List of Publications by Year in descending order

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131	11,398	47006 47	30087
papers	citations	h-index	g-index
135	135	135	7438
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Characterization of 260 Isolates of Aspergillus Section Flavi Obtained from Sesame Seeds in Punjab, Pakistan. Toxins, 2022, 14, 117.	3.4	4
2	The velvet-activated putative C6 transcription factor VadZ regulates development and sterigmatocystin production in Aspergillus nidulans. Fungal Biology, 2022, , .	2.5	5
3	Analysis of E.U. Rapid Alert System (RASFF) Notifications for Aflatoxins in Exported U.S. Food and Feed Products for 2010–2019. Toxins, 2021, 13, 90.	3.4	22
4	Transcriptomic, Protein-DNA Interaction, and Metabolomic Studies of VosA, VelB, and WetA in Aspergillus nidulans Asexual Spores. MBio, 2021, 12, .	4.1	29
5	Characterization of the mbsA Gene Encoding a Putative APSES Transcription Factor in Aspergillus fumigatus. International Journal of Molecular Sciences, 2021, 22, 3777.	4.1	9
6	The DUG Pathway Governs Degradation of Intracellular Glutathione in Aspergillus nidulans. Applied and Environmental Microbiology, 2021, 87, .	3.1	6
7	Characterization of the asexual developmental genes brlA and wetA in Monascus ruber M7. Fungal Genetics and Biology, 2021, 151, 103564.	2.1	18
8	AtfA-Independent Adaptation to the Toxic Heavy Metal Cadmium in Aspergillus nidulans. Microorganisms, 2021, 9, 1433.	3.6	10
9	The putative sensor histidine kinase VadJ coordinates development and sterigmatocystin production in Aspergillus nidulans. Journal of Microbiology, 2021, 59, 746-752.	2.8	4
10	Transcriptomic and Functional Studies of the RGS Protein Rax1 in Aspergillus fumigatus. Pathogens, 2020, 9, 36.	2.8	3
11	More P450s Are Involved in Secondary Metabolite Biosynthesis in Streptomyces Compared to Bacillus, Cyanobacteria, and Mycobacterium. International Journal of Molecular Sciences, 2020, 21, 4814.	4.1	20
12	Velvet activated McrA plays a key role in cellular and metabolic development in Aspergillus nidulans. Scientific Reports, 2020, 10, 15075.	3.3	6
13	Heterotrimeric G-Protein Signalers and RGSs in Aspergillus fumigatus. Pathogens, 2020, 9, 902.	2.8	16
14	Comparative Analysis, Structural Insights, and Substrate/Drug Interaction of CYP128A1 in Mycobacterium tuberculosis. International Journal of Molecular Sciences, 2020, 21, 4816.	4.1	7
15	The role of the VosA-repressed dnjA gene in development and metabolism in Aspergillus species. Current Genetics, 2020, 66, 621-633.	1.7	15
16	The velvet Regulator VosA Governs Survival and Secondary Metabolism of Sexual Spores in Aspergillus nidulans. Genes, 2020, 11, 103.	2.4	15
17	A Liquid Chromatographic Method for Rapid and Sensitive Analysis of Aflatoxins in Laboratory Fungal Cultures. Toxins, 2020, 12, 93.	3.4	15
18	Comprehensive Analyses of Cytochrome P450 Monooxygenases and Secondary Metabolite Biosynthetic Gene Clusters in Cyanobacteria. International Journal of Molecular Sciences, 2020, 21, 656.	4.1	21

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19	Increased Cd ²⁺ biosorption capability of <i>Aspergillus nidulans</i> elicited by <i>crpA</i> deletion. Journal of Basic Microbiology, 2020, 60, 574-584.	3.3	7
20	The Putative APSES Transcription Factor RgdA Governs Growth, Development, Toxigenesis, and Virulence in Aspergillus fumigatus. MSphere, 2020, 5, .	2.9	13
21	Effects of Different G-Protein α-Subunits on Growth, Development and Secondary Metabolism of Monascus ruber M7. Frontiers in Microbiology, 2019, 10, 1555.	3.5	17
22	RgsA Attenuates the PKA Signaling, Stress Response, and Virulence in the Human Opportunistic Pathogen Aspergillus fumigatus. International Journal of Molecular Sciences, 2019, 20, 5628.	4.1	10
23	The velvet repressed vidA gene plays a key role in governing development in Aspergillus nidulans. Journal of Microbiology, 2019, 57, 893-899.	2.8	13
24	RgsD negatively controls development, toxigenesis, stress response, and virulence in Aspergillus fumigatus. Scientific Reports, 2019, 9, 811.	3.3	15
25	Distribution and Diversity of Cytochrome P450 Monooxygenases in the Fungal Class Tremellomycetes. International Journal of Molecular Sciences, 2019, 20, 2889.	4.1	12
26	Cytochrome P450 Monooxygenase-Mediated Metabolic Utilization of Benzo[a]Pyrene by Aspergillus Species. MBio, 2019, 10, .	4.1	22
27	Cytochrome P450 Monooxygenase CYP139 Family Involved in the Synthesis of Secondary Metabolites in 824 Mycobacterial Species. International Journal of Molecular Sciences, 2019, 20, 2690.	4.1	13
28	Developmental Decisions in Aspergillus nidulans. , 2019, , 63-80.		10
28		3.3	10
	Developmental Decisions in Aspergillus nidulans. , 2019, , 63-80. Similarities, variations, and evolution of cytochrome P450s in Streptomyces versus Mycobacterium.	3.3	
29	Developmental Decisions in Aspergillus nidulans. , 2019, , 63-80. Similarities, variations, and evolution of cytochrome P450s in Streptomyces versus Mycobacterium. Scientific Reports, 2019, 9, 3962. Disturbance in biosynthesis of arachidonic acid impairs the sexual development of the onion blight		28
30	Developmental Decisions in Aspergillus nidulans. , 2019, , 63-80. Similarities, variations, and evolution of cytochrome P450s in Streptomyces versus Mycobacterium. Scientific Reports, 2019, 9, 3962. Disturbance in biosynthesis of arachidonic acid impairs the sexual development of the onion blight pathogen Stemphylium eturmiunum. Current Genetics, 2019, 65, 759-771. Comparative analyses and structural insights of the novel cytochrome P450 fusion protein family	1.7	28
29 30 31	Developmental Decisions in Aspergillus nidulans. , 2019, , 63-80. Similarities, variations, and evolution of cytochrome P450s in Streptomyces versus Mycobacterium. Scientific Reports, 2019, 9, 3962. Disturbance in biosynthesis of arachidonic acid impairs the sexual development of the onion blight pathogen Stemphylium eturmiunum. Current Genetics, 2019, 65, 759-771. Comparative analyses and structural insights of the novel cytochrome P450 fusion protein family CYP5619 in Oomycetes. Scientific Reports, 2018, 8, 6597. Controlling aflatoxin contamination and propagation of Aspergillus flavus by a soy-fermenting	1.7 3.3	28 3 4
29 30 31 32	Developmental Decisions in Aspergillus nidulans., 2019,, 63-80. Similarities, variations, and evolution of cytochrome P450s in Streptomyces versus Mycobacterium. Scientific Reports, 2019, 9, 3962. Disturbance in biosynthesis of arachidonic acid impairs the sexual development of the onion blight pathogen Stemphylium eturmiunum. Current Genetics, 2019, 65, 759-771. Comparative analyses and structural insights of the novel cytochrome P450 fusion protein family CYP5619 in Oomycetes. Scientific Reports, 2018, 8, 6597. Controlling aflatoxin contamination and propagation of Aspergillus flavus by a soy-fermenting Aspergillus oryzae strain. Scientific Reports, 2018, 8, 16871.	1.7 3.3 3.3	28 3 4 66
30 31 32 33	Developmental Decisions in Aspergillus nidulans., 2019, , 63-80. Similarities, variations, and evolution of cytochrome P450s in Streptomyces versus Mycobacterium. Scientific Reports, 2019, 9, 3962. Disturbance in biosynthesis of arachidonic acid impairs the sexual development of the onion blight pathogen Stemphylium eturmiunum. Current Genetics, 2019, 65, 759-771. Comparative analyses and structural insights of the novel cytochrome P450 fusion protein family CYP5619 in Oomycetes. Scientific Reports, 2018, 8, 6597. Controlling aflatoxin contamination and propagation of Aspergillus flavus by a soy-fermenting Aspergillus oryzae strain. Scientific Reports, 2018, 8, 16871. Comparative Analyses of Cytochrome P450s and Those Associated with Secondary Metabolism in Bacillus Species. International Journal of Molecular Sciences, 2018, 19, 3623. Efficient Adsorption on Benzoyl and Stearoyl Cellulose to Remove Phenanthrene and Pyrene from	1.7 3.3 3.3 4.1	28 3 4 66

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37	Blooming of Unusual Cytochrome P450s by Tandem Duplication in the Pathogenic Fungus Conidiobolus coronatus. International Journal of Molecular Sciences, 2018, 19, 1711.	4.1	15
38	Lipid Biosynthesis as an Antifungal Target. Journal of Fungi (Basel, Switzerland), 2018, 4, 50.	3.5	61
39	Bioremediation and microbial metabolism of benzo(a)pyrene. Molecular Microbiology, 2018, 109, 433-444.	2.5	59
40	Systematic Dissection of the Evolutionarily Conserved WetA Developmental Regulator across a Genus of Filamentous Fungi. MBio, 2018, 9, .	4.1	68
41	Characterization and regulated naproxen release of hydroxypropyl cyclosophoraose-pullulan microspheres. Journal of Industrial and Engineering Chemistry, 2017, 48, 108-118.	5.8	28
42	Comparative genomics reveals high biological diversity and specific adaptations in the industrially and medically important fungal genus Aspergillus. Genome Biology, 2017, 18, 28.	8.8	417
43	High molecular weight genomic DNA mini-prep for filamentous fungi. Fungal Genetics and Biology, 2017, 104, 1-5.	2.1	17
44	Characterization of the rax1 gene encoding a putative regulator of G protein signaling in Aspergillus fumigatus. Biochemical and Biophysical Research Communications, 2017, 487, 426-432.	2.1	13
45	Diversity, Application, and Synthetic Biology of Industrially Important Aspergillus Fungi. Advances in Applied Microbiology, 2017, 100, 161-202.	2.4	114
46	MybA, a transcription factor involved in conidiation and conidial viability of the human pathogen <i>Aspergillus fumigatus </i> . Molecular Microbiology, 2017, 105, 880-900.	2.5	31
47	Occurrence, Toxicity, and Analysis of Major Mycotoxins in Food. International Journal of Environmental Research and Public Health, 2017, 14, 632.	2.6	763
48	Characteristics of a Regulator of G-Protein Signaling (RGS) rgsC in Aspergillus fumigatus. Frontiers in Microbiology, 2017, 8, 2058.	3.5	19
49	Transcriptome-Based Modeling Reveals that Oxidative Stress Induces Modulation of the AtfA-Dependent Signaling Networks in <i>Aspergillus nidulans</i> Linternational Journal of Genomics, 2017, 2017, 1-14.	1.6	18
50	WetA bridges cellular and chemical development in Aspergillus flavus. PLoS ONE, 2017, 12, e0179571.	2.5	48
51	The role of VosA/VelB-activated developmental gene vadA in Aspergillus nidulans. PLoS ONE, 2017, 12, e0177099.	2.5	25
52	Negative regulation and developmental competence in Aspergillus. Scientific Reports, 2016, 6, 28874.	3.3	77
53	Characterization of the aodA, dnmA, mnSOD and pimA genes in Aspergillus nidulans. Scientific Reports, 2016, 6, 20523.	3.3	26
54	Aspergillus fumigatus spore proteomics and genetics reveal that VeA represses DefA-mediated DNA damage response. Journal of Proteomics, 2016, 148, 26-35.	2.4	2

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55	Molecular evolutionary dynamics of cytochrome P450 monooxygenases across kingdoms: Special focus on mycobacterial P450s. Scientific Reports, 2016, 6, 33099.	3.3	61
56	Enhancing bio-availability of β-naphthoflavone by supramolecular complexation with 6,6′-thiobis(methylene)-β-cyclodextrin dimer. Carbohydrate Polymers, 2016, 151, 40-50.	10.2	12
57	Chemically functionalized silica gel with alkynyl terminated monolayers as an efficient new material for removal of mercury ions from water. Journal of Industrial and Engineering Chemistry, 2016, 35, 376-382.	5.8	32
58	Developmental regulators in Aspergillus fumigatus. Journal of Microbiology, 2016, 54, 223-231.	2.8	58
59	1 Molecular Biology of Asexual Sporulation in Filamentous Fungi. , 2016, , 3-19.		6
60	Characterization of gprK Encoding a Putative Hybrid G-Protein-Coupled Receptor in Aspergillus fumigatus. PLoS ONE, 2016, 11, e0161312.	2.5	32
61	Velvet Regulators in Aspergillus spp Microbiology and Biotechnology Letters, 2016, 44, 409-419.	0.4	37
62	Gâ€protein coupled receptorâ€mediated nutrient sensing and developmental control in <scp><i>A</i></scp> <i>spergillus nidulans</i>	2.5	31
63	The WOPR Domain Protein OsaA Orchestrates Development in Aspergillus nidulans. PLoS ONE, 2015, 10, e0137554.	2.5	25
64	Genome-Wide Annotation and Comparative Analysis of Cytochrome P450 Monooxygenases in Basidiomycete Biotrophic Plant Pathogens. PLoS ONE, 2015, 10, e0142100.	2.5	46
65	Proteomic analyses reveal the key roles of BrlA and AbaA in biogenesis of gliotoxin in Aspergillus fumigatus. Biochemical and Biophysical Research Communications, 2015, 463, 428-433.	2.1	25
66	Genetic Control of Asexual Development in Aspergillus fumigatus. Advances in Applied Microbiology, 2015, 90, 93-107.	2.4	27
67	\hat{l}^3 -Glutamyl transpeptidase (GgtA) of Aspergillus nidulans is not necessary for bulk degradation of glutathione. Archives of Microbiology, 2015, 197, 285-297.	2.2	6
68	Diversity and evolution of cytochrome P450 monooxygenases in Oomycetes. Scientific Reports, 2015, 5, 11572.	3.3	24
69	Novel magnetic nanoparticles coated by benzene- and \hat{l}^2 -cyclodextrin-bearing dextran, and the sorption of polycyclic aromatic hydrocarbon. Carbohydrate Polymers, 2015, 133, 221-228.	10.2	25
70	Epigenetics of Fungal Secondary Metabolism Related Genes. Fungal Biology, 2015, , 29-42.	0.6	4
71	Velvet-mediated repression of \hat{l}^2 -glucan synthesis in Aspergillus nidulans spores. Scientific Reports, 2015, 5, 10199.	3.3	41
72	Removal of methyl violet dye by adsorption onto N-benzyltriazole derivatized dextran. RSC Advances, 2015, 5, 34327-34334.	3.6	34

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73	Core oxidative stress response in Aspergillus nidulans. BMC Genomics, 2015, 16, 478.	2.8	45
74	Hydroxypropyl cyclic \hat{l}^2 -(1â†'2)-d-glucans and epichlorohydrin \hat{l}^2 -cyclodextrin dimers as effective carbohydrate-solubilizers for polycyclic aromatic hydrocarbons. Carbohydrate Research, 2015, 401, 82-88.	2.3	6
75	Cytochrome P450 monooxygenase analysis in free-living and symbiotic microalgae Coccomyxa sp. C-169 and Chlorella sp. NC64A. Algae, 2015, 30, 233-239.	2.3	4
76	NsdD Is a Key Repressor of Asexual Development in <i>Aspergillus nidulans</i> . Genetics, 2014, 197, 159-173.	2.9	71
77	Intermolecular complexation of low-molecular-weight succinoglycans directs solubility enhancement of pindolol. Carbohydrate Polymers, 2014, 106, 101-108.	10.2	15
78	Mild, Selective Oxidation of Aromatic Alcohols Using $font^2$ (font-Cyclodextrin-Functionalized Glass Microparticles: Characterization, Stability, and Application. Synthetic Communications, 2014, 44, 589-599.	2.1	1
79	Upstream Regulation of Mycotoxin Biosynthesis. Advances in Applied Microbiology, 2014, 86, 251-278.	2.4	21
80	Fungal Cytochrome P450 Monooxygenases: Their Distribution, Structure, Functions, Family Expansion, and Evolutionary Origin. Genome Biology and Evolution, 2014, 6, 1620-1634.	2. 5	179
81	VelC Positively Controls Sexual Development in Aspergillus nidulans. PLoS ONE, 2014, 9, e89883.	2.5	69
82	Antimicrobial Properties of Glass Surface Functionalized with Silver-doped Terminal-alkynyl Monolayers. Bulletin of the Korean Chemical Society, 2014, 35, 39-44.	1.9	2
83	Comparative proteomic analyses reveal that FlbA down-regulates gliT expression and SOD activity in Aspergillus fumigatus. Journal of Proteomics, 2013, 87, 40-52.	2.4	17
84	The Velvet Family of Fungal Regulators Contains a DNA-Binding Domain Structurally Similar to NF-κB. PLoS Biology, 2013, 11, e1001750.	5.6	121
85	Expression and Activity of Catalases Is Differentially Affected by GpaA (Ga) and FlbA (Regulator of G) Tj ETQq1 1 (0.784314	rgBT /Overloo
86	GÎ ² -Like CpcB Plays a Crucial Role for Growth and Development of Aspergillus nidulans and Aspergillus fumigatus. PLoS ONE, 2013, 8, e70355.	2.5	28
87	The Putative Guanine Nucleotide Exchange Factor RicA Mediates Upstream Signaling for Growth and Development in Aspergillus. Eukaryotic Cell, 2012, 11, 1399-1412.	3.4	25
88	Characterization of the <i>velvet</i> regulators in <i><scp>A</scp>spergillus fumigatus</i> Molecular Microbiology, 2012, 86, 937-953.	2.5	84
89	Genetic control of asexual sporulation in filamentous fungi. Current Opinion in Microbiology, 2012, 15, 669-677.	5.1	331
90	Multi-Copy Genetic Screen in Aspergillus nidulans. Methods in Molecular Biology, 2012, 944, 183-190.	0.9	29

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91	Investigation of In Vivo Protein Interactions in Aspergillus Spores. Methods in Molecular Biology, 2012, 944, 251-257.	0.9	2
92	Antifungal activity of extracellular hydrolases produced by autolysing Aspergillus nidulans cultures. Journal of Microbiology, 2012, 50, 849-854.	2.8	10
93	The Role, Interaction and Regulation of the Velvet Regulator VelB in Aspergillus nidulans. PLoS ONE, 2012, 7, e45935.	2.5	101
94	The small molecular mass antifungal protein of <i>Penicillium chrysogenum</i> $\hat{a}\in \hat{a}$ a mechanism of action oriented review. Journal of Basic Microbiology, 2011, 51, 561-571.	3.3	35
95	Extracellular proteinase formation in carbon starving <i>Aspergillus nidulans</i> cultures – physiological function and regulation. Journal of Basic Microbiology, 2011, 51, 625-634.	3.3	17
96	AbaA and WetA govern distinct stages of Aspergillus fumigatus development. Microbiology (United) Tj ETQq0 0	0 rgBT /Ov	verlock 10 Tf 5
97	The choC gene encoding a putative phospholipid methyltransferase is essential for growth and development in Aspergillus nidulans. Current Genetics, 2010, 56, 283-296.	1.7	19
98	FlbC is a putative nuclear C2H2 transcription factor regulating development in Aspergillus nidulans. Molecular Microbiology, 2010, 77, 1203-1219.	2.5	138
99	Aspergillus fumigatus <i>flbB</i> Encodes Two Basic Leucine Zipper Domain (bZIP) Proteins Required for Proper Asexual Development and Gliotoxin Production. Eukaryotic Cell, 2010, 9, 1711-1723.	3.4	48
100	LaeA Control of Velvet Family Regulatory Proteins for Light-Dependent Development and Fungal Cell-Type Specificity. PLoS Genetics, 2010, 6, e1001226.	3.5	233
101	Characterization of the developmental regulator FlbE in Aspergillus fumigatus and Aspergillus nidulans. Fungal Genetics and Biology, 2010, 47, 981-993.	2.1	72
102	Regulation of Development in <i>Aspergillus nidulans</i> and <i>Aspergillus fumigatus</i> Mycobiology, 2010, 38, 229.	1.7	108
103	Differential Roles of the ChiB Chitinase in Autolysis and Cell Death of <i>Aspergillus nidulans</i> Eukaryotic Cell, 2009, 8, 738-746.	3.4	80
104	$G\hat{l}^2\hat{l}^3$ -mediated growth and developmental control in Aspergillus fumigatus. Current Genetics, 2009, 55, 631-641.	1.7	30
105	VelB/VeA/LaeA Complex Coordinates Light Signal with Fungal Development and Secondary Metabolism. Science, 2008, 320, 1504-1506.	12.6	843
106	Basic-Zipper-Type Transcription Factor FlbB Controls Asexual Development in <i>Aspergillus nidulans</i> . Eukaryotic Cell, 2008, 7, 38-48.	3.4	97
107	A Novel Regulator Couples Sporogenesis and Trehalose Biogenesis in Aspergillus nidulans. PLoS ONE, 2007, 2, e970.	2.5	215
108	Growth and Developmental Control in the Model and Pathogenic Aspergilli. Eukaryotic Cell, 2006, 5, 1577-1584.	3.4	80

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109	G-protein and cAMP-mediated signaling in aspergilli: A genomic perspective. Fungal Genetics and Biology, 2006, 43, 490-502.	2.1	131
110	FluG-Dependent Asexual Development in Aspergillus nidulans Occurs via Derepression. Genetics, 2006, 172, 1535-1544.	2.9	102
111	Upstream and Downstream Regulation of Asexual Development in Aspergillus fumigatus. Eukaryotic Cell, 2006, 5, 1585-1595.	3.4	134
112	The Phosducin-Like Protein PhnA Is Required for $G\hat{I}^2\hat{I}^3$ -Mediated Signaling for Vegetative Growth, Developmental Control, and Toxin Biosynthesis in Aspergillus nidulans. Eukaryotic Cell, 2006, 5, 400-410.	3.4	59
113	Heterotrimeric G protein signaling and RGSs in Aspergillus nidulans. Journal of Microbiology, 2006, 44, 145-54.	2.8	110
114	Genomic sequence of the pathogenic and allergenic filamentous fungus Aspergillus fumigatus. Nature, 2005, 438, 1151-1156.	27.8	1,272
115	The pkaB Gene Encoding the Secondary Protein Kinase A Catalytic Subunit Has a Synthetic Lethal Interaction with pkaA and Plays Overlapping and Opposite Roles in Aspergillus nidulans. Eukaryotic Cell, 2005, 4, 1465-1476.	3.4	57
116	The Heterotrimeric G-Protein $GanB(\hat{l}_{\pm})$ - $SfaD(\hat{l}^{2})$ - $GpgA(\hat{l}^{3})$ Is a Carbon Source Sensor Involved in Early cAMP-Dependent Germination in Aspergillus nidulans. Genetics, 2005, 171, 71-80.	2.9	118
117	Multiple Roles of a Heterotrimeric G-Protein \hat{I}^3 -Subunit in Governing Growth and Development of Aspergillus nidulans. Genetics, 2005, 171, 81-89.	2.9	64
118	Regulation of Secondary Metabolism in Filamentous Fungi. Annual Review of Phytopathology, 2005, 43, 437-458.	7.8	454
119	A putative G protein-coupled receptor negatively controls sexual development in Aspergillus nidulans. Molecular Microbiology, 2004, 51, 1333-1345.	2.5	97
120	Regulators of G-protein signalling in Aspergillus nidulans: RgsA downregulates stress response and stimulates asexual sporulation through attenuation of GanB (\widehat{Gl}) signalling. Molecular Microbiology, 2004, 53, 529-540.	2.5	114
121	The gprA and gprB genes encode putative G protein-coupled receptors required for self-fertilization in Aspergillus nidulans. Molecular Microbiology, 2004, 53, 1611-1623.	2.5	103
122	Double-joint PCR: a PCR-based molecular tool for gene manipulations in filamentous fungi. Fungal Genetics and Biology, 2004, 41, 973-981.	2.1	1,072
123	Suppressor Mutations Bypass the Requirement of <i>fluG</i> for Asexual Sporulation and Sterigmatocystin Production in <i>Aspergillus nidulans</i> Genetics, 2003, 165, 1083-1093.	2.9	63
124	The nsdD gene encodes a putative GATA-type transcription factor necessary for sexual development of Aspergillus nidulans. Molecular Microbiology, 2001, 41, 299-309.	2.5	200
125	Extragenic Suppressors of Loss-of-Function Mutations in the Aspergillus FlbA Regulator of G-Protein Signaling Domain Protein. Genetics, 1999, 151, 97-105.	2.9	43
126	Coordinate control of secondary metabolite production and asexual sporulation in Aspergillus nidulans. Current Opinion in Microbiology, 1998, 1, 674-677.	5.1	67

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127	Asexual Sporulation in <i>Aspergillus nidulans</i> i>Nicrobiology and Molecular Biology Reviews, 1998, 62, 35-54.	6.6	645
128	Dominant mutations affecting both sporulation and sterigmatocystin biosynthesis in Aspergillus nidulans. Current Genetics, 1997, 32, 218-224.	1.7	44
129	Conservation of structure and function of the aflatoxin regulatory geneaflR fromAspergillus nidulans and A. flavus. Current Genetics, 1996, 29, 549-555.	1.7	236
130	Conservation of structure and function of the aflatoxin regulatory gene aflR from Aspergillus nidulans and A. flavus. Current Genetics, 1996, 29, 549-555.	1.7	28
131	Regulation of <i>Aspergillus </i> Conidiation., 0,, 557-576.		23