Tanxiao Huang

List of Publications by Year
in descending order

11 papers	499 citations	1163117 8 h-index	1281871 11 g-index
11 all docs	11 docs citations	11 times ranked	1278 citing authors

#	ARTICLE	IF	CITATIONS
1	The genomic characteristics of RET fusion positive tumors in Chinese non-small cell lung cancer (NSCLC) patients. Journal of Cancer Research and Clinical Oncology, 2023, 149, 1019-1028.	2.5	2
2	Metastatic Low-Grade Sarcoma with *CARS*-*ALK* Fusion Dramatically Responded to Multiple ALK Tyrosine Kinase Inhibitors: A Case Report with Comprehensive Genomic Analysis. Oncologist, 2021, 26, e524-e529.	3.7	6
3	The Genomic Characteristics of ALK Fusion Positive Tumors in Chinese NSCLC Patients. Frontiers in Oncology, 2020, 10, 726.	2.8	21
4	Circulating Tumor DNA Is Capable of Monitoring the Therapeutic Response and Resistance in Advanced Colorectal Cancer Patients Undergoing Combined Target and Chemotherapy. Frontiers in Oncology, 2020, 10, 466.	2.8	22
5	Targeted deep sequencing helps distinguish independent primary tumors from intrapulmonary metastasis for lung cancer diagnosis. Journal of Cancer Research and Clinical Oncology, 2020, 146, 2359-2367.	2.5	9
6	Comprehensive analysis of POLE and POLD1 Gene Variations identifies cancer patients potentially benefit from immunotherapy in Chinese population. Scientific Reports, 2019, 9, 15767.	3.3	34
7	Gencore: an efficient tool to generate consensus reads for error suppressing and duplicate removing of NGS data. BMC Bioinformatics, 2019, 20, 606.	2.6	43
8	Sequencing of circulating tumor DNA for dynamic monitoring of gene mutations in advanced non-small cell lung cancer. Oncology Letters, 2018, 15, 3726-3734.	1.8	6
9	Detection of Urothelial Bladder Carcinoma via Microfluidic Immunoassay and Single-Cell DNA Copy-Number Alteration Analysis of Captured Urinary-Exfoliated Tumor Cells. Cancer Research, 2018, 78, 4073-4085.	0.9	34
10	GeneFuse: detection and visualization of target gene fusions from DNA sequencing data. International Journal of Biological Sciences, 2018, 14, 843-848.	6.4	36
11	AfterQC: automatic filtering, trimming, error removing and quality control for fastq data. BMC Bioinformatics, 2017, 18, 80.	2.6	286