

Andreas Bachmair

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

1,279
citations

567281

15
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552781

26
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all docs

28
docs citations

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times ranked

1772
citing authors

#	ARTICLE	IF	CITATIONS
1	Nitric Oxide Sensing in Plants Is Mediated by Proteolytic Control of Group VII ERF Transcription Factors. <i>Molecular Cell</i> , 2014, 53, 369-379.	9.7	312
2	The eukaryotic N-end rule pathway: conserved mechanisms and diverse functions. <i>Trends in Cell Biology</i> , 2014, 24, 603-611.	7.9	171
3	The Scope, Functions, and Dynamics of Posttranslational Protein Modifications. <i>Annual Review of Plant Biology</i> , 2019, 70, 119-151.	18.7	158
4	Substrates Related to Chromatin and to RNA-Dependent Processes Are Modified by Arabidopsis SUMO Isoforms That Differ in a Conserved Residue with Influence on Desumoylation. <i>Plant Physiology</i> , 2009, 149, 1529-1540.	4.8	91
5	Update on sumoylation: defining core components of the plant SUMO conjugation system by phylogenetic comparison. <i>New Phytologist</i> , 2012, 195, 23-31.	7.3	75
6	Arabidopsis PIAL1 and 2 Promote SUMO Chain Formation as E4-Type SUMO Ligases and Are Involved in Stress Responses and Sulfur Metabolism. <i>Plant Cell</i> , 2014, 26, 4547-4560.	6.6	73
7	Distinct branches of the N-end rule pathway modulate the plant immune response. <i>New Phytologist</i> , 2019, 221, 988-1000.	7.3	59
8	SUMOylation represses SnRK1 signaling in Arabidopsis. <i>Plant Journal</i> , 2016, 85, 120-133.	5.7	56
9	Distinct roles for Arabidopsis SUMO protease ESD4 and its closest homolog ELS1. <i>Planta</i> , 2011, 233, 63-73.	3.2	52
10	Interplay between phosphorylation and SUMOylation events determines CESTA protein fate in brassinosteroid signalling. <i>Nature Communications</i> , 2014, 5, 4687.	12.8	46
11	Extensive Analysis of GmFTL and GmCOL Expression in Northern Soybean Cultivars in Field Conditions. <i>PLoS ONE</i> , 2015, 10, e0136601.	2.5	27
12	Ubiquitin Lys 63 chains – second-most abundant, but poorly understood in plants. <i>Frontiers in Plant Science</i> , 2014, 5, 15.	3.6	26
13	Protein sumoylation and phosphorylation intersect in Arabidopsis signaling. <i>Plant Journal</i> , 2017, 91, 505-517.	5.7	25
14	Sumoylation and phosphorylation: hidden and overt links. <i>Journal of Experimental Botany</i> , 2018, 69, 4583-4590.	4.8	24
15	Revised nomenclature and functional overview of the ULP gene family of plant deSUMOylating proteases. <i>Journal of Experimental Botany</i> , 2018, 69, 4505-4509.	4.8	20
16	Small Ubiquitin-Like Modifier Conjugating Enzyme with Active Site Mutation Acts as Dominant Negative Inhibitor of SUMO Conjugation in Arabidopsis. <i>Journal of Integrative Plant Biology</i> , 2013, 55, 75-82.	8.5	16
17	SUMO chain formation relies on the amino-terminal region of SUMO-conjugating enzyme and has dedicated substrates in plants. <i>Biochemical Journal</i> , 2018, 475, 61-74.	3.7	10
18	Cellular Control of Protein Turnover via the Modification of the Amino Terminus. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3545.	4.1	8

#	ARTICLE	IF	CITATIONS
19	How cells coordinate waste removal through their major proteolytic pathways. <i>Nature Cell Biology</i> , 2015, 17, 841-842.	10.3	7
20	Virus-like particle formation and translational start site choice of the plant retrotransposon Tto1. <i>Virology</i> , 2008, 373, 437-446.	2.4	6
21	A synthetic biology approach allows inducible retrotransposition in whole plants. <i>Systems and Synthetic Biology</i> , 2010, 4, 133-138.	1.0	6
22	SUMO Chain Formation by Plant Enzymes. <i>Methods in Molecular Biology</i> , 2016, 1450, 97-105.	0.9	4
23	Seedling Germination: Seedlings Follow Sunshine and Fresh Air. <i>Current Biology</i> , 2015, 25, R565-R566.	3.9	2
24	A Yeast-Based Functional Assay to Study Plant N-Degron – N-Recognin Interactions. <i>Frontiers in Plant Science</i> , 2021, 12, 806129.	3.6	2
25	Deletion analysis of the 3' long terminal repeat sequence of plant retrotransposon Tto1 identifies 125 base pairs redundancy as sufficient for first strand transfer. <i>Virology</i> , 2011, 412, 75-82.	2.4	1
26	Transcriptome, metabolome and suppressor analysis reveal an essential role for the ubiquitin-proteasome system in seedling chloroplast development. <i>BMC Plant Biology</i> , 2022, 22, 183.	3.6	1
27	Expression and Purification of the Arabidopsis E4 SUMO Ligases PIAL1 and PIAL2. <i>Bio-protocol</i> , 2015, 5, .	0.4	0