

Christopher J F Cameron

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9138396/publications.pdf>

Version: 2024-02-01

11
papers

320
citations

1684188

5
h-index

1474206

9
g-index

17
all docs

17
docs citations

17
times ranked

498
citing authors

#	ARTICLE	IF	CITATIONS
1	Functional annotation of human long noncoding RNAs via molecular phenotyping. <i>Genome Research</i> , 2020, 30, 1060-1072.	5.5	109
2	RADICL-seq identifies general and cell type-specific principles of genome-wide RNA-chromatin interactions. <i>Nature Communications</i> , 2020, 11, 1018.	12.8	98
3	PAM multiplicity marks genomic target sites as inhibitory to CRISPR-Cas9 editing. <i>Nature Communications</i> , 2015, 6, 10124.	12.8	52
4	HIFI: estimating DNA-DNA interaction frequency from Hi-C data at restriction-fragment resolution. <i>Genome Biology</i> , 2020, 21, 11.	8.8	24
5	Double-Stranded Biotinylated Donor Enhances Homology-Directed Repair in Combination with Cas9 Monoavidin in Mammalian Cells. <i>CRISPR Journal</i> , 2018, 1, 414-430.	2.9	12
6	2C-ChIP: measuring chromatin immunoprecipitation signal from defined genomic regions with deep sequencing. <i>BMC Genomics</i> , 2019, 20, 162.	2.8	4
7	Nodal modulator (NOMO) is required to sustain endoplasmic reticulum morphology. <i>Journal of Biological Chemistry</i> , 2021, 297, 100937.	3.4	4
8	LAMPS: an analysis pipeline for sequence-specific ligation-mediated amplification reads. <i>BMC Research Notes</i> , 2020, 13, 273.	1.4	3
9	Neural grammar networks for toxicology. , 2010, , .		1
10	Classifying and scoring of molecules with the NGN: new datasets, significance tests, and generalization. <i>BMC Bioinformatics</i> , 2010, 11, S4.	2.6	0
11	Profiling Chromatin Landscape at High Resolution and Throughput with 2C-ChIP. <i>Methods in Molecular Biology</i> , 2021, 2157, 127-157.	0.9	0