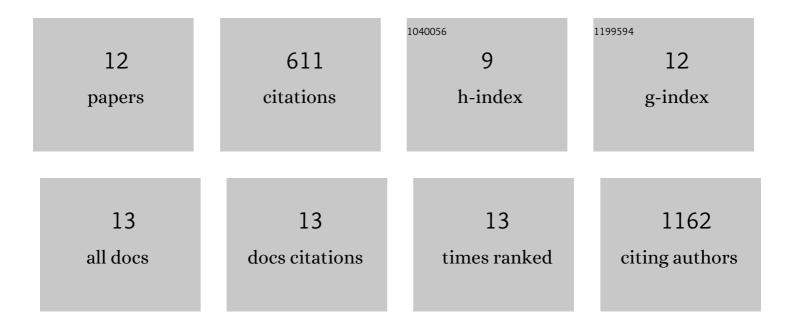
Sara Ten Have

List of Publications by Year in descending order

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SADA TEN HAVE

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