

# Heribert Hirt

## List of Publications by Year in descending order

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233  
papers

36,006  
citations

4942

84  
h-index

3394

183  
g-index

240  
all docs

240  
docs citations

240  
times ranked

27686  
citing authors

#	ARTICLE	IF	CITATIONS
1	In vivo identification of putative CPK5 substrates in <i>Arabidopsis thaliana</i> . <i>Plant Science</i> , 2022, 314, 111121.	1.7	6
2	Beat the heat: plant- and microbe-mediated strategies for crop thermotolerance. <i>Trends in Plant Science</i> , 2022, 27, 802-813.	4.3	43
3	ROS homeostasis mediated by MPK4 and SUMM2 determines synergid cell death. <i>Nature Communications</i> , 2022, 13, 1746.	5.8	18
4	The Lysâ€œmotif receptor <sc><i>LYK4</i></sc> mediates <i>Enterobacter</i> sp. <sc>SA187</sc> triggered salt tolerance in <i>Arabidopsis thaliana</i>. <i>Environmental Microbiology</i> , 2022, 24, 223-239.	1.8	4
5	Analysis of the <i>Arabidopsis&lt;i&gt; coilin&lt;/i&gt;</i> mutant reveals a positive role of AtCOILIN in plant immunity. <i>Plant Physiology</i> , 2022, 190, 745-761.	2.3	6
6	Root endophyte induced plant thermotolerance by constitutive chromatin modification at heat stress memory gene loci. <i>EMBO Reports</i> , 2021, 22, e51049.	2.0	71
7	Complete Genome Sequence of <i>Cellulomonas</i> sp. JZ18, a Root Endophytic Bacterium Isolated from the Perennial Desert Tussock-Grass <i>Panicum turgidum</i> . <i>Current Microbiology</i> , 2021, 78, 1135-1141.	1.0	4
8	The Seed Development Factors TT2 and MYB5 Regulate Heat Stress Response in <i>Arabidopsis</i> . <i>Genes</i> , 2021, 12, 746.	1.0	13
9	G3BPs in Plant Stress. <i>Frontiers in Plant Science</i> , 2021, 12, 680710.	1.7	6
10	Stomatal regulation: Role of H <sub>2</sub> S-induced persulfidation in ABA signaling. <i>Molecular Plant</i> , 2021, 14, 858-860.	3.9	8
11	Polycomb-dependent differential chromatin compartmentalization determines gene coregulation in <i>Arabidopsis</i>. <i>Genome Research</i> , 2021, 31, 1230-1244.	2.4	36
12	Multiple strategies of plant colonization by beneficial endophytic <sc><i>Enterobacter</i> sp. <sc>SA187</sc>. <i>Environmental Microbiology</i> , 2021, 23, 6223-6240.	1.8	10
13	Chromatin phosphoproteomics unravels a function for AT-hook motif nuclear localized protein AHL13 in PAMP-triggered immunity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	25
14	Development, validation, and application of an HPLC-MS/MS method for quantification of oxidized fatty acids in plants. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2021, 1186, 123006.	1.2	2
15	Coordinated bacterial and plant sulfur metabolism in <i>Enterobacter</i> sp. SA187â€œinduced plant salt stress tolerance. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	40
16	CATION-CHLORIDE CO-TRANSPORTER 1 (CCC1) Mediates Plant Resistance against <i>Pseudomonas syringae</i>. <i>Plant Physiology</i> , 2020, 182, 1052-1065.	2.3	7
17	Desert Microbes for Boosting Sustainable Agriculture in Extreme Environments. <i>Frontiers in Microbiology</i> , 2020, 11, 1666.	1.5	87
18	GCN5 modulates salicylic acid homeostasis by regulating H3K14ac levels at the 5â€œ and 3â€œ ends of its target genes. <i>Nucleic Acids Research</i> , 2020, 48, 5953-5966.	6.5	44

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19	Tailoring plant-associated microbial inoculants in agriculture: a roadmap for successful application. <i>Journal of Experimental Botany</i> , 2020, 71, 3878-3901.	2.4	118
20	Complete genome sequence of the endophytic bacterium <i>Cellulosimicrobium</i> sp. JZ28 isolated from the root endosphere of the perennial desert tussock grass <i>Panicum turgidum</i> . <i>Archives of Microbiology</i> , 2020, 202, 1563-1569.	1.0	8
21	Complete Genome Sequence of <i>Paenibacillus</i> sp. JZ16, a Plant Growth Promoting Root Endophytic Bacterium of the Desert Halophyte <i>Zygophyllum Simplex</i> . <i>Current Microbiology</i> , 2020, 77, 1097-1103.	1.0	15
22	Wheat chromatin architecture is organized in genome territories and transcription factories. <i>Genome Biology</i> , 2020, 21, 104.	3.8	99
23	Nanofabrication of Isoporous Membranes for Cell Fractionation. <i>Scientific Reports</i> , 2020, 10, 6138.	1.6	22
24	Wounding and Insect Feeding Trigger Two Independent MAPK Pathways with Distinct Regulation and Kinetics. <i>Plant Cell</i> , 2020, 32, 1988-2003.	3.1	61
25	Role of MPK4 in pathogen-associated molecular pattern-triggered alternative splicing in <i>Arabidopsis</i> . <i>PLoS Pathogens</i> , 2020, 16, e1008401.	2.1	38
26	Genome Insights of the Plant-Growth Promoting Bacterium <i>Cronobacter muytjensii</i> JZ38 With Volatile-Mediated Antagonistic Activity Against <i>Phytophthora infestans</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 369.	1.5	39
27	Healthy soils for healthy plants for healthy humans. <i>EMBO Reports</i> , 2020, 21, e51069.	2.0	60
28	Role of MPK4 in pathogen-associated molecular pattern-triggered alternative splicing in <i>Arabidopsis</i> . , 2020, 16, e1008401.		0
29	Role of MPK4 in pathogen-associated molecular pattern-triggered alternative splicing in <i>Arabidopsis</i> . , 2020, 16, e1008401.		0
30	Role of MPK4 in pathogen-associated molecular pattern-triggered alternative splicing in <i>Arabidopsis</i> . , 2020, 16, e1008401.		0
31	Role of MPK4 in pathogen-associated molecular pattern-triggered alternative splicing in <i>Arabidopsis</i> . , 2020, 16, e1008401.		0
32	Role of MPK4 in pathogen-associated molecular pattern-triggered alternative splicing in <i>Arabidopsis</i> . , 2020, 16, e1008401.		0
33	Phosphorylation regulates the activity of INDETERMINATE-DOMAIN (IDD/BIRD) proteins in response to diverse environmental conditions. <i>Plant Signaling and Behavior</i> , 2019, 14, e1642037.	1.2	7
34	<i>Piriformospora indica</i> alters Na <sup>+</sup> /K <sup>+</sup> homeostasis, antioxidant enzymes and LeNHX1 expression of greenhouse tomato grown under salt stress. <i>Scientia Horticulturae</i> , 2019, 256, 108532.	1.7	97
35	The Polycomb protein <i>LHP1</i> regulates <i>Arabidopsis thaliana</i> stress responses through the repression of the MYC-dependent branch of immunity. <i>Plant Journal</i> , 2019, 100, 1118-1131.	2.8	52
36	Mining biosynthetic gene clusters in <i>Virgibacillus</i> genomes. <i>BMC Genomics</i> , 2019, 20, 696.	1.2	7

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37	INDETERMINATE-DOMAIN 4 (IDD4) coordinates immune responses with plant-growth in <i>Arabidopsis thaliana</i> . <i>PLoS Pathogens</i> , 2019, 15, e1007499.	2.1	17
38	OXI1 and DAD Regulate Light-Induced Cell Death Antagonistically through Jasmonate and Salicylate Levels. <i>Plant Physiology</i> , 2019, 180, 1691-1708.	2.3	30
39	A Chimeric IDD4 Repressor Constitutively Induces Immunity in <i>Arabidopsis</i> via the Modulation of Salicylic Acid and Jasmonic Acid Homeostasis. <i>Plant and Cell Physiology</i> , 2019, 60, 1536-1555.	1.5	17
40	MAP 4K4 associates with BIK 1 to regulate plant innate immunity. <i>EMBO Reports</i> , 2019, 20, e47965.	2.0	22
41	Comparative genomics study reveals Red Sea <i>Bacillus</i> with characteristics associated with potential microbial cell factories (MCFs). <i>Scientific Reports</i> , 2019, 9, 19254.	1.6	6
42	Bioprospecting desert plant <i>Bacillus</i> endophytic strains for their potential to enhance plant stress tolerance. <i>Scientific Reports</i> , 2019, 9, 18154.	1.6	69
43	Phylogenetically diverse endophytic bacteria from desert plants induce transcriptional changes of tissue-specific ion transporters and salinity stress in <i>Arabidopsis thaliana</i> . <i>Plant Science</i> , 2019, 280, 228-240.	1.7	33
44	The Lamin-Like LITTLE NUCLEI 1 (LINC1) Regulates Pattern-Triggered Immunity and Jasmonic Acid Signaling. <i>Frontiers in Plant Science</i> , 2019, 10, 1639.	1.7	26
45	Plant Immunity: The MTI-ETI Model and Beyond. <i>Current Issues in Molecular Biology</i> , 2019, 30, 39-58.	1.0	31
46	Metaorganisms in extreme environments: do microbes play a role in organismal adaptation?. <i>Zoology</i> , 2018, 127, 1-19.	0.6	194
47	Quantitative Phosphoproteomic Analysis Reveals Shared and Specific Targets of <i>Arabidopsis</i> Mitogen-Activated Protein Kinases (MAPKs) MPK3, MPK4, and MPK6. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 61-80.	2.5	80
48	Desert plant bacteria reveal host influence and beneficial plant growth properties. <i>PLoS ONE</i> , 2018, 13, e0208223.	1.1	76
49	The Trihelix transcription factor GT2-like 1 (GTL1) promotes salicylic acid metabolism, and regulates bacterial-triggered immunity. <i>PLoS Genetics</i> , 2018, 14, e1007708.	1.5	41
50	Plant Immunity: From Signaling to Epigenetic Control of Defense. <i>Trends in Plant Science</i> , 2018, 23, 833-844.	4.3	198
51	Modify the Histone to Win the Battle: Chromatin Dynamics in Plant-Pathogen Interactions. <i>Frontiers in Plant Science</i> , 2018, 9, 355.	1.7	106
52	Nuclear Signaling of Plant MAPKs. <i>Frontiers in Plant Science</i> , 2018, 9, 469.	1.7	168
53	Boosting Alfalfa ( <i>Medicago sativa</i> L.) Production With Rhizobacteria From Various Plants in Saudi Arabia. <i>Frontiers in Microbiology</i> , 2018, 9, 477.	1.5	35
54	In silico exploration of Red Sea <i>Bacillus</i> genomes for natural product biosynthetic gene clusters. <i>BMC Genomics</i> , 2018, 19, 382.	1.2	17

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55	Ethylene induced plant stress tolerance by <i>Enterobacter</i> sp. SA187 is mediated by 2-oxo-4-methylthiobutyric acid production. <i>PLoS Genetics</i> , 2018, 14, e1007273.	1.5	95
56	Quantification of Root Colonizing Bacteria. <i>Bio-protocol</i> , 2018, 8, .	0.2	3
57	The <i>Arabidopsis</i> homolog of human G3BP1 is a key regulator of stomatal and apoplastic immunity. <i>Life Science Alliance</i> , 2018, 1, e201800046.	1.3	16
58	Draft Genome Sequence of the Plant Growth-Promoting Rhizobacterium <i>Acinetobacter radioresistens</i> Strain SA188 Isolated from the Desert Plant <i>Indigofera argentea</i> . <i>Genome Announcements</i> , 2017, 5, .	0.8	5
59	Draft Genome Sequence of the Plant Growth-Promoting <i>Pseudomonas punonensis</i> Strain D1-6 Isolated from the Desert Plant <i>Erodium hirtum</i> in Jordan. <i>Genome Announcements</i> , 2017, 5, .	0.8	9
60	Constitutively Active <i>Arabidopsis</i> MAP Kinase 3 Triggers Defense Responses Involving Salicylic Acid and SUMM2 Resistance Protein. <i>Plant Physiology</i> , 2017, 174, 1238-1249.	2.3	57
61	A high quality <i>Arabidopsis</i> transcriptome for accurate transcript-level analysis of alternative splicing. <i>Nucleic Acids Research</i> , 2017, 45, 5061-5073.	6.5	262
62	Draft Genome Sequence of Plant Growth-Promoting <i>Micrococcus luteus</i> Strain K39 Isolated from <i>Cyperus conglomeratus</i> in Saudi Arabia. <i>Genome Announcements</i> , 2017, 5, .	0.8	11
63	Review: Mitogen-Activated Protein Kinases in nutritional signaling in <i>Arabidopsis</i> . <i>Plant Science</i> , 2017, 260, 101-108.	1.7	70
64	Plant-Specific Histone Deacetylases HDT1/2 Regulate <i>GIBBERELLIN 2-OXIDASE2</i> Expression to Control <i>Arabidopsis</i> Root Meristem Cell Number. <i>Plant Cell</i> , 2017, 29, 2183-2196.	3.1	69
65	Constitutive activity of the <i>Arabidopsis</i> MAP Kinase 3 confers resistance to <i>Pseudomonas syringae</i> and drives robust immune responses. <i>Plant Signaling and Behavior</i> , 2017, 12, e1356533.	1.2	14
66	Challenges Faced in Field Application of Phosphate-Solubilizing Bacteria. , 2017, , 125-143.		12
67	The <i>Arabidopsis</i> SWI/SNF protein BAF60 mediates seedling growth control by modulating DNA accessibility. <i>Genome Biology</i> , 2017, 18, 114.	3.8	53
68	The heat-shock protein/chaperone network and multiple stress resistance. <i>Plant Biotechnology Journal</i> , 2017, 15, 405-414.	4.1	513
69	Draft Genome Sequence of <i>Ochrobactrum intermedium</i> Strain SA148, a Plant Growth-Promoting Desert Rhizobacterium. <i>Genome Announcements</i> , 2017, 5, .	0.8	5
70	Draft Genome Sequence of <i>Enterobacter</i> sp. Sa187, an Endophytic Bacterium Isolated from the Desert Plant <i>Indigofera argentea</i> . <i>Genome Announcements</i> , 2017, 5, .	0.8	5
71	Complete Genome Sequence Analysis of <i>Enterobacter</i> sp. SA187, a Plant Multi-Stress Tolerance Promoting Endophytic Bacterium. <i>Frontiers in Microbiology</i> , 2017, 8, 2023.	1.5	83
72	MAPK-triggered chromatin reprogramming by histone deacetylase in plant innate immunity. <i>Genome Biology</i> , 2017, 18, 131.	3.8	73

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73	Draft Genome Sequence of <i>Halomonas elongata</i> Strain K4, an Endophytic Growth-Promoting Bacterium Enhancing Salinity Tolerance <i>In Planta</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	13
74	Plant Growth Promoting Rhizobacteria and Silicon Synergistically Enhance Salinity Tolerance of Mung Bean. <i>Frontiers in Plant Science</i> , 2016, 7, 876.	1.7	178
75	Convergence of Multiple MAP3Ks on MKK3 Identifies a Set of Novel Stress MAPK Modules. <i>Frontiers in Plant Science</i> , 2016, 07, 1941.	1.7	35
76	Draft Genome Sequence of the Phosphate-Solubilizing Bacterium <i>Pseudomonas argentinensis</i> Strain SA190 Isolated from the Desert Plant <i>Indigofera argentea</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	9
77	Draft Genome Sequence of the Plant Growth-Promoting <i>Cupriavidus gilardii</i> Strain JZ4 Isolated from the Desert Plant <i>Tribulus terrestris</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	17
78	The Role of MAPK Modules and ABA during Abiotic Stress Signaling. <i>Trends in Plant Science</i> , 2016, 21, 677-685.	4.3	326
79	Aquaporins Link ROS Signaling to Plant Immunity. <i>Plant Physiology</i> , 2016, 171, 1540-1540.	2.3	15
80	Interview with Heribert Hirt. <i>Trends in Plant Science</i> , 2016, 21, 1-2.	4.3	5
81	LHP1 Regulates H3K27me3 Spreading and Shapes the Three-Dimensional Conformation of the Arabidopsis Genome. <i>PLoS ONE</i> , 2016, 11, e0158936.	1.1	97
82	Bacterial Rhizosphere Biodiversity from Several Pioneer Desert Sand Plants Near Jizan, Saudi Arabia. <i>The Open Conference Proceedings Journal</i> , 2016, 7, 70-79.	0.6	4
83	A SWI/SNF Chromatin Remodelling Protein Controls Cytokinin Production through the Regulation of Chromatin Architecture. <i>PLoS ONE</i> , 2015, 10, e0138276.	1.1	25
84	Plant-Microbe Interactions and Water Management in Arid and Saline Soils. , 2015, , 265-276.		20
85	Signaling Mechanisms in Pattern-Triggered Immunity (PTI). <i>Molecular Plant</i> , 2015, 8, 521-539.	3.9	750
86	Identification and characterization of an ABA-activated MAP kinase cascade in <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 2015, 82, 232-244.	2.8	187
87	Plant MAPK cascades: Just rapid signaling modules?. <i>Plant Signaling and Behavior</i> , 2015, 10, e1062197.	1.2	23
88	<i>Salmonella enterica</i> induces and subverts the plant immune system. <i>Frontiers in Microbiology</i> , 2014, 5, 141.	1.5	31
89	<i>Salmonella enterica</i> Flagellin Is Recognized via FLS2 and Activates PAMP-Triggered Immunity in <i>Arabidopsis thaliana</i> . <i>Molecular Plant</i> , 2014, 7, 657-674.	3.9	75
90	The <i>Salmonella</i> effector protein SpvC, a phosphothreonine lyase is functional in plant cells. <i>Frontiers in Microbiology</i> , 2014, 5, 548.	1.5	27

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91	Proteomic and phosphoproteomic analyses of chromatin-associated proteins from <i>Arabidopsis thaliana</i> . <i>Proteomics</i> , 2014, 14, 2141-2155.	1.3	18
92	Salt-induced subcellular kinase relocation and seedling susceptibility caused by overexpression of Medicago SIMKK in <i>Arabidopsis</i> . <i>Journal of Experimental Botany</i> , 2014, 65, 2335-2350.	2.4	37
93	The role of ABA and MAPK signaling pathways in plant abiotic stress responses. <i>Biotechnology Advances</i> , 2014, 32, 40-52.	6.0	528
94	The BAF60 Subunit of the SWI/SNF Chromatin-Remodeling Complex Directly Controls the Formation of a Gene Loop at <i>FLOWERING LOCUS C</i> in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 26, 538-551.	3.1	82
95	Phosphorylation-dependent regulation of plant chromatin and chromatin-associated proteins. <i>Proteomics</i> , 2014, 14, 2127-2140.	1.3	26
96	Identification of Novel PAMP-Triggered Phosphorylation and Dephosphorylation Events in <i>Arabidopsis thaliana</i> by Quantitative Phosphoproteomic Analysis. <i>Journal of Proteome Research</i> , 2014, 13, 2137-2151.	1.8	44
97	Auxin efflux by PIN-FORMED proteins is activated by two different protein kinases, D6 PROTEIN KINASE and PINOID. <i>ELife</i> , 2014, 3, .	2.8	205
98	Functional analysis of <i>Arabidopsis</i> immune-related MAPKs uncovers a role for MPK3 as negative regulator of inducible defences. <i>Genome Biology</i> , 2014, 15, R87.	13.9	137
99	Protein Complexes Characterization in <i>Arabidopsis thaliana</i> by Tandem Affinity Purification Coupled to Mass Spectrometry Analysis. <i>Methods in Molecular Biology</i> , 2014, 1171, 237-250.	0.4	3
100	The role of the kinase <i>OXI1</i> in cadmium- and copper-induced molecular responses in <i>Arabidopsis thaliana</i> . <i>Plant, Cell and Environment</i> , 2013, 36, 1228-1238.	2.8	50
101	Rhizosphere Microbes as Essential Partners for Plant Stress Tolerance. <i>Molecular Plant</i> , 2013, 6, 242-245.	3.9	220
102	Improvement of stress tolerance in plants by genetic manipulation of mitogen-activated protein kinases. <i>Biotechnology Advances</i> , 2013, 31, 118-128.	6.0	124
103	New checkpoints in stomatal defense. <i>Trends in Plant Science</i> , 2013, 18, 295-297.	4.3	52
104	Constitutively active MPK4 helps to clarify its role in plant immunity. <i>Plant Signaling and Behavior</i> , 2013, 8, e22991.	1.2	7
105	An Abscisic Acid-Independent Oxylipin Pathway Controls Stomatal Closure and Immune Defense in <i>Arabidopsis</i> . <i>PLoS Biology</i> , 2013, 11, e1001513.	2.6	239
106	Brassinosteroid-regulated GSK3/Shaggy-like Kinases Phosphorylate Mitogen-activated Protein (MAP) Kinase Kinases, Which Control Stomata Development in <i>Arabidopsis thaliana</i> . <i>Journal of Biological Chemistry</i> , 2013, 288, 7519-7527.	1.6	152
107	Dual function of MIPS1 as a metabolic enzyme and transcriptional regulator. <i>Nucleic Acids Research</i> , 2013, 41, 2907-2917.	6.5	35
108	Regulation of the heat stress response in <i>Arabidopsis</i> by MPK6-targeted phosphorylation of the heat stress factor HsfA2. <i>PeerJ</i> , 2013, 1, e59.	0.9	106

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109	Constitutively Active Mitogen-Activated Protein Kinase Versions Reveal Functions of <i>Arabidopsis</i> MPK4 in Pathogen Defense Signaling. <i>Plant Cell</i> , 2012, 24, 4281-4293.	3.1	163
110	Automated Phosphopeptide Identification Using Multiple MS/MS Fragmentation Modes. <i>Journal of Proteome Research</i> , 2012, 11, 5695-5703.	1.8	16
111	Plants as alternative hosts for Salmonella. <i>Trends in Plant Science</i> , 2012, 17, 245-249.	4.3	92
112	Role of AGC kinases in plant growth and stress responses. <i>Cellular and Molecular Life Sciences</i> , 2012, 69, 3259-3267.	2.4	45
113	The <i>Arabidopsis</i> protein kinase Pto-interacting 1 is a common target of the oxidative signal-inducible 1 and mitogen-activated protein kinases. <i>FEBS Journal</i> , 2011, 278, 1126-1136.	2.2	50
114	Isolation and characterization of plant protein complexes by mass spectrometry. <i>Proteomics</i> , 2011, 11, 1824-1833.	1.3	23
115	Linking the proteins' Elucidation of proteome-scale networks using mass spectrometry. <i>Mass Spectrometry Reviews</i> , 2011, 30, 268-297.	2.8	23
116	Conservation of Salmonella Infection Mechanisms in Plants and Animals. <i>PLoS ONE</i> , 2011, 6, e24112.	1.1	114
117	The OX11 Kinase Pathway Mediates Piriformospora indica-Induced Growth Promotion in Arabidopsis. <i>PLoS Pathogens</i> , 2011, 7, e1002051.	2.1	126
118	AGC kinases in plant development and defense. <i>Plant Signaling and Behavior</i> , 2011, 6, 1030-1033.	1.2	24
119	New insights into an old story: Agrobacterium-induced tumour formation in plants by plant transformation. <i>EMBO Journal</i> , 2010, 29, 1021-1032.	3.5	216
120	Bioinformatic and Systems Biology Tools to Generate Testable Models of Signaling Pathways and Their Targets. <i>Plant Physiology</i> , 2010, 152, 460-469.	2.3	17
121	The MAP Kinase MPK4 Is Required for Cytokinesis in <i>Arabidopsis thaliana</i> . <i>Plant Cell</i> , 2010, 22, 3778-3790.	3.1	185
122	Mechanism of MAPK-targeted gene expression unraveled in plants. <i>Cell Cycle</i> , 2010, 9, 18-19.	1.3	6
123	Transgenerational Stress Memory Is Not a General Response in Arabidopsis. <i>PLoS ONE</i> , 2009, 4, e5202.	1.1	142
124	VIP1 response elements mediate mitogen-activated protein kinase 3-induced stress gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 18414-18419.	3.3	128
125	A Major Role of the MEK1/MKK1/2-MPK4 Pathway in ROS Signalling. <i>Molecular Plant</i> , 2009, 2, 120-137.	3.9	250
126	Disentangling the Complexity of Mitogen-Activated Protein Kinases and Reactive Oxygen Species Signaling. <i>Plant Physiology</i> , 2009, 149, 606-615.	2.3	120



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127	MAP KINASE PHOSPHATASE1 and PROTEIN TYROSINE PHOSPHATASE1 Are Repressors of Salicylic Acid Synthesis and SNC1-Mediated Responses in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2009, 21, 2884-2897.	3.1	216
128	MAPK cascade signalling networks in plant defence. <i>Current Opinion in Plant Biology</i> , 2009, 12, 421-426.	3.5	612
129	Protein tyrosine phosphorylation in plants: more abundant than expected?. <i>Trends in Plant Science</i> , 2009, 14, 71-76.	4.3	87
130	Possible involvement of MAP kinase pathways in acquired metal-tolerance induced by heat in plants. <i>Planta</i> , 2008, 228, 499-509.	1.6	24
131	Protein networking: insights into global functional organization of proteomes. <i>Proteomics</i> , 2008, 8, 799-816.	1.3	74
132	Towards functional phosphoproteomics by mapping differential phosphorylation events in signaling networks. <i>Proteomics</i> , 2008, 8, 4453-4465.	1.3	51
133	Site-Specific Phosphorylation Profiling of <i>Arabidopsis</i> Proteins by Mass Spectrometry and Peptide Chip Analysis. <i>Journal of Proteome Research</i> , 2008, 7, 2458-2470.	1.8	139
134	<i>Arabidopsis</i> MAPKs: a complex signalling network involved in multiple biological processes. <i>Biochemical Journal</i> , 2008, 413, 217-226.	1.7	652
135	The Dark Side of the Salad: <i>Salmonella typhimurium</i> Overcomes the Innate Immune Response of <i>Arabidopsis thaliana</i> and Shows an Endopathogenic Lifestyle. <i>PLoS ONE</i> , 2008, 3, e2279.	1.1	142
136	The PP2C-Type Phosphatase AP2C1, Which Negatively Regulates MPK4 and MPK6, Modulates Innate Immunity, Jasmonic Acid, and Ethylene Levels in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2007, 19, 2213-2224.	3.1	302
137	The <i>Arabidopsis</i> Mitogen-Activated Protein Kinase Kinase MKK3 Is Upstream of Group C Mitogen-Activated Protein Kinases and Participates in Pathogen Signaling. <i>Plant Cell</i> , 2007, 19, 3266-3279.	3.1	234
138	The MAP Kinase Kinase MKK2 Affects Disease Resistance in <i>Arabidopsis</i> . <i>Molecular Plant-Microbe Interactions</i> , 2007, 20, 589-596.	1.4	108
139	Using phosphoproteomics to reveal signalling dynamics in plants. <i>Trends in Plant Science</i> , 2007, 12, 404-411.	4.3	63
140	Trojan Horse Strategy in <i>Agrobacterium</i> Transformation: Abusing MAPK Defense Signaling. <i>Science</i> , 2007, 318, 453-456.	6.0	251
141	A plastid-localized glycogen synthase kinase $\epsilon$ 3 modulates stress tolerance and carbohydrate metabolism. <i>Plant Journal</i> , 2007, 49, 1076-1090.	2.8	70
142	The BRI1-Associated Kinase 1, BAK1, Has a Brassinolide-Independent Role in Plant Cell-Death Control. <i>Current Biology</i> , 2007, 17, 1116-1122.	1.8	356
143	Activation of members of a MAPK module in $\beta$ -glucan elicitor-mediated non-host resistance of soybean. <i>Planta</i> , 2007, 225, 1559-1571.	1.6	41
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