

Szymon AwiÅewski

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3356729/publications.pdf>

Version: 2024-02-01

14
papers

2,016
citations

759233

12
h-index

1058476

14
g-index

14
all docs

14
docs citations

14
times ranked

2502
citing authors

#	ARTICLE	IF	CITATIONS
1	Single seeds exhibit transcriptional heterogeneity during secondary dormancy induction. <i>Plant Physiology</i> , 2022, 190, 211-225.	4.8	12
2	Long Noncoding RNAs in Plants. <i>Annual Review of Plant Biology</i> , 2021, 72, 245-271.	18.7	83
3	Light Regulates Plant Alternative Splicing through the Control of Transcriptional Elongation. <i>Molecular Cell</i> , 2019, 73, 1066-1074.e3.	9.7	102
4	Arabidopsis SWI/SNF chromatin remodeling complex binds both promoters and terminators to regulate gene expression. <i>Nucleic Acids Research</i> , 2017, 45, gkw1273.	14.5	58
5	Developmental transitions in Arabidopsis are regulated by antisense RNAs resulting from bidirectionally transcribed genes. <i>RNA Biology</i> , 2017, 14, 838-842.	3.1	4
6	Antisense transcription represses <i>Arabidopsis</i> seed dormancy QTL <i>DOG1</i> to regulate drought tolerance. <i>EMBO Reports</i> , 2017, 18, 2186-2196.	4.5	42
7	Alternative Polyadenylation of the Sense Transcript Controls Antisense Transcription of DELAY OF GERMINATION 1 in Arabidopsis. <i>Molecular Plant</i> , 2017, 10, 1349-1352.	8.3	24
8	Control of seed dormancy in <i>Arabidopsis</i> by a <i>cis</i> -acting noncoding antisense transcript. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E7846-E7855.	7.1	113
9	Seed Dormancy in Arabidopsis Is Controlled by Alternative Polyadenylation of <i>DOG1</i> . <i>Plant Physiology</i> , 2016, 170, 947-955.	4.8	72
10	A specialized histone H1 variant is required for adaptive responses to complex abiotic stress and related DNA methylation in Arabidopsis. <i>Plant Physiology</i> , 2015, 169, pp.00493.2015.	4.8	101
11	<i>NTR1</i> is required for transcription elongation checkpoints at alternative exons in <i>Arabidopsis</i> . <i>EMBO Journal</i> , 2015, 34, 544-558.	7.8	52
12	Targeted 3' Processing of Antisense Transcripts Triggers <i>Arabidopsis FLC</i> Chromatin Silencing. <i>Science</i> , 2010, 327, 94-97.	12.6	435
13	Cold-induced silencing by long antisense transcripts of an Arabidopsis Polycomb target. <i>Nature</i> , 2009, 462, 799-802.	27.8	802
14	Small RNA-mediated chromatin silencing directed to the 3' region of the Arabidopsis gene encoding the developmental regulator, FLC. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 3633-3638.	7.1	116