Yulong Wei

List of Publications by Year in descending order

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Version: 2024-02-01

1125271 1306789 12 184 7 13 citations g-index h-index papers 14 14 14 269 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	Conservation of griseofulvin genes in the <code><i>gsf</i>gene</code> cluster among fungal genomes. G3: Genes, Genomes, Genetics, 2022, 12, .	0.8	5
2	In Silico Molecular Dynamics of Griseofulvin and Its Derivatives Revealed Potential Therapeutic Applications for COVID-19. International Journal of Molecular Sciences, 2022, 23, 6889.	1.8	6
3	Predicting mammalian species at risk of being infected by SARS-CoV-2 from an ACE2 perspective. Scientific Reports, 2021, 11, 1702.	1.6	22
4	Applications of Protein Secondary Structure Algorithms in SARS-CoV-2 Research. Journal of Proteome Research, 2021, 20, 1457-1463.	1.8	3
5	Does Saccharomyces cerevisiae Require Specific Post-Translational Silencing against Leaky Translation of Haclup?. Microorganisms, 2021, 9, 620.	1.6	1
6	Coronavirus genomes carry the signatures of their habitats. PLoS ONE, 2020, 15, e0244025.	1.1	25
7	Unique Shine–Dalgarno Sequences in Cyanobacteria and Chloroplasts Reveal Evolutionary Differences in Their Translation Initiation. Genome Biology and Evolution, 2019, 11, 3194-3206.	1.1	10
8	An improved estimation of tRNA expression to better elucidate the coevolution between tRNA abundance and codon usage in bacteria. Scientific Reports, 2019, 9, 3184.	1.6	28
9	RNA-Seq-Based Analysis Reveals Heterogeneity in Mature 16S rRNA 3′ Termini and Extended Anti-Shine-Dalgarno Motifs in Bacterial Species. G3: Genes, Genomes, Genetics, 2018, 8, 3973-3979.	0.8	4
10	The Role of +4U as an Extended Translation Termination Signal in Bacteria. Genetics, 2017, 205, 539-549.	1.2	22
11	Elucidating the 16S rRNA 3′ boundaries and defining optimal SD/aSD pairing in Escherichia coli and Bacillus subtilis using RNA-Seq data. Scientific Reports, 2017, 7, 17639.	1.6	22
12	Coevolution between Stop Codon Usage and Release Factors in Bacterial Species. Molecular Biology and Evolution, 2016, 33, 2357-2367.	3.5	29