

Sergey Shilov

List of Publications by Year in descending order

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Version: 2024-02-01

9
papers

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2257263

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2053342

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#	ARTICLE	IF	CITATIONS
1	Complete Genome Sequences of Gram-Positive Opportunistic Pathogens Isolated from Hospitals in Almaty, Kazakhstan. <i>Microbiology Resource Announcements</i> , 2022, , e0009322.	0.3	2
2	Comparison of Transcriptional Responses and Metabolic Alterations in Three Multidrug-Resistant Model Microorganisms, <i>Staphylococcus aureus</i> ATCC BAA-39, <i>Escherichia coli</i> ATCC BAA-196, and <i>Acinetobacter baumannii</i> ATCC BAA-1790, on Exposure to Iodine-Containing Nano-micelle Drug FS-1. <i>MSystems</i> , 2021, 6, .	1.7	2
3	Complete Genome Sequences of Gram-Negative Opportunistic Pathogens Isolated in Hospitals in Almaty, Kazakhstan. <i>Microbiology Resource Announcements</i> , 2021, 10, e0097421.	0.3	5
4	Transcriptomics and methylomics study on the effect of iodine-containing drug FS-1 on ATCC BAA-196. <i>Future Microbiology</i> , 2021, 16, 1063-1085.	1.0	0
5	The Effect of Iodine-Containing Nano-Micelles, FS-1, on Antibiotic Resistance, Gene Expression and Epigenetic Modifications in the Genome of Multidrug Resistant MRSA Strain <i>Staphylococcus aureus</i> ATCC BAA-39. <i>Frontiers in Microbiology</i> , 2020, 11, 581660.	1.5	9
6	Complete Genome Sequence of Collection Strain <i>Acinetobacter baumannii</i> ATCC BAA-1790, Used as a Model To Study the Antibiotic Resistance Reversion Induced by Iodine-Containing Complexes. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	6
7	GENE EXPRESSION PROFILING OF MULTI-DRUG RESISTANT E. COLI AFTER EXPOSURE BY NANOMOLECULAR IODINE-CONTAINING COMPLEX. <i>News of the National Academy of Sciences of the Republic of Kazakhstan Series of Biological and Medical</i> , 2020, 4, 10-18.	0.0	4
8	Assembly of Complete Genome Sequences of Negative-Control and Experimental Strain Variants of <i>Staphylococcus aureus</i> ATCC BAA-39 Selected under the Effect of the Drug FS-1, Which Induces Antibiotic Resistance Reversion. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	7
9	Complete Genome Sequence of a Multidrug-Resistant Strain, <i>Escherichia coli</i> ATCC BAA-196, as a Model for Studying Induced Antibiotic Resistance Reversion. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	5