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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular determinants of response to PD-L1 blockade across tumor types. Nature Communications, 2021, 12, 3969.	12.8	79
2	Anti–PD-L1 Treatment Results in Functional Remodeling of the Macrophage Compartment. Cancer Research, 2019, 79, 1493-1506.	0.9	118
3	Coexpression of Inhibitory Receptors Enriches for Activated and Functional CD8+ T Cells in Murine Syngeneic Tumor Models. Cancer Immunology Research, 2019, 7, 963-976.	3.4	36
4	Therapeutic resistance and susceptibility is shaped by cooperative multi-compartment tumor adaptation. Cell Death and Differentiation, 2019, 26, 2416-2429.	11.2	25
5	TGFβ attenuates tumour response to PD-L1 blockade by contributing to exclusion of T cells. Nature, 2018, 554, 544-548.	27.8	3,359
6	Clinical activity and molecular correlates of response to atezolizumab alone or in combination with bevacizumab versus sunitinib in renal cell carcinoma. Nature Medicine, 2018, 24, 749-757.	30.7	900
7	Atezolizumab in patients with locally advanced and metastatic urothelial carcinoma who have progressed following treatment with platinum-based chemotherapy: a single-arm, multicentre, phase 2 trial. Lancet, The, 2016, 387, 1909-1920.	13.7	3,077
8	Immune biomarkers associated with clinical benefit from atezolizumab (MPDL3280a; anti-PD-L1) in advanced urothelial bladder cancer (UBC). , 2015, 3, .		12
9	Whole genome sequences of 2 octogenarians with sustained cognitive abilities. Neurobiology of Aging, 2015, 36, 1435-1438.	3.1	1
10	Rituximab Efficiently Depletes Increased CD20-Expressing T Cells in Multiple Sclerosis Patients. Journal of Immunology, 2014, 193, 580-586.	0.8	223
11	The autoimmune disease-associated transcription factors EOMES and TBX21 are dysregulated in multiple sclerosis and define a molecular subtype of disease. Clinical Immunology, 2014, 151, 16-24.	3.2	49
12	Blood RNA profiling in a large cohort of multiple sclerosis patients and healthy controls. Human Molecular Genetics, 2013, 22, 4194-4205.	2.9	81
13	A Genome-Wide RNA Interference Screen Identifies Caspase 4 as a Factor Required for Tumor Necrosis Factor Alpha Signaling. Molecular and Cellular Biology, 2012, 32, 3372-3381.	2.3	36
14	In depth comparison of an individual's DNA and its lymphoblastoid cell line using whole genome sequencing. BMC Genomics, 2012, 13, 477.	2.8	34
15	An RNAi screen identifies USP2 as a factor required for TNFâ€Î±â€induced NFâ€ÎºB signaling. International Journal of Cancer, 2011, 129, 607-618.	5.1	49