Wei Gao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5056955/publications.pdf

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19	421	11	19
papers	citations	h-index	g-index
19	19	19	487 citing authors
all docs	docs citations	times ranked	

#	Article	IF	Citations
1	Isolation and identification of pigments from oyster mushrooms with black, yellow and pink caps. Food Chemistry, 2022, 372, 131171.	8.2	15
2	Characterization of the wheat VQ protein family and expression of candidate genes associated with seed dormancy and germination. BMC Plant Biology, 2022, 22, 119.	3.6	7
3	Fine Mapping and Functional Analysis of the Gene <i>PcTYR</i> , Involved in Control of Cap Color of Pleurotus cornucopiae. Applied and Environmental Microbiology, 2022, 88, e0217321.	3.1	6
4	Identification of the Wheat (TriticumÂaestivum) IQD Gene Family and an Expression Analysis of Candidate Genes Associated with Seed Dormancy and Germination. International Journal of Molecular Sciences, 2022, 23, 4093.	4.1	3
5	Genetic Linkage and Physical Mapping for an Oyster Mushroom (<i>Pleurotus cornucopiae</i>) and Quantitative Trait Locus Analysis for Cap Color. Applied and Environmental Microbiology, 2021, 87, e0095321.	3.1	8
6	Identification and expression analysis of candidate genes related to seed dormancy and germination in the wheat GATA family. Plant Physiology and Biochemistry, 2021, 169, 343-359.	5.8	6
7	Identification of the wheat C3H gene family and expression analysis of candidates associated with seed dormancy and germination. Plant Physiology and Biochemistry, 2020, 156, 524-537.	5.8	18
8	Identification and Validation of New Stable QTLs for Grain Weight and Size by Multiple Mapping Models in Common Wheat. Frontiers in Genetics, 2020, $11,584859$.	2.3	8
9	Expression patterns of two pal genes of Pleurotus ostreatus across developmental stages and under heat stress. BMC Microbiology, 2019, 19, 231.	3.3	30
10	A genetic linkage map of Pleurotus tuoliensis integrated with physical mapping of the de novo sequenced genome and the mating type loci. BMC Genomics, 2018, 19, 18.	2.8	34
11	Genome-Wide Characterization and Expression Analyses of Pleurotus ostreatus MYB Transcription Factors during Developmental Stages and under Heat Stress Based on de novo Sequenced Genome. International Journal of Molecular Sciences, 2018, 19, 2052.	4.1	36
12	Cloning, purification and characterization of trehalose-6-phosphate synthase from Pleurotus tuoliensis. Peerl, 2018, 6, e5230.	2.0	12
13	Developments in breeding of Agaricus bisporus var. bisporus: progress made and technical and legal hurdles to take. Applied Microbiology and Biotechnology, 2017, 101, 1819-1829.	3.6	49
14	Differential Expression Patterns of Pleurotus ostreatus Catalase Genes during Developmental Stages and under Heat Stress. Genes, 2017, 8, 335.	2.4	36
15	A detailed analysis of the recombination landscape of the button mushroom Agaricus bisporus var. bisporus. Fungal Genetics and Biology, 2016, 93, 35-45.	2.1	75
16	The famous cultivated mushroom Bailinggu is a separate species of the Pleurotus eryngii species complex. Scientific Reports, 2016, 6, 33066.	3.3	21
17	Multi-trait QTL analysis for agronomic and quality characters of Agaricus bisporus (button) Tj ETQq1 1 0.78431	4 rgBT /Ov	erlock 10 Tf 5

Quantitative trait locus mapping for bruising sensitivity and cap color of Agaricus bisporus (button) Tj ETQq $0\,0\,0\,$ rgBT /Overlock $10\,$ Tf $5\,$ Cuantitative trait locus mapping for bruising sensitivity and cap color of Agaricus bisporus (button) Tj ETQq $0\,0\,0\,$ rgBT /Overlock $10\,$ Tf $5\,$ Cuantitative trait locus mapping for bruising sensitivity and cap color of Agaricus bisporus (button) Tj ETQq $0\,0\,0\,$ rgBT /Overlock $10\,$ Tf $5\,$ Cuantitative trait locus mapping for bruising sensitivity and cap color of Agaricus bisporus (button) Tj ETQq $0\,0\,0\,$ rgBT /Overlock $10\,$ Tf $10\,$ Tf

18

#	Article	IF	CITATIONS
19	Genetic Variation and Combining Ability Analysis of Bruising Sensitivity in Agaricus bisporus. PLoS ONE, 2013, 8, e76826.	2.5	17