

Gerhard H Braus

List of Publications by Year in descending order

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Version: 2024-02-01

248
papers

21,792
citations

22099

59
h-index

11030

137
g-index

263
all docs

263
docs citations

263
times ranked

29511
citing authors

#	ARTICLE	IF	CITATIONS
1	Vacuole fragmentation depends on a novel Atg18-containing retromer-complex. <i>Autophagy</i> , 2023, 19, 278-295.	4.3	13
2	Biosynthesis of Antibacterial Iron-Chelating Tropolones in <i>Aspergillus nidulans</i> as Response to Glycopeptide-Producing Streptomycetes. <i>Frontiers in Fungal Biology</i> , 2022, 2, .	0.9	8
3	SEED LIPID DROPLET PROTEIN1, SEED LIPID DROPLET PROTEIN2, and LIPID DROPLET PLASMA MEMBRANE ADAPTOR mediate lipid droplet-plasma membrane tethering. <i>Plant Cell</i> , 2022, 34, 2424-2448.	3.1	12
4	Adhesion as a Focus in <i>Trichoderma</i> -Root Interactions. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 372.	1.5	6
5	Multi-omics analysis of xylem sap uncovers dynamic modulation of poplar defenses by ammonium and nitrate. <i>Plant Journal</i> , 2022, 111, 282-303.	2.8	11
6	Design of typical genes for heterologous gene expression. <i>Scientific Reports</i> , 2022, 12, .	1.6	0
7	Novel Fus3- and Ste12-interacting protein FsiA activates cell fusion-related genes in both Ste12-dependent and -independent manners in Ascomycete filamentous fungi. <i>Molecular Microbiology</i> , 2021, 115, 723-738.	1.2	6
8	The velvet protein Vel1 controls initial plant root colonization and conidia formation for xylem distribution in <i>Verticillium</i> wilt. <i>PLoS Genetics</i> , 2021, 17, e1009434.	1.5	20
9	Draft Genome Sequence of <i>Saccharomyces cerevisiae</i> LW2591Y, a Laboratory Strain for <i>In Vivo</i> Multigene Assemblies. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.3	1
10	DEAD-box RNA helicase Dbp4/DDX10 is an enhancer of α -synuclein toxicity and oligomerization. <i>PLoS Genetics</i> , 2021, 17, e1009407.	1.5	19
11	Unfolded Protein Response and Scaffold Independent Pheromone MAP Kinase Signaling Control <i>Verticillium dahliae</i> Growth, Development, and Plant Pathogenesis. <i>Journal of Fungi (Basel, Switzerland)</i> 10:1075-1091 (2021) https://doi.org/10.3390/jof10071075 / https://pubmed.ncbi.nlm.nih.gov/3511075/	0.784314	10
12	Identification of Two Novel Peptides That Inhibit α -Synuclein Toxicity and Aggregation. <i>Frontiers in Molecular Neuroscience</i> , 2021, 14, 659926.	1.4	8
13	<i>Arabidopsis thaliana</i> EARLY RESPONSIVE TO DEHYDRATION 7 Localizes to Lipid Droplets via Its Senescence Domain. <i>Frontiers in Plant Science</i> , 2021, 12, 658961.	1.7	16
14	<i>Pseudomonas</i> Strains Induce Transcriptional and Morphological Changes and Reduce Root Colonization of <i>Verticillium</i> spp.. <i>Frontiers in Microbiology</i> , 2021, 12, 652468.	1.5	6
15	A 20-kb lineage-specific genomic region tames virulence in pathogenic amphidiploid <i>Verticillium longisporum</i> . <i>Molecular Plant Pathology</i> , 2021, 22, 939-953.	2.0	6
16	LDIP cooperates with SEIPIN and LDAP to facilitate lipid droplet biogenesis in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2021, 33, 3076-3103.	3.1	31
17	The role of <i>Aspergillus nidulans</i> polo-like kinase PlkA in microtubule-organizing center control. <i>Journal of Cell Science</i> , 2021, 134, .	1.2	3
18	α -Synuclein Decreases the Abundance of Proteasome Subunits and Alters Ubiquitin Conjugates in Yeast Cells, 2021, 10, 2229.	1.8	5

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19	Dynamic and Reversible Aggregation of the Human CAP Superfamily Member GAPR-1 in Protein Inclusions in <i>Saccharomyces cerevisiae</i> . <i>Journal of Molecular Biology</i> , 2021, 433, 167162.	2.0	2
20	Guidelines for the use and interpretation of assays for monitoring autophagy (4th) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702 Td (edition 4.3)	4.3	1,430
21	Secondary metabolites of <i>H₂O₂</i> cells mediate protection of fungal reproductive and overwintering structures against fungivorous animals. <i>ELife</i> , 2021, 10, .	2.8	7
22	The Nma1 protein promotes long distance transport mediated by early endosomes in <i>Ustilago maydis</i> . <i>Molecular Microbiology</i> , 2021, , .	1.2	1
23	A J Domain Protein Functions as a Histone Chaperone to Maintain Genome Integrity and the Response to DNA Damage in a Human Fungal Pathogen. <i>MBio</i> , 2021, 12, e0327321.	1.8	2
24	Production of the Fragrance Geraniol in Peroxisomes of a Product-Tolerant Baker's Yeast. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020, 8, 582052.	2.0	22
25	The High Osmolarity Glycerol Mitogen-Activated Protein Kinase regulates glucose catabolite repression in filamentous fungi. <i>PLoS Genetics</i> , 2020, 16, e1008996.	1.5	15
26	<i>Verticillium longisporum</i> Elicits Media-Dependent Secretome Responses With Capacity to Distinguish Between Plant-Related Environments. <i>Frontiers in Microbiology</i> , 2020, 11, 1876.	1.5	18
27	<i>H₂O₂</i> Cells of <i>Aspergillus nidulans</i> with Nuclear Storage and Developmental Backup Functions Are Reminiscent of Multipotent Stem Cells. <i>MBio</i> , 2020, 11, .	1.8	9
28	The Novel J-Domain Protein Mrj1 Is Required for Mitochondrial Respiration and Virulence in <i>Cryptococcus neoformans</i> . <i>MBio</i> , 2020, 11, .	1.8	15
29	Identification of Low-Abundance Lipid Droplet Proteins in Seeds and Seedlings. <i>Plant Physiology</i> , 2020, 182, 1326-1345.	2.3	44
30	The COP9 signalosome mediates the Spt23 regulated fatty acid desaturation and ergosterol biosynthesis. <i>FASEB Journal</i> , 2020, 34, 4870-4889.	0.2	10
31	The Vta1 transcriptional regulator is required for microsclerotia melanization in <i>Verticillium dahliae</i> . <i>Fungal Biology</i> , 2020, 124, 490-500.	1.1	13
32	Growing a circular economy with fungal biotechnology: a white paper. <i>Fungal Biology and Biotechnology</i> , 2020, 7, 5.	2.5	228
33	8 Coordination of Fungal Secondary Metabolism and Development. , 2020, , 173-205.		2
34	Title is missing!. , 2020, 16, e1008996.		0
35	Title is missing!. , 2020, 16, e1008996.		0
36	Title is missing!. , 2020, 16, e1008996.		0

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37	Title is missing!. , 2020, 16, e1008996.		0
38	NBR1 is involved in selective pexophagy in filamentous ascomycetes and can be functionally replaced by a tagged version of its human homolog. <i>Autophagy</i> , 2019, 15, 78-97.	4.3	18
39	COP9 Signalosome Interaction with UspA/Usp15 Deubiquitinase Controls VeA-Mediated Fungal Multicellular Development. <i>Biomolecules</i> , 2019, 9, 238.	1.8	15
40	Integration of Fungus-Specific CandA-C1 into a Trimeric CandA Complex Allowed Splitting of the Gene for the Conserved Receptor Exchange Factor of CullinA E3 Ubiquitin Ligases in <i>Aspergilli</i> . <i>MBio</i> , 2019, 10, .	1.8	9
41	Antimicrobial propensity of ultrananocrystalline diamond films with embedded silver nanodroplets. <i>Diamond and Related Materials</i> , 2019, 93, 168-178.	1.8	10
42	Yeast-Based Screens to Target Alpha-Synuclein Toxicity. <i>Methods in Molecular Biology</i> , 2019, 1948, 145-156.	0.4	4
43	Broad Substrate-Specific Phosphorylation Events Are Associated With the Initial Stage of Plant Cell Wall Recognition in <i>Neurospora crassa</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 2317.	1.5	25
44	Genome sequencing of evolved aspergilli populations reveals robust genomes, transversions in <i>A. flavus</i> , and sexual aberrancy in non-homologous end-joining mutants. <i>BMC Biology</i> , 2019, 17, 88.	1.7	18
45	<i>Verticillium dahliae</i> transcription factors Som1 and Vta3 control microsclerotia formation and sequential steps of plant root penetration and colonisation to induce disease. <i>New Phytologist</i> , 2019, 221, 2138-2159.	3.5	50
46	Cytoplasmic retention and degradation of a mitotic inducer enable plant infection by a pathogenic fungus. <i>ELife</i> , 2019, 8, .	2.8	7
47	Fungal Morphogenesis, from the Polarized Growth of Hyphae to Complex Reproduction and Infection Structures. <i>Microbiology and Molecular Biology Reviews</i> , 2018, 82, .	2.9	231
48	Fluorescent pseudomonads pursue media-dependent strategies to inhibit growth of pathogenic <i>Verticillium</i> fungi. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 817-831.	1.7	6
49	Importance of Stress Response Mechanisms in Filamentous Fungi for Agriculture and Industry. , 2018, , 189-222.		2
50	The trehalose protective mechanism during thermal stress in <i>Saccharomyces cerevisiae</i> : the roles of Ath1 and Agt1. <i>FEMS Yeast Research</i> , 2018, 18, .	1.1	37
51	Response to Comment on "Sterilizing immunity in the lung relies on targeting fungal apoptosis-like programmed cell death" <i>Science</i> , 2018, 360, .	6.0	1
52	Velvet domain protein VosA represses the zinc cluster transcription factor SclB regulatory network for <i>Aspergillus nidulans</i> asexual development, oxidative stress response and secondary metabolism. <i>PLoS Genetics</i> , 2018, 14, e1007511.	1.5	29
53	Heavy Metal-Induced Expression of PcaA Provides Cadmium Tolerance to <i>Aspergillus fumigatus</i> and Supports Its Virulence in the <i>Galleria mellonella</i> Model. <i>Frontiers in Microbiology</i> , 2018, 9, 744.	1.5	26
54	Sumoylation Protects Against β^2 -Synuclein Toxicity in Yeast. <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 94.	1.4	11

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55	A novel STRIPAK complex component mediates hyphal fusion and fruiting body development in filamentous fungi. <i>Molecular Microbiology</i> , 2018, 110, 513-532.	1.2	19
56	PUX10 Is a Lipid Droplet-Localized Scaffold Protein That Interacts with CELL DIVISION CYCLE48 and Is Involved in the Degradation of Lipid Droplet Proteins. <i>Plant Cell</i> , 2018, 30, 2137-2160.	3.1	78
57	Regulation of <i>Aspergillus nidulans</i> CreA-Mediated Catabolite Repression by the F-Box Proteins Fbx23 and Fbx47. <i>MBio</i> , 2018, 9, .	1.8	70
58	Sem1 links proteasome stability and specificity to multicellular development. <i>PLoS Genetics</i> , 2018, 14, e1007141.	1.5	15
59	Comparative genomics reveals high biological diversity and specific adaptations in the industrially and medically important fungal genus <i>Aspergillus</i> . <i>Genome Biology</i> , 2017, 18, 28.	3.8	417
60	<i>Arabidopsis</i> lipid droplet-associated protein (LDAP) interacting protein (LDIP) influences lipid droplet size and neutral lipid homeostasis in both leaves and seeds. <i>Plant Journal</i> , 2017, 92, 1182-1201.	2.8	71
61	Capturing the Asc1p/Receptor for Activated C Kinase 1 (RACK1) Microenvironment at the Head Region of the 40S Ribosome with Quantitative BioID in Yeast. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 2199-2218.	2.5	63
62	The truncated NLR protein TIR1 is a MOS6/IMPORTIN13 interaction partner required for plant immunity. <i>Plant Journal</i> , 2017, 92, 808-821.	2.8	43
63	Sterilizing immunity in the lung relies on targeting fungal apoptosis-like programmed cell death. <i>Science</i> , 2017, 357, 1037-1041.	6.0	92
64	MybA, a transcription factor involved in conidiation and conidial viability of the human pathogen <i>Aspergillus fumigatus</i> . <i>Molecular Microbiology</i> , 2017, 105, 880-900.	1.2	31
65	Proteomic profiling of the antifungal drug response of <i>Aspergillus fumigatus</i> to voriconazole. <i>International Journal of Medical Microbiology</i> , 2017, 307, 398-408.	1.5	12
66	BcXYG1, a Secreted Xyloglucanase from <i>Botrytis cinerea</i> , Triggers Both Cell Death and Plant Immune Responses. <i>Plant Physiology</i> , 2017, 175, 438-456.	2.3	102
67	Analysis of the lipid body proteome of the oleaginous alga <i>Lobosphaera incisa</i> . <i>BMC Plant Biology</i> , 2017, 17, 98.	1.6	44
68	Asc1p/RACK1 Connects Ribosomes to Eukaryotic Phosphosignaling. <i>Molecular and Cellular Biology</i> , 2017, 37, .	1.1	29
69	The DenA/DEN1 Interacting Phosphatase DipA Controls Septa Positioning and Phosphorylation-Dependent Stability of Cytoplasmatic DenA/DEN1 during Fungal Development. <i>PLoS Genetics</i> , 2016, 12, e1005949.	1.5	18
70	SCF Ubiquitin Ligase F-box Protein Fbx15 Controls Nuclear Co-repressor Localization, Stress Response and Virulence of the Human Pathogen <i>Aspergillus fumigatus</i> . <i>PLoS Pathogens</i> , 2016, 12, e1005899.	2.1	60
71	Current challenges of research on filamentous fungi in relation to human welfare and a sustainable bio-economy: a white paper. <i>Fungal Biology and Biotechnology</i> , 2016, 3, 6.	2.5	208
72	Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). <i>Autophagy</i> , 2016, 12, 1-222.	4.3	4,701

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73	<i>CHK2</i> and <i>BRCA1</i> tumor-suppressor axis restrains oncogenic Aurora-A kinase to ensure proper mitotic microtubule assembly. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 1817-1822.	3.3	51
74	Changes of global gene expression and secondary metabolite accumulation during light-dependent <i>Aspergillus nidulans</i> development. <i>Fungal Genetics and Biology</i> , 2016, 87, 30-53.	0.9	56
75	Yeast reveals similar molecular mechanisms underlying alpha- and beta-synuclein toxicity. <i>Human Molecular Genetics</i> , 2016, 25, 275-290.	1.4	29
76	The devil is in the details: comparison between COP9 signalosome (CSN) and the LID of the 26S proteasome. <i>Current Genetics</i> , 2016, 62, 129-136.	0.8	12
77	The putative oncogene CEP72 inhibits the mitotic function of BRCA1 and induces chromosomal instability. <i>Oncogene</i> , 2016, 35, 2398-2406.	2.6	22
78	<i>Bacillus thuringiensis</i> and <i>Bacillus weihenstephanensis</i> Inhibit the Growth of Phytopathogenic <i>Verticillium</i> Species. <i>Frontiers in Microbiology</i> , 2016, 7, 2171.	1.5	74
79	C-Terminal Tyrosine Residue Modifications Modulate the Protective Phosphorylation of Serine 129 of α -Synuclein in a Yeast Model of Parkinson's Disease. <i>PLoS Genetics</i> , 2016, 12, e1006098.	1.5	49
80	In vitro Deneddylation Assay. <i>Bio-protocol</i> , 2016, 6, .	0.2	2
81	Transcription Factor SomA Is Required for Adhesion, Development and Virulence of the Human Pathogen <i>Aspergillus fumigatus</i> . <i>PLoS Pathogens</i> , 2015, 11, e1005205.	2.1	57
82	Draft Genome Sequence of the Phenazine-Producing <i>Pseudomonas fluorescens</i> Strain 2-79. <i>Genome Announcements</i> , 2015, 3, .	0.8	5
83	Genetically Encoding Lysine Modifications on Histone H4. <i>ACS Chemical Biology</i> , 2015, 10, 939-944.	1.6	46
84	Dissecting the function of the different chitin synthases in vegetative growth and sexual development in <i>Neurospora crassa</i> . <i>Fungal Genetics and Biology</i> , 2015, 75, 30-45.	0.9	52
85	Posttranslational Modifications and Clearing of α -Synuclein Aggregates in Yeast. <i>Biomolecules</i> , 2015, 5, 617-634.	1.8	33
86	Draft Genome Sequence of the Beneficial Rhizobacterium <i>Pseudomonas fluorescens</i> DSM 8569, a Natural Isolate of Oilseed Rape (<i>Brassica napus</i>). <i>Genome Announcements</i> , 2015, 3, .	0.8	2
87	One Juliet and four Romeos: VeA and its methyltransferases. <i>Frontiers in Microbiology</i> , 2015, 6, 1.	1.5	1,444
88	The <i>SrkA</i> Kinase Is Part of the <i>SakA</i> Mitogen-Activated Protein Kinase Interactome and Regulates Stress Responses and Development in <i>Aspergillus nidulans</i> . <i>Eukaryotic Cell</i> , 2015, 14, 495-510.	3.4	66
89	Integration of the catalytic subunit activates deneddylase activity <i>in vivo</i> as final step in fungal COP9 signalosome assembly. <i>Molecular Microbiology</i> , 2015, 97, 110-124.	1.2	18
90	RNAseq analysis of <i>Aspergillus fumigatus</i> in blood reveals a just wait and see resting stage behavior. <i>BMC Genomics</i> , 2015, 16, 640.	1.2	25

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91	Systematic Comparison of the Effects of Alpha-synuclein Mutations on Its Oligomerization and Aggregation. <i>PLoS Genetics</i> , 2014, 10, e1004741.	1.5	168
92	Manipulation of fungal development as source of novel secondary metabolites for biotechnology. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 8443-8455.	1.7	43
93	Interplay between Sumoylation and Phosphorylation for Protection against α -Synuclein Inclusions. <i>Journal of Biological Chemistry</i> , 2014, 289, 31224-31240.	1.6	63
94	Infections with the vascular pathogens <i>Verticillium longisporum</i> and <i>Verticillium dahliae</i> induce distinct disease symptoms and differentially affect drought stress tolerance of <i>Arabidopsis thaliana</i> . <i>Environmental and Experimental Botany</i> , 2014, 108, 23-37.	2.0	38
95	<i>Verticillium</i> transcription activator of adhesion <i>Vta2</i> suppresses microsclerotia formation and is required for systemic infection of plant roots. <i>New Phytologist</i> , 2014, 202, 565-581.	3.5	92
96	<i>Verticillium dahliae</i> VdTHI4, involved in thiazole biosynthesis, stress response and DNA repair functions, is required for vascular disease induction in tomato. <i>Environmental and Experimental Botany</i> , 2014, 108, 14-22.	2.0	40
97	A novel <i>Arabidopsis</i> CHITIN ELICITOR RECEPTOR KINASE 1 (CERK1) mutant with enhanced pathogen-induced cell death and altered receptor processing. <i>New Phytologist</i> , 2014, 204, 955-967.	3.5	55
98	α -Synuclein interacts with the switch region of Rab8a in a Ser129 phosphorylation-dependent manner. <i>Neurobiology of Disease</i> , 2014, 70, 149-161.	2.1	84
99	Establishing a versatile Golden Gate cloning system for genetic engineering in fungi. <i>Fungal Genetics and Biology</i> , 2014, 62, 1-10.	0.9	102
100	Membrane-Bound Methyltransferase Complex VapA-VipC-VapB Guides Epigenetic Control of Fungal Development. <i>Developmental Cell</i> , 2014, 29, 406-420.	3.1	63
101	Molecular diagnosis to discriminate pathogen and apathogen species of the hybrid <i>Verticillium longisporum</i> on the oilseed crop <i>Brassica napus</i> . <i>Applied Microbiology and Biotechnology</i> , 2013, 97, 4467-4483.	1.7	30
102	conF and conJ contribute to conidia germination and stress response in the filamentous fungus <i>Aspergillus nidulans</i> . <i>Fungal Genetics and Biology</i> , 2013, 56, 42-53.	0.9	31
103	A structural model of PpoA derived from SAXS-analysis—Implications for substrate conversion. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2013, 1831, 1449-1457.	1.2	9
104	Interplay of the fungal sumoylation network for control of multicellular development. <i>Molecular Microbiology</i> , 2013, 90, 1125-1145.	1.2	40
105	The Velvet Family of Fungal Regulators Contains a DNA-Binding Domain Structurally Similar to NF- κ B. <i>PLoS Biology</i> , 2013, 11, e1001750.	2.6	121
106	Control of Multicellular Development by the Physically Interacting Deneddylases DEN1/DenA and COP9 Signalosome. <i>PLoS Genetics</i> , 2013, 9, e1003275.	1.5	42
107	RACK1/Asc1p, a Ribosomal Node in Cellular Signaling. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 87-105.	2.5	37
108	Mutual Cross Talk between the Regulators Hac1 of the Unfolded Protein Response and Gcn4 of the General Amino Acid Control of <i>Saccharomyces cerevisiae</i> . <i>Eukaryotic Cell</i> , 2013, 12, 1142-1154.	3.4	17

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109	The Cpc1 Regulator of the Cross-Pathway Control of Amino Acid Biosynthesis Is Required for Pathogenicity of the Vascular Pathogen <i>Verticillium longisporum</i> . <i>Molecular Plant-Microbe Interactions</i> , 2013, 26, 1312-1324.	1.4	55
110	The <i>Aspergillus nidulans</i> MAPK Module AnSte11-Ste50-Ste7-Fus3 Controls Development and Secondary Metabolism. <i>PLoS Genetics</i> , 2012, 8, e1002816.	1.5	182
111	Transcriptional Activation and Production of Tryptophan-Derived Secondary Metabolites in <i>Arabidopsis</i> Roots Contributes to the Defense against the Fungal Vascular Pathogen <i>Verticillium longisporum</i> . <i>Molecular Plant</i> , 2012, 5, 1389-1402.	3.9	120
112	Aggregate Clearance of β -Synuclein in <i>Saccharomyces cerevisiae</i> Depends More on Autophagosome and Vacuole Function Than on the Proteasome. <i>Journal of Biological Chemistry</i> , 2012, 287, 27567-27579.	1.6	66
113	Characterization of the velvet regulators in <i>Aspergillus fumigatus</i> . <i>Molecular Microbiology</i> , 2012, 86, 937-953.	1.2	84
114	The Plant Host <i>Brassica napus</i> Induces in the Pathogen <i>Verticillium longisporum</i> the Expression of Functional Catalase Peroxidase Which Is Required for the Late Phase of Disease. <i>Molecular Plant-Microbe Interactions</i> , 2012, 25, 569-581.	1.4	55
115	Breaking the Silence: Protein Stabilization Uncovers Silenced Biosynthetic Gene Clusters in the Fungus <i>Aspergillus nidulans</i> . <i>Applied and Environmental Microbiology</i> , 2012, 78, 8234-8244.	1.4	64
116	Molecular characterization of the <i>Aspergillus nidulans</i> fbxA encoding an F-box protein involved in xylanase induction. <i>Fungal Genetics and Biology</i> , 2012, 49, 130-140.	0.9	29
117	Fungal S-adenosylmethionine synthetase and the control of development and secondary metabolism in <i>Aspergillus nidulans</i> . <i>Fungal Genetics and Biology</i> , 2012, 49, 443-454.	0.9	25
118	Identification of Protein Complexes from Filamentous Fungi with Tandem Affinity Purification. <i>Methods in Molecular Biology</i> , 2012, 944, 191-205.	0.4	37
119	Structure-functional analysis of the <i>Dictyoglomus</i> cell envelope. <i>Systematic and Applied Microbiology</i> , 2012, 35, 279-290.	1.2	10
120	The COP9 signalosome counteracts the accumulation of cullin SCF ubiquitin E3 RING ligases during fungal development. <i>Molecular Microbiology</i> , 2012, 83, 1162-1177.	1.2	40
121	Coordination of secondary metabolism and development in fungi: the velvet family of regulatory proteins. <i>FEMS Microbiology Reviews</i> , 2012, 36, 1-24.	3.9	477
122	Comparative genomics of citric-acid-producing <i>Aspergillus niger</i> ATCC 1015 versus enzyme-producing CBS 513.88. <i>Genome Research</i> , 2011, 21, 885-897.	2.4	329
123	A Feedback Circuit between Transcriptional Activation and Self-Destruction of Gcn4 Separates Its Metabolic and Morphogenic Response in Diploid Yeasts. <i>Journal of Molecular Biology</i> , 2011, 405, 909-925.	2.0	8
124	5'UTR: Identification and Analysis of Translationally Regulative Untranslated Regions in Amino Acid Starved Yeast Cells. <i>Molecular and Cellular Proteomics</i> , 2011, 10, M110.003350.	2.5	5
125	Recruitment of the inhibitor Cand1 to the cullin substrate adaptor site mediates interaction to the neddylation site. <i>Molecular Biology of the Cell</i> , 2011, 22, 153-164.	0.9	27
126	Silencing of Vlaro2 for chorismate synthase revealed that the phytopathogen <i>Verticillium longisporum</i> induces the cross-pathway control in the xylem. <i>Applied Microbiology and Biotechnology</i> , 2010, 85, 1961-1976.	1.7	62

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127	The COP9 signalosome mediates transcriptional and metabolic response to hormones, oxidative stress protection and cell wall rearrangement during fungal development. <i>Molecular Microbiology</i> , 2010, 78, 964-979.	1.2	81
128	LaeA Control of Velvet Family Regulatory Proteins for Light-Dependent Development and Fungal Cell-Type Specificity. <i>PLoS Genetics</i> , 2010, 6, e1001226.	1.5	233
129	Dissection of mitotic functions of the yeast cyclin Clb2. <i>Cell Cycle</i> , 2010, 9, 2611-2619.	1.3	7
130	Fungal development and the COP9 signalosome. <i>Current Opinion in Microbiology</i> , 2010, 13, 672-676.	2.3	74
131	Spotlight on <i>Aspergillus nidulans</i> photosensory systems. <i>Fungal Genetics and Biology</i> , 2010, 47, 900-908.	0.9	138
132	Degradation of <i>Saccharomyces cerevisiae</i> Transcription Factor Gcn4 Requires a C-Terminal Nuclear Localization Signal in the Cyclin Pcl5. <i>Eukaryotic Cell</i> , 2009, 8, 496-510.	3.4	9
133	The Yeast HtrA Orthologue Ynm3 Is a Protease with Chaperone Activity that Aids Survival Under Heat Stress. <i>Molecular Biology of the Cell</i> , 2009, 20, 68-77.	0.9	32
134	The protein kinase ImeB is required for light-mediated inhibition of sexual development and for mycotoxin production in <i>Aspergillus nidulans</i> . <i>Molecular Microbiology</i> , 2009, 71, 1278-1295.	1.2	42
135	Pre-fibrillar α -synuclein variants with impaired β -structure increase neurotoxicity in Parkinson's disease models. <i>EMBO Journal</i> , 2009, 28, 3256-3268.	3.5	411
136	The 2008 update of the <i>Aspergillus nidulans</i> genome annotation: A community effort. <i>Fungal Genetics and Biology</i> , 2009, 46, S2-S13.	0.9	99
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