

Sara Ten Have

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2763822/publications.pdf>

Version: 2024-02-01

12
papers

611
citations

1040056

9
h-index

1199594

12
g-index

13
all docs

13
docs citations

13
times ranked

1162
citing authors

#	ARTICLE	IF	CITATIONS
1	A high quality Arabidopsis transcriptome for accurate transcript-level analysis of alternative splicing. <i>Nucleic Acids Research</i> , 2017, 45, 5061-5073.	14.5	262
2	Mass spectrometry-based immuno-precipitation proteomics – The user's guide. <i>Proteomics</i> , 2011, 11, 1153-1159.	2.2	76
3	tRNA-Derived RNA Fragments Associate with Human Multisynthetase Complex (MSC) and Modulate Ribosomal Protein Translation. <i>Journal of Proteome Research</i> , 2017, 16, 413-420.	3.7	72
4	Quantitative Profiling of the Human Substantia Nigra Proteome from Laser-capture Microdissected FFPE Tissue. <i>Molecular and Cellular Proteomics</i> , 2020, 19, 839-851.	3.8	39
5	Phosphorylation of Elp1 by Hrr25 Is Required for Elongator-Dependent tRNA Modification in Yeast. <i>PLoS Genetics</i> , 2015, 11, e1004931.	3.5	38
6	Quantitative analysis of the tomato nuclear proteome during <i>Phytophthora capsici</i> infection unveils regulators of immunity. <i>New Phytologist</i> , 2017, 215, 309-322.	7.3	29
7	Self-oligomerization regulates stability of survival motor neuron protein isoforms by sequestering an SCF ^{Slmb} degron. <i>Molecular Biology of the Cell</i> , 2018, 29, 96-110.	2.1	27
8	EMSY expression affects multiple components of the skin barrier with relevance to atopic dermatitis. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 144, 470-481.	2.9	23
9	Composition of the Survival Motor Neuron (SMN) Complex in <i>Drosophila melanogaster</i> . <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 491-503.	1.8	16
10	Functional and proteomic analysis of a full thickness filaggrin-deficient skin organoid model. <i>Wellcome Open Research</i> , 2019, 4, 134.	1.8	13
11	The Ndc80 complex targets Bod1 to human mitotic kinetochores. <i>Open Biology</i> , 2017, 7, 170099.	3.6	8
12	SILAC-based quantitative proteomic analysis of <i>Drosophila</i> gastrula stage embryos mutant for fibroblast growth factor signalling. <i>Fly</i> , 2020, 14, 10-28.	1.7	7