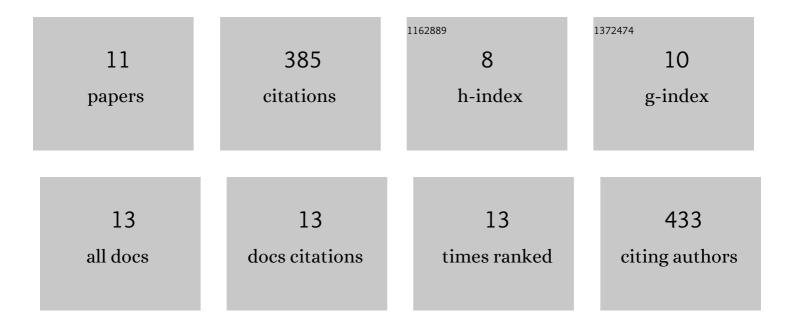
## Jia-Jun Qiu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8038471/publications.pdf Version: 2024-02-01



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