Guangan Hu

List of Publications by Year in descending order

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623734 839539 20 964 14 18 citations h-index g-index papers 21 21 21 1813 all docs docs citations times ranked citing authors

#	Article	IF	Citations
1	Ovarian Cancer Ascites Inhibits Transcriptional Activation of NK Cells Partly through CA125. Journal of Immunology, 2022, 208, 2227-2238.	0.8	6
2	Memory-like NK cells armed with a neoepitope-specific CAR exhibit potent activity against NPM1 mutated acute myeloid leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119 , .	7.1	44
3	High-throughput phenotypic screen and transcriptional analysis identify new compounds and targets for macrophage reprogramming. Nature Communications, 2021, 12, 773.	12.8	62
4	MFSD7C switches mitochondrial ATP synthesis to thermogenesis in response to heme. Nature Communications, 2020, 11, 4837.	12.8	21
5	LILRB3 (ILT5) is a myeloid cell checkpoint that elicits profound immunomodulation. JCI Insight, 2020, 5, .	5.0	26
6	Vitamin D3-vitamin D receptor axis suppresses pulmonary emphysema by maintaining alveolar macrophage homeostasis and function. EBioMedicine, 2019, 45, 563-577.	6.1	26
7	Induction and Therapeutic Targeting of Human NPM1c+ Myeloid Leukemia in the Presence of Autologous Immune System in Mice. Journal of Immunology, 2019, 202, 1885-1894.	0.8	7
8	Cyclophosphamide Enhances Cancer Antibody Immunotherapy in the Resistant Bone Marrow Niche by Modulating Macrophage Fcl³R Expression. Cancer Immunology Research, 2019, 7, 1876-1890.	3.4	23
9	Differentiation of Club Cells to Alveolar Epithelial Cells In Vitro. Scientific Reports, 2017, 7, 41661.	3.3	44
10	Antibiotics induce polarization of pleural macrophages to M2-like phenotype in patients with tuberculous pleuritis. Scientific Reports, 2017, 7, 14982.	3.3	14
11	Integrated analysis of the Plasmodium species transcriptome. EBioMedicine, 2016, 7, 255-266.	6.1	55
12	Information-dense analysis for information-dense understanding. Translational Cancer Research, 2016, 5, S1078-S1081.	1.0	0
13	The Dawn of Systems Cancer Immunotherapy. Archives of Surgical Oncology, 2015, 01, .	0.1	O
14	A genome-wide regulatory network identifies key transcription factors for memory CD8+ T-cell development. Nature Communications, 2013, 4, 2830.	12.8	103
15	Defining species specific genome differences in malaria parasites. BMC Genomics, 2010, 11, 128.	2.8	12
16	Transcriptional profiling of growth perturbations of the human malaria parasite Plasmodium falciparum. Nature Biotechnology, 2010, 28, 91-98.	17.5	196
17	Crystal Structure of the FK506 Binding Domain of <i>Plasmodium falciparum</i> FKBP35 in Complex with FK506. Biochemistry, 2008, 47, 5951-5961.	2.5	35
18	The transcriptome of <i>Plasmodium vivax</i> reveals divergence and diversity of transcriptional regulation in malaria parasites. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 16290-16295.	7.1	234

#	Article	IF	CITATIONS
19	Selection of long oligonucleotides for gene expression microarrays using weighted rank-sum strategy. BMC Bioinformatics, 2007, 8, 350.	2.6	50
20	Identification of One Intron Loss and Phylogenetic Evolution of Dfak Gene in the Drosophila melanogaster Species Group. Genetica, 2005, 125, 223-230.	1.1	6