

Chenyue Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

331
citations

1040056

9
h-index

888059

17
g-index

24
all docs

24
docs citations

24
times ranked

607
citing authors

#	ARTICLE	IF	CITATIONS
1	Differences in Stage of Cancer at Diagnosis, Treatment, and Survival by Race and Ethnicity Among Leading Cancer Types. <i>JAMA Network Open</i> , 2020, 3, e202950.	5.9	71
2	Na ⁺ /K ⁺ -ATPase β 1 subunit, a novel therapeutic target for hepatocellular carcinoma. <i>Oncotarget</i> , 2015, 6, 28183-28193.	1.8	39
3	Prognostic nutritional index serves as a predicative marker of survival and associates with systemic inflammatory response in metastatic intrahepatic cholangiocarcinoma. <i>OncoTargets and Therapy</i> , 2016, Volume 9, 6417-6423.	2.0	33
4	Serum liver enzymes serve as prognostic factors in patients with intrahepatic cholangiocarcinoma. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 1441-1449.	2.0	33
5	TERT mutations correlate with higher TMB value and unique tumor microenvironment and may be a potential biomarker for anti-CTLA4 treatment. <i>Cancer Medicine</i> , 2020, 9, 7151-7160.	2.8	33
6	CD44, a marker of cancer stem cells, is positively correlated with PD-L1 expression and immune cells infiltration in lung adenocarcinoma. <i>Cancer Cell International</i> , 2020, 20, 583.	4.1	31
7	Integrative analysis of genomic alteration, immune cells infiltration and prognosis of lung squamous cell carcinoma (LUSC) to identify smoking-related biomarkers. <i>International Immunopharmacology</i> , 2020, 89, 107053.	3.8	11
8	LHPP suppresses tumorigenesis of intrahepatic cholangiocarcinoma by inhibiting the TGF β 2/smad signaling pathway. <i>International Journal of Biochemistry and Cell Biology</i> , 2021, 132, 105845.	2.8	11
9	Identification and Validation of a Tumor Microenvironment-Related Gene Signature in Hepatocellular Carcinoma Prognosis. <i>Frontiers in Genetics</i> , 2021, 12, 717319.	2.3	10
10	KEAP1-NFE2L2 ^{mutant} NSCLC and Immune Checkpoint Inhibitors: A Large Database Analysis. <i>Journal of Thoracic Oncology</i> , 2020, 15, e85-e86.	1.1	9
11	Construction of immune-related and prognostic lncRNA clusters and identification of their immune and genomic alterations characteristics in lung adenocarcinoma samples. <i>Aging</i> , 2020, 12, 9868-9881.	3.1	9
12	Patients With BRAF-Mutant NSCLC May Not Benefit From Immune Checkpoint Inhibitors: A Population-Based Study. <i>JTO Clinical and Research Reports</i> , 2020, 1, 100006.	1.1	8
13	Male patients with TERT mutation may be more likely to benefit from immunotherapy, especially for melanoma. <i>Aging</i> , 2020, 12, 17288-17294.	3.1	7
14	Genetic alternations and immune characteristics in patients with small cell lung cancer. <i>Cancer Communications</i> , 2021, 41, 1075-1078.	9.2	6
15	Prognostic values of common clinical parameters in advanced pancreatic ductal adenocarcinoma: a large multicenter cohort study of ten years. <i>Discovery Medicine</i> , 2018, 25, 91-98.	0.5	6
16	Regulation of immune microenvironment may enable MET-altered NSCLC patients to benefit from immune checkpoint inhibitors. <i>Lung Cancer</i> , 2021, 154, 221-223.	2.0	4
17	Is chronic hepatitis B infection a protective factor for the progression of advanced pancreatic ductal adenocarcinoma? An analysis from a large multicenter cohort study. <i>Oncotarget</i> , 2016, 7, 85603-85612.	1.8	3
18	Genomic Variations and Immune-Related Features of TMB, PD-L1 Expression and CD8 ⁺ T Cell Infiltration in Chinese Pulmonary Sarcomatoid Carcinoma. <i>International Journal of General Medicine</i> , 2022, Volume 15, 4209-4220.	1.8	3

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19	Loss of heterozygosity related to TMB and TNB may predict PFS for patients with SCLC received the first line setting. <i>Journal of Translational Medicine</i> , 2021, 19, 385.	4.4	2
20	The source of the tumor tissue should be taken into consideration when distinguishing tumor mutational burden scores. <i>Lung Cancer</i> , 2021, 154, 214-215.	2.0	1
21	Identification of a Gene Signature Closely Related to Immunosuppressive Tumour Microenvironment Predicting Prognosis of Patients in EGFR Mutant Lung Adenocarcinoma. <i>Frontiers in Oncology</i> , 2021, 11, 732841.	2.8	1
22	Therapeutic utility of Lung-MAP: ushering into an era of genomic and biomarker-driven clinical trials. <i>Signal Transduction and Targeted Therapy</i> , 2021, 6, 141.	17.1	0
23	The negative relationship between patients with NSCLC harbored STK11/KEAP1 copy number variation and immune microenvironment infiltration. <i>Journal of Translational Medicine</i> , 2021, 19, 259.	4.4	0