Zhengyan Kan

List of Publications by Year in descending order

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933264 1199470 3,078 14 10 12 citations g-index h-index papers 15 15 15 7775 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Genome-wide survey of recurrent HBV integration in hepatocellular carcinoma. Nature Genetics, 2012, 44, 765-769.	9.4	785
2	Single-cell RNA-seq enables comprehensive tumour and immune cell profiling in primary breast cancer. Nature Communications, 2017, 8, 15081.	5.8	743
3	The BioMart community portal: an innovative alternative to large, centralized data repositories. Nucleic Acids Research, 2015, 43, W589-W598.	6.5	682
4	Whole-genome sequencing identifies recurrent mutations in hepatocellular carcinoma. Genome Research, 2013, 23, 1422-1433.	2.4	457
5	Multi-omics profiling of younger Asian breast cancers reveals distinctive molecular signatures. Nature Communications, 2018, 9, 1725.	5 . 8	122
6	Chemotherapy induces dynamic immune responses in breast cancers that impact treatment outcome. Nature Communications, 2020, 11, 6175.	5.8	92
7	Decoding complex patterns of genomic rearrangement in hepatocellular carcinoma. Genomics, 2014, 103, 189-203.	1.3	49
8	Cooperation Between Distinct Cancer Driver Genes Underlies Intertumor Heterogeneity in Hepatocellular Carcinoma. Gastroenterology, 2020, 159, 2203-2220.e14.	0.6	47
9	OASIS: web-based platform for exploring cancer multi-omics data. Nature Methods, 2016, 13, 9-10.	9.0	45
10	Immune gene expression profiling reveals heterogeneity in luminal breast tumors. Breast Cancer Research, 2019, 21, 147.	2.2	43
11	Prospective longitudinal multi-omics study of palbociclib resistance in hormone receptor+/HER2-metastatic breast cancer Journal of Clinical Oncology, 2021, 39, 1013-1013.	0.8	7
12	Genomic characteristics of breast cancer to predict response of neoadjuvant chemotherapy and long-term prognosis Journal of Clinical Oncology, 2021, 39, 557-557.	0.8	6
13	APPLICATIONS OF GENETICS, GENOMICS AND BIOINFORMATICS IN DRUG DISCOVERY., 2018,,.		0
14	Abstract P1-08-07: Prediction model of the response of neoadjuvant chemotherapy and long term survival according to multi-omic profiling in cooperation with clinicopathologic features in patients with breast cancer. Cancer Research, 2022, 82, P1-08-07-P1-08-07.	0.4	0