

SÃ©bastien Aubourg

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2727839/publications.pdf>

Version: 2024-02-01

10
papers

2,420
citations

1040056

9
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

3554
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-Wide Analysis of Arabidopsis Pentatricopeptide Repeat Proteins Reveals Their Essential Role in Organelle Biogenesis [W]. <i>Plant Cell</i> , 2004, 16, 2089-2103.	6.6	1,132
2	High-quality de novo assembly of the apple genome and methylome dynamics of early fruit development. <i>Nature Genetics</i> , 2017, 49, 1099-1106.	21.4	693
3	Versatile Gene-Specific Sequence Tags for Arabidopsis Functional Genomics: Transcript Profiling and Reverse Genetics Applications. <i>Genome Research</i> , 2004, 14, 2176-2189.	5.5	282
4	CATdb: a public access to Arabidopsis transcriptome data from the URGV-CATMA platform. <i>Nucleic Acids Research</i> , 2007, 36, D986-D990.	14.5	160
5	The SCOOP12 peptide regulates defense response and root elongation in <i>Arabidopsis thaliana</i> . <i>Journal of Experimental Botany</i> , 2019, 70, 1349-1365.	4.8	59
6	Exploration of plant genomes in the FLAGdb++ environment. <i>Plant Methods</i> , 2011, 7, 8.	4.3	45
7	GEM2Net: from gene expression modeling to -omics networks, a new CATdb module to investigate <i>Arabidopsis thaliana</i> genes involved in stress response. <i>Nucleic Acids Research</i> , 2015, 43, D1010-D1017.	14.5	14
8	Analysis of CATMA transcriptome data identifies hundreds of novel functional genes and improves gene models in the Arabidopsis genome. <i>BMC Genomics</i> , 2007, 8, 401.	2.8	12
9	The peptide SCOOP12 acts on reactive oxygen species homeostasis to modulate cell division and elongation in Arabidopsis primary root. <i>Journal of Experimental Botany</i> , 2022, 73, 6115-6132.	4.8	12
10	The MIK2/SCOOP Signaling System Contributes to Arabidopsis Resistance Against Herbivory by Modulating Jasmonate and Indole Glucosinolate Biosynthesis. <i>Frontiers in Plant Science</i> , 2022, 13, 852808.	3.6	11