

Nomeda Kuisiene

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

226
citations

1163117

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334
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | The cave microbiome as a source for drug discovery: Reality or pipe dream?. <i>Biochemical Pharmacology</i> , 2017, 134, 18-34. | 4.4 | 69 |
| 2 | 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 |
| 3 | <i>Bacillus butanolivorans</i> sp. nov., a species with industrial application for the remediation of n-butanol. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 505-509. | 1.7 | 26 |
| 4 | Phylogenetic, Inter, and Intraspecific Sequence Analysis of spo0A Gene of the Genus <i>Geobacillus</i> . <i>Current Microbiology</i> , 2009, 58, 547-553. | 2.2 | 15 |
| 5 | Down into the Earth: microbial diversity of the deepest cave of the world. <i>Biologia (Poland)</i> , 2015, 70, 989-1002. | 1.5 | 12 |
| 6 | 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 |
| 7 | 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 |
| 8 | Collagen-like sequences encoded by extremophilic and extremotolerant bacteria. <i>Genomics</i> , 2020, 112, 2271-2281. | 2.9 | 9 |
| 9 | 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 |
| 10 | 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 |
| 11 | 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 |
| 12 | Genetic Characterization of Multidrug-Resistant <i>E. coli</i> Isolates from Bloodstream Infections in Lithuania. <i>Microorganisms</i> , 2022, 10, 449. | 3.6 | 5 |
| 13 | 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 |
| 14 | Evaluation of the Inter- and Intrahospital Spread of Multidrug Resistant Gram-Negative Bacteria in Lithuanian Hospitals. <i>Microbial Drug Resistance</i> , 2019, 25, 326-335. | 2.0 | 3 |
| 15 | 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 |
| 16 | Gene expression and activity analysis of the first thermophilic U32 peptidase. <i>Open Life Sciences</i> , 2012, 7, 587-595. | 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 | 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 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | 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 |
| 20 | Bioamperometric Systems with Fructose Dehydrogenase From <i>Gluconobacter japonicus</i> for D-Galactose Monitoring. <i>Electroanalysis</i> , 2021, 33, 1393-1397. | 2.9 | 1 |
| 21 | Unusual intragenomic and interspecific variability of <i>Geobacillus</i> 16S-23S rRNA internal transcribed spacers. <i>Open Life Sciences</i> , 2008, 3, 396-406. | 1.4 | 0 |