

Junyu Long

List of Publications by Year in descending order

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Version: 2024-02-01

55
papers

1,686
citations

430754

18
h-index

345118

36
g-index

57
all docs

57
docs citations

57
times ranked

1882
citing authors

#	ARTICLE	IF	CITATIONS
1	A real-world study of the efficacy and safety of anti-PD-1 antibodies plus lenvatinib in patients with advanced gallbladder cancer. <i>Cancer Immunology, Immunotherapy</i> , 2022, , 1.	2.0	8
2	A mutation-based gene set predicts survival benefit after immunotherapy across multiple cancers and reveals the immune response landscape. <i>Genome Medicine</i> , 2022, 14, 20.	3.6	26
3	Lenvatinib Beyond First-Line Therapy in Patients With Advanced Biliary Tract Carcinoma. <i>Frontiers in Oncology</i> , 2022, 12, 785535.	1.3	6
4	eRNAs Identify Immune Microenvironment Patterns and Provide a Novel Prognostic Tool in Acute Myeloid Leukemia. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, 877117.	1.6	1
5	Lenvatinib plus PD-1 inhibitors in 378 unresectable hepatocellular carcinoma: A large real-world study from two centers.. <i>Journal of Clinical Oncology</i> , 2022, 40, e16155-e16155.	0.8	2
6	Genomic characterization and translational immunotherapy of microsatellite instability-high (MSI-H) in cholangiocarcinoma.. <i>Journal of Clinical Oncology</i> , 2022, 40, 4101-4101.	0.8	0
7	Transcriptional landscape of cholangiocarcinoma revealed by weighted gene coexpression network analysis. <i>Briefings in Bioinformatics</i> , 2021, 22, .	3.2	46
8	The Efficacy and Safety of Apatinib Plus Camrelizumab in Patients With Previously Treated Advanced Biliary Tract Cancer: A Prospective Clinical Study. <i>Frontiers in Oncology</i> , 2021, 11, 646979.	1.3	23
9	Construction and validation of a prognostic signature using CNV-driven genes for hepatocellular carcinoma. <i>Annals of Translational Medicine</i> , 2021, 9, 765-765.	0.7	3
10	Comprehensive Analysis of Autophagy-Associated lncRNAs Reveal Potential Prognostic Prediction in Pancreatic Cancer. <i>Frontiers in Oncology</i> , 2021, 11, 596573.	1.3	7
11	Cell-free DNA copy number variations predict efficacy of immune checkpoint inhibitor-based therapy in hepatobiliary cancers. , 2021, 9, e001942.		22
12	Polygenic risk score: A promising predictor for hepatocellular carcinoma in the population with non-alcoholic fatty liver disease. <i>Journal of Hepatology</i> , 2021, 74, 1493-1494.	1.8	3
13	Targeted Next-Generation Sequencing Combined With Circulating-Free DNA Deciphers Spatial Heterogeneity of Resected Multifocal Hepatocellular Carcinoma. <i>Frontiers in Immunology</i> , 2021, 12, 673248.	2.2	4
14	Identification of a Novel Tumor Microenvironment Prognostic Signature for Advanced-Stage Serous Ovarian Cancer. <i>Cancers</i> , 2021, 13, 3343.	1.7	14
15	Association of chronic liver disease with the prognosis of COVID-19 patients. <i>Journal of Hepatology</i> , 2021, , .	1.8	1
16	Comprehensive exploration of tumor mutational burden and immune infiltration in diffuse glioma. <i>International Immunopharmacology</i> , 2021, 96, 107610.	1.7	9
17	Identification of NOTCH4 mutation as a response biomarker for immune checkpoint inhibitor therapy. <i>BMC Medicine</i> , 2021, 19, 154.	2.3	32
18	Downstaging and resection of hepatocellular carcinoma in patients with extrahepatic metastases after stereotactic therapy. <i>Hepatobiliary Surgery and Nutrition</i> , 2021, 10, 434-442.	0.7	28

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19	Genomic characterization of co-existing neoplasia and carcinoma lesions reveals distinct evolutionary paths of gallbladder cancer. <i>Nature Communications</i> , 2021, 12, 4753.	5.8	12
20	Racial/Ethnic Differences in the 21-Gene Recurrence Score Assay Among Women With Breast Cancer. <i>JAMA Oncology</i> , 2021, 7, 1248.	3.4	0
21	Genomic instability in lower-grade glioma: Prediction of prognosis based on lncRNA and immune infiltration. <i>Molecular Therapy - Oncolytics</i> , 2021, 22, 431-443.	2.0	6
22	Apatinib as non-first-line treatment in patients with Intrahepatic Cholangiocarcinoma. <i>Journal of Cancer</i> , 2021, 12, 1555-1562.	1.2	7
23	Mutational spectrum and precision oncology for biliary tract carcinoma. <i>Theranostics</i> , 2021, 11, 4585-4598.	4.6	39
24	Identification of TMB, CD8 T cell abundance, and homologous repair pathway mutation frequency as predictors of the benefit-toxicity ratio of anti-PD-1/PD-L1 therapy. <i>Clinical and Translational Medicine</i> , 2021, 11, e598.	1.7	0
25	Differential genes and scoring criteria among immunogenomic clusters of lower-grade gliomas. <i>International Immunopharmacology</i> , 2021, 101, 108376.	1.7	4
26	Gut microbiome is associated with the clinical response to anti-PD-1 based immunotherapy in hepatobiliary cancers. , 2021, 9, e003334.		101
27	Identification and Validation of Constructing the Prognostic Model With Four DNA Methylation-Driven Genes in Pancreatic Cancer. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 709669.	1.8	3
28	Application of the Lung Immune Prognostic Index From Research to Clinical Practice. <i>JAMA Oncology</i> , 2020, 6, 299.	3.4	0
29	Comprehensive analysis of tumour mutation burden and the immune microenvironment in hepatocellular carcinoma. <i>International Immunopharmacology</i> , 2020, 89, 107135.	1.7	17
30	Expression of p42.3 in non-small cell lung cancer. <i>Annals of Translational Medicine</i> , 2020, 8, 819-819.	0.7	7
31	Pembrolizumab combined with lenvatinib as non-first-line therapy in patients with refractory biliary tract carcinoma. <i>Hepatobiliary Surgery and Nutrition</i> , 2020, 9, 414-424.	0.7	93
32	Development and Validation of a Prognostic Nomogram for Gastric Cancer Based on DNA Methylation-Driven Differentially Expressed Genes. <i>International Journal of Biological Sciences</i> , 2020, 16, 1153-1165.	2.6	45
33	Comparing the efficacy and safety of second-line therapies for advanced hepatocellular carcinoma: a network meta-analysis of phase III trials. <i>Therapeutic Advances in Gastroenterology</i> , 2020, 13, 175628482093248.	1.4	10
34	Dyslipidemia Might Be Associated with an Increased Risk of Osteoarthritis. <i>BioMed Research International</i> , 2020, 2020, 1-9.	0.9	20
35	CeRNA regulatory network-based analysis to study the roles of noncoding RNAs in the pathogenesis of intrahepatic cholangiocellular carcinoma. <i>Aging</i> , 2020, 12, 1047-1086.	1.4	27
36	T lymphocytes in hepatocellular carcinoma immune microenvironment: insights into human immunology and immunotherapy. <i>American Journal of Cancer Research</i> , 2020, 10, 4585-4606.	1.4	8

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37	CK20 and lymph node involvement predict adverse outcome of malignant intraductal papillary neoplasm of the bile duct. <i>Histology and Histopathology</i> , 2020, 35, 449-456.	0.5	2
38	Construction and Investigation of a lncRNA-Associated ceRNA Regulatory Network in Cholangiocarcinoma. <i>Frontiers in Oncology</i> , 2019, 9, 649.	1.3	32
39	DNA methylation-driven genes for constructing diagnostic, prognostic, and recurrence models for hepatocellular carcinoma. <i>Theranostics</i> , 2019, 9, 7251-7267.	4.6	99
40	Transcriptomic analysis and identification of prognostic biomarkers in cholangiocarcinoma. <i>Oncology Reports</i> , 2019, 42, 1833-1842.	1.2	16
41	Alterations in DNA Damage Repair Genes in Primary Liver Cancer. <i>Clinical Cancer Research</i> , 2019, 25, 4701-4711.	3.2	74
42	Construction and comprehensive analysis of a ceRNA network to reveal potential prognostic biomarkers for hepatocellular carcinoma. <i>Cancer Cell International</i> , 2019, 19, 90.	1.8	90
43	Combination regimens with PD-1/PD-L1 immune checkpoint inhibitors for gastrointestinal malignancies. <i>Journal of Hematology and Oncology</i> , 2019, 12, 42.	6.9	58
44	Development and validation of a TP53-associated immune prognostic model for hepatocellular carcinoma. <i>EBioMedicine</i> , 2019, 42, 363-374.	2.7	257
45	Comprehensive analysis of a ceRNA network reveals potential prognostic cytoplasmic lncRNAs involved in HCC progression. <i>Journal of Cellular Physiology</i> , 2019, 234, 18837-18848.	2.0	103
46	The diagnostic and prognostic role of RhoA in hepatocellular carcinoma. <i>Aging</i> , 2019, 11, 5158-5172.	1.4	16
47	Precision oncology for gallbladder cancer: insights from genetic alterations and clinical practice. <i>Annals of Translational Medicine</i> , 2019, 7, 467-467.	0.7	24
48	Metabolic syndrome and the risk of cholangiocarcinoma: a hospital-based case–control study in China. <i>Cancer Management and Research</i> , 2018, Volume 10, 3849-3855.	0.9	13
49	A four-gene-based prognostic model predicts overall survival in patients with hepatocellular carcinoma. <i>Journal of Cellular and Molecular Medicine</i> , 2018, 22, 5928-5938.	1.6	128
50	Classification of gallbladder cancer by assessment of CD8+ TIL and PD-L1 expression. <i>BMC Cancer</i> , 2018, 18, 766.	1.1	42
51	PD-1/PD-L blockade in gastrointestinal cancers: lessons learned and the road toward precision immunotherapy. <i>Journal of Hematology and Oncology</i> , 2017, 10, 146.	6.9	77
52	Identification of hub genes involved in the development of hepatocellular carcinoma by transcriptome sequencing. <i>Oncotarget</i> , 2017, 8, 60358-60367.	0.8	8
53	Systematic review and meta-analysis: cholecystectomy and the risk of cholangiocarcinoma. <i>Oncotarget</i> , 2017, 8, 59648-59657.	0.8	13
54	Development and Validation of a Prognostic Nomogram for Gastric Cancer Based on DNA Methylation-Driven Genes. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0

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55	Lenvatinib Plus PD-1 Blockade in Refractory or Unresectable Biliary Tract Carcinoma: Insights from Real-World Evidences. SSRN Electronic Journal, 0, , .	0.4	0