Qiuying Yang

List of Publications by Year in descending order

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932766 1199166 12 585 10 12 citations h-index g-index papers 12 12 12 871 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	RNA Interference in Fungi: Pathways, Functions, and Applications. Eukaryotic Cell, 2011, 10, 1148-1155.	3.4	191
2	AhSSK1, a novel SKP1-like protein that interacts with the S-locus F-box protein SLF. Plant Journal, 2006, 46, 780-793.	2.8	83
3	The DNA/RNA-Dependent RNA Polymerase QDE-1 Generates Aberrant RNA and dsRNA for RNAi in a Process Requiring Replication Protein A and a DNA Helicase. PLoS Biology, 2010, 8, e1000496.	2.6	71
4	Molecular Cytogenetic Characterization of the Antirrhinum majus GenomeSequence data from this article have been deposited with the EMBL/GenBank Data Libraries under the accession nos. AY630561 (for BAC 5E10) and AY6305612 (for BAC 36D21) Genetics, 2005, 169, 325-335.	1.2	42
5	Diversity of centromeric repeats in two closely related wild rice species, Oryza officinalis and Oryza rhizomatis. Molecular Genetics and Genomics, 2006, 275, 421-430.	1.0	38
6	Transcription of the Major Neurospora crassa microRNA–Like Small RNAs Relies on RNA Polymerase III. PLoS Genetics, 2013, 9, e1003227.	1.5	38
7	Mechanism of siRNA production from repetitive DNA. Genes and Development, 2015, 29, 526-537.	2.7	34
8	Heterochromatic and genetic features are consistent with recombination suppression of the self-incompatibility locus in Antirrhinum. Plant Journal, 2007, 51, 140-151.	2.8	30
9	Genetic features of a pollen-part mutation suggest an inhibitory role for the Antirrhinum pollen self-incompatibility determinant. Plant Molecular Biology, 2009, 70, 499-509.	2.0	30
10	Histone H3K56 Acetylation Is Required for Quelling-induced Small RNA Production through Its Role in Homologous Recombination. Journal of Biological Chemistry, 2014, 289, 9365-9371.	1.6	16
11	Digital Inventory of Arabidopsis Transcripts Revealed by 61 RNA Sequencing Samples. Plant Physiology, 2014, 166, 869-878.	2.3	8
12	Cytological characterization of the tandem repetitive sequences and their methylation status in the Antirrhinum majus genome. Genomics, 2008, 92, 107-114.	1.3	4