

Fidel RamÃ rez

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6770405/publications.pdf>

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14

papers

6,662

citations

840776

11

h-index

1125743

13

g-index

17

all docs

17

docs citations

17

times ranked

14051

citing authors

#	ARTICLE		IF	CITATIONS
1	CD206+ tumor-associated macrophages cross-present tumor antigen and drive antitumor immunity. <i>JCI Insight</i> , 2022, 7, .		5.0	29
2	miR449 Protects Airway Regeneration by Controlling AURKA/HDAC6-Mediated Ciliary Disassembly. <i>International Journal of Molecular Sciences</i> , 2022, 23, 7749.		4.1	1
3	pyGenomeTracks: reproducible plots for multivariate genomic datasets. <i>Bioinformatics</i> , 2021, 37, 422-423.		4.1	237
4	Aprovechamiento de residuos en la construcciÃ³n de galpones como alternativa de sostenibilidad en el corregimiento El Prodigio, en San Luis, Antioquia-Colombia. <i>Tecno LÃ³gicas</i> , 2021, 24, e1830.		0.3	0
5	A single-cell RNA-sequencing training and analysis suite using the Galaxy framework. <i>GigaScience</i> , 2020, 9, .		6.4	14
6	Hi-C guided assemblies reveal conserved regulatory topologies on X and autosomes despite extensive genome shuffling. <i>Genes and Development</i> , 2019, 33, 1591-1612.		5.9	43
7	CAPRI enables comparison of evolutionarily conserved RNA interacting regions. <i>Nature Communications</i> , 2019, 10, 2682.		12.8	39
8	High-resolution TADs reveal DNA sequences underlying genome organization in flies. <i>Nature Communications</i> , 2018, 9, 189.		12.8	652
9	RELACS nuclei barcoding enables high-throughput ChIP-seq. <i>Communications Biology</i> , 2018, 1, 214.		4.4	30
10	A comprehensive analysis of 195 DNA methylomes reveals shared and cell-specific features of partially methylated domains. <i>Genome Biology</i> , 2018, 19, 150.		8.8	71
11	Community-Driven Data Analysis Training for Biology. <i>Cell Systems</i> , 2018, 6, 752-758.e1.		6.2	141
12	Galaxy HiCExplorer: a web server for reproducible Hi-C data analysis, quality control and visualization. <i>Nucleic Acids Research</i> , 2018, 46, W11-W16.		14.5	168
13	deepTools2: a next generation web server for deep-sequencing data analysis. <i>Nucleic Acids Research</i> , 2016, 44, W160-W165.		14.5	5,157
14	High-Affinity Sites Form an Interaction Network to Facilitate Spreading of the MSL Complex across the X Chromosome in Drosophila. <i>Molecular Cell</i> , 2015, 60, 146-162.		9.7	70