

Bang Feng

List of Publications by Year in descending order

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Version: 2024-02-01

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papers

878
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516710

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docs citations

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times ranked

695
citing authors

#	ARTICLE	IF	CITATIONS
1	Evolutionary innovations through gain and loss of genes in the ectomycorrhizal Boletales. <i>New Phytologist</i> , 2022, 233, 1383-1400.	7.3	19
2	Elevation Matters More than Season in Shaping the Heterogeneity of Soil and Root Associated Ectomycorrhizal Fungal Community. <i>Microbiology Spectrum</i> , 2022, 10, e0195021.	3.0	15
3	Life Cycle and Phylogeography of True Truffles. <i>Genes</i> , 2022, 13, 145.	2.4	2
4	<i>Cantharellus albus</i> , a striking new species from Southwest China. <i>Phytotaxa</i> , 2020, 470, 133-144.	0.3	6
5	African origin and global distribution patterns: Evidence inferred from phylogenetic and biogeographical analyses of ectomycorrhizal fungal genus <i>Strobilomyces</i> . <i>Journal of Biogeography</i> , 2018, 45, 201-212.	3.0	28
6	Studies on diversity of higher fungi in Yunnan, southwestern China: A review. <i>Plant Diversity</i> , 2018, 40, 165-171.	3.7	26
7	Using mating-type loci to improve taxonomy of the <i>Tuber indicum</i> complex, and discovery of a new species, <i>T. longispinosum</i> . <i>PLoS ONE</i> , 2018, 13, e0193745.	2.5	13
8	Ecological and physical barriers shape genetic structure of the Alpine porcini (<i>Boletus reticulocephus</i>). <i>Mycorrhiza</i> , 2017, 27, 261-272.	2.8	10
9	Genetic diversity and breeding history of Winter Mushroom (<i>Flammulina velutipes</i>) in China uncovered by genomic SSR markers. <i>Gene</i> , 2016, 591, 227-235.	2.2	34
10	Multilocus phylogenetic analyses reveal unexpected abundant diversity and significant disjunct distribution pattern of the Hedgehog Mushrooms (<i>Hydnum</i> L.). <i>Scientific Reports</i> , 2016, 6, 25586.	3.3	29
11	Drainage isolation and climate change-driven population expansion shape the genetic structures of <i>Tuber indicum</i> complex in the Hengduan Mountains region. <i>Scientific Reports</i> , 2016, 6, 21811.	3.3	29
12	Four new genera of the fungal family Boletaceae. <i>Fungal Diversity</i> , 2016, 81, 1-24.	12.3	61
13	Porcini mushrooms (<i>Boletus</i> sect. <i>Boletus</i>) from China. <i>Fungal Diversity</i> , 2016, 81, 189-212.	12.3	36
14	The genus <i>Imleria</i> (Boletaceae) in East Asia. <i>Phytotaxa</i> , 2014, 191, 81.	0.3	17
15	Molecular phylogenetic analyses redefine seven major clades and reveal 22 new generic clades in the fungal family Boletaceae. <i>Fungal Diversity</i> , 2014, 69, 93-115.	12.3	183
16	Molecular phylogeny of <i>Caloboletus</i> (Boletaceae) and a new species in East Asia. <i>Mycological Progress</i> , 2014, 13, 1127.	1.4	25
17	The taxonomic foundation, species circumscription and continental endemisms of <i>Singerocybe</i> : evidence from morphological and molecular data. <i>Mycologia</i> , 2014, 106, 1015-1026.	1.9	16
18	Molecular phylogeny and taxonomy of the genus <i>Veloporphyrellus</i> . <i>Mycologia</i> , 2014, 106, 291-306.	1.9	35

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19	Borofutus, a new genus of Boletaceae from tropical Asia: phylogeny, morphology and taxonomy. <i>Fungal Diversity</i> , 2013, 58, 215-226.	12.3	66
20	New species and distinctive geographical divergences of the genus <i>Sparassis</i> (Basidiomycota): evidence from morphological and molecular data. <i>Mycological Progress</i> , 2013, 12, 445-454.	1.4	26
21	DNA Sequence Analyses Reveal Abundant Diversity, Endemism and Evidence for Asian Origin of the Porcini Mushrooms. <i>PLoS ONE</i> , 2012, 7, e37567.	2.5	79
22	<i>Zangia</i> , a new genus of Boletaceae supported by molecular and morphological evidence. <i>Fungal Diversity</i> , 2011, 49, 125-143.	12.3	86
23	<i>Ovipoculum album</i> , a new anamorph with gelatinous cupulate bulbiferous conidiomata from China and with affinities to the Auriculariales (Basidiomycota). <i>Fungal Diversity</i> , 2010, 43, 55-65.	12.3	16
24	Genetic Diversity of Dahongjun, the Commercially Important "Big Red Mushroom" from Southern China. <i>PLoS ONE</i> , 2010, 5, e10684.	2.5	21