Andreas E Muller

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Analysis of histone acetyltransferase and histone deacetylase families of Arabidopsis thaliana suggests functional diversification of chromatin modification among multicellular eukaryotes. Nucleic Acids Research, 2002, 30, 5036-5055.	14.5	672
2	Flowering time control and applications in plant breeding. Trends in Plant Science, 2009, 14, 563-573.	8.8	480
3	Effectiveness of RNA interference in transgenic plants. FEBS Letters, 2004, 566, 223-228.	2.8	188
4	The Role of a Pseudo-Response Regulator Gene in Life Cycle Adaptation and Domestication of Beet. Current Biology, 2012, 22, 1095-1101.	3.9	135
5	Sugar beet contains a large CONSTANS-LIKE gene family including a CO homologue that is independent of the early-bolting (B) gene locus. Journal of Experimental Botany, 2008, 59, 2735-2748.	4.8	118
6	Flowering time variation in oilseed rape (Brassica napus L.) is associated with allelic variation in the FRIGIDA homologue BnaA.FRI.a. Journal of Experimental Botany, 2011, 62, 5641-5658.	4.8	114
7	Palindromic sequences and A+T-rich DNA elements promote illegitimate recombination in Nicotiana tabacum. Journal of Molecular Biology, 1999, 291, 29-46.	4.2	64
8	Siteâ€directed mutagenesis in bread and durum wheat via pollination by <i>cas9</i> /guide RNAâ€ŧransgenic maize used as haploidy inducer. Plant Biotechnology Journal, 2020, 18, 2376-2378.	8.3	48
9	Conservation and divergence of autonomous pathway genes in the flowering regulatory network of Beta vulgaris. Journal of Experimental Botany, 2011, 62, 3359-3374.	4.8	41
10	A repetitive DNA fragment carrying a hot spot for de novo DNA methylation enhances expression variegation in tobacco and petunia. Plant Journal, 1995, 8, 919-932.	5.7	38
11	A survey of EMS-induced biennial Beta vulgaris mutants reveals a novel bolting locus which is unlinked to the bolting gene B. Theoretical and Applied Genetics, 2010, 121, 1117-1131.	3.6	36
12	The transformation booster sequence from Petunia hybrida is a retrotransposon derivative that binds to the nuclear scaffold. Molecular Genetics and Genomics, 1995, 247, 614-622.	2.4	34
13	The FLC-like gene BvFL1 is not a major regulator of vernalization response in biennial beets. Frontiers in Plant Science, 2014, 5, 146.	3.6	33
14	Genetic analysis of bolting after winter in sugar beet (Beta vulgaris L.). Theoretical and Applied Genetics, 2014, 127, 2479-2489.	3.6	32
15	EcoTILLING in Beta vulgaris reveals polymorphisms in the FLC-like gene BvFL1that are associated with annuality and winter hardiness. BMC Plant Biology, 2013, 13, 52.	3.6	31
16	Bolting and flowering control in sugar beet: relationships and effects of gibberellin, the bolting gene B and vernalization. AoB PLANTS, 2010, 2010, plq012.	2.3	27
17	A Detailed Analysis of the BR1 Locus Suggests a New Mechanism for Bolting after Winter in Sugar Beet (Beta vulgaris L.). Frontiers in Plant Science, 2016, 7, 1662.	3.6	26
18	Analysis of hypermethylation in the RPS element suggests a signal function for short inverted repeats in de novo methylation. Plant Molecular Biology, 2002, 48, 383-399.	3.9	20

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19	Genetic identification of a novel bolting locus in Beta vulgaris which promotes annuality independently of the bolting gene B. Molecular Breeding, 2012, 29, 989-998.	2.1	16
20	A Paragenetic Perspective on Integration of RNA Silencing into the Epigenome and Its Role in the Biology of Higher Plants. Cold Spring Harbor Symposia on Quantitative Biology, 2006, 71, 481-485.	1.1	15
21	Microhomologies between T-DNA ends and target sites often occur in inverted orientation and may be responsible for the high frequency of T-DNA-associated inversions. Plant Cell Reports, 2007, 26, 617-630.	5.6	14
22	Genetics and Genomics of Flowering Time Regulation in Sugar Beet. , 2014, , 3-26.		9
23	BvPRR7 is a cold responsive gene with a clock function in beet. Biologia Plantarum, 2016, 60, 95-104.	1.9	8
24	QTL for delayed bolting after winter detected in leaf beet (<i>BetaÂvulgaris</i> L.). Plant Breeding, 2017, 136, 237-244.	1.9	7
25	Gene Silencing in Plants: Transgenes as Targets and Effectors. Biotechnology in Agriculture and Forestry, 2010, , 79-101.	0.2	4
26	Control and Silencing of Transgene Expression. , 0, , .		3
27	Motywy dziaÅ,alnoÅ›ci charytatywnej w KoÅ›ciele starożytnym. Teologia Praktyczna, 2019, , 9-20. 	0.0	0