

Julian S Tonti-Filippini

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8839146/publications.pdf>

Version: 2024-02-01

13
papers

9,939
citations

686830

13
h-index

1125271

13
g-index

14
all docs

14
docs citations

14
times ranked

14745
citing authors

#	ARTICLE	IF	CITATIONS
1	Human DNA methylomes at base resolution show widespread epigenomic differences. <i>Nature</i> , 2009, 462, 315-322.	13.7	4,063
2	Highly Integrated Single-Base Resolution Maps of the Epigenome in Arabidopsis. <i>Cell</i> , 2008, 133, 523-536.	13.5	2,229
3	Global Epigenomic Reconfiguration During Mammalian Brain Development. <i>Science</i> , 2013, 341, 1237905.	6.0	1,609
4	Experimental Analysis of the Arabidopsis Mitochondrial Proteome Highlights Signaling and Regulatory Components, Provides Assessment of Targeting Prediction Programs, and Indicates Plant-Specific Mitochondrial Proteins [W]. <i>Plant Cell</i> , 2004, 16, 241-256.	3.1	550
5	SUBA: the Arabidopsis Subcellular Database. <i>Nucleic Acids Research</i> , 2007, 35, D213-D218.	6.5	394
6	A Link between RNA Metabolism and Silencing Affecting Arabidopsis Development. <i>Developmental Cell</i> , 2008, 14, 854-866.	3.1	394
7	Large scale genome skimming from herbarium material for accurate plant identification and phylogenomics. <i>Plant Methods</i> , 2020, 16, 1.	1.9	197
8	What can we do with 1000 plastid genomes?. <i>Plant Journal</i> , 2017, 90, 808-818.	2.8	165
9	Combining Experimental and Predicted Datasets for Determination of the Subcellular Location of Proteins in Arabidopsis. <i>Plant Physiology</i> , 2005, 139, 598-609.	2.3	120
10	Characterization of the Preprotein and Amino Acid Transporter Gene Family in Arabidopsis. <i>Plant Physiology</i> , 2007, 143, 199-212.	2.3	94
11	The Expansion and Diversification of Pentatricopeptide Repeat RNA-Editing Factors in Plants. <i>Molecular Plant</i> , 2020, 13, 215-230.	3.9	71
12	Plastome-Wide Rearrangements and Gene Losses in Carnivorous Droseraceae. <i>Genome Biology and Evolution</i> , 2019, 11, 472-485.	1.1	40
13	STAR: an integrated solution to management and visualization of sequencing data. <i>Bioinformatics</i> , 2013, 29, 3204-3210.	1.8	13