

Yunhao Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5721531/publications.pdf>

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9
papers

1,360
citations

1163117

8
h-index

1474206

9
g-index

9
all docs

9
docs citations

9
times ranked

1590
citing authors

#	ARTICLE	IF	CITATIONS
1	Nanopore sequencing technology, bioinformatics and applications. <i>Nature Biotechnology</i> , 2021, 39, 1348-1365.	17.5	521
2	MED25 connects enhancerâ€“promoter looping and MYC2-dependent activation of jasmonate signalling. <i>Nature Plants</i> , 2019, 5, 616-625.	9.3	82
3	Single-molecule long-read sequencing reveals the chromatin basis of gene expression. <i>Genome Research</i> , 2019, 29, 1329-1342.	5.5	46
4	iASPP mediates p53 selectivity through a modular mechanism fine-tuning DNA recognition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 17470-17479.	7.1	20
5	E-C coupling structural protein junctophilin-2 encodes a stress-adaptive transcription regulator. <i>Science</i> , 2018, 362, .	12.6	78
6	A Statistical Method for Observing Personal Diploid Methylomes and Transcriptomes with Single-Molecule Real-Time Sequencing. <i>Genes</i> , 2018, 9, 460.	2.4	2
7	IDP-ASE: haplotyping and quantifying allele-specific expression at the gene and gene isoform level by hybrid sequencing. <i>Nucleic Acids Research</i> , 2017, 45, e32-e32.	14.5	42
8	Comprehensive comparison of Pacific Biosciences and Oxford Nanopore Technologies and their applications to transcriptome analysis. <i>F1000Research</i> , 2017, 6, 100.	1.6	366
9	Comprehensive comparison of Pacific Biosciences and Oxford Nanopore Technologies and their applications to transcriptome analysis. <i>F1000Research</i> , 2017, 6, 100.	1.6	203