BalÃ;zs Vajna

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5078961/publications.pdf Version: 2024-02-01



ΒΛΙ Ã:75 ΜΛΙΝΙΛ

#	Article	IF	CITATIONS
1	Distribution of bacterial single cell parameters and their estimation from turbidity detection times. Food Microbiology, 2022, 104, 103972.	4.2	0
2	Evaluating the combined effect of biochar and PGPR inoculants on the bacterial community in acidic sandy soil. Applied Soil Ecology, 2021, 160, 103856.	4.3	12
3	Aboveâ€ground parts of white grapevine <scp><i>Vitis vinifera</i></scp> cv. Furmint share core members of the fungal microbiome. Environmental Microbiology Reports, 2021, 13, 509-520.	2.4	23
4	Succession and potential role of bacterial communities during <i>Pleurotus ostreatus</i> production. FEMS Microbiology Ecology, 2021, 97, .	2.7	6
5	Grazing pressure-induced shift in planktonic bacterial communities with the dominance of acIII-A1 actinobacterial lineage in soda pans. Scientific Reports, 2020, 10, 19871.	3.3	12
6	Whole genome sequence analysis of Cupriavidus campinensis S14E4C, a heavy metal resistant bacterium. Molecular Biology Reports, 2020, 47, 3973-3985.	2.3	16
7	Relationships Between Chemical Defenses of Common Toad (Bufo bufo) Tadpoles and Bacterial Community Structure of their Natural Aquatic Habitat. Journal of Chemical Ecology, 2020, 46, 534-543.	1.8	3
8	Critical point analysis and biocide treatment in a microbiologically contaminated water purification system of a power plant. SN Applied Sciences, 2019, 1, 1.	2.9	9
9	Monitoring of soil microbial inoculants and their impact on maize (Zea mays L.) rhizosphere using T-RFLP molecular fingerprint method. Applied Soil Ecology, 2019, 138, 233-244.	4.3	15
10	Effect of Long-Term Cropping Systems on the Diversity of the Soil Bacterial Communities. Agronomy, 2019, 9, 878.	3.0	16
11	Fertoeibacter niger gen. nov., sp. nov. a novel alkaliphilic bacterium of the family Rhodobacteraceae. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	1.7	8
12	Bacterial–fungal interactions: ecology, mechanisms and challenges. FEMS Microbiology Reviews, 2018, 42, 335-352.	8.6	468
13	Differences in planktonic microbial communities associated with three types of macrophyte stands in a shallow lake. FEMS Microbiology Ecology, 2018, 94, .	2.7	16
14	Texture and type of polymer fiber carrier determine bacterial colonization and biofilm properties in wastewater treatment. Chemical Engineering Journal, 2015, 264, 824-834.	12.7	42
15	Characterisation of the large-scale production process of oyster mushroom (Pleurotus ostreatus) with the analysis of succession and spatial heterogeneity of lignocellulolytic enzyme activities. Fungal Biology, 2015, 119, 1354-1363.	2.5	20
16	One-year monitoring of meta-cleavage dioxygenase gene expression and microbial community dynamics reveals the relevance of subfamily I.2.C extradiol dioxygenases in hypoxic, BTEX-contaminated groundwater. Systematic and Applied Microbiology, 2013, 36, 339-350.	2.8	33
17	Remarkable impact of PAHs and TPHs on the richness and diversity of bacterial species in surface soils exposed to long-term hydrocarbon pollution. World Journal of Microbiology and Biotechnology, 2013, 29, 1989-2002.	3.6	54
18	Thermus composti sp. nov., isolated from oyster mushroom compost. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 1486-1490.	1.7	27

BalÃizs Vajna

#	Article	IF	CITATIONS
19	An Improved Sequence-aided T-RFLP Analysis of Bacterial Succession During Oyster Mushroom Substrate Preparation. Microbial Ecology, 2012, 64, 702-713.	2.8	24
20	Ferrate treatment for inactivation of bacterial community in municipal secondary effluent. Bioresource Technology, 2012, 107, 116-121.	9.6	41
21	Diversity and seasonal dynamics of the photoautotrophic picoplankton in Lake Balaton (Hungary). Aquatic Microbial Ecology, 2011, 63, 273-287.	1.8	27
22	Microbial community structure changes during oyster mushroom substrate preparation. Applied Microbiology and Biotechnology, 2010, 86, 367-375.	3.6	26
23	Diversity and activity of cultivable aerobic planktonic bacteria of a saline Lake located in Sovata, Romania. Folia Microbiologica, 2010, 55, 461-466.	2.3	15
24	DGGE and T-RFLP Analysis of Bacterial Succession during Mushroom Compost Production and Sequence-aided T-RFLP Profile of Mature Compost. Microbial Ecology, 2009, 57, 522-533.	2.8	90
25	Phenotypic characterization and molecular taxonomic studies on Bacillus and related isolates from Phragmites australis periphyton. Aquatic Botany, 2007, 86, 243-252.	1.6	15
26	Genetic Variability ofgagandenvRegions of HIV Type 1 Strains Circulating in Slovenia. AIDS Research and Human Retroviruses, 2006, 22, 109-113.	1.1	5
27	Increased incidence of genetic human prion disease in Hungary. Neurology, 2005, 65, 1666-1669.	1.1	12