

Nomeda Kuisiene

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

226
citations

1163117

8
h-index

1058476

14
g-index

21
all docs

21
docs citations

21
times ranked

334
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic Characterization of Multidrug-Resistant <i>E. coli</i> Isolates from Bloodstream Infections in Lithuania. <i>Microorganisms</i> , 2022, 10, 449.	3.6	5
2	Diversity of Polyketide Synthases and Nonribosomal Peptide Synthetases Revealed Through Metagenomic Analysis of a Deep Oligotrophic Cave. <i>Microbial Ecology</i> , 2021, 81, 110-121.	2.8	8
3	Multidrug-Resistant <i>Acinetobacter baumannii</i> Genetic Characterization and Spread in Lithuania in 2014, 2016, and 2018. <i>Life</i> , 2021, 11, 151.	2.4	2
4	Bioamperometric Systems with Fructose Dehydrogenase From <i>Gluconobacter japonicus</i> for D-Tagatose Monitoring. <i>Electroanalysis</i> , 2021, 33, 1393-1397.	2.9	1
5	Collagen-like sequences encoded by extremophilic and extremotolerant bacteria. <i>Genomics</i> , 2020, 112, 2271-2281.	2.9	9
6	Genome Mining and Characterization of Biosynthetic Gene Clusters in Two Cave Strains of <i>Paenibacillus</i> sp.. <i>Frontiers in Microbiology</i> , 2020, 11, 612483.	3.5	9
7	First genotypic characterization of toxigenic <i>Clostridioides difficile</i> in Lithuanian hospitals reveals the prevalence of the hypervirulent ribotype 027/ST1. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2019, 38, 1953-1959.	2.9	1
8	Screening and Transcriptional Analysis of Polyketide Synthases and Non-ribosomal Peptide Synthetases in Bacterial Strains From Krubera-Voronja Cave. <i>Frontiers in Microbiology</i> , 2019, 10, 2149.	3.5	9
9	Evaluation of the Inter- and Intra-hospital Spread of Multidrug Resistant Gram-Negative Bacteria in Lithuanian Hospitals. <i>Microbial Drug Resistance</i> , 2019, 25, 326-335.	2.0	3
10	Draft genome sequence of pectic polysaccharide-degrading moderate thermophilic bacterium <i>Geobacillus thermodenitrificans</i> DSM 101594. <i>Brazilian Journal of Microbiology</i> , 2017, 48, 7-8.	2.0	6
11	The cave microbiome as a source for drug discovery: Reality or pipe dream?. <i>Biochemical Pharmacology</i> , 2017, 134, 18-34.	4.4	69
12	Combining prebiotics with probiotic bacteria can enhance bacterial growth and secretion of bacteriocins. <i>International Journal of Biological Macromolecules</i> , 2016, 89, 669-676.	7.5	32
13	Characterization of antimicrobial activity of culturable bacteria isolated from Krubera-Voronja Cave. <i>International Journal of Speleology</i> , 2016, 45, 275-287.	1.0	12
14	Down into the Earth: microbial diversity of the deepest cave of the world. <i>Biologia (Poland)</i> , 2015, 70, 989-1002.	1.5	12
15	Characterization of a Novel Thermostable Oligopeptidase from <i>Geobacillus thermoleovorans</i> DSM 15325. <i>Journal of Microbiology and Biotechnology</i> , 2015, 25, 1070-1083.	2.1	4
16	Identification of proteins involved in starch and polygalacturonic acid degradation using LC/MS. <i>Open Life Sciences</i> , 2014, 9, 708-716.	1.4	1
17	Characterization and evaluation of tandem repeats for the identification of <i>Geobacillus</i> . <i>Open Life Sciences</i> , 2013, 8, 549-560.	1.4	1
18	Gene expression and activity analysis of the first thermophilic U32 peptidase. <i>Open Life Sciences</i> , 2012, 7, 587-595.	1.4	1

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19	Phylogenetic, Inter, and Intraspecific Sequence Analysis of spo0A Gene of the Genus Geobacillus. Current Microbiology, 2009, 58, 547-553.	2.2	15
20	Unusual intragenomic and interspecific variability of Geobacillus 16S-23S rRNA internal transcribed spacers. Open Life Sciences, 2008, 3, 396-406.	1.4	0
21	Bacillus butanolivorans sp. nov., a species with industrial application for the remediation of n-butanol. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 505-509.	1.7	26