Ephraim Kenigsberg

List of Publications by Year in descending order

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759055 1125617 2,424 14 12 13 citations h-index g-index papers 17 17 17 4146 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Ulcerative colitis is characterized by a plasmablast-skewed humoral response associated with disease activity. Nature Medicine, 2022, 28, 766-779.	15.2	70
2	A myeloid–stromal niche and gp130 rescue in NOD2-driven Crohn's disease. Nature, 2021, 593, 275-281.	13.7	65
3	Intestinal Host Response to SARS-CoV-2 Infection and COVID-19 Outcomes in Patients With Gastrointestinal Symptoms. Gastroenterology, 2021, 160, 2435-2450.e34.	0.6	118
4	Tissue-resident macrophages provide a pro-tumorigenic niche to early NSCLC cells. Nature, 2021, 595, 578-584.	13.7	284
5	Abstract 64: Characterization of molecular and spatial diversity of macrophages in hepatocellular carcinoma., 2021,,.		1
6	685â€Characterization of molecular and spatial diversity of macrophages in hepatocellular carcinoma. , 2021, 9, A713-A713.		0
7	Single-cell analysis of human non-small cell lung cancer lesions refines tumor classification and patient stratification. Cancer Cell, 2021, 39, 1594-1609.e12.	7.7	151
8	Serum Amyloid A Proteins Induce Pathogenic Th17 Cells and Promote Inflammatory Disease. Cell, 2020, 180, 79-91.e16.	13.5	243
9	DNA methylation disruption reshapes the hematopoietic differentiation landscape. Nature Genetics, 2020, 52, 378-387.	9.4	154
10	A conserved dendritic-cell regulatory program limits antitumour immunity. Nature, 2020, 580, 257-262.	13.7	476
11	Altering Sphingolipid Metabolism Attenuates Cell Death and Inflammatory Response After Myocardial Infarction. Circulation, 2020, 141, 916-930.	1.6	84
12	Single-Cell Analysis of Crohn's Disease Lesions Identifies a Pathogenic Cellular Module Associated with Resistance to Anti-TNF Therapy. Cell, 2019, 178, 1493-1508.e20.	13.5	519
13	Prioritizing Crohn's disease genes by integrating association signals with gene expression implicates monocyte subsets. Genes and Immunity, 2019, 20, 577-588.	2.2	16
14	MARS-seq2.0: an experimental and analytical pipeline for indexed sorting combined with single-cell RNA sequencing. Nature Protocols, 2019, 14, 1841-1862.	5.5	200