

COVID-19 immune features revealed by a large-scale sin

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Identification of Distinct Immune Cells Associated with Various Clinical Presentations of COVID-19. SSRN Electronic Journal, 0, , .	0.4	0
2	scAdapt: virtual adversarial domain adaptation network for single cell RNA-seq data classification across platforms and species. Briefings in Bioinformatics, 2021, 22, .	3.2	13
7	Next-generation crop engineering. Nature Plants, 2021, 7, 241-241.	4.7	3
9	BET inhibition blocks inflammation-induced cardiac dysfunction and SARS-CoV-2 infection. Cell, 2021, 184, 2167-2182.e22.	13.5	131
13	Circulating Calprotectin as a Biomarker of COVID-19 Severity. Expert Review of Clinical Immunology, 2021, 17, 431-443.	1.3	70
14	Single-cell multi-omics analysis of the immune response in COVID-19. Nature Medicine, 2021, 27, 904-916.	15.2	452
15	COVID-19: complexity of disease severity revealed by systemic and localized single cell immune atlas. Signal Transduction and Targeted Therapy, 2021, 6, 156.	7.1	1
19	Multidisciplinary approach for post-acute COVID-19 syndrome: time to break down the walls. European Respiratory Journal, 2021, 58, 2101090.	3.1	18
20	Transcriptional Changes in CD16+ Monocytes May Contribute to the Pathogenesis of COVID-19. Frontiers in Immunology, 2021, 12, 665773.	2.2	20
22	Integrative genomics analysis reveals a 21q22.11 locus contributing risk to COVID-19. Human Molecular Genetics, 2021, 30, 1247-1258.	1.4	28
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27	A cohort autopsy study defines COVID-19 systemic pathogenesis. Cell Research, 2021, 31, 836-846.	5.7	93
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39	An Impaired Inflammatory and Innate Immune Response in COVID-19. Molecules and Cells, 2021, 44, 384-391.	1.0	13

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40	Exploring the utility of extracellular vesicles in ameliorating viral infection-associated inflammation, cytokine storm and tissue damage. <i>Translational Oncology</i> , 2021, 14, 101095.	1.7	23
42	CRISPR Technique Incorporated with Single-Cell RNA Sequencing for Studying Hepatitis B Infection. <i>Analytical Chemistry</i> , 2021, 93, 10756-10761.	3.2	9
43	Multimodal single-cell omics analysis identifies epithelium-immune cell interactions and immune vulnerability associated with sex differences in COVID-19. <i>Signal Transduction and Targeted Therapy</i> , 2021, 6, 292.	7.1	13
45	Interactomes of SARS-CoV-2 and human coronaviruses reveal host factors potentially affecting pathogenesis. <i>EMBO Journal</i> , 2021, 40, e107776.	3.5	53
48	Natural killer cells and unconventional T cells in COVID-19. <i>Current Opinion in Virology</i> , 2021, 49, 176-182.	2.6	28
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