

# Robert T Jones

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11290109/publications.pdf>

Version: 2024-02-01

12  
papers

2,209  
citations

933447

10  
h-index

1281871

11  
g-index

12  
all docs

12  
docs citations

12  
times ranked

4772  
citing authors

#	ARTICLE	IF	CITATIONS
1	TRIM28 is a transcriptional activator of the mutant TERT promoter in human bladder cancer. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	24
2	Inhibition of the CCL2 receptor, CCR2, enhances tumor response to immune checkpoint therapy. Communications Biology, 2020, 3, 720.	4.4	82
3	ROS1 Gene Rearrangements Are Associated With an Elevated Risk of Peridiagnosis Thromboembolic Events. Journal of Thoracic Oncology, 2019, 14, 596-605.	1.1	56
4	Targeting DDR2 enhances tumor response to anti-PD-1 immunotherapy. Science Advances, 2019, 5, eaav2437.	10.3	92
5	Two methods of prediction signatures. Nature Reviews Urology, 2018, 15, 340-342.	3.8	0
6	Pharmacogenomics. Urologic Clinics of North America, 2016, 43, 77-86.	1.8	9
7	Genomic Characterization of Brain Metastases Reveals Branched Evolution and Potential Therapeutic Targets. Cancer Discovery, 2015, 5, 1164-1177.	9.4	821
8	Cross-reactivity of the BRAF VE1 antibody with epitopes in axonemal dyneins leads to staining of cilia. Modern Pathology, 2015, 28, 596-606.	5.5	55
9	Angiomatous meningiomas have a distinct genetic profile with multiple chromosomal polysomies including polysomy of chromosome 5. Oncotarget, 2014, 5, 10596-10606.	1.8	65
10	Sporadic hemangioblastomas are characterized by cryptic VHL inactivation. Acta Neuropathologica Communications, 2014, 2, 167.	5.2	65
11	Exome sequencing identifies BRAF mutations in papillary craniopharyngiomas. Nature Genetics, 2014, 46, 161-165.	21.4	408
12	Genomic sequencing of meningiomas identifies oncogenic SMO and AKT1 mutations. Nature Genetics, 2013, 45, 285-289.	21.4	532