Hong Zou

List of Publications by Year
in descending order

43 papers	737 citations	567281 15 h-index	580821 25 g-index
45 all docs	45 docs citations	45 times ranked	1457 citing authors

#	Article	IF	Citations
1	SR-B1 and CD10 combined immunoprofile for differential diagnosis of metastatic clear cell renal cell carcinoma and clear cell carcinoma of the ovary. Journal of Molecular Histology, 2021, 52, 539-544.	2.2	4
2	Synergistic Inhibition of Drug-Resistant Colon Cancer Growth with PI3K/mTOR Dual Inhibitor BEZ235 and Nano-Emulsioned Paclitaxel via Reducing Multidrug Resistance and Promoting Apoptosis. International Journal of Nanomedicine, 2021, Volume 16, 2173-2186.	6.7	24
3	IFITM1, CD10, SMA, and h-caldesmon as a helpful combination in differential diagnosis between endometrial stromal tumor and cellular leiomyoma. BMC Cancer, 2021, 21, 1047.	2.6	9
4	Biological and prognostic value of ETV5 in high-grade serous ovarian cancer. Journal of Ovarian Research, 2021, 14, 149.	3.0	1
5	Long non‐coding RNA MIR31HG as a prognostic predictor for malignant cancers: A meta‐and bioinformatics analysis. Journal of Clinical Laboratory Analysis, 2021, , e24082.	2.1	6
6	Immune Infiltration in Gastric Cancer Microenvironment and Its Clinical Significance. Frontiers in Cell and Developmental Biology, 2021, 9, 762029.	3.7	10
7	ETV5-mediated upregulation of lncRNA CTBP1-DT as a ceRNA facilitates HGSOC progression by regulating miR-188-5p/MAP3K3 axis. Cell Death and Disease, 2021, 12, 1146.	6.3	9
8	Small-cell variant renal oncocytoma: Case report on its clinicopathological and genetic characteristics and literature review. Gene, 2020, 730, 144266.	2.2	5
9	Prognostic value of cripto-1 expression in non-small-cell lung cancer patients: a systematic review and meta-analysis. Biomarkers in Medicine, 2020, 14, 317-329.	1.4	2
10	Bioinformatics-based screening of key genes for transformation of liver cirrhosis to hepatocellular carcinoma. Journal of Translational Medicine, 2020, 18, 40.	4.4	39
11	Identification of hub genes associated with esophageal cancer progression using bioinformatics analysis. Oncology Letters, 2020, 20, 214.	1.8	1
12	Identification of hub genes associated with esophageal cancer progression using bioinformatics analysis. Oncology Letters, 2020, 20, 1-1.	1.8	3
13	Prognostic impact of tumor-associated macrophage infiltration in esophageal cancer: a meta-analysis. Future Oncology, 2019, 15, 2303-2317.	2.4	47
14	Overexpression of MAP3K3 promotes tumour growth through activation of the NF-κB signalling pathway in ovarian carcinoma. Scientific Reports, 2019, 9, 8401.	3.3	13
15	Cytomegalovirus infective gastritis in an immunocompetent host misdiagnosed as malignancy on upper gastrointestinal endoscopy: a case report and review of literature. Human Pathology, 2019, 92, 107-112.	2.0	11
16	Laser capture microdissection for detecting the expression of epithelial‐mesenchymal transition‐related genes in epithelial and spindle cells of paraffin‐embedded formalin‐fixed biphasic synovial sarcoma. Clinical and Experimental Pharmacology and Physiology, 2018, 45, 675-682.	1.9	0
17	Ovarian microcystic stromal tumor with undetermined potential: case study with molecular analysis and literature review. Human Pathology, 2018, 78, 171-176.	2.0	23
18	Clinicopathological significance of Bmi-1 overexpression in esophageal cancer: a meta-analysis. Biomarkers in Medicine, 2018, 12, 71-81.	1.4	5

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19	Exploring the Histogenesis and Diagnostic Strategy Using Immunoassay and RT-PCR in Alveolar Soft Part Sarcoma. Pathology and Oncology Research, 2018, 24, 593-600.	1.9	6
20	Matrix metalloproteinase-14 induces epithelial-to-mesenchymal transition in synovial sarcoma. Human Pathology, 2018, 80, 201-209.	2.0	9
21	Overexpression of Polo-like kinase1 (PLK1) in chondrosarcoma and its implications for cancer progression. International Journal of Clinical and Experimental Pathology, 2018, 11, 1707-1711.	0.5	0
22	Prognostic value of the Micro<scp>RNA</scp>‑29 family in multiple human cancers: A meta‑analysis and systematic review. Clinical and Experimental Pharmacology and Physiology, 2017, 44, 441-454.	1.9	37
23	The increased number of tumor-associated macrophage is associated with overexpression of VEGF-C, plays an important role in Kazakh ESCC invasion and metastasis. Experimental and Molecular Pathology, 2017, 102, 15-21.	2.1	30
24	The clinicopathological parameters significance of CD133 and Nestin in epithelial ovarian cancer: a meta-analysis. Future Oncology, 2017, 13, 2555-2570.	2.4	7
25	The expression profile and clinicopathological significance of Notch1 in patients with colorectal cancer: a meta-analysis. Future Oncology, 2017, 13, 2103-2118.	2.4	8
26	Evaluation of expression of cancer stem cell markers and fusion gene in synovial sarcoma: Insights into histogenesis and pathogenesis. Oncology Reports, 2017, 37, 3351-3360.	2.6	16
27	Transforming growth factor-β1 signaling promotes epithelial-mesenchymal transition-like phenomena, cell motility, and cell invasion in synovial sarcoma cells. PLoS ONE, 2017, 12, e0182680.	2.5	16
28	CD163 as a marker of M2 macrophage, contribute to predict aggressiveness and prognosis of Kazakh esophageal squamous cell carcinoma. Oncotarget, 2017, 8, 21526-21538.	1.8	114
29	Synergistic inhibition of colon cancer cell growth with nanoemulsion-loaded paclitaxel and Pl3K/mTOR dual inhibitor BEZ235 through apoptosis. International Journal of Nanomedicine, 2016, 11, 1947.	6.7	28
30	Membrane type 1-matrix metalloproteinase induces epithelial-to-mesenchymal transition in esophageal squamous cell carcinoma: Observations from clinical and in vitro analyses. Scientific Reports, 2016, 6, 22179.	3.3	45
31	Effect of TGF-β1 on the Migration and Recruitment of Mesenchymal Stem Cells after Vascular Balloon Injury: Involvement of Matrix Metalloproteinase-14. Scientific Reports, 2016, 6, 21176.	3.3	28
32	Relationship between microvessel density and cancer stem cells in tumor angiogenesis: a meta-analysis. Biomarkers in Medicine, 2016, 10, 919-932.	1.4	7
33	Identification of potential mutations and genomic alterations in the epithelial and spindle cell components of biphasic synovial sarcomas using a human exome SNP chip. BMC Medical Genomics, 2015, 8, 69.	1.5	15
34	Meta-Analysis of the Effect of Mesenchymal Stem Cell Transplantation on Vascular Remodeling after Carotid Balloon Injury in Animal Models. PLoS ONE, 2015, 10, e0120082.	2.5	3
35	Clinicopathological significance of ALDH1A1 in lung, colorectal, and breast cancers: a meta-analysis. Biomarkers in Medicine, 2015, 9, 777-790.	1.4	16
36	Down-Regulated E-Cadherin Expression Is Associated with Poor Five-Year Overall Survival in Bone and Soft Tissue Sarcoma: Results of a Meta-Analysis. PLoS ONE, 2015, 10, e0121448.	2.5	21

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37	Chromosomal imbalances revealed in primary renal cell carcinomas by comparative genomic hybridization. International Journal of Clinical and Experimental Pathology, 2015, 8, 3636-47.	0.5	7
38	Papillary renal cell carcinoma: a clinicopathological and whole-genome exon sequencing study. International Journal of Clinical and Experimental Pathology, 2015, 8, 8311-35.	0.5	16
39	Chromophobe renal cell carcinoma with and without sarcomatoid change: a clinicopathological, comparative genomic hybridization, and whole-exome sequencing study. American Journal of Translational Research (discontinued), 2015, 7, 2482-99.	0.0	5
40	Analysis of Molecular Cytogenetic Alteration in Rhabdomyosarcoma by Array Comparative Genomic Hybridization. PLoS ONE, 2014, 9, e94924.	2.5	20
41	TGF-Î²1/Smad Signaling Pathway Regulates Epithelial-to-Mesenchymal Transition in Esophageal Squamous Cell Carcinoma: In Vitro and Clinical Analyses of Cell Lines and Nomadic Kazakh Patients from Northwest Xinjiang, China. PLoS ONE, 2014, 9, e112300.	2.5	54
42	Xp11 translocation renal cell carcinoma in adults: a clinicopathological and comparative genomic hybridization study. International Journal of Clinical and Experimental Pathology, 2014, 7, 236-45.	0.5	10
43	Unclassified renal cell carcinoma: a clinicopathological, comparative genomic hybridization, and whole-genome exon sequencing study. International Journal of Clinical and Experimental Pathology, 2014, 7, 3865-75.	0.5	4