

Ad Wiebenga

List of Publications by Year in descending order

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26
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4795
citing authors

#	ARTICLE	IF	CITATIONS
1	Macroalgae Derived Fungi Have High Abilities to Degrade Algal Polymers. <i>Microorganisms</i> , 2020, 8, 52.	1.6	24
2	Colonies of the fungus <i>Aspergillus niger</i> are highly differentiated to adapt to local carbon source variation. <i>Environmental Microbiology</i> , 2020, 22, 1154-1166.	1.8	15
3	A comparative genomics study of 23 <i>Aspergillus</i> species from section Flavi. <i>Nature Communications</i> , 2020, 11, 1106.	5.8	125
4	Transcriptome analysis of <i>Aspergillus niger</i> xlnR and xkiA mutants grown on corn Stover and soybean hulls reveals a highly complex regulatory network. <i>BMC Genomics</i> , 2019, 20, 853.	1.2	5
5	Genomic and Genetic Insights Into a Cosmopolitan Fungus, <i>Paecilomyces variotii</i> (Eurotiales). <i>Frontiers in Microbiology</i> , 2018, 9, 3058.	1.5	35
6	Investigation of inter- and intraspecies variation through genome sequencing of <i>Aspergillus</i> section Nigri. <i>Nature Genetics</i> , 2018, 50, 1688-1695.	9.4	160
7	Physiological background of the remarkably high Cd ²⁺ tolerance of the <i>Aspergillus fumigatus</i> Af293 strain. <i>Journal of Basic Microbiology</i> , 2018, 58, 957-967.	1.8	10
8	Secretion of small proteins is species-specific within <i>Aspergillus</i> sp. <i>Microbial Biotechnology</i> , 2017, 10, 323-329.	2.0	23
9	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
10	Genetic transformation of the white-rot fungus <i>Dichomitus squalens</i> using a new commercial protoplasting cocktail. <i>Journal of Microbiological Methods</i> , 2017, 143, 38-43.	0.7	12
11	High resolution visualization and exo-proteomics reveal the physiological role of XlnR and AraR in plant biomass colonization and degradation by <i>Aspergillus niger</i> . <i>Environmental Microbiology</i> , 2017, 19, 4587-4598.	1.8	6
12	Expansion of Signal Transduction Pathways in Fungi by Extensive Genome Duplication. <i>Current Biology</i> , 2016, 26, 1577-1584.	1.8	175
13	<i>Penicillium subrubescens</i> is a promising alternative for <i>Aspergillus niger</i> in enzymatic plant biomass saccharification. <i>New Biotechnology</i> , 2016, 33, 834-841.	2.4	27
14	Protease and lipase activities of fungal and bacterial strains derived from an artisanal raw ewe's milk cheese. <i>International Journal of Food Microbiology</i> , 2016, 237, 17-27.	2.1	72
15	Closely related fungi employ diverse enzymatic strategies to degrade plant biomass. <i>Biotechnology for Biofuels</i> , 2015, 8, 107.	6.2	111
16	Genetic Interaction of <i>Aspergillus nidulans</i> galR, xlnR and araR in Regulating D-Galactose and L-Arabinose Release and Catabolism Gene Expression. <i>PLoS ONE</i> , 2015, 10, e0143200.	1.1	41
17	<i>Aspergillus niger</i> RhaR, a regulator involved in l-rhamnose release and catabolism. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 5531-40.	1.7	56
18	Genomic and transcriptomic analysis of <i>Laccaria bicolor</i> CAZome reveals insights into polysaccharides remodelling during symbiosis establishment. <i>Fungal Genetics and Biology</i> , 2014, 72, 168-181.	0.9	81

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19	The genome of the white-rot fungus <i>Pycnoporus cinnabarinus</i> : a basidiomycete model with a versatile arsenal for lignocellulosic biomass breakdown. <i>BMC Genomics</i> , 2014, 15, 486.	1.2	91
20	Carbohydrate utilization and metabolism is highly differentiated in <i>Agaricus bisporus</i> . <i>BMC Genomics</i> , 2013, 14, 663.	1.2	35
21	Comparative genomics of the white-rot fungi, <i>Phanerochaete carnos</i> a and <i>P. chrysosporium</i> , to elucidate the genetic basis of the distinct wood types they colonize. <i>BMC Genomics</i> , 2012, 13, 444.	1.2	125
22	Comparative genomics of <i>Ceriporiopsis subvermispora</i> and <i>Phanerochaete chrysosporium</i> provide insight into selective ligninolysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 5458-5463.	3.3	259
23	The Paleozoic Origin of Enzymatic Lignin Decomposition Reconstructed from 31 Fungal Genomes. <i>Science</i> , 2012, 336, 1715-1719.	6.0	1,424
24	Carbohydrate-active enzymes from the zygomycete fungus <i>Rhizopus oryzae</i> : a highly specialized approach to carbohydrate degradation depicted at genome level. <i>BMC Genomics</i> , 2011, 12, 38.	1.2	105
25	The Plant Cell Wall "Decomposing Machinery Underlies the Functional Diversity of Forest Fungi. <i>Science</i> , 2011, 333, 762-765.	6.0	512
26	An enzyme cocktail for efficient protoplast formation in <i>Aspergillus niger</i> . <i>Journal of Microbiological Methods</i> , 2009, 76, 305-306.	0.7	59