

Jia-Jun Qiu

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

385
citations

1162889

8
h-index

1372474

10
g-index

13
all docs

13
docs citations

13
times ranked

433
citing authors

#	ARTICLE	IF	CITATIONS
1	PredictProtein - Predicting Protein Structure and Function for 29 Years. <i>Nucleic Acids Research</i> , 2021, 49, W535-W540.	6.5	135
2	ProNA2020 predicts protein-DNA, protein-RNA, and protein-protein binding proteins and residues from sequence. <i>Journal of Molecular Biology</i> , 2020, 432, 2428-2443.	2.0	67
3	Long non-coding RNA LINC01296 is a potential prognostic biomarker in patients with colorectal cancer. <i>Tumor Biology</i> , 2015, 36, 7175-7183.	0.8	60
4	Long non-coding RNA MALAT1 protects preterm infants with bronchopulmonary dysplasia by inhibiting cell apoptosis. <i>BMC Pulmonary Medicine</i> , 2017, 17, 199.	0.8	35
5	Identification and functional analysis of long non-coding RNAs in human and mouse early embryos based on single-cell transcriptome data. <i>Oncotarget</i> , 2016, 7, 61215-61228.	0.8	27
6	Dysfunctions of mitochondria in close association with strong perturbation of long noncoding RNAs expression in down syndrome. <i>International Journal of Biochemistry and Cell Biology</i> , 2017, 92, 115-120.	1.2	19
7	Network-based protein-protein interaction prediction method maps perturbations of cancer interactome. <i>PLoS Genetics</i> , 2021, 17, e1009869.	1.5	13
8	Adrenomedullin regulated by miRNA-574-3p protects premature infants with bronchopulmonary dysplasia. <i>Bioscience Reports</i> , 2020, 40, .	1.1	12
9	RNA editing regulates lncRNA splicing in human early embryo development. <i>PLoS Computational Biology</i> , 2021, 17, e1009630.	1.5	6
10	Protein-protein and protein-nucleic acid binding residues important for common and rare sequence variants in human. <i>BMC Bioinformatics</i> , 2020, 21, 452.	1.2	4
11	Genome-wide hypermethylation is closely associated with abnormal expression of genes involved in neural development in induced pluripotent stem cells derived from a Down syndrome mouse model. <i>Cell Biology International</i> , 2021, 45, 1383-1392.	1.4	0