular Evolution</i> , 2002, 54, 42-53.	1.8	64
17	<i>tie-dyed1</i> Regulates Carbohydrate Accumulation in Maize Leaves. <i>Plant Physiology</i> , 2006, 142, 1511-1522.	4.8	59
18	Three <i>FLOWERING LOCUS T</i> -like genes function as potential florigens and mediate photoperiod response in sorghum. <i>New Phytologist</i> , 2016, 210, 946-959.	7.3	59

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19	Molecular Analysis of Two Pyruvate Dehydrogenase Kinases from Maize. <i>Journal of Biological Chemistry</i> , 1998, 273, 26618-26623.	3.4	41
20	Regional mutagenesis using Dissociation in maize. <i>Methods</i> , 2009, 49, 248-254.	3.8	40
21	Beyond flowering time. <i>Plant Signaling and Behavior</i> , 2011, 6, 1267-1270.	2.4	39
22	The Dihydrolipoamide S-Acetyltransferase Subunit of the Mitochondrial Pyruvate Dehydrogenase Complex from Maize Contains a Single Lipoyl Domain. <i>Journal of Biological Chemistry</i> , 1999, 274, 21769-21775.	3.4	33
23	Characterization of a gene from <i>Zea mays</i> related to the <i>Arabidopsis</i> flowering-time gene LUMINIDEPENDENS. <i>Plant Molecular Biology</i> , 2000, 44, 107-122.	3.9	31
24	Over-expression of the photoperiod response regulator ZmCCT10 modifies plant architecture, flowering time and inflorescence morphology in maize. <i>PLoS ONE</i> , 2019, 14, e0203728.	2.5	30
25	The Maize <i>Hairy Sheath Frayed1</i> (<i>Hsf1</i>) Mutation Alters Leaf Patterning through Increased Cytokinin Signaling. <i>Plant Cell</i> , 2020, 32, 1501-1518.	6.6	30
26	A Pectin Methylesterase ZmPme3 Is Expressed in Gametophyte factor1-s (<i>Ga1-s</i>) Silks and Maps to that Locus in Maize (<i>Zea mays</i> L.). <i>Frontiers in Plant Science</i> , 2017, 8, 1926.	3.6	28
27	The Maize Floral Transition. , 2009, , 41-55.		27
28	A maize mutant with decreased capacity to accumulate chloroplast protein synthesis elongation factor (EF-Tu) displays reduced tolerance to heat stress. <i>Plant Science</i> , 2004, 167, 1367-1374.	3.6	24
29	Genetic and molecular analysis of a three-component transposable-element system in maize. <i>Molecular Genetics and Genomics</i> , 1993, 237-237, 105-112.	2.4	16
30	ZMPP2, a novel type-2C protein phosphatase from maize. <i>Journal of Experimental Botany</i> , 2001, 52, 1739-1740.	4.8	4
31	Putting the Function in Maize Genomics. <i>Plant Genome</i> , 2009, 2, .	2.8	1
32	Understanding and Manipulation of the Flowering Network and the Perfection of Seed Quality. , 2010, , 167-198.		1