Pauline E Jullien

List of Publications by Year in descending order

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758635 752256 19 1,883 12 20 h-index citations g-index papers 25 25 25 1978 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Reprogramming of DNA Methylation in Pollen Guides Epigenetic Inheritance via Small RNA. Cell, 2012, 151, 194-205.	13.5	506
2	Maintenance of DNA Methylation during the Arabidopsis Life Cycle Is Essential for Parental Imprinting. Plant Cell, 2006, 18, 1360-1372.	3.1	264
3	Retinoblastoma and Its Binding Partner MSI1 Control Imprinting in Arabidopsis. PLoS Biology, 2008, 6, e194.	2.6	220
4	DNA Methylation Dynamics during Sexual Reproduction in Arabidopsis thaliana. Current Biology, 2012, 22, 1825-1830.	1.8	214
5	Polycomb Group Complexes Self-Regulate Imprinting of the Polycomb Group Gene MEDEA in Arabidopsis. Current Biology, 2006, 16, 486-492.	1.8	194
6	The Female Gametophyte and the Endosperm Control Cell Proliferation and Differentiation of the Seed Coat in Arabidopsis. Plant Cell, 2007, 18, 3491-3501.	3.1	111
7	Gamete-specific epigenetic mechanisms shape genomic imprinting. Current Opinion in Plant Biology, 2009, 12, 637-642.	3.5	64
8	Parental Genome Dosage Imbalance Deregulates Imprinting in Arabidopsis. PLoS Genetics, 2010, 6, e1000885.	1.5	63
9	Movement and differential consumption of short interfering RNA duplexes underlie mobile RNA interference. Nature Plants, 2020, 6, 789-799.	4.7	54
10	DNA methylation reprogramming during plant sexual reproduction?. Trends in Genetics, 2010, 26, 394-399.	2.9	42
11	DNA Methylation Influences the Expression of <i>DICER-LIKE4</i> Isoforms, Which Encode Proteins of Alternative Localization and Function. Plant Cell, 2016, 28, 2786-2804.	3.1	41
12	Functional characterization of Arabidopsis ARGONAUTE 3 in reproductive tissues. Plant Journal, 2020, 103, 1796-1809.	2.8	22
13	Asymmetric expression of Argonautes in reproductive tissues. Plant Physiology, 2022, 188, 38-43.	2.3	17
14	The miRNome function transitions from regulating developmental genes to transposable elements during pollen maturation. Plant Cell, 2022, 34, 784-801.	3.1	17
15	A complex of <i>Arabidopsis</i> DRB proteins can impair dsRNA processing. Rna, 2017, 23, 782-797.	1.6	13
16	Analysis of CATMA transcriptome data identifies hundreds of novel functional genes and improves gene models in the Arabidopsis genome. BMC Genomics, 2007, 8, 401.	1.2	12
17	Evolution of CG Methylation Maintenance Machinery in Plants. Epigenomes, 2021, 5, 19.	0.8	11
18	The Diversity of Plant Small RNAs Silencing Mechanisms. Chimia, 2019, 73, 362.	0.3	5

#	Article	IF	CITATIONS
19	DNA Methyltransferase 3 (MET3) is regulated by Polycomb group complex during Arabidopsis endosperm development. Plant Reproduction, 2022, 35, 141-151.	1.3	4