## Henghui Zhang

List of Publications by Year in descending order

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82 papers 1,344 citations

394421 19 h-index 31 g-index

88 all docs 88 docs citations

88 times ranked 2374 citing authors

#	Article	IF	Citations
1	Peritumoural neutrophils negatively regulate adaptive immunity via the PD-L1/PD-1 signalling pathway in hepatocellular carcinoma. Journal of Experimental and Clinical Cancer Research, 2015, 34, 141.	8.6	186
2	S100A4 promotes liver fibrosis via activation of hepatic stellate cells. Journal of Hepatology, 2015, 62, 156-164.	3.7	133
3	VEGF-D: a novel biomarker for detection of COVID-19 progression. Critical Care, 2020, 24, 373.	5.8	77
4	Diagnostic value of circulating cell-free DNA levels for hepatocellular carcinoma. International Journal of Infectious Diseases, 2018, 67, 92-97.	3.3	66
5	PD-L1 expression and tumor mutational burden status for prediction of response to chemotherapy and targeted therapy in non-small cell lung cancer. Journal of Experimental and Clinical Cancer Research, 2019, 38, 193.	8.6	61
6	Tumor copy-number alterations predict response to immune-checkpoint-blockade in gastrointestinal cancer., 2020, 8, e000374.		43
7	Mutational spectrum and precision oncology for biliary tract carcinoma. Theranostics, 2021, 11, 4585-4598.	10.0	39
8	Storm of soluble immune checkpoints associated with disease severity of COVID-19. Signal Transduction and Targeted Therapy, 2020, 5, 192.	17.1	38
9	Massive PD-L1 and CD8 double positive TILs characterize an immunosuppressive microenvironment with high mutational burden in lung cancer., 2021, 9, e002356.		35
10	Persistent High Percentage of HLA-DR+CD38high CD8+ T Cells Associated With Immune Disorder and Disease Severity of COVID-19. Frontiers in Immunology, 2021, 12, 735125.	4.8	35
11	Sendai Virus Induces Persistent Olfactory Dysfunction in a Murine Model of PVOD via Effects on Apoptosis, Cell Proliferation, and Response to Odorants. PLoS ONE, 2016, 11, e0159033.	2.5	34
12	Gutâ€derived lymphocyte recruitment to liver and induce liver injury in nonâ€alcoholic fatty liver disease mouse model. Journal of Gastroenterology and Hepatology (Australia), 2016, 31, 676-684.	2.8	31
13	Molecular profiles and tumor mutational burden analysis in Chinese patients with gynecologic cancers. Scientific Reports, 2018, 8, 8990.	3.3	31
14	NOTCH3 is a Prognostic Factor and Is Correlated With Immune Tolerance in Gastric Cancer. Frontiers in Oncology, 2020, 10, 574937.	2.8	29
15	DNA Damage Repair Gene Mutations Are Indicative of a Favorable Prognosis in Colorectal Cancer Treated With Immune Checkpoint Inhibitors. Frontiers in Oncology, 2020, 10, 549777.	2.8	26
16	The transcription factor RFX5 is a transcriptional activator of the TPP1 gene in hepatocellular carcinoma. Oncology Reports, 2017, 37, 289-296.	2.6	25
17	Mutational Landscape of cfDNA Identifies Distinct Molecular Features Associated With Therapeutic Response to First-Line Platinum-Based Doublet Chemotherapy in Patients with Advanced NSCLC. Theranostics, 2017, 7, 4753-4762.	10.0	25
18	Serological Markers Associated With Response to Immune Checkpoint Blockade in Metastatic Gastrointestinal Tract Cancer. JAMA Network Open, 2019, 2, e197621.	5.9	25

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19	Targeted Therapy with Anlotinib for a Patient with an Oncogenic <i>FGFR3</i> - <i>TACC3</i> Fusion and Recurrent Glioblastoma. Oncologist, 2021, 26, 173-177.	3.7	23
20	Prediction of immune checkpoint inhibition with immune oncology-related gene expression in gastrointestinal cancer using a machine learning classifier., 2020, 8, e000631.		22
21	Cell-free DNA copy number variations predict efficacy of immune checkpoint inhibitor-based therapy in hepatobiliary cancers., 2021, 9, e001942.		22
22	Deep sequencing of hepatitis B virus basal core promoter and precore mutants in HBeAg-positive chronic hepatitis B patients. Scientific Reports, 2015, 5, 17950.	3.3	21
23	The frequency and inter-relationship of PD-L1 expression and tumour mutational burden across multiple types of advanced solid tumours in China. Experimental Hematology and Oncology, 2020, 9, 17.	5.0	21
24	A genomic mutation signature predicts the clinical outcomes of immunotherapy and characterizes immunophenotypes in gastrointestinal cancer. Npj Precision Oncology, 2021, 5, 36.	5.4	20
25	Peripheral cytokine levels as predictive biomarkers of benefit from immune checkpoint inhibitors in cancer therapy. Biomedicine and Pharmacotherapy, 2020, 129, 110457.	<b>5.</b> 6	19
26	Genomic features and tumor immune microenvironment alteration in NSCLC treated with neoadjuvant PD-1 blockade. Npj Precision Oncology, 2022, 6, 2.	5.4	17
27	Dexamethasone affects mouse olfactory mucosa gene expression and attenuates genes related to neurite outgrowth. International Forum of Allergy and Rhinology, 2015, 5, 907-918.	2.8	16
28	The mutational pattern of homologous recombination-related (HRR) genes in Chinese colon cancer and its relevance to immunotherapy responses. Aging, 2021, 13, 2365-2378.	3.1	15
29	The CD68+ macrophages to CD8+ T-cell ratio is associated with clinical outcomes in hepatitis B virus (HBV)-related hepatocellular carcinoma. Hpb, 2021, 23, 1061-1071.	0.3	12
30	Germline HLA-B evolutionary divergence influences the efficacy of immune checkpoint blockade therapy in gastrointestinal cancer. Genome Medicine, 2021, 13, 175.	8.2	12
31	PTCH1 mutation promotes antitumor immunity and the response to immune checkpoint inhibitors in colorectal cancer patients. Cancer Immunology, Immunotherapy, 2022, 71, 111-120.	4.2	11
32	Abnormal CD4 + T helper (Th) 1 cells and activated memory B cells are associated with type III asymptomatic mixed cryoglobulinemia in HCV infection. Virology Journal, 2015, 12, 100.	3.4	10
33	Molecular analysis of cellâ€free DNA identifies distinct molecular features in patients with chemosensitive and chemorefractory small cell lung cancer. Cancer Communications, 2019, 39, 1-5.	9.2	10
34	Prognostic impact of gene copy number instability and tumor mutation burden in patients with resectable gastric cancer. Cancer Communications, 2020, 40, 63-66.	9.2	10
35	Pathologic complete response to preoperative immunotherapy in a lung adenocarcinoma patient with bone metastasis: A case report. Thoracic Cancer, 2020, 11, 1094-1098.	1.9	10
36	A Machine Learning Approach Yields a Multiparameter Prognostic Marker in Liver Cancer. Cancer Immunology Research, 2021, 9, 337-347.	3.4	10

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37	EPHA5 mutations predict survival after immunotherapy in lung adenocarcinoma. Aging, 2021, 13, 598-618.	3.1	10
38	Characterization of the immune microenvironment in brain metastases from different solid tumors. Cancer Medicine, 2020, 9, 2299-2308.	2.8	9
39	Durable benefit from immunotherapy and accompanied lupus erythematosus in pancreatic adenocarcinoma with DNA repair deficiency., 2020, 8, e000463.		8
40	Application of plasma circulating cell-free DNA detection to the molecular diagnosis of hepatocellular carcinoma. American Journal of Translational Research (discontinued), 2019, 11, 1428-1445.	0.0	8
41	The mutation profiles of cell-free DNA in patients with oesophageal squamous cell carcinoma who were responsive and non-responsive to neoadjuvant chemotherapy. Journal of Thoracic Disease, 2020, 12, 4274-4283.	1.4	7
42	An Immunogram for an Individualized Assessment of the Antitumor Immune Response in Patients With Hepatocellular Carcinoma. Frontiers in Oncology, 2020, 10, 1189.	2.8	6
43	Short-term response to immune-chemotherapy and immune features of a ceritinib-resistant patient with <i>ROS1</i> -rearranged lung adenocarcinoma., 2021, 9, e001967.		6
44	Genetic and immune characteristics of sentinel lymph node metastases and multiple lymph node metastases compared to their matched primary breast tumours. EBioMedicine, 2021, 71, 103542.	6.1	6
45	The mutation of homologous recombination repair genetics is a potential biomarker for immunotherapy in microsatellite stable colon cancer Journal of Clinical Oncology, 2020, 38, 4076-4076.	1.6	6
46	The reduction in CD8+PD-1+ T cells in liver histological tissue is related to Pegylated IFN-α therapy outcomes in chronic hepatitis B patients. BMC Infectious Diseases, 2020, 20, 590.	2.9	4
47	Identification of immune checkpoint and cytokine signatures associated with the response to immune checkpoint blockade in gastrointestinal cancers. Cancer Immunology, Immunotherapy, 2021, 70, 2669-2679.	4.2	4
48	The mutational landscape of MSI-H and MSS colorectal cancer Journal of Clinical Oncology, 2019, 37, e15122-e15122.	1.6	4
49	Identifying tumor promoting genomic alterations in tumor-associated fibroblasts via retrovirus-insertional mutagenesis. Oncotarget, 2017, 8, 97231-97245.	1.8	4
50	Alterations in DNA damage response and repair genes as potential biomarkers for immune checkpoint blockade in gastrointestinal cancer. Cancer Biology and Medicine, 2022, 19, 1139-1149.	3.0	4
51	FOXP3 expression in FOXP3+ tumor cells promotes hepatocellular cells metastasis. Translational Cancer Research, 2020, 9, 5868-5881.	1.0	4
52	Case Report: Sarcoid-Like Reactions and Tertiary Lymphoid Structures Following Dual Checkpoint Inhibition in a Patient with Early-Stage Lung Adenocarcinoma. Frontiers in Immunology, 2022, 13, 794217.	4.8	4
53	Clinical Response to Neoadjuvant Immunotherapy Combined with Targeted Therapy and Chemotherapy in Oral Squamous Cell Carcinoma: Experience in Three Patients. OncoTargets and Therapy, 2022, Volume 15, 353-359.	2.0	4
54	Characterization of Hyperprogression After Immunotherapy in a Lung Adenocarcinoma Patient WithÂStrong Expression of Programmed Death LigandÂ1. Journal of Thoracic Oncology, 2020, 15, e4-e8.	1.1	3

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55	Genomic profile and immune microenvironment in patients with relapsed stage IA lung adenocarcinoma. Translational Oncology, 2021, 14, 100942.	3.7	3
56	Durable Complete Response to Pembrolizumab in Esophageal Squamous Cell Carcinoma With Divergent Microsatellite Status: A Case Report. Frontiers in Oncology, 2021, 11, 767957.	2.8	3
57	Abnormal phenotypic features of IgM+B cell subsets in patients with chronic hepatitis C virus infection. Experimental and Therapeutic Medicine, 2017, 14, 1846-1852.	1.8	2
58	Clinical Outcome and Molecular Analysis of a Chinese Patient With Lung Adenocarcinoma Harboring Rare EGFR Mutation V834L. Journal of Thoracic Oncology, 2018, 13, e189-e191.	1.1	2
59	Molecular Alterations in Circulating Cell-Free DNA in Patients with Colorectal Adenoma or Carcinoma Cancer Management and Research, 2020, Volume 12, 5159-5167.	1.9	2
60	Ki-67 versus MammaPrint/BluePrint for assessing luminal type breast cancer Journal of Clinical Oncology, 2020, 38, e13673-e13673.	1.6	2
61	Tumor copy number alteration (CNA) burden as a prognostic factor for overall survival in Chinese gastric cancers Journal of Clinical Oncology, 2019, 37, e15555-e15555.	1.6	2
62	Partial recovery of disturbed V-J pairing profiles of T-cell receptor in people living with HIV receiving long-term antiretroviral therapy. Science China Life Sciences, 2021, 64, 152-161.	4.9	1
63	Single cell transcriptome revealed tumor associated antigen (TAA) profile in lung adenocarcinoma (LUAD). Biomarker Research, 2021, 9, 41.	6.8	1
64	Comparative molecular profiling of distant metastatic and non-distant metastatic lung adenocarcinoma Neoplasma, 2021, 68, 253-261.	1.6	1
65	The mutational landscape of circulating cell free DNA in patients with esophageal squamous cell carcinoma response and non-response to neoadjuvant chemotherapy Journal of Clinical Oncology, 2018, 36, e16082-e16082.	1.6	1
66	Association of HLA class I genotype with outcomes of gastrointestinal cancer patients with immunotherapy Journal of Clinical Oncology, 2020, 38, e16551-e16551.	1.6	1
67	Immune Effective Score as a Predictor of Response to Neoadjuvant Trastuzumab Therapy and a Prognostic Indicator for HER2-Positive Breast Cancer. Current Oncology, 2022, 29, 283-293.	2.2	1
68	Soluble Immune-Related Proteins as New Candidate Serum Biomarkers for the Diagnosis and Progression of Lymphangioleiomyomatosis. Frontiers in Immunology, 2022, 13, 844914.	4.8	1
69	Features in genomics and tumor immune microenvironment in NSCLC treated with neoadjuvant PD-1 blockade Journal of Clinical Oncology, 2021, 39, 9063-9063.	1.6	0
70	PD-L1 expression and TMB status in newly diagnosed metastatic non-small cell lung cancer and their effect on prognosis after EGFR-TKI or platinum-based chemotherapy Journal of Clinical Oncology, 2018, 36, e24294-e24294.	1.6	0
71	Molecular profiles and tumor mutational burden analysis in Chinese patients with gynecologic cancers Journal of Clinical Oncology, 2018, 36, e17543-e17543.	1.6	0
72	Association of high copy number instability (CNI) score with prognosis in patients with gastric cancer after surgical resection Journal of Clinical Oncology, 2019, 37, e15552-e15552.	1.6	0

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73	The mutational profile analysis of different response to neoadjuvant chemoradiation therapy in local advanced esophageal squamous cell cancer patients Journal of Clinical Oncology, 2019, 37, e15560-e15560.	1.6	0
74	The cross talk between the molecular alterations and tumor immunity in the microenvironment in non-small-cell lung carcinoma Journal of Clinical Oncology, 2019, 37, e20043-e20043.	1.6	0
75	The mutational profile analysis of extramural vascular invasion in rectal cancer Journal of Clinical Oncology, 2019, 37, e15128-e15128.	1.6	O
76	Prediction of hepatocellular carcinoma patient survival using machine learning classification rules Journal of Clinical Oncology, 2019, 37, e15649-e15649.	1.6	0
77	Mutation of DNA damage repair genes confers an immune-privileged tumor microenvironment in colorectal cancer with a prognostic value Journal of Clinical Oncology, 2020, 38, 4080-4080.	1.6	0
78	Effect of TP53 mutation on antitumor immunity and responsiveness to immunotherapy in colorectal cancer Journal of Clinical Oncology, 2020, 38, e16014-e16014.	1.6	0
79	Application of MammaPrint test on Chinese patient with breast cancer Journal of Clinical Oncology, 2020, 38, e13671-e13671.	1.6	0
80	The association of CD8 <sup>+</sup> T-cell infiltration and PD-L1 expression with prognosis of Chinese pulmonary large cell neuroendocrine carcinoma Journal of Clinical Oncology, 2020, 38, e21003-e21003.	1.6	0
81	PD-1 or PD-L1 co-localizes with immune cells for prediction of prognosis in non-small cell lung cancer Journal of Clinical Oncology, 2020, 38, e21029-e21029.	1.6	0
82	Distinct transcriptional profiles in plasma exosomes associated with recurrence of nasopharyngeal carcinoma patients with standard treatment Journal of Clinical Oncology, 2020, 38, 6534-6534.	1.6	0