Makoto Kurachi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5842691/publications.pdf

Version: 2024-02-01

25 papers 7,822 citations

430442 18 h-index 25 g-index

25 all docs

25 docs citations

25 times ranked

13460 citing authors

#	Article	IF	CITATIONS
1	Molecular and cellular insights into T cell exhaustion. Nature Reviews Immunology, 2015, 15, 486-499.	10.6	3,159
2	Epigenetic stability of exhausted T cells limits durability of reinvigoration by PD-1 blockade. Science, 2016, 354, 1160-1165.	6.0	939
3	The epigenetic landscape of T cell exhaustion. Science, 2016, 354, 1165-1169.	6.0	694
4	Bioenergetic Insufficiencies Due to Metabolic Alterations Regulated by the Inhibitory Receptor PD-1 Are an Early Driver of CD8 + T Cell Exhaustion. Immunity, 2016, 45, 358-373.	6.6	560
5	Developmental Relationships of Four Exhausted CD8+ T Cell Subsets Reveals Underlying Transcriptional and Epigenetic Landscape Control Mechanisms. Immunity, 2020, 52, 825-841.e8.	6.6	497
6	TCF-1-Centered Transcriptional Network Drives an Effector versus Exhausted CD8ÂT Cell-Fate Decision. Immunity, 2019, 51, 840-855.e5.	6.6	409
7	The transcription factor BATF operates as an essential differentiation checkpoint in early effector CD8+ T cells. Nature Immunology, 2014, 15, 373-383.	7.0	289
8	CXCR6 regulates localization of tissue-resident memory CD8 T cells to the airways. Journal of Experimental Medicine, 2019, 216, 2748-2762.	4.2	216
9	Janus kinase inhibition lessens inflammation and ameliorates disease in murine models of hemophagocytic lymphohistiocytosis. Blood, 2016, 127, 1666-1675.	0.6	207
10	CD8+ T cell exhaustion. Seminars in Immunopathology, 2019, 41, 327-337.	2.8	169
11	Lineage-Determining Transcription Factor TCF-1 Initiates the Epigenetic Identity of T Cells. Immunity, 2018, 48, 243-257.e10.	6.6	164
12	InÂvivo CD8+ TÂcell CRISPR screening reveals control by Fli1 in infection and cancer. Cell, 2021, 184, 1262-1280.e22.	13.5	107
13	Long-Term Persistence of Exhausted CD8ÂT Cells in Chronic Infection Is Regulated by MicroRNA-155. Cell Reports, 2018, 23, 2142-2156.	2.9	84
14	miR-150 Regulates Memory CD8ÂT Cell Differentiation via c-Myb. Cell Reports, 2017, 20, 2584-2597.	2.9	70
15	Group 1 Innate Lymphoid Cell Lineage Identity Is Determined by a cis-Regulatory Element Marked by a Long Non-coding RNA. Immunity, 2017, 47, 435-449.e8.	6.6	57
16	Inhibitory signaling sustains a distinct early memory CD8 $<$ sup $>+<$ /sup $>$ T cell precursor that is resistant to DNA damage. Science Immunology, 2021, 6, .	5.6	52
17	Optimized retroviral transduction of mouse T cells for in vivo assessment of gene function. Nature Protocols, 2017, 12, 1980-1998.	5. 5	47
18	Hidden Caveat of Inducible Cre Recombinase. Immunity, 2019, 51, 591-592.	6.6	23

#	Article	IF	CITATIONS
19	Batf-mediated epigenetic control of effector CD8 ⁺ T cell differentiation. Science Immunology, 2022, 7, eabi4919.	5.6	19
20	Trib1 regulates T cell differentiation during chronic infection by restraining the effector program. Journal of Experimental Medicine, 2020, 217, .	4.2	15
21	Transient Depletion of CD4+ Cells Induces Remodeling of the TCR Repertoire in Gastrointestinal Cancer. Cancer Immunology Research, 2021, 9, 624-636.	1.6	13
22	MCPIP1 reduces HBV-RNA by targeting its epsilon structure. Scientific Reports, 2020, 10, 20763.	1.6	10
23	Generation of tumor antigen-specific murine CD8+ T cells with enhanced anti-tumor activity via highly efficient CRISPR/Cas9 genome editing. International Immunology, 2018, 30, 141-154.	1.8	9
24	MicroRNA-29a attenuates CD8 T cell exhaustion and induces memory-like CD8 T cells during chronic infection. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2106083119.	3.3	7
25	Leukocyte cell-derived chemotaxin 2 is an antiviral regulator acting through the proto-oncogene MET. Nature Communications, 2022, 13, .	5.8	6