

Olga Mf Pontes

List of Publications by Year in descending order

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36
papers

6,436
citations

186265

28
h-index

377865

34
g-index

56
all docs

56
docs citations

56
times ranked

6618
citing authors

#	ARTICLE	IF	CITATIONS
1	Gateway-compatible vectors for plant functional genomics and proteomics. <i>Plant Journal</i> , 2006, 45, 616-629.	5.7	1,658
2	Plant Nuclear RNA Polymerase IV Mediates siRNA and DNA Methylation-Dependent Heterochromatin Formation. <i>Cell</i> , 2005, 120, 613-622.	28.9	602
3	The Arabidopsis Chromatin-Modifying Nuclear siRNA Pathway Involves a Nucleolar RNA Processing Center. <i>Cell</i> , 2006, 126, 79-92.	28.9	399
4	An ARGONAUTE4-Containing Nuclear Processing Center Colocalized with Cajal Bodies in Arabidopsis thaliana. <i>Cell</i> , 2006, 126, 93-106.	28.9	350
5	A Concerted DNA Methylation/Histone Methylation Switch Regulates rRNA Gene Dosage Control and Nucleolar Dominance. <i>Molecular Cell</i> , 2004, 13, 599-609.	9.7	336
6	Chromosomal locus rearrangements are a rapid response to formation of the allotetraploid Arabidopsis suecica genome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 18240-18245.	7.1	251
7	An SNF2 Protein Associated with Nuclear RNA Silencing and the Spread of a Silencing Signal between Cells in Arabidopsis. <i>Plant Cell</i> , 2007, 19, 1507-1521.	6.6	251
8	An RNA polymerase II- and AGO4-associated protein acts in RNA-directed DNA methylation. <i>Nature</i> , 2010, 465, 106-109.	27.8	228
9	A Histone Acetyltransferase Regulates Active DNA Demethylation in Arabidopsis. <i>Science</i> , 2012, 336, 1445-1448.	12.6	224
10	An Effector of RNA-Directed DNA Methylation in Arabidopsis Is an ARGONAUTE 4- and RNA-Binding Protein. <i>Cell</i> , 2009, 137, 498-508.	28.9	220
11	Erasure of histone acetylation by Arabidopsis HDA6 mediates large-scale gene silencing in nucleolar dominance. <i>Genes and Development</i> , 2006, 20, 1283-1293.	5.9	219
12	VIM1, a methylcytosine-binding protein required for centromeric heterochromatinization. <i>Genes and Development</i> , 2007, 21, 267-277.	5.9	167
13	ROS3 is an RNA-binding protein required for DNA demethylation in Arabidopsis. <i>Nature</i> , 2008, 455, 1259-1262.	27.8	150
14	Multimegabase Silencing in Nucleolar Dominance Involves siRNA-Directed DNA Methylation and Specific Methylcytosine-Binding Proteins. <i>Molecular Cell</i> , 2008, 32, 673-684.	9.7	144
15	Mechanisms of HDA6-mediated rRNA gene silencing: suppression of intergenic Pol II transcription and differential effects on maintenance versus siRNA-directed cytosine methylation. <i>Genes and Development</i> , 2010, 24, 1119-1132.	5.9	143
16	NRPD4, a protein related to the RPB4 subunit of RNA polymerase II, is a component of RNA polymerases IV and V and is required for RNA-directed DNA methylation. <i>Genes and Development</i> , 2009, 23, 318-330.	5.9	126
17	JMJ14, a JmjC domain protein, is required for RNA silencing and cell-to-cell movement of an RNA silencing signal in Arabidopsis. <i>Genes and Development</i> , 2010, 24, 986-991.	5.9	116
18	siRNA and miRNA processing: new functions for Cajal bodies. <i>Current Opinion in Genetics and Development</i> , 2008, 18, 197-203.	3.3	103

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19	A conserved transcriptional regulator is required for RNA-directed DNA methylation and plant development. <i>Genes and Development</i> , 2009, 23, 2717-2722.	5.9	92
20	Natural variation in nucleolar dominance reveals the relationship between nucleolus organizer chromatin topology and rRNA gene transcription in <i>Arabidopsis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 11418-11423.	7.1	85
21	A DNA 3' Phosphatase Functions in Active DNA Demethylation in <i>Arabidopsis</i> . <i>Molecular Cell</i> , 2012, 45, 357-370.	9.7	81
22	Posttranscriptional gene silencing in nuclei. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 409-414.	7.1	80
23	Postembryonic Establishment of Megabase-Scale Gene Silencing in Nucleolar Dominance. <i>PLoS ONE</i> , 2007, 2, e1157.	2.5	69
24	RNA Polymerase V Functions in <i>Arabidopsis</i> Interphase Heterochromatin Organization Independently of the 24-nt siRNA-Directed DNA Methylation Pathway. <i>Molecular Plant</i> , 2009, 2, 700-710.	8.3	63
25	Metal A and Metal B Sites of Nuclear RNA Polymerases Pol IV and Pol V Are Required for siRNA-Dependent DNA Methylation and Gene Silencing. <i>PLoS ONE</i> , 2009, 4, e4110.	2.5	51
26	Heterochromatic siRNAs and DDM1 Independently Silence Aberrant 5S rDNA Transcripts in <i>Arabidopsis</i> . <i>PLoS ONE</i> , 2009, 4, e5932.	2.5	42
27	A subgroup of SGS3-like proteins act redundantly in RNA-directed DNA methylation. <i>Nucleic Acids Research</i> , 2012, 40, 4422-4431.	14.5	40
28	Intersection of Small RNA Pathways in <i>Arabidopsis thaliana</i> Sub-Nuclear Domains. <i>PLoS ONE</i> , 2013, 8, e65652.	2.5	40
29	DNA Topoisomerase I Promotes Transcriptional Silencing of Transposable Elements through DNA Methylation and Histone Lysine 9 Dimethylation in <i>Arabidopsis</i> . <i>PLoS Genetics</i> , 2014, 10, e1004446.	3.5	26
30	Extra views on RNA-dependent DNA methylation and MBD6-dependent heterochromatin formation in nucleolar dominance. <i>Nucleus</i> , 2010, 1, 254-259.	2.2	20
31	IDN2 Interacts with RPA and Facilitates DNA Double-Strand Break Repair by Homologous Recombination in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2017, 29, 589-599.	6.6	19
32	Extra views on RNA-dependent DNA methylation and MBD6-dependent heterochromatin formation in nucleolar dominance. <i>Nucleus</i> , 2010, 1, 254-259.	2.2	15
33	The cytological and molecular role of DOMAINS REARRANGED METHYLTRANSFERASE3 in RNA-dependent DNA methylation of <i>Arabidopsis thaliana</i> . <i>BMC Research Notes</i> , 2014, 7, 721.	1.4	13
34	Heterochromatin: condense or excise. <i>Nature Cell Biology</i> , 2007, 9, 19-20.	10.3	12
35	Connecting the dots of RNA-directed DNA methylation in <i>Arabidopsis thaliana</i> . <i>Chromosome Research</i> , 2014, 22, 225-240.	2.2	0
36	Roles of non-coding, Pol IV-dependent RNAs in megabase-scale silencing and long-range heterochromatin interactions in <i>Arabidopsis</i> . <i>FASEB Journal</i> , 2008, 22, 534.3.	0.5	0