

Asset Daniyarov

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

94
citations

1937685

4
h-index

1588992

8
g-index

11
all docs

11
docs citations

11
times ranked

95
citing authors

#	ARTICLE	IF	CITATIONS
1	Use of retrotransposon-derived genetic markers to analyse genomic variability in plants. <i>Functional Plant Biology</i> , 2019, 46, 15.	2.1	45
2	Genetic analysis of the grapevine genotypes of the Russian <i>Vitis</i> ampelographic collection using iPBS markers. <i>Genetica</i> , 2019, 147, 91-101.	1.1	28
3	Determinants of resistance in <i>Bacteroides fragilis</i> strain BFR_KZ01 isolated from a patient with peritonitis in Kazakhstan. <i>Journal of Global Antimicrobial Resistance</i> , 2021, 25, 1-4.	2.2	7
4	Genomic Analysis of Multidrug-Resistant <i>Mycobacterium tuberculosis</i> Strains From Patients in Kazakhstan. <i>Frontiers in Genetics</i> , 2021, 12, 683515.	2.3	5
5	Whole genome sequence data of <i>Mycobacterium tuberculosis</i> XDR strain, isolated from patient in Kazakhstan. <i>Data in Brief</i> , 2020, 33, 106416.	1.0	3
6	Whole-genome sequencing data of Kazakh individuals. <i>BMC Research Notes</i> , 2021, 14, 45.	1.4	2
7	re-Searcher: GUI-based bioinformatics tool for simplified genomics data mining of VCF files. <i>PeerJ</i> , 2021, 9, e11333.	2.0	2
8	Meta-Analysis of Esophageal Cancer Transcriptomes Using Independent Component Analysis. <i>Frontiers in Genetics</i> , 2021, 12, 683632.	2.3	2
9	THE CASE OF PRE-EXTENSIVELY DRUG-RESISTANT TUBERCULOSIS DETECTION BY USING THE WHOLE GENOME SEQUENCING <i>M.TUBERCULOSIS</i> . <i>Chest</i> , 2020, 157, A18.	0.8	0
10	Analysis of <i>Bacteroides fragilis</i> Clinical Strains Isolated in Kazakhstan. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.6	0
11	Draft Genome Sequence of <i>Lactobacillus salivarius</i> KZ-NCB, Isolated from Chicken Cecum. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	0