

# Dorothee Nickles

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11619693/publications.pdf>

Version: 2024-02-01

15  
papers

8,079  
citations

687363

13  
h-index

1058476

14  
g-index

15  
all docs

15  
docs citations

15  
times ranked

13332  
citing authors

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