Nai Tran-Dinh

List of Publications by Year
in descending order

17	639	759233 12	888059 17
papers	citations	h-index	g-index

17	17	17	1061
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The Coal Seam Microbiome (CSMB) reference set, a lingua franca for the microbial coal-to-methane community. International Journal of Coal Geology, 2018, 186, 41-50.	5.0	46
2	Gamarada debralockiae gen. nov. sp. nov.â€"the genome of the most widespread Australian ericoid mycorrhizal fungus. Mycorrhiza, 2018, 28, 379-389.	2.8	9
3	First evidence of Pezoloma ericae in Australia: using the Biomes of Australia Soil Environments (BASE) to explore the Australian phylogeography of known ericoid mycorrhizal and root-associated fungi. Mycorrhiza, 2017, 27, 587-594.	2.8	15
4	Genomic insights into the carbohydrate catabolism of Cairneyella variabilis gen. nov. sp. nov., the first reports from a genome of an ericoid mycorrhizal fungus from the southern hemisphere. Mycorrhiza, 2016, 26, 345-352.	2.8	18
5	Fungal identification using a Bayesian classifier and the Warcup training set of internal transcribed spacer sequences. Mycologia, 2016, 108, 1-5.	1.9	178
6	Genomic and chemical insights into biosurfactant production by the mangrove-derived strain Bacillus safensis CCMA-560. Applied Microbiology and Biotechnology, 2015, 99, 3155-3167.	3.6	30
7	Selection of non-toxigenic strains ofAspergillus flavusfor biocontrol of aflatoxins in maize in Thailand. Biocontrol Science and Technology, 2014, 24, 652-661.	1.3	21
8	Draft Genome Sequence of Clostridium sporogenes PA 3679, the Common Nontoxigenic Surrogate for Proteolytic Clostridium botulinum. Journal of Bacteriology, 2012, 194, 1631-1632.	2.2	25
9	Clostridium sporogenes PA 3679 and Its Uses in the Derivation of Thermal Processing Schedules for Low-Acid Shelf-Stable Foods and as a Research Model for Proteolytic Clostridium botulinum. Journal of Food Protection, 2012, 75, 779-792.	1.7	35
10	Survey of Vietnamese Peanuts, Corn and Soil for the Presence of Aspergillus flavus and Aspergillus parasiticus. Mycopathologia, 2009, 168, 257-268.	3.1	38
11	Utility of Microsatellite Markers and Amplified Fragment Length Polymorphism in the Study of Potentially Ochratoxigenic Black Aspergilli. Current Microbiology, 2008, 57, 348-355.	2.2	13
12	Isolation and characterization of polymorphic microsatellite markers for Alternaria alternata. Molecular Ecology Notes, 2006, 6, 405-407.	1.7	13
13	Isolation and characterization of six polymorphic microsatellite loci in Aspergillus niger. Molecular Ecology Notes, 2005, 5, 375-377.	1.7	9
14	The Development of Genetic Markers from Fungal Genome Initiatives. Applied Mycology and Biotechnology, 2004, 4, 1-27.	0.3	8
15	Xanthones from a microfungus of the genus Xylaria. Phytochemistry, 2004, 65, 2373-2378.	2.9	72
16	Complete Genomic Sequence of SfV, a Serotype-Converting Temperate Bacteriophage of Shigellaflexneri. Journal of Bacteriology, 2002, 184, 1974-1987.	2.2	83
17	Characterization of microsatellite loci in the aflatoxigenic fungi Aspergillus flavus and Aspergillus parasiticus. Molecular Ecology, 2000, 9, 2170-2172.	3.9	26