Pavel Tarlykov

List of Publications by Year in descending order

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Version: 2024-02-01

933447 839539 24 326 10 citations h-index papers

g-index 25 25 25 458 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	Damage-Induced Mutation Clustering in Gram-Positive Bacteria: Preliminary Data. Symmetry, 2022, 14, 1431.	2.2	O
2	Analysis of Bacteroides fragilis Clinical Strains Isolated in Kazakhstan. Microbiology Resource Announcements, $2021,10,.$	0.6	O
3	Determinants of resistance in Bacteroides fragilis strain BFR_KZ01 isolated from a patient with peritonitis in Kazakhstan. Journal of Global Antimicrobial Resistance, 2021, 25, 1-4.	2.2	7
4	Mitochondrial DNA analysis of ancient sheep from Kazakhstan: evidence for early sheep introduction. Heliyon, 2021, 7, e08011.	3.2	2
5	Draft Genome Sequence of a Bacteroides fragilis Strain Isolated from Peritoneal Fluid of a Patient from Kazakhstan. Microbiology Resource Announcements, 2020, 9, .	0.6	0
6	The medieval Mongolian roots of Y-chromosomal lineages from South Kazakhstan. BMC Genetics, 2020, 21, 87.	2.7	15
7	Development and validation of hybrid Brillouin-Raman spectroscopy for non-contact assessment of mechano-chemical properties of urine proteins as biomarkers of kidney diseases. BMC Nephrology, 2020, 21, 229.	1.8	13
8	Draft Genome Sequence of an Extensively Drug-Resistant Mycobacterium tuberculosis Clinical Isolate, 3485_MTB, from Nur-Sultan, Kazakhstan. Microbiology Resource Announcements, 2020, 9, .	0.6	3
9	Genomic analysis of Latin American-Mediterranean family of Mycobacterium tuberculosis clinical strains from Kazakhstan. Memorias Do Instituto Oswaldo Cruz, 2020, 115, e200215.	1.6	5
10	Genomic characterization of MDR/XDR-TB in Kazakhstan by a combination of high-throughput methods predominantly shows the ongoing transmission of L2/Beijing 94–32 central Asian/Russian clusters. BMC Infectious Diseases, 2019, 19, 553.	2.9	10
11	Genetic Characterization of Kazakh Native Sheep Breeds Using Mitochondrial DNA. OnLine Journal of Biological Sciences, 2018, 18, 341-348.	0.4	5
12	Topokaryotyping demonstrates single cell variability and stress dependent variations in nuclear envelope associated domains. Nucleic Acids Research, 2018, 46, e135-e135.	14.5	3
13	Epidemiology of Brucellosis and Genetic Diversity of Brucella abortus in Kazakhstan. PLoS ONE, 2016, 11, e0167496.	2.5	31
14	Genetic risk factors for restenosis after percutaneous coronary intervention in Kazakh population. Human Genomics, 2016, 10, 15.	2.9	8
15	ZNF555 protein binds to transcriptional activator site of 4qA allele and <i>ANT1</i> ii>implication in Facioscapulohumeral dystrophy. Nucleic Acids Research, 2015, 43, 8227-8242.	14.5	15
16	Genetic diversity of Brucella abortus and Brucella melitensis in Kazakhstan using MLVA-16. Infection, Genetics and Evolution, 2015, 34, 173-180.	2.3	36
17	ABO Blood Group Genotyping by Real-time PCR in Kazakh Population. Central Asian Journal of Global Health, 2014, 3, 177.	0.6	1
18	PUB-NChIPâ€""in vivo biotinylation―approach to study chromatin in proximity to a protein of interest. Genome Research, 2013, 23, 331-340.	5 . 5	27

#	Article	IF	CITATION
19	Mitochondrial and Y-chromosomal profile of the Kazakh population from East Kazakhstan. Croatian Medical Journal, 2013, 54, 17-24.	0.7	29
20	Draft Genome Sequence of the Live Vaccine Strain Brucella abortus 82. Genome Announcements, 2013, $1, \ldots$	0.8	7
21	Draft Genome Sequence of Rhodococcus erythropolis DN1, a Crude Oil Biodegrader. Genome Announcements, 2013, 1, .	0.8	12
22	Proteomic Analysis of <i>Sulfolobus solfataricus</i> during <i>Sulfolobus</i> Turreted Icosahedral Virus Infection. Journal of Proteome Research, 2012, 11, 1420-1432.	3.7	26
23	Something Old, Something New, Something Borrowed; How the Thermoacidophilic Archaeon Sulfolobus solfataricus Responds to Oxidative Stress. PLoS ONE, 2009, 4, e6964.	2.5	70
24	Crystal and molecular structure of tigogenin. Russian Journal of Applied Chemistry, 2006, 79, 1371-1373.	0.5	1