

Ilya S Korotetskiy

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

199
citations

1477746

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1125271

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all docs

29
docs citations

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times ranked

247
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Newcastle disease outbreaks in Kazakhstan and Kyrgyzstan during 1998, 2000, 2001, 2003, 2004, and 2005 were caused by viruses of the genotypes VIIb and VIId. <i>Virus Genes</i> , 2009, 39, 94-101. | 0.7 | 47 |
| 2 | Genomic Insight into Mechanisms of Reversion of Antibiotic Resistance in Multidrug Resistant <i>Mycobacterium tuberculosis</i> Induced by a Nanomolecular Iodine-Containing Complex FS-1. <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 151. | 1.8 | 26 |
| 3 | Role of the horizontal gene exchange in evolution of pathogenic <i>Mycobacteria</i> . <i>BMC Evolutionary Biology</i> , 2015, 15, S2. | 3.2 | 23 |
| 4 | Molecular Characterization of Virulent Newcastle Disease Virus Isolates from Chickens during the 1998 NDV Outbreak in Kazakhstan. <i>Virus Genes</i> , 2005, 31, 13-20. | 0.7 | 17 |
| 5 | Characterization of pigeon paramyxoviruses (Newcastle disease virus) isolated in Kazakhstan in 2005. <i>Virologica Sinica</i> , 2012, 27, 93-99. | 1.2 | 11 |
| 6 | 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 |
| 7 | 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 |
| 8 | 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 |
| 9 | ISOLATION AND CHARACTERIZATION ISOLATES OF NOSOCOMIAL INFECTIONS. <i>The Bulletin</i> , 2019, 5, 199-209. | 0.0 | 6 |
| 10 | Complete Genome Sequence of Multidrug-Resistant Clinical Isolate <i>Mycobacterium tuberculosis</i> 187.0, Used To Study the Effect of Drug Susceptibility Reversion by the New Medicinal Drug FS-1. <i>Genome Announcements</i> , 2015, 3, . | 0.8 | 5 |
| 11 | Clade-Specific Distribution of Antibiotic Resistance Mutations in the Population of <i>Mycobacterium tuberculosis</i> - Prospects for Drug Resistance Reversion. , 2018, , . | | 5 |
| 12 | 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 |
| 13 | 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 |
| 14 | 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 |
| 15 | Constraints of Drug Resistance in <i>Mycobacterium tuberculosis</i> - Prospects for Pharmacological Reversion of Susceptibility to Antibiotics. <i>The Open Conference Proceedings Journal</i> , 2017, 8, 33-43. | 0.6 | 3 |
| 16 | The Effect of Antibiotic-Resistant and Sensitive <i>Escherichia coli</i> on the Production of Pro-Inflammatory Cytokine Response by Human Peripheral Blood Mononuclear Cells. <i>Journal of Clinical & Cellular Immunology</i> , 2017, 8, . | 1.5 | 3 |
| 17 | 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 |
| 18 | Coding-Complete Genome Sequence of Swine Influenza Virus Isolate A/Swine/Karaganda/04/2020 (H1N1) from Kazakhstan. <i>Microbiology Resource Announcements</i> , 2021, 10, e0078621. | 0.3 | 2 |

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|----|--|-----|-----------|
| 19 | Complete Genome Sequences of Gram-Positive Opportunistic Pathogens Isolated from Hospitals in Almaty, Kazakhstan. Microbiology Resource Announcements, 2022, , e0009322. | 0.3 | 2 |
| 20 | Genetic Changes in Influenza a Virus Genes Responsible for Formation of Drug Resistance Phenotype. Journal of Human Virology & Retrovirology, 2016, 3, . | 0.1 | 1 |
| 21 | Impact Genetic Characterization of H5N1 Avian Influenza Virus Isolated from Dead Mute Swan (Cygnus Tj ETQq1 1 0.784314 rgBT /C | 0.6 | 0 |
| 22 | STUDY ON ANTIVIRAL ACTIVITY OF COORDINATION COMPOUND BASED ON MOLECULAR IODINE AGAINST INFLUENZA A VIRUS. , 2017, , . | | 0 |
| 23 | NEW SEMIORGANIC IODINE COMPLEX, ITS STRUCTURE AND BIOLOGICAL ACTIVITY. Series Chemistry and Technology, 2019, 1, 70-77. | 0.1 | 0 |
| 24 | Transcriptomics and methylomics study on the effect of iodine-containing drug FS-1 on ATCC BAA-196. Future Microbiology, 2021, 16, 1063-1085. | 1.0 | 0 |