

Andrew C Warner

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/489564/publications.pdf>

Version: 2024-02-01

11
papers

401
citations

1307594

7
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

1360
citing authors

#	ARTICLE	IF	CITATIONS
1	TCR-engineered T cells targeting E7 for patients with metastatic HPV-associated epithelial cancers. <i>Nature Medicine</i> , 2021, 27, 419-425.	30.7	156
2	Distribution and Functional Consequences of Somatic MAP2K1 Variants in Affected Skin Associated with Bone Lesions in Melorheostosis. <i>Journal of Investigative Dermatology</i> , 2021, 141, 688-692.e11.	0.7	3
3	Prostate cancer evolution from multilineage primary to single lineage metastases with implications for liquid biopsy. <i>Nature Communications</i> , 2020, 11, 5070.	12.8	44
4	Naturally Acquired Mouse Kidney Parvovirus Infection Produces a Persistent Interstitial Nephritis in Immunocompetent Laboratory Mice. <i>Veterinary Pathology</i> , 2020, 57, 915-925.	1.7	16
5	Characteristics of Breast Ducts in Normal-Risk and High-risk Women and Their Relationship to Ductal Cytologic Atypia. <i>Cancer Prevention Research</i> , 2020, 13, 1027-1036.	1.5	2
6	Cancer targeting by TCR gene-engineered T cells directed against Kita-Kyushu Lung Cancer Antigen-1. , 2019, 7, 229.		27
7	Laser Microdissection Workflow for Isolating Nucleic Acids from Fixed and Frozen Tissue Samples. <i>Methods in Molecular Biology</i> , 2018, 1723, 33-93.	0.9	3
8	Gammaherpesvirus infection and malignant disease in rhesus macaques experimentally infected with SIV or SHIV. <i>PLoS Pathogens</i> , 2018, 14, e1007130.	4.7	10
9	Telomere Length and Survival of Patients with Hepatocellular Carcinoma in the United States. <i>PLoS ONE</i> , 2016, 11, e0166828.	2.5	10
10	An Improved Breast Epithelial Sampling Method for Molecular Profiling and Biomarker Analysis in Women at Risk for Breast Cancer. <i>Breast Cancer: Basic and Clinical Research</i> , 2015, 9, BCBCR.S23577.	1.1	7
11	Characterizing the genetic basis of methylome diversity in histologically normal human lung tissue. <i>Nature Communications</i> , 2014, 5, 3365.	12.8	123