

# You Guo

## List of Publications by Year in descending order

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Version: 2024-02-01

14  
papers

429  
citations

759233

12  
h-index

1058476

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g-index

16  
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16  
docs citations

16  
times ranked

352  
citing authors

#	ARTICLE	IF	CITATIONS
1	A qualitative transcriptional signature for the early diagnosis of colorectal cancer. <i>Cancer Science</i> , 2019, 110, 3225-3234.	3.9	21
2	A rank-based algorithm of differential expression analysis for small cell line data with statistical control. <i>Briefings in Bioinformatics</i> , 2019, 20, 482-491.	6.5	23
3	A qualitative signature for early diagnosis of hepatocellular carcinoma based on relative expression orderings. <i>Liver International</i> , 2018, 38, 1812-1819.	3.9	33
4	A qualitative signature for predicting pathological response to neoadjuvant chemoradiation in locally advanced rectal cancers. <i>Radiotherapy and Oncology</i> , 2018, 129, 149-153.	0.6	20
5	Quantitative or qualitative transcriptional diagnostic signatures? A case study for colorectal cancer. <i>BMC Genomics</i> , 2018, 19, 99.	2.8	36
6	Shared liver-like transcriptional characteristics in liver metastases and corresponding primary colorectal tumors. <i>Journal of Cancer</i> , 2018, 9, 1500-1505.	2.5	7
7	Evaluating hepatocellular carcinoma cell lines for tumour samples using within-sample relative expression orderings of genes. <i>Liver International</i> , 2017, 37, 1688-1696.	3.9	21
8	Robust transcriptional signatures for low-input RNA samples based on relative expression orderings. <i>BMC Genomics</i> , 2017, 18, 913.	2.8	45
9	Circumvent the uncertainty in the applications of transcriptional signatures to tumor tissues sampled from different tumor sites. <i>Oncotarget</i> , 2017, 8, 30265-30275.	1.8	72
10	Identifying reproducible cancer-associated highly expressed genes with important functional significances using multiple datasets. <i>Scientific Reports</i> , 2016, 6, 36227.	3.3	2
11	Differential expression analysis for individual cancer samples based on robust within-sample relative gene expression orderings across multiple profiling platforms. <i>Oncotarget</i> , 2016, 7, 68909-68920.	1.8	63
12	Common DNA methylation alterations of Alzheimer's disease and aging in peripheral whole blood. <i>Oncotarget</i> , 2016, 7, 19089-19098.	1.8	25
13	An individualized prognostic signature and multi-omics distinction for early stage hepatocellular carcinoma patients with surgical resection. <i>Oncotarget</i> , 2016, 7, 24097-24110.	1.8	37
14	Identifying clinically relevant drug resistance genes in drug-induced resistant cancer cell lines and post-chemotherapy tissues. <i>Oncotarget</i> , 2015, 6, 41216-41227.	1.8	24