

Stefan Simm

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

985
citations

516710

16
h-index

552781

26
g-index

27
all docs

27
docs citations

27
times ranked

1504
citing authors

#	ARTICLE	IF	CITATIONS
1	Enhanced pro-apoptosis gene signature following the activation of TAp63 ^Δ in oocytes upon γ irradiation. <i>Cell Death and Disease</i> , 2022, 13, 204.	6.3	5
2	Cloning and Functional Characterization of Dog OCT1 and OCT2: Another Step in Exploring Species Differences in Organic Cation Transporters. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5100.	4.1	1
3	Macrophages Are Polarized toward an Inflammatory Phenotype by their Aged Microenvironment in the Human Skin. <i>Journal of Investigative Dermatology</i> , 2022, 142, 3136-3145.e11.	0.7	5
4	Insertion of plastidic β -barrel proteins into the outer envelopes of plastids involves an intermembrane space intermediate formed with Toc75-V/OEP80. <i>Plant Cell</i> , 2021, 33, 1657-1681.	6.6	15
5	Effect of thermospermine on expression profiling of different gene using massive analysis of cDNA ends (MACE) and vascular maintenance in Arabidopsis. <i>Physiology and Molecular Biology of Plants</i> , 2021, 27, 577-586.	3.1	3
6	miRNAs involved in transcriptome remodeling during pollen development and heat stress response in <i>Solanum lycopersicum</i> . <i>Scientific Reports</i> , 2020, 10, 10694.	3.3	22
7	The Existence and Localization of Nuclear snoRNAs in <i>Arabidopsis thaliana</i> Revisited. <i>Plants</i> , 2020, 9, 1016.	3.5	14
8	Transcriptional Basis for Differential Thermosensitivity of Seedlings of Various Tomato Genotypes. <i>Genes</i> , 2020, 11, 655.	2.4	5
9	Toc75 Δ /OEP80 is processed during translocation into chloroplasts, and the membrane-embedded form exposes its POTRA domain to the intermembrane space. <i>FEBS Open Bio</i> , 2020, 10, 444-454.	2.3	14
10	Functional diversification of tomato HsfA1 factors is based on DNA binding domain properties. <i>Gene</i> , 2019, 714, 143985.	2.2	20
11	HEATSTER: A Database and Web Server for Identification and Classification of Heat Stress Transcription Factors in Plants. <i>Bioinformatics and Biology Insights</i> , 2019, 13, 117793221882136.	2.0	26
12	The repressor and co-activator HsfB1 regulates the major heat stress transcription factors in tomato. <i>Plant, Cell and Environment</i> , 2019, 42, 874-890.	5.7	63
13	Regulation of two GTPases Toc159 and Toc34 in the translocon of the outer envelope of chloroplasts. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2019, 1867, 627-636.	2.3	14
14	Identification of the TXNIP IRES and characterization of the impact of regulatory IRES trans-acting factors. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2018, 1861, 147-157.	1.9	12
15	The coupling of transcriptome and proteome adaptation during development and heat stress response of tomato pollen. <i>BMC Genomics</i> , 2018, 19, 447.	2.8	68
16	Alternative splicing in tomato pollen in response to heat stress. <i>DNA Research</i> , 2017, 24, dsw051.	3.4	55
17	Reducing Cytoplasmic Polyamine Oxidase Activity in Arabidopsis Increases Salt and Drought Tolerance by Reducing Reactive Oxygen Species Production and Increasing Defense Gene Expression. <i>Frontiers in Plant Science</i> , 2016, 7, 214.	3.6	46
18	HsfA2 Controls the Activity of Developmentally and Stress-Regulated Heat Stress Protection Mechanisms in Tomato Male Reproductive Tissues. <i>Plant Physiology</i> , 2016, 170, 2461-2477.	4.8	148

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19	50Âyears of amino acid hydrophobicity scales: revisiting the capacity for peptide classification. <i>Biological Research</i> , 2016, 49, 31.	3.4	77
20	Survey of Genes Involved in Biosynthesis, Transport, and Signaling of Phytohormones with Focus on <i>Solanum lycopersicum</i> . <i>Bioinformatics and Biology Insights</i> , 2016, 10, BBI.S38425.	2.0	21
21	The membrane proteome of male gametophyte in <i>Solanum lycopersicum</i> . <i>Journal of Proteomics</i> , 2016, 131, 48-60.	2.4	25
22	Identification and Expression Analysis of Ribosome Biogenesis Factor Co-orthologs in <i>Solanum lycopersicum</i> . <i>Bioinformatics and Biology Insights</i> , 2015, 9, BBI.S20751.	2.0	62
23	Chaperone network composition in <i>Solanum lycopersicum</i> explored by transcriptome profiling and microarray meta-analysis. <i>Plant, Cell and Environment</i> , 2015, 38, 693-709.	5.7	71
24	The composition of the global and feature specific cyanobacterial core-genomes. <i>Frontiers in Microbiology</i> , 2015, 6, 219.	3.5	38
25	Defining the Core Proteome of the Chloroplast Envelope Membranes. <i>Frontiers in Plant Science</i> , 2013, 4, 11.	3.6	75
26	40S Ribosome Biogenesis Co-Factors Are Essential for Gametophyte and Embryo Development. <i>PLoS ONE</i> , 2013, 8, e54084.	2.5	74
27	Relevance and Regulation of Alternative Splicing in Plant Heat Stress Response: Current Understanding and Future Directions. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	6