

Alejandro J Brenes

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

661
citations

933447

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1199594

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21
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21
docs citations

21
times ranked

1321
citing authors

#	ARTICLE	IF	CITATIONS
1	Nrf2 activation reprograms macrophage intermediary metabolism and suppresses the type I interferon response. <i>IScience</i> , 2022, 25, 103827.	4.1	51
2	type I IFN, prothrombotic hyperinflammatory neutrophil signature is distinct for COVID-19 ARDS. <i>Wellcome Open Research</i> , 2021, 6, 38.	1.8	29
3	Erosion of human X chromosome inactivation causes major remodeling of the iPSC proteome. <i>Cell Reports</i> , 2021, 35, 109032.	6.4	23
4	type I IFN, prothrombotic hyperinflammatory neutrophil signature is distinct for COVID-19 ARDS. <i>Wellcome Open Research</i> , 2021, 6, 38.	1.8	35
5	Tissue environment, not ontogeny, defines murine intestinal intraepithelial T lymphocytes. <i>ELife</i> , 2021, 10, .	6.0	14
6	Population-scale proteome variation in human induced pluripotent stem cells. <i>ELife</i> , 2020, 9, .	6.0	40
7	Multibatch TMT Reveals False Positives, Batch Effects and Missing Values. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 1967-1980.	3.8	128
8	Quantitative analysis of T cell proteomes and environmental sensors during T cell differentiation. <i>Nature Immunology</i> , 2019, 20, 1542-1554.	14.5	152
9	The Encyclopedia of Proteome Dynamics: the KinoViewer. <i>Bioinformatics</i> , 2019, 35, 1441-1442.	4.1	9
10	Antigen receptor control of methionine metabolism in T cells. <i>ELife</i> , 2019, 8, .	6.0	132
11	The Encyclopedia of Proteome Dynamics: a big data ecosystem for (prote)omics. <i>Nucleic Acids Research</i> , 2018, 46, D1202-D1209.	14.5	19
12	Proteome-wide analysis of protein abundance and turnover remodelling during oncogenic transformation of human breast epithelial cells. <i>Wellcome Open Research</i> , 2018, 3, 51.	1.8	18