

Rongbin Zheng

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11497331/publications.pdf>

Version: 2024-02-01

10
papers

1,884
citations

1163117

8
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

2901
citing authors

#	ARTICLE	IF	CITATIONS
1	ViMIC: a database of human disease-related virus mutations, integration sites and cis-effects. <i>Nucleic Acids Research</i> , 2022, 50, D918-D927.	14.5	15
2	TISCH: a comprehensive web resource enabling interactive single-cell transcriptome visualization of tumor microenvironment. <i>Nucleic Acids Research</i> , 2021, 49, D1420-D1430.	14.5	481
3	Determinants of transcription factor regulatory range. <i>Nature Communications</i> , 2020, 11, 2472.	12.8	28
4	Cistrome Data Browser and Toolkit: analyzing human and mouse genomic data using compendia of ChIP-seq and chromatin accessibility data. <i>Quantitative Biology</i> , 2020, 8, 267-276.	0.5	11
5	Lisa: inferring transcriptional regulators through integrative modeling of public chromatin accessibility and ChIP-seq data. <i>Genome Biology</i> , 2020, 21, 32.	8.8	161
6	GIDB: a knowledge database for the automated curation and multidimensional analysis of molecular signatures in gastrointestinal cancer. <i>Database: the Journal of Biological Databases and Curation</i> , 2019, 2019, .	3.0	3
7	Cistrome-GO: a web server for functional enrichment analysis of transcription factor ChIP-seq peaks. <i>Nucleic Acids Research</i> , 2019, 47, W206-W211.	14.5	76
8	Cistrome Data Browser: expanded datasets and new tools for gene regulatory analysis. <i>Nucleic Acids Research</i> , 2019, 47, D729-D735.	14.5	527
9	Cistrome Cancer: A Web Resource for Integrative Gene Regulation Modeling in Cancer. <i>Cancer Research</i> , 2017, 77, e19-e22.	0.9	130
10	Cistrome Data Browser: a data portal for ChIP-Seq and chromatin accessibility data in human and mouse. <i>Nucleic Acids Research</i> , 2017, 45, D658-D662.	14.5	451