

Julian S Tonti-Filippini

List of Publications by Citations

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Version: 2024-04-20

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

12
papers

8,152
citations

12
h-index

14
g-index

14
ext. papers

9,218
ext. citations

17.2
avg, IF

5.16
L-index

#	Paper	IF	Citations
12	Human DNA methylomes at base resolution show widespread epigenomic differences. <i>Nature</i> , 2009 , 462, 315-22	50.4	3401
11	Highly integrated single-base resolution maps of the epigenome in Arabidopsis. <i>Cell</i> , 2008 , 133, 523-36	56.2	1896
10	Global epigenomic reconfiguration during mammalian brain development. <i>Science</i> , 2013 , 341, 1237905	33.3	1283
9	Experimental analysis of the Arabidopsis mitochondrial proteome highlights signaling and regulatory components, provides assessment of targeting prediction programs, and indicates plant-specific mitochondrial proteins. <i>Plant Cell</i> , 2004 , 16, 241-56	11.6	461
8	SUBA: the Arabidopsis Subcellular Database. <i>Nucleic Acids Research</i> , 2007 , 35, D213-8	20.1	364
7	A link between RNA metabolism and silencing affecting Arabidopsis development. <i>Developmental Cell</i> , 2008 , 14, 854-66	10.2	328
6	Combining experimental and predicted datasets for determination of the subcellular location of proteins in Arabidopsis. <i>Plant Physiology</i> , 2005 , 139, 598-609	6.6	116
5	Characterization of the preprotein and amino acid transporter gene family in Arabidopsis. <i>Plant Physiology</i> , 2007 , 143, 199-212	6.6	83
4	Large scale genome skimming from herbarium material for accurate plant identification and phylogenomics. <i>Plant Methods</i> , 2020 , 16, 1	5.8	61
3	The Expansion and Diversification of Pentatricopeptide Repeat RNA-Editing Factors in Plants. <i>Molecular Plant</i> , 2020 , 13, 215-230	14.4	28
2	Plastome-Wide Rearrangements and Gene Losses in Carnivorous Droseraceae. <i>Genome Biology and Evolution</i> , 2019 , 11, 472-485	3.9	23
1	STAR: an integrated solution to management and visualization of sequencing data. <i>Bioinformatics</i> , 2013 , 29, 3204-10	7.2	9