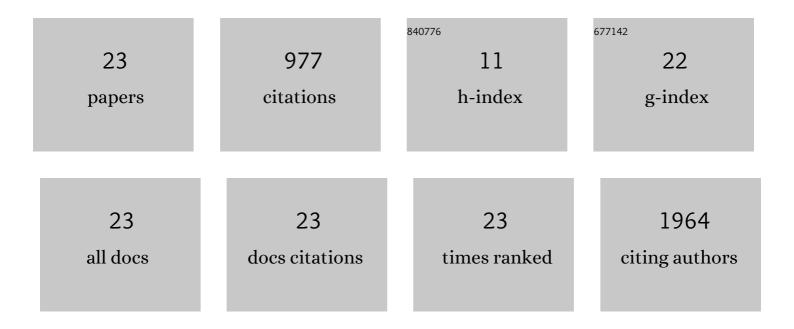
Jia-jian Zhou

List of Publications by Year in descending order

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Ιιλ-ΙΙΛΝ ΖΗΟΙΙ

#	Article	IF	CITATIONS
1	Two related female patients with familial facial pigmentary demarcation lines. Journal of Cosmetic Dermatology, 2022, 21, 5279-5281.	1.6	0
2	Engineered Cas12a-Plus nuclease enables gene editing with enhanced activity and specificity. BMC Biology, 2022, 20, 91.	3.8	15
3	Reciprocal Regulation between IncRNA ANRIL and p15 in Steroid-Induced Glaucoma. Cells, 2022, 11, 1468.	4.1	11
4	Large scale RNA-binding proteins/LncRNAs interaction analysis to uncover lncRNA nuclear localization mechanisms. Briefings in Bioinformatics, 2021, 22, .	6.5	19
5	Tag-seq: a convenient and scalable method for genome-wide specificity assessment of CRISPR/Cas nucleases. Communications Biology, 2021, 4, 830.	4.4	10
6	Cabazitaxel suppresses colorectal cancer cell growth via enhancing the p53 antitumor pathway. FEBS Open Bio, 2021, 11, 3032-3050.	2.3	3
7	Immune Cell Infiltration Analysis Demonstrates Excessive Mast Cell Activation in Psoriasis. Frontiers in Immunology, 2021, 12, 773280.	4.8	18
8	β-Catenin safeguards the ground state of mousepluripotency by strengthening the robustness of the transcriptional apparatus. Science Advances, 2020, 6, eaba1593.	10.3	10
9	PARMAP: A Pan-Genome-Based Computational Framework for Predicting Antimicrobial Resistance. Frontiers in Microbiology, 2020, 11, 578795.	3.5	10
10	Hippo-YAP signaling controls lineage differentiation of mouse embryonic stem cells through modulating the formation of super-enhancers. Nucleic Acids Research, 2020, 48, 7182-7196.	14.5	41
11	Elevated H3K27ac in aged skeletal muscle leads to increase in extracellular matrix and fibrogenic conversion of muscle satellite cells. Aging Cell, 2019, 18, e12996.	6.7	35
12	MyoD induced enhancer RNA interacts with hnRNPL to activate target gene transcription during myogenic differentiation. Nature Communications, 2019, 10, 5787.	12.8	70
13	SKmDB: an integrated database of next generation sequencing information in skeletal muscle. Bioinformatics, 2019, 35, 847-855.	4.1	2
14	lncFunTK: a toolkit for functional annotation of long noncoding RNAs. Bioinformatics, 2018, 34, 3415-3416.	4.1	12
15	Capturing the interactome of newly transcribed RNA. Nature Methods, 2018, 15, 213-220.	19.0	170
16	PCGF5 is required for neural differentiation of embryonic stem cells. Nature Communications, 2018, 9, 1463.	12.8	60
17	Bioinformatics for Novel Long Intergenic Noncoding RNA (lincRNA) Identification in Skeletal Muscle Cells. Methods in Molecular Biology, 2017, 1556, 355-362.	0.9	3
18	MyoD- and FoxO3-mediated hotspot interaction orchestrates super-enhancer activity during myogenic differentiation. Nucleic Acids Research, 2017, 45, 8785-8805.	14.5	51

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#	Article	IF	CITATIONS
19	LncFunNet: an integrated computational framework for identification of functional long noncoding RNAs in mouse skeletal muscle cells. Nucleic Acids Research, 2017, 45, e108-e108.	14.5	43
20	Multi-step formation, evolution, and functionalization of new cytoplasmic male sterility genes in the plant mitochondrial genomes. Cell Research, 2017, 27, 130-146.	12.0	65
21	Transcriptome analysis demonstrate widespread differential expression of long noncoding RNAs involve in Larimichthys crocea immune response. Fish and Shellfish Immunology, 2016, 51, 1-8.	3.6	41
22	An Intracellular Laccase is Responsible for the Epicatechin Mediated Anthocyanin Degradation in Litchi Fruit Pericarp. Plant Physiology, 2015, 169, pp.00359.2015.	4.8	78
23	The genome of the leaf-cutting ant <i>Acromyrmex echinatior</i> suggests key adaptations to advanced social life and fungus farming. Genome Research, 2011, 21, 1339-1348.	5.5	210