Peng Qi

List of Publications by Year in descending order

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623734 794594 1,737 20 14 19 citations g-index h-index papers 20 20 20 2869 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	BRAF, C-KIT, and NRAS mutations correlated with different clinicopathological features: an analysis of 691 melanoma patients from a single center. Annals of Translational Medicine, 2022, 10, 31-31.	1.7	3
2	Clinical validation of a 90-gene expression test for tumor tissue of origin diagnosis: a large-scale multicenter study of 1417 patients. Journal of Translational Medicine, 2022, 20, 114.	4.4	7
3	Concordance of the 21-gene assay between core needle biopsy and resection specimens in early breast cancer patients. Breast Cancer Research and Treatment, 2021, 186, 327-342.	2.5	14
4	Performance of Automated Dissection on Formalin-Fixed Paraffin-Embedded Tissue Sections for the 21-Gene Recurrence Score Assay. Technology in Cancer Research and Treatment, 2020, 19, 153303382096076.	1.9	5
5	Development and Clinical Validation of a 90-Gene Expression Assay for Identifying Tumor Tissue Origin. Journal of Molecular Diagnostics, 2020, 22, 1139-1150.	2.8	13
6	E2F1 induces LSINCT5 transcriptional activity and promotes gastric cancer progression by affecting the epithelial-mesenchymal transition. Cancer Management and Research, 2018, Volume 10, 2563-2571.	1.9	28
7	PTTG3P promotes gastric tumour cell proliferation and invasion and is an indicator of poor prognosis. Journal of Cellular and Molecular Medicine, 2017, 21, 3360-3371.	3.6	42
8	A Positive Feedback Loop of IncRNA- <i>PVT1</i> and FOXM1 Facilitates Gastric Cancer Growth and Invasion. Clinical Cancer Research, 2017, 23, 2071-2080.	7.0	210
9	Response to Tyrosine Kinase Inhibitors in Lung Adenocarcinoma with the Rare Epidermal Growth Factor Receptor Mutation S768I: a Retrospective Analysis and Literature Review. Targeted Oncology, 2017, 12, 81-88.	3.6	19
10	Circulating Long RNAs in Serum Extracellular Vesicles: Their Characterization and Potential Application as Biomarkers for Diagnosis of Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1158-1166.	2.5	175
11	Circulating long non-coding RNAs in cancer: current status and future perspectives. Molecular Cancer, 2016, 15, 39.	19.2	250
12	Pituitary tumor-transforming gene-1 serves as an independent prognostic biomarker for gastric cancer. Gastric Cancer, 2016, 19, 107-115.	5. 3	28
13	Long non-coding RNAs LOC285194, RP11-462C24.1 and Nbla12O61 in serum provide a new approach for distinguishing patients with colorectal cancer from healthy controls. Oncotarget, 2016, 7, 70769-70778.	1.8	34
14	Circulating <scp>CUDR</scp> , <scp>LSINCT</scp> â€5 and <scp>PTENP</scp> 1 long noncoding <scp>RNA</scp> s in sera distinguish patients with gastric cancer from healthy controls. International Journal of Cancer, 2015, 137, 1128-1135.	5.1	143
15	Reciprocal repression between TUSC7 and miR-23b in gastric cancer. International Journal of Cancer, 2015, 137, 1269-1278.	5.1	82
16	Down-regulation of ncRAN, a long non-coding RNA, contributes to colorectal cancer cell migration and invasion and predicts poor overall survival for colorectal cancer patients. Molecular Carcinogenesis, 2015, 54, 742-750.	2.7	61
17	Long non-coding RNAs in cancer invasion and metastasis. Modern Pathology, 2015, 28, 4-13.	5. 5	71
18	Long non-coding RNAs in colorectal cancer: implications for pathogenesis and clinical application. Modern Pathology, 2014, 27, 1310-1320.	5. 5	101

#	Article	IF	CITATIONS
19	The long non-coding RNAs, a new cancer diagnostic and therapeutic gold mine. Modern Pathology, 2013, 26, 155-165.	5.5	449
20	Clinicopathological, molecular and prognostic characteristics of cancer of unknown primary in China: An analysis of 1420 cases. Cancer Medicine, 0 , , .	2.8	2