Shuai Jiang

List of Publications by Year
in descending order

12 papers	765 citations	1307594 7 h-index	1281871 11 g-index
13 all docs	13 docs citations	13 times ranked	936 citing authors

#	Article	IF	Citations
1	Database Resources of the National Genomics Data Center, China National Center for Bioinformation in 2022. Nucleic Acids Research, 2022, 50, D27-D38.	14.5	364
2	CompoDynamics: a comprehensive database for characterizing sequence composition dynamics. Nucleic Acids Research, 2022, 50, D962-D969.	14.5	6
3	Quantitative model suggests both intrinsic and contextual features contribute to the transcript coding ability determination in cells. Briefings in Bioinformatics, 2022, 23, .	6.5	0
4	LncExpDB: an expression database of human long non-coding RNAs. Nucleic Acids Research, 2021, 49, D962-D968.	14.5	55
5	The Global Landscape of SARS-CoV-2 Genomes, Variants, and Haplotypes in 2019nCoVR. Genomics, Proteomics and Bioinformatics, 2020, 18, 749-759.	6.9	88
6	Multi-omics annotation of human long non-coding RNAs. Biochemical Society Transactions, 2020, 48, 1545-1556.	3.4	6
7	An online coronavirus analysis platform from the National Genomics Data Center. Zoological Research, 2020, 41, 705-708.	2.1	62
8	An expanded landscape of human long noncoding RNA. Nucleic Acids Research, 2019, 47, 7842-7856.	14.5	92
9	GenTree, an integrated resource for analyzing the evolution and function of primate-specific coding genes. Genome Research, 2019, 29, 682-696.	5.5	67
10	Systematic identification and annotation of multiple-variant compound effects at transcription factor binding sites in human genome. Journal of Genetics and Genomics, 2018, 45, 373-379.	3.9	5
11	Accurately annotate compound effects of genetic variants using a context-sensitive framework. Nucleic Acids Research, 2017, 45, e82-e82.	14.5	9
12	LocExpress: a web server for efficiently estimating expression of novel transcripts. BMC Genomics, 2016, 17, 1023.	2.8	6