

Judy Callis

List of Publications by Year in descending order

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47
papers

5,290
citations

159585

30
h-index

223800

46
g-index

151
all docs

151
docs citations

151
times ranked

5506
citing authors

#	ARTICLE	IF	CITATIONS
1	Functional Analysis of the RING-Type Ubiquitin Ligase Family of Arabidopsis. <i>Plant Physiology</i> , 2005, 137, 13-30.	4.8	524
2	Interactions of the COP9 Signalosome with the E3 Ubiquitin Ligase SCFTIR1 in Mediating Auxin Response. <i>Science</i> , 2001, 292, 1379-1382.	12.6	451
3	Ubiquitin, Hormones and Biotic Stress in Plants. <i>Annals of Botany</i> , 2007, 99, 787-822.	2.9	432
4	Genome Analysis and Functional Characterization of the E2 and RING-Type E3 Ligase Ubiquitination Enzymes of Arabidopsis. <i>Plant Physiology</i> , 2005, 139, 1597-1611.	4.8	365
5	KEEP ON GOING, a RING E3 Ligase Essential for Arabidopsis Growth and Development, Is Involved in Abscisic Acid Signaling. <i>Plant Cell</i> , 2007, 18, 3415-3428.	6.6	347
6	The Arabidopsis Aux/IAA Protein Family Has Diversified in Degradation and Auxin Responsiveness. <i>Plant Cell</i> , 2006, 18, 699-714.	6.6	265
7	Rapid Degradation of Auxin/Indoleacetic Acid Proteins Requires Conserved Amino Acids of Domain II and Is Proteasome Dependent. <i>Plant Cell</i> , 2001, 13, 2349-2360.	6.6	260
8	The Ubiquitination Machinery of the Ubiquitin System. <i>The Arabidopsis Book</i> , 2014, 12, e0174.	0.5	260
9	Degradation of Aux/IAA proteins is essential for normal auxin signalling. <i>Plant Journal</i> , 2000, 21, 553-562.	5.7	254
10	The intron of Arabidopsis thaliana polyubiquitin genes is conserved in location and is a quantitative determinant of chimeric gene expression. <i>Plant Molecular Biology</i> , 1993, 21, 895-906.	3.9	226
11	Protein degradation in signaling. <i>Current Opinion in Plant Biology</i> , 2000, 3, 381-386.	7.1	183
12	Regulation of Cullin RING Ligases. <i>Annual Review of Plant Biology</i> , 2008, 59, 467-489.	18.7	175
13	Arabidopsis Has Two Redundant Cullin3 Proteins That Are Essential for Embryo Development and That Interact with RBX1 and BTB Proteins to Form Multisubunit E3 Ubiquitin Ligase Complexes in Vivo. <i>Plant Cell</i> , 2005, 17, 1180-1195.	6.6	153
14	Ubiquitin ligases mediate growth and development by promoting protein death. <i>Current Opinion in Plant Biology</i> , 2007, 10, 624-632.	7.1	150
15	Independent modulation of Arabidopsis thaliana polyubiquitin mRNAs in different organs and in response to environmental changes. <i>Plant Journal</i> , 1997, 11, 1017-1027.	5.7	120
16	<sc>ABA</sc> and the ubiquitin E3 ligase <sc>KEEP ON GOING</sc> affect proteolysis of the <i><sc>A</sc>rabadopsis thaliana</i> transcription factors <sc>ABF</sc>1 and <sc>ABF</sc>3. <i>Plant Journal</i> , 2013, 75, 965-976.	5.7	114
17	Related to Ubiquitin 1 and 2 Are Redundant and Essential and Regulate Vegetative Growth, Auxin Signaling, and Ethylene Production in Arabidopsis. <i>Plant Cell</i> , 2004, 16, 2418-2432.	6.6	79
18	The Rub Family of Ubiquitin-like Proteins. <i>Journal of Biological Chemistry</i> , 1998, 273, 34976-34982.	3.4	78

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19	The RING E3 Ligase KEEP ON GOING Modulates JASMONATE ZIM-DOMAIN12 Stability. <i>Plant Physiology</i> , 2015, 169, 1405-1417.	4.8	76
20	The plastid-localized pfkB-type carbohydrate kinases FRUCTOKINASE-LIKE 1 and 2 are essential for growth and development of <i>Arabidopsis thaliana</i> . <i>BMC Plant Biology</i> , 2012, 12, 102.	3.6	70
21	Ubiquitin on the Move: The Ubiquitin Modification System Plays Diverse Roles in the Regulation of Endoplasmic Reticulum- and Plasma Membrane-Localized Proteins. <i>Plant Physiology</i> , 2012, 160, 56-64.	4.8	58
22	Polypeptide tags, ubiquitous modifiers for plant protein regulation. , 1999, 41, 435-442.		55
23	Acceleration of Aux/IAA proteolysis is specific for auxin and independent of AXR1. <i>Plant Journal</i> , 2003, 35, 285-294.	5.7	53
24	Degradation of the auxin response factor ARF1. <i>Plant Journal</i> , 2008, 54, 118-128.	5.7	48
25	Engineering in vivo instability of firefly luciferase and <i>Escherichia coli</i> beta-glucuronidase in higher plants using recognition elements from the ubiquitin pathway. <i>Plant Molecular Biology</i> , 1998, 37, 337-347.	3.9	45
26	Isolation and Characterization of <i>cul1-7</i> , a Recessive Allele of <i>CULLIN1</i> That Disrupts SCF Function at the C Terminus of CUL1 in <i>Arabidopsis thaliana</i> . <i>Genetics</i> , 2009, 181, 945-963.	2.9	41
27	Identification and biochemical characterization of the fructokinase gene family in <i>Arabidopsis thaliana</i> . <i>BMC Plant Biology</i> , 2017, 17, 83.	3.6	40
28	The Ubiquitin E3 Ligase LOSS OF GDU2 Is Required for GLUTAMINE DUMPER1-Induced Amino Acid Secretion in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2012, 158, 1628-1642.	4.8	39
29	Lysine Residues Are Not Required for Proteasome-Mediated Proteolysis of the Auxin/Indole Acidic Acid Protein IAA1. <i>Plant Physiology</i> , 2015, 168, 708-720.	4.8	39
30	A role for phospholipase A in auxin-regulated gene expression. <i>FEBS Letters</i> , 2007, 581, 4205-4211.	2.8	36
31	Histidine-Tagged Ubiquitin Substitutes for Wild-Type Ubiquitin in <i>Saccharomyces cerevisiae</i> and Facilitates Isolation and Identification of in Vivo Substrates of the Ubiquitin Pathway. <i>Analytical Biochemistry</i> , 2000, 282, 54-64.	2.4	34
32	The ubiquitin system affects agronomic plant traits. <i>Journal of Biological Chemistry</i> , 2020, 295, 13940-13955.	3.4	32
33	High Performance Liquid Chromatography Resolution of Ubiquitin Pathway Enzymes from Wheat Germ. <i>Plant Physiology</i> , 1990, 94, 710-716.	4.8	28
34	Selective auxin agonists induce specific AUX/IAA protein degradation to modulate plant development. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 6463-6472.	7.1	23
35	BRIZ1 and BRIZ2 Proteins Form a Heteromeric E3 Ligase Complex Required for Seed Germination and Post-germination Growth in <i>Arabidopsis thaliana</i> . <i>Journal of Biological Chemistry</i> , 2010, 285, 37070-37081.	3.4	20
36	Identification of the Plant Ribokinase and Discovery of a Role for <i>Arabidopsis</i> Ribokinase in Nucleoside Metabolism. <i>Journal of Biological Chemistry</i> , 2016, 291, 22572-22582.	3.4	20

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37	A model for the evolution of polyubiquitin genes from the study of <i>Arabidopsis thaliana</i> ecotypes. <i>Plant Molecular Biology</i> , 1997, 34, 745-758.	3.9	16
38	AXR1-ECR1 and AXL1-ECR1 heterodimeric RUB-activating enzymes diverge in function in <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 2011, 75, 515-526.	3.9	16
39	Functional conservation between mammalian MGRN1 and plant LOG2 ubiquitin ligases. <i>FEBS Letters</i> , 2013, 587, 3400-3405.	2.8	15
40	Broadening the impact of plant science through innovative, integrative, and inclusive outreach. <i>Plant Direct</i> , 2021, 5, e00316.	1.9	14
41	Preparation, Characterization, and Use of Tagged Ubiquitins. <i>Methods in Enzymology</i> , 2005, 399, 51-64.	1.0	9
42	Control of Amino Acid Homeostasis by a Ubiquitin Ligase-Coactivator Protein Complex. <i>Journal of Biological Chemistry</i> , 2017, 292, 3827-3840.	3.4	7
43	<i>Arabidopsis</i> fructokinase-like protein associations are regulated by ATP. <i>Biochemical Journal</i> , 2017, 474, 1789-1801.	3.7	7
44	A genetic screen for mutants defective in IAA1-LUC degradation in <i>Arabidopsis thaliana</i> reveals an important requirement for TOPOISOMERASE6 in auxin physiology. <i>Plant Signaling and Behavior</i> , 2014, 9, e972207.	2.4	4
45	Recovery of DDB1a (DAMAGED DNA BINDING PROTEIN1a) in a Screen to Identify Novel RUB-Modified Proteins in <i>Arabidopsis thaliana</i> . <i>Molecular Plant</i> , 2012, 5, 1163-1166.	8.3	3
46	The <i>Arabidopsis thaliana</i> E3 Ubiquitin Ligase BRIZ Functions in Abscisic Acid Response. <i>Frontiers in Plant Science</i> , 2021, 12, 641849.	3.6	3
47	Factors that affect protein abundance of a positive regulator of abscisic acid signalling, the basic leucine zipper transcription factor ABRE-binding factor 2 (ABF2). <i>Plant Direct</i> , 2021, 5, e00330.	1.9	2