Yanjie Liu

List of Publications by Year in descending order

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ΥΛΝΠΕΙΠΙ

#	Article	IF	CITATIONS
1	Distinct Roles of Velvet Complex in the Development, Stress Tolerance, and Secondary Metabolism in Pestalotiopsis microspora, a Taxol Producer. Genes, 2018, 9, 164.	2.4	22
2	The AMP-Activated Protein Kinase Homolog Snf1 Concerts Carbon Utilization, Conidia Production and the Biosynthesis of Secondary Metabolites in the Taxol-Producer Pestalotiopsis microspora. Genes, 2018, 9, 59.	2.4	19
3	Gα-cAMP/PKA pathway positively regulates pigmentation, chaetoglobosin A biosynthesis and sexual development in Chaetomium globosum. PLoS ONE, 2018, 13, e0195553.	2.5	14
4	Regulation of the Cα-cAMP/PKA signaling pathway in cellulose utilization of Chaetomium globosum. Microbial Cell Factories, 2018, 17, 160.	4.0	13
5	Amino Acid Sensor Kinase Gcn2 Is Required for Conidiation, Secondary Metabolism, and Cell Wall Integrity in the Taxol-Producer Pestalotiopsis microspora. Frontiers in Microbiology, 2017, 8, 1879.	3.5	9
6	A Gene Cluster for the Biosynthesis of Dibenzodioxocinons in the Endophyte Pestalotiopsis microspora, a Taxol Producer. Journal of Microbiology and Biotechnology, 2019, 29, 1570-1579.	2.1	7
7	Transcription Factors Pmr1 and Pmr2 Cooperatively Regulate Melanin Biosynthesis, Conidia Development and Secondary Metabolism in Pestalotiopsis microspora. Journal of Fungi (Basel,) Tj ETQq1 1 0.784	-3 194argBT	/Overlock 10
8	A Fungal Diterpene Synthase Is Responsible for Sterol Biosynthesis for Growth. Frontiers in Microbiology, 2020, 11, 1426.	3.5	3
9	Deletion of a Rare Fungal PKS CgPKS11 Promotes Chaetoglobosin A Biosynthesis, Yet Defers the Growth and Development of Chaetomium globosum. Journal of Fungi (Basel, Switzerland), 2021, 7, 750.	3.5	3
10	An effective method to produce 7-epitaxol from taxol in HCO3–. Bioorganic and Medicinal Chemistry Letters, 2020, 30, 127285.	2.2	2
11	Acquiring novel chemicals by overexpression of a transcription factor DibT in the dibenzodioxocinone biosynthetic cluster in Pestalotiopsis microspora. Microbiological Research, 2022, 257, 126977.	5.3	1