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List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

2,489
citations

279798

23
h-index

377865

34
g-index

41
all docs

41
docs citations

41
times ranked

5931
citing authors

#	ARTICLE	IF	CITATIONS
1	The chromatin remodeller ATRX facilitates diverse nuclear processes, in a stochastic manner, in both heterochromatin and euchromatin. <i>Nature Communications</i> , 2022, 13, .	12.8	20
2	Multi-Modal Characterization of Monocytes in Idiopathic Pulmonary Fibrosis Reveals a Primed Type I Interferon Immune Phenotype. <i>Frontiers in Immunology</i> , 2021, 12, 623430.	4.8	34
3	IL-6 effector function of group 2 innate lymphoid cells (ILC2) is NOD2 dependent. <i>Science Immunology</i> , 2021, 6, .	11.9	8
4	BET inhibition disrupts transcription but retains enhancer-promoter contact. <i>Nature Communications</i> , 2021, 12, 223.	12.8	84
5	Single-cell atlas of colonic CD8+ T cells in ulcerative colitis. <i>Nature Medicine</i> , 2020, 26, 1480-1490.	30.7	126
6	DOT1L inhibition reveals a distinct subset of enhancers dependent on H3K79 methylation. <i>Nature Communications</i> , 2019, 10, 2803.	12.8	99
7	Reconstruction of the Global Neural Crest Gene Regulatory Network In Vivo. <i>Developmental Cell</i> , 2019, 51, 255-276.e7.	7.0	108
8	Transcriptomic profiling of the myeloma bone-lining niche reveals BMP signalling inhibition to improve bone disease. <i>Nature Communications</i> , 2019, 10, 4533.	12.8	46
9	Cell-intrinsic depletion of Aml1-ETO-expressing pre-leukemic hematopoietic stem cells by K-Ras activating mutation. <i>Haematologica</i> , 2019, 104, 2215-2224.	3.5	9
10	Canonical Notch signaling is dispensable for adult steady-state and stress myelo-erythropoiesis. <i>Blood</i> , 2018, 131, 1712-1719.	1.4	14
11	Single-cell analysis reveals the continuum of human lympho-myeloid progenitor cells. <i>Nature Immunology</i> , 2018, 19, 85-97.	14.5	193
12	Does osteogenic potential of clonal human bone marrow mesenchymal stem/stromal cells correlate with their vascular supportive ability?. <i>Stem Cell Research and Therapy</i> , 2018, 9, 351.	5.5	6
13	SCL/TAL1 cooperates with Polycomb RYBP-PRC1 to suppress alternative lineages in blood-fated cells. <i>Nature Communications</i> , 2018, 9, 5375.	12.8	29
14	From Pioneer to Repressor: Bimodal foxd3 Activity Dynamically Remodels Neural Crest Regulatory Landscape In Vivo. <i>Developmental Cell</i> , 2018, 47, 608-628.e6.	7.0	92
15	Impact of spliceosome mutations on RNA splicing in myelodysplasia: dysregulated genes/pathways and clinical associations. <i>Blood</i> , 2018, 132, 1225-1240.	1.4	168
16	MLL-AF4 binds directly to a BCL-2 specific enhancer and modulates H3K27Acetylation. <i>Experimental Hematology</i> , 2017, 47, 64-75.	0.4	25
17	MLL-AF4 Spreading Identifies Binding Sites that Are Distinct from Super-Enhancers and that Govern Sensitivity to DOT1L Inhibition in Leukemia. <i>Cell Reports</i> , 2017, 18, 482-495.	6.4	69
18	Hepcidin is regulated by promoter-associated histone acetylation and HDAC3. <i>Nature Communications</i> , 2017, 8, 403.	12.8	45

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19	Niche-mediated depletion of the normal hematopoietic stem cell reservoir by Flt3-ITD-induced myeloproliferation. <i>Journal of Experimental Medicine</i> , 2017, 214, 2005-2021.	8.5	43
20	CD1a presentation of endogenous antigens by group 2 innate lymphoid cells. <i>Science Immunology</i> , 2017, 2, .	11.9	57
21	M1-like monocytes are a major immunological determinant of severity in previously healthy adults with life-threatening influenza. <i>JCI Insight</i> , 2017, 2, e91868.	5.0	59
22	The U2AF1S34F mutation induces lineage-specific splicing alterations in myelodysplastic syndromes. <i>Journal of Clinical Investigation</i> , 2017, 127, 2206-2221.	8.2	69
23	A Comprehensive Analysis of the Impact of HIV on HCV Immune Responses and Its Association with Liver Disease Progression in a Unique Plasma Donor Cohort. <i>PLoS ONE</i> , 2016, 11, e0158037.	2.5	2
24	CTAS: a CT score to quantify disease activity in pulmonary sarcoidosis. <i>Thorax</i> , 2016, 71, 1161-1163.	5.6	26
25	Transforming Growth Factor β 2 Drives Hemogenic Endothelium Programming and the Transition to Hematopoietic Stem Cells. <i>Developmental Cell</i> , 2016, 38, 358-370.	7.0	75
26	Interferon-induced transmembrane protein-3 rs12252-C is associated with rapid progression of acute HIV-1 infection in Chinese MSM cohort. <i>Aids</i> , 2015, 29, 889-894.	2.2	45
27	Interaction between p53 Mutation and a Somatic HDMX Biomarker Better Defines Metastatic Potential in Breast Cancer. <i>Cancer Research</i> , 2015, 75, 698-708.	0.9	13
28	IGF-1R inhibition induces schedule-dependent sensitization of human melanoma to temozolomide. <i>Oncotarget</i> , 2015, 6, 39877-39890.	1.8	20
29	A Polymorphic p53 Response Element in KIT Ligand Influences Cancer Risk and Has Undergone Natural Selection. <i>Cell</i> , 2013, 155, 410-422.	28.9	115
30	The Inheritance of p53. , 2013, , 25-46.		0
31	Alternate Splicing of the p53 Inhibitor HDMX Offers a Superior Prognostic Biomarker than p53 Mutation in Human Cancer. <i>Cancer Research</i> , 2012, 72, 4074-4084.	0.9	58
32	Effect of Five Genetic Variants Associated with Lung Function on the Risk of Chronic Obstructive Lung Disease, and Their Joint Effects on Lung Function. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2011, 184, 786-795.	5.6	128
33	A Comprehensive Evaluation of Potential Lung Function Associated Genes in the SpiroMeta General Population Sample. <i>PLoS ONE</i> , 2011, 6, e19382.	2.5	56
34	Genome-wide association study identifies five loci associated with lung function. <i>Nature Genetics</i> , 2010, 42, 36-44.	21.4	518
35	Integration of single-cell RNA-Seq and CyTOF data characterises heterogeneity of rare cell subpopulations. <i>F1000Research</i> , 0, 11, 560.	1.6	0