

JosÃ© Alquicira-HernÃ¡ndez

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6726014/publications.pdf>

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9
papers

1,531
citations

1163117

8
h-index

1372567

10
g-index

14
all docs

14
docs citations

14
times ranked

2735
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-cell eQTL mapping identifies cell type-specific genetic control of autoimmune disease. <i>Science</i> , 2022, 376, eabf3041.	12.6	171
2	<i>Nebulosa</i> recovers single-cell gene expression signals by kernel density estimation. <i>Bioinformatics</i> , 2021, 37, 2485-2487.	4.1	133
3	Large-scale cis- and trans-eQTL analyses identify thousands of genetic loci and polygenic scores that regulate blood gene expression. <i>Nature Genetics</i> , 2021, 53, 1300-1310.	21.4	590
4	No evidence that plasmablasts transdifferentiate into developing neutrophils in severe COVID-19 disease. <i>Clinical and Translational Immunology</i> , 2021, 10, e1308.	3.8	10
5	Benchmarking of cell type deconvolution pipelines for transcriptomics data. <i>Nature Communications</i> , 2020, 11, 5650.	12.8	207
6	Programmatic access to bacterial regulatory networks with <i>regutools</i> . <i>Bioinformatics</i> , 2020, 36, 4532-4534.	4.1	4
7	ascend: R package for analysis of single-cell RNA-seq data. <i>GigaScience</i> , 2019, 8, .	6.4	36
8	scPred: accurate supervised method for cell-type classification from single-cell RNA-seq data. <i>Genome Biology</i> , 2019, 20, 264.	8.8	263
9	Human splicing diversity and the extent of unannotated splice junctions across human RNA-seq samples on the Sequence Read Archive. <i>Genome Biology</i> , 2016, 17, 266.	8.8	94