Pavel Tarlykov

List of Publications by Year
in descending order

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24	326	933447	839539
papers	citations	10	18
		h-index	g-index

25	25	25	458
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Something Old, Something New, Something Borrowed; How the Thermoacidophilic Archaeon Sulfolobus solfataricus Responds to Oxidative Stress. PLoS ONE, 2009, 4, e6964.	2.5	70
2	Genetic diversity of Brucella abortus and Brucella melitensis in Kazakhstan using MLVA-16. Infection, Genetics and Evolution, 2015, 34, 173-180.	2.3	36
3	Epidemiology of Brucellosis and Genetic Diversity of Brucella abortus in Kazakhstan. PLoS ONE, 2016, 11, e0167496.	2.5	31
4	Mitochondrial and Y-chromosomal profile of the Kazakh population from East Kazakhstan. Croatian Medical Journal, 2013, 54, 17-24.	0.7	29
5	PUB-NChIPâ€"â€œin vivo biotinylationâ€•approach to study chromatin in proximity to a protein of interest. Genome Research, 2013, 23, 331-340.	5.5	27
6	Proteomic Analysis of Sulfolobus solfataricus during Sulfolobus Turreted Icosahedral Virus Infection. Journal of Proteome Research, 2012, 11, 1420-1432.	3.7	26
7	ZNF555 protein binds to transcriptional activator site of 4qA allele andANT1: potential implication in Facioscapulohumeral dystrophy. Nucleic Acids Research, 2015, 43, 8227-8242.	14.5	15
8	The medieval Mongolian roots of Y-chromosomal lineages from South Kazakhstan. BMC Genetics, 2020, 21, 87.	2.7	15
9	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
10	Draft Genome Sequence of Rhodococcus erythropolis DN1, a Crude Oil Biodegrader. Genome Announcements, 2013, 1, .	0.8	12
11	Genomic characterization of MDR/XDR-TB in Kazakhstan by a combination of high-throughput methods predominantly shows the ongoing transmission of L2/Beijing 94â€"32 central Asian/Russian clusters. BMC Infectious Diseases, 2019, 19, 553.	2.9	10
12	Genetic risk factors for restenosis after percutaneous coronary intervention in Kazakh population. Human Genomics, 2016, 10, 15.	2.9	8
13	Draft Genome Sequence of the Live Vaccine Strain Brucella abortus 82. Genome Announcements, 2013, 1, .	0.8	7
14	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
15	Genetic Characterization of Kazakh Native Sheep Breeds Using Mitochondrial DNA. OnLine Journal of Biological Sciences, 2018, 18, 341-348.	0.4	5
16	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
17	Topokaryotyping demonstrates single cell variability and stress dependent variations in nuclear envelope associated domains. Nucleic Acids Research, 2018, 46, e135-e135.	14.5	3
18	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

#	Article	IF	Citations
19	Mitochondrial DNA analysis of ancient sheep from Kazakhstan: evidence for early sheep introduction. Heliyon, 2021, 7, e08011.	3.2	2
20	Crystal and molecular structure of tigogenin. Russian Journal of Applied Chemistry, 2006, 79, 1371-1373.	0.5	1
21	ABO Blood Group Genotyping by Real-time PCR in Kazakh Population. Central Asian Journal of Global Health, 2014, 3, 177.	0.6	1
22	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
23	Analysis of Bacteroides fragilis Clinical Strains Isolated in Kazakhstan. Microbiology Resource Announcements, 2021, 10, .	0.6	0
24	Damage-Induced Mutation Clustering in Gram-Positive Bacteria: Preliminary Data. Symmetry, 2022, 14, 1431.	2.2	0