

Quentin Bayard

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

15
papers

469
citations

8
h-index

16
g-index

16
ext. papers

771
ext. citations

11.8
avg, IF

3.13
L-index

#	Paper	IF	Citations
15	Mutational signatures reveal the dynamic interplay of risk factors and cellular processes during liver tumorigenesis. <i>Nature Communications</i> , 2017 , 8, 1315	17.4	135
14	Analysis of Liver Cancer Cell Lines Identifies Agents With Likely Efficacy Against Hepatocellular Carcinoma and Markers of Response. <i>Gastroenterology</i> , 2019 , 157, 760-776	13.3	77
13	Cyclin A2/E1 activation defines a hepatocellular carcinoma subclass with a rearrangement signature of replication stress. <i>Nature Communications</i> , 2018 , 9, 5235	17.4	73
12	Clinical Impact of Genomic Diversity From Early to Advanced Hepatocellular Carcinoma. <i>Hepatology</i> , 2020 , 71, 164-182	11.2	62
11	Adeno-associated virus in the liver: natural history and consequences in tumour development. <i>Gut</i> , 2020 , 69, 737-747	19.2	36
10	Palimpsest: an R package for studying mutational and structural variant signatures along clonal evolution in cancer. <i>Bioinformatics</i> , 2018 , 34, 3380-3381	7.2	26
9	BAP1 mutations define a homogeneous subgroup of hepatocellular carcinoma with fibrolamellar-like features and activated PKA. <i>Journal of Hepatology</i> , 2020 , 72, 924-936	13.4	18
8	Telomere length is key to hepatocellular carcinoma diversity and telomerase addiction is an actionable therapeutic target. <i>Journal of Hepatology</i> , 2021 , 74, 1155-1166	13.4	11
7	Recurrent chromosomal rearrangements of , and activating JAK/STAT pathway in inflammatory hepatocellular adenomas. <i>Gut</i> , 2020 , 69, 1667-1676	19.2	8
6	Multi-site tumor sampling highlights molecular intra-tumor heterogeneity in malignant pleural mesothelioma. <i>Genome Medicine</i> , 2021 , 13, 113	14.4	7
5	DNA Methylation Signatures Reveal the Diversity of Processes Remodeling Hepatocellular Carcinoma Methylomes. <i>Hepatology</i> , 2021 , 74, 816-834	11.2	7
4	Integrated Genomic Analysis Identifies Driver Genes and Cisplatin-Resistant Progenitor Phenotype in Pediatric Liver Cancer. <i>Cancer Discovery</i> , 2021 , 11, 2524-2543	24.4	4
3	High Impact: The Role of Promiscuous Binding Sites in Polypharmacology. <i>Molecules</i> , 2019 , 24,	4.8	3
2	Common genetic variation in alcohol-related hepatocellular carcinoma: a case-control genome-wide association study.. <i>Lancet Oncology</i> , 2022 , 23, 161-171	21.7	1
1	Structure, Dynamics, and Impact of Replication Stress-Induced Structural Variants in Hepatocellular Carcinoma.. <i>Cancer Research</i> , 2022 , 82, 1470-1481	10.1	