Yang Liao

List of Publications by Year in descending order

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Version: 2024-02-01

		279487	329751
39	24,562	23	37
papers	citations	h-index	g-index
43	43	43	49897
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	featureCounts: an efficient general purpose program for assigning sequence reads to genomic features. Bioinformatics, 2014, 30, 923-930.	1.8	16,839
2	The Subread aligner: fast, accurate and scalable read mapping by seed-and-vote. Nucleic Acids Research, 2013, 41, e108-e108.	6.5	2,389
3	The R package Rsubread is easier, faster, cheaper and better for alignment and quantification of RNA sequencing reads. Nucleic Acids Research, 2019, 47, e47-e47.	6.5	1,744
4	Hobit and Blimp1 instruct a universal transcriptional program of tissue residency in lymphocytes. Science, 2016, 352, 459-463.	6.0	721
5	The transcriptional regulators IRF4, BATF and IL-33 orchestrate development and maintenance of adipose tissue–resident regulatory T cells. Nature Immunology, 2015, 16, 276-285.	7.0	442
6	Transcription Factor IRF4 Promotes CD8+ T Cell Exhaustion and Limits the Development of Memory-like T Cells during Chronic Infection. Immunity, 2017, 47, 1129-1141.e5.	6.6	335
7	Transcriptional profiling of mouse B cell terminal differentiation defines a signature for antibody-secreting plasma cells. Nature Immunology, 2015, 16, 663-673.	7.0	332
8	Blimp-1 controls plasma cell function through the regulation of immunoglobulin secretion and the unfolded protein response. Nature Immunology, 2016, 17, 323-330.	7.0	310
9	A molecular threshold for effector CD8+ T cell differentiation controlled by transcription factors Blimp-1 and T-bet. Nature Immunology, 2016, 17, 422-432.	7. O	145
10	Sex-specific adipose tissue imprinting of regulatory T cells. Nature, 2020, 579, 581-585.	13.7	141
11	c-Maf-dependent Treg cell control of intestinal TH17 cells and IgA establishes host–microbiota homeostasis. Nature Immunology, 2019, 20, 471-481.	7. 0	138
12	Assessing technical performance in differential gene expression experiments with external spike-in RNA control ratio mixtures. Nature Communications, 2014, 5, 5125.	5.8	122
13	The TNF Receptor Superfamily-NF-κB Axis Is Critical to Maintain Effector Regulatory T Cells in Lymphoid and Non-lymphoid Tissues. Cell Reports, 2017, 20, 2906-2920.	2.9	115
14	Effector Regulatory T Cell Differentiation and Immune Homeostasis Depend on the Transcription Factor Myb. Immunity, 2017, 46, 78-91.	6.6	83
15	Pax5 loss imposes a reversible differentiation block in B-progenitor acute lymphoblastic leukemia. Genes and Development, 2014, 28, 1337-1350.	2.7	73
16	Environmental sensing by mature B cells is controlled by the transcription factors PU.1 and SpiB. Nature Communications, 2017, 8, 1426.	5.8	71
17	IMiDs prime myeloma cells for daratumumab-mediated cytotoxicity through loss of Ikaros and Aiolos. Blood, 2018, 132, 2166-2178.	0.6	65
18	Attenuation of TCR-induced transcription by Bach2 controls regulatory T cell differentiation and homeostasis. Nature Communications, 2020, 11, 252.	5.8	59

#	Article	IF	Citations
19	Dynamic changes in Id3 and E-protein activity orchestrate germinal center and plasma cell development. Journal of Experimental Medicine, 2016, 213, 1095-1111.	4.2	53
20	TCF-1 limits the formation of Tc17 cells via repression of the MAF–RORγt axis. Journal of Experimental Medicine, 2019, 216, 1682-1699.	4.2	48
21	IRF4 Activity Is Required in Established Plasma Cells to Regulate Gene Transcription and Mitochondrial Homeostasis. Cell Reports, 2019, 29, 2634-2645.e5.	2.9	47
22	An update on using CRISPR/Cas9 in the one-cell stage mouse embryo for generating complex mutant alleles. Cell Death and Differentiation, 2017, 24, 1821-1822.	5.0	38
23	Transcription Factor T-bet in B Cells Modulates Germinal Center Polarization and Antibody Affinity Maturation in Response to Malaria. Cell Reports, 2019, 29, 2257-2269.e6.	2.9	36
24	Read trimming is not required for mapping and quantification of RNA-seq reads at the gene level. NAR Genomics and Bioinformatics, 2020, 2, Iqaa068.	1.5	22
25	Impact of RNA-seq data analysis algorithms on gene expression estimation and downstream prediction. Scientific Reports, 2020, 10, 17925.	1.6	18
26	The transcription factor IRF4 represses proapoptotic BMF and BIM to licence multiple myeloma survival. Leukemia, 2021, 35, 2114-2118.	3.3	18
27	Mining the Plasma Cell Transcriptome for Novel Cell Surface Proteins. International Journal of Molecular Sciences, 2018, 19, 2161.	1.8	17
28	Tailored NEOadjuvant epirubicin, cyclophosphamide and Nanoparticle Albumin-Bound paclitaxel for breast cancer: The phase II NEONAB trialâ€"Clinical outcomes and molecular determinants of response. PLoS ONE, 2019, 14, e0210891.	1.1	13
29	Mining Micro-blogs: Opportunities and Challenges. , 2012, , 129-159.		13
30	Quantitative proteomic profiling reveals novel regionâ€specific markers in the adult mouse brain. Proteomics, 2014, 14, 241-261.	1.3	12
31	Plasmacytoid dendritic cell heterogeneity is defined by CXCL10 expression following TLR7 stimulation. Immunology and Cell Biology, 2018, 96, 1083-1094.	1.0	12
32	A point mutation in the <i>Ncr1</i> signal peptide impairs the development of innate lymphoid cell subsets. Oncolmmunology, 2018, 7, e1475875.	2.1	9
33	Molecular profiling reveals features of clinical immunity and immunosuppression in asymptomatic <i>P. falciparum</i> malaria. Molecular Systems Biology, 2022, 18, e10824.	3.2	9
34	Impact of gene annotation choice on the quantification of RNA-seq data. BMC Bioinformatics, 2022, 23, 107.	1.2	8
35	Epithelial de-differentiation triggered by co-ordinate epigenetic inactivation of the EHF and CDX1 transcription factors drives colorectal cancer progression. Cell Death and Differentiation, 2022, 29, 2288-2302.	5.0	6
36	ScoreFinder: A method for collaborative quality inference on user-generated content., 2010,,.		2

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#	Article	IF	CITATIONS
37	Dynamic changes in Id3 and E-protein activity orchestrate germinal center and plasma cell development. Journal of Cell Biology, 2016, 213, 2135OIA110.	2.3	1
38	Decentralisation of ScoreFinder: A Framework for Credibility Management on User-Generated Contents. Lecture Notes in Computer Science, 2010, , 272-282.	1.0	1
39	ScoreTree: A Decentralised Framework for Credibility Management of User-Generated Content. Lecture Notes in Computer Science, 2011, , 249-256.	1.0	0