

# Andreas Bachmair

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1307783/publications.pdf>

Version: 2024-02-01

27  
papers

1,279  
citations

567281

15  
h-index

552781

26  
g-index

28  
all docs

28  
docs citations

28  
times ranked

1772  
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	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
3	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
4	Distinct branches of the N"end rule pathway modulate the plant immune response. <i>New Phytologist</i> , 2019, 221, 988-1000.	7.3	59
5	The Scope, Functions, and Dynamics of Posttranslational Protein Modifications. <i>Annual Review of Plant Biology</i> , 2019, 70, 119-151.	18.7	158
6	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
7	Sumoylation and phosphorylation: hidden and overt links. <i>Journal of Experimental Botany</i> , 2018, 69, 4583-4590.	4.8	24
8	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
9	Protein sumoylation and phosphorylation intersect in Arabidopsis signaling. <i>Plant Journal</i> , 2017, 91, 505-517.	5.7	25
10	<sc>SUMO</sc>ylation represses Sn<sc>RK</sc>1 signaling in Arabidopsis. <i>Plant Journal</i> , 2016, 85, 120-133.	5.7	56
11	SUMO Chain Formation by Plant Enzymes. <i>Methods in Molecular Biology</i> , 2016, 1450, 97-105.	0.9	4
12	Seedling Germination: Seedlings Follow Sunshine and"Fresh Air. <i>Current Biology</i> , 2015, 25, R565-R566.	3.9	2
13	How cells coordinate waste removal through their major proteolytic pathways. <i>Nature Cell Biology</i> , 2015, 17, 841-842.	10.3	7
14	Extensive Analysis of GmFTL and GmCOL Expression in Northern Soybean Cultivars in Field Conditions. <i>PLoS ONE</i> , 2015, 10, e0136601.	2.5	27
15	Expression and Purification of the Arabidopsis E4 SUMO Ligases PIAL1 and PIAL2. <i>Bio-protocol</i> , 2015, 5, .	0.4	0
16	Ubiquitin Lys 63 chains " second-most abundant, but poorly understood in plants. <i>Frontiers in Plant Science</i> , 2014, 5, 15.	3.6	26
17	<i>Arabidopsis</i> 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
18	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

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19	Interplay between phosphorylation and SUMOylation events determines CESTA protein fate in brassinosteroid signalling. <i>Nature Communications</i> , 2014, 5, 4687.	12.8	46
20	The eukaryotic N-end rule pathway: conserved mechanisms and diverse functions. <i>Trends in Cell Biology</i> , 2014, 24, 603-611.	7.9	171
21	Small Ubiquitin-Like Modifier Conjugating Enzyme with Active Site Mutation Acts as Dominant Negative Inhibitor of SUMO Conjugation in <i>Arabidopsis</i> . <i>Journal of Integrative Plant Biology</i> , 2013, 55, 75-82.	8.5	16
22	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
23	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
24	Distinct roles for <i>Arabidopsis</i> SUMO protease ESD4 and its closest homolog ELS1. <i>Planta</i> , 2011, 233, 63-73.	3.2	52
25	A synthetic biology approach allows inducible retrotransposition in whole plants. <i>Systems and Synthetic Biology</i> , 2010, 4, 133-138.	1.0	6
26	Substrates Related to Chromatin and to RNA-Dependent Processes Are Modified by <i>Arabidopsis</i> SUMO Isoforms That Differ in a Conserved Residue with Influence on Desumoylation. <i>Plant Physiology</i> , 2009, 149, 1529-1540.	4.8	91
27	Virus-like particle formation and translational start site choice of the plant retrotransposon Tto1. <i>Virology</i> , 2008, 373, 437-446.	2.4	6