

Ying Wan

List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

1,081
citations

471509

17
h-index

414414

32
g-index

36
all docs

36
docs citations

36
times ranked

2136
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular dissection of the miR-17-92 cluster's critical dual roles in promoting Th1 responses and preventing inducible Treg differentiation. <i>Blood</i> , 2011, 118, 5487-5497.	1.4	270
2	Identification of a serum microRNA expression signature for detection of lung cancer, involving miR-23b, miR-221, miR-148b and miR-423-3p. <i>Lung Cancer</i> , 2017, 114, 6-11.	2.0	67
3	Diversity index of mucosal resident T lymphocyte repertoire predicts clinical prognosis in gastric cancer. <i>Oncotmunology</i> , 2015, 4, e1001230.	4.6	57
4	MeCP2 Reinforces STAT3 Signaling and the Generation of Effector CD4 ⁺ T Cells by Promoting miR-124-Mediated Suppression of SOCS5. <i>Science Signaling</i> , 2014, 7, ra25.	3.6	55
5	Dax1 and Nanog act in parallel to stabilize mouse embryonic stem cells and induced pluripotency. <i>Nature Communications</i> , 2014, 5, 5042.	12.8	55
6	The GTPase Rab3b/3c-positive recycling vesicles are involved in cross-presentation in dendritic cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 15801-15806.	7.1	53
7	Forced miR-146a expression causes autoimmune lymphoproliferative syndrome in mice via downregulation of Fas in germinal center B cells. <i>Blood</i> , 2013, 121, 4875-4883.	1.4	52
8	Analysis of the Rab GTPase Interactome in Dendritic Cells Reveals Anti-microbial Functions of the Rab32 Complex in Bacterial Containment. <i>Immunity</i> , 2016, 44, 422-437.	14.3	42
9	CD8+NKT-like cells regulate the immune response by killing antigen-bearing DCs. <i>Scientific Reports</i> , 2015, 5, 14124.	3.3	33
10	TCR repertoire and CDR3 motif analyses depict the role of $\gamma\delta$ T cells in Ankylosing spondylitis. <i>EBioMedicine</i> , 2019, 47, 414-426.	6.1	32
11	Association of CD8+ T lymphocyte repertoire spreading with the severity of DRESS syndrome. <i>Scientific Reports</i> , 2015, 5, 9913.	3.3	27
12	Rab25 upregulation correlates with the proliferation, migration, and invasion of renal cell carcinoma. <i>Biochemical and Biophysical Research Communications</i> , 2015, 458, 745-750.	2.1	27
13	UHRF1 is required for basal stem cell proliferation in response to airway injury. <i>Cell Discovery</i> , 2017, 3, 17019.	6.7	27
14	MicroRNA-146a promotes IgE class switch in B cells via upregulating 14-3-3 σ expression. <i>Molecular Immunology</i> , 2017, 92, 180-189.	2.2	26
15	miR-21a in exosomes from Lewis lung carcinoma cells accelerates tumor growth through targeting PDCD4 to enhance expansion of myeloid-derived suppressor cells. <i>Oncogene</i> , 2020, 39, 6354-6369.	5.9	23
16	Collaboration between Distinct Rab Small GTPase Trafficking Circuits Mediates Bacterial Clearance from the Bladder Epithelium. <i>Cell Host and Microbe</i> , 2017, 22, 330-342.e4.	11.0	22
17	Proteome screening of pleural effusions identifies IL1A as a diagnostic biomarker for non-small cell lung cancer. <i>Biochemical and Biophysical Research Communications</i> , 2015, 457, 177-182.	2.1	21
18	Depletion of Rab32 decreases intracellular lipid accumulation and induces lipolysis through enhancing ATGL expression in hepatocytes. <i>Biochemical and Biophysical Research Communications</i> , 2016, 471, 492-496.	2.1	20

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19	Glimpse of natural selection of long-lived T-cell clones in healthy life. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 9858-9863.	7.1	19
20	Type I Interferon Therapy Limits CNS Autoimmunity by Inhibiting CXCR3-Mediated Trafficking of Pathogenic Effector T Cells. Cell Reports, 2019, 28, 486-497.e4.	6.4	19
21	The tumor microenvironment disarms CD8 ⁺ T lymphocyte function via a miR-26a-EZH2 axis. Oncoimmunology, 2016, 5, e1245267.	4.6	15
22	MicroRNA-146a Overexpression Impairs the Positive Selection during T Cell Development. Frontiers in Immunology, 2017, 8, 2006.	4.8	15
23	Nac1 promotes self-renewal of embryonic stem cells through direct transcriptional regulation of c-Myc. Oncotarget, 2017, 8, 47607-47618.	1.8	15
24	The Dynll1-Cox4i1 Complex Regulates Intracellular Pathogen Clearance via Release of Mitochondrial Reactive Oxygen Species. Infection and Immunity, 2020, 88, .	2.2	12
25	Conversion of effector CD4 ⁺ T cells to a CD8 ⁺ MHC II-recognizing lineage. Cellular and Molecular Immunology, 2021, 18, 150-161.	10.5	12
26	The kinase p38 β functions in dendritic cells to regulate Th2-cell differentiation and allergic inflammation. , 2022, 19, 805-819.		12
27	Streamlined Low-Input Transcriptomics through EASY-RNAseq. Journal of Molecular Biology, 2019, 431, 5075-5085.	4.2	9
28	MiR-26a targets EphA2 to resist intracellular Listeria monocytogenes in macrophages. Molecular Immunology, 2020, 128, 69-78.	2.2	8
29	TIPS: trajectory inference of pathway significance through pseudotime comparison for functional assessment of single-cell RNAseq data. Briefings in Bioinformatics, 2021, 22, .	6.5	8
30	VisTCR: An Interactive Software for T Cell Repertoire Sequencing Data Analysis. Frontiers in Genetics, 2020, 11, 771.	2.3	7
31	Rab32-related antimicrobial pathway is involved in the progression of dextran sodium sulfate-induced colitis. FEBS Open Bio, 2018, 8, 1658-1668.	2.3	6
32	TCR repertoire characteristics predict clinical response to adoptive CTL therapy against nasopharyngeal carcinoma. Oncoimmunology, 2021, 10, 1955545.	4.6	6
33	Mapping Cell Phenomics with Multiparametric Flow Cytometry Assays. Phenomics, 2022, 2, 272-281.	2.9	5
34	Hsa-miR-31 Governs T-Cell Homeostasis in HIV Protection via IFN- β -Stat1-T-Bet Axis. Frontiers in Immunology, 2021, 12, 771279.	4.8	3
35	Mapping the spatial distribution of T cells in repertoire dimension. Molecular Immunology, 2021, 138, 161-171.	2.2	1