

Jessica Zucman-Rossi

List of Publications by Year in descending order

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Version: 2024-02-01

394
papers

50,040
citations

3325

91
h-index

1705

213
g-index

420
all docs

420
docs citations

420
times ranked

51377
citing authors

#	ARTICLE	IF	CITATIONS
1	Signatures of mutational processes in human cancer. <i>Nature</i> , 2013, 500, 415-421.	13.7	8,060
2	Hepatocellular carcinoma. <i>Nature Reviews Disease Primers</i> , 2021, 7, 6.	18.1	2,757
3	International network of cancer genome projects. <i>Nature</i> , 2010, 464, 993-998.	13.7	2,114
4	Hepatocellular carcinoma. <i>Nature Reviews Disease Primers</i> , 2016, 2, 16018.	18.1	1,863
5	Gene fusion with an ETS DNA-binding domain caused by chromosome translocation in human tumours. <i>Nature</i> , 1992, 359, 162-165.	13.7	1,724
6	Exome sequencing of hepatocellular carcinomas identifies new mutational signatures and potential therapeutic targets. <i>Nature Genetics</i> , 2015, 47, 505-511.	9.4	1,372
7	Alteration in a new gene encoding a putative membrane-organizing protein causes neuro-fibromatosis type 2. <i>Nature</i> , 1993, 363, 515-521.	13.7	1,351
8	Integrated analysis of somatic mutations and focal copy-number changes identifies key genes and pathways in hepatocellular carcinoma. <i>Nature Genetics</i> , 2012, 44, 694-698.	9.4	1,229
9	Transcriptome classification of HCC is related to gene alterations and to new therapeutic targets. <i>Hepatology</i> , 2007, 45, 42-52.	3.6	1,034
10	The Ewing Family of Tumors – A Subgroup of Small-Round-Cell Tumors Defined by Specific Chimeric Transcripts. <i>New England Journal of Medicine</i> , 1994, 331, 294-299.	13.9	1,010
11	Genetic Landscape and Biomarkers of Hepatocellular Carcinoma. <i>Gastroenterology</i> , 2015, 149, 1226-1239.e4.	0.6	980
12	Genotype-phenotype correlation in hepatocellular adenoma: New classification and relationship with HCC. <i>Hepatology</i> , 2006, 43, 515-524.	3.6	733
13	Mechanisms of HBV-induced hepatocellular carcinoma. <i>Journal of Hepatology</i> , 2016, 64, S84-S101.	1.8	664
14	MicroRNA profiling in hepatocellular tumors is associated with clinical features and oncogene/tumor suppressor gene mutations. <i>Hepatology</i> , 2008, 47, 1955-1963.	3.6	634
15	Toward understanding and exploiting tumor heterogeneity. <i>Nature Medicine</i> , 2015, 21, 846-853.	15.2	604
16	Hepatocellular adenoma subtype classification using molecular markers and immunohistochemistry. <i>Hepatology</i> , 2007, 46, 740-748.	3.6	554
17	Ewing sarcoma 11;22 translocation produces a chimeric transcription factor that requires the DNA-binding domain encoded by FLI1 for transformation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1993, 90, 5752-5756.	3.3	538
18	A gp130-IL6-STAT3-YAP module links inflammation to epithelial regeneration. <i>Nature</i> , 2015, 519, 57-62.	13.7	528

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19	Histological subtypes of hepatocellular carcinoma are related to gene mutations and molecular tumour classification. <i>Journal of Hepatology</i> , 2017, 67, 727-738.	1.8	525
20	High frequency of telomerase reverse-transcriptase promoter somatic mutations in hepatocellular carcinoma and preneoplastic lesions. <i>Nature Communications</i> , 2013, 4, 2218.	5.8	513
21	EWS and ATF-1 gene fusion induced by t(12;22) translocation in malignant melanoma of soft parts. <i>Nature Genetics</i> , 1993, 4, 341-345.	9.4	483
22	Frequent in-frame somatic deletions activate gp130 in inflammatory hepatocellular tumours. <i>Nature</i> , 2009, 457, 200-204.	13.7	437
23	Hepatocellular adenoma management and phenotypic classification: The Bordeaux experience. <i>Hepatology</i> , 2009, 50, 481-489.	3.6	394
24	Recurrent AAV2-related insertional mutagenesis in human hepatocellular carcinomas. <i>Nature Genetics</i> , 2015, 47, 1187-1193.	9.4	387
25	EASL Clinical Practice Guidelines on the management of benign liver tumours. <i>Journal of Hepatology</i> , 2016, 65, 386-398.	1.8	372
26	DNA methylation-based prognosis and epdrivers in hepatocellular carcinoma. <i>Hepatology</i> , 2015, 61, 1945-1956.	3.6	367
27	GNAS-activating mutations define a rare subgroup of inflammatory liver tumors characterized by STAT3 activation. <i>Journal of Hepatology</i> , 2012, 56, 184-191.	1.8	354
28	Genomic portrait of resectable hepatocellular carcinomas: Implications of <i>RB1</i> and <i>FGF19</i> aberrations for patient stratification. <i>Hepatology</i> , 2014, 60, 1972-1982.	3.6	345
29	Bi-allelic inactivation of TCF1 in hepatic adenomas. <i>Nature Genetics</i> , 2002, 32, 312-315.	9.4	333
30	Molecular and histological correlations in liver cancer. <i>Journal of Hepatology</i> , 2019, 71, 616-630.	1.8	308
31	Genetics of hepatocellular tumors. <i>Oncogene</i> , 2006, 25, 3778-3786.	2.6	304
32	A Hepatocellular Carcinoma 5-Gene Score Associated With Survival of Patients After Liver Resection. <i>Gastroenterology</i> , 2013, 145, 176-187.	0.6	302
33	Molecular Classification of Hepatocellular Adenoma Associates With Risk Factors, Bleeding, and Malignant Transformation. <i>Gastroenterology</i> , 2017, 152, 880-894.e6.	0.6	290
34	Cloning and characterization of the Ewing's sarcoma and peripheral neuroepithelioma t(11;22) translocation breakpoints. <i>Genes Chromosomes and Cancer</i> , 1992, 5, 271-277.	1.5	284
35	Hepatocellular adenomas: Magnetic resonance imaging features as a function of molecular pathological classification. <i>Hepatology</i> , 2008, 48, 808-818.	3.6	277
36	Telomerase reverse transcriptase promoter mutation is an early somatic genetic alteration in the transformation of premalignant nodules in hepatocellular carcinoma on cirrhosis. <i>Hepatology</i> , 2014, 60, 1983-1992.	3.6	268

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37	Hepatocellular Benign Tumors—From Molecular Classification to Personalized Clinical Care. <i>Gastroenterology</i> , 2013, 144, 888-902.	0.6	251
38	Molecular Subtypes of Clear Cell Renal Cell Carcinoma Are Associated with Sunitinib Response in the Metastatic Setting. <i>Clinical Cancer Research</i> , 2015, 21, 1329-1339.	3.2	250
39	chCC—CCA: Consensus terminology for primary liver carcinomas with both hepatocytic and cholangiocytic differentiation. <i>Hepatology</i> , 2018, 68, 113-126.	3.6	244
40	Genomic Profiling of Hepatocellular Adenomas Reveals Recurrent FRK-Activating Mutations and the Mechanisms of Malignant Transformation. <i>Cancer Cell</i> , 2014, 25, 428-441.	7.7	240
41	Molecular pathogenesis of focal nodular hyperplasia and hepatocellular adenoma. <i>Journal of Hepatology</i> , 2008, 48, 163-170.	1.8	235
42	Differential effects of inactivated Axin1 and activated β -catenin mutations in human hepatocellular carcinomas. <i>Oncogene</i> , 2007, 26, 774-780.	2.6	230
43	Mutational signatures reveal the dynamic interplay of risk factors and cellular processes during liver tumorigenesis. <i>Nature Communications</i> , 2017, 8, 1315.	5.8	228
44	Genotype—phenotype correlation of CTNNB1 mutations reveals different β -catenin activity associated with liver tumor progression. <i>Hepatology</i> , 2016, 64, 2047-2061.	3.6	222
45	Intra-tumoral tertiary lymphoid structures are associated with a low risk of early recurrence of hepatocellular carcinoma. <i>Journal of Hepatology</i> , 2019, 70, 58-65.	1.8	219
46	Somatic mutations activating STAT3 in human inflammatory hepatocellular adenomas. <i>Journal of Experimental Medicine</i> , 2011, 208, 1359-1366.	4.2	218
47	High Incidence of Activating <i>TERT</i> Promoter Mutations in Meningiomas Undergoing Malignant Progression. <i>Brain Pathology</i> , 2014, 24, 184-189.	2.1	209
48	Clinical, Morphologic, and Molecular Features Defining So-Called Telangiectatic Focal Nodular Hyperplasias of the Liver. <i>Gastroenterology</i> , 2005, 128, 1211-1218.	0.6	207
49	A MYC—aurora kinase A protein complex represents an actionable drug target in p53-altered liver cancer. <i>Nature Medicine</i> , 2016, 22, 744-753.	15.2	207
50	PNPLA3 gene in liver diseases. <i>Journal of Hepatology</i> , 2016, 65, 399-412.	1.8	205
51	Oncogenic conversion of a novel orphan nuclear receptor by chromosome translocation. <i>Human Molecular Genetics</i> , 1995, 4, 2219-2226.	1.4	190
52	Tissue metabolomics of hepatocellular carcinoma: Tumor energy metabolism and the role of transcriptomic classification. <i>Hepatology</i> , 2013, 58, 229-238.	3.6	172
53	Pathological diagnosis of liver cell adenoma and focal nodular hyperplasia: Bordeaux update. <i>Journal of Hepatology</i> , 2007, 46, 521-527.	1.8	170
54	Familia liver adenomatosis associated with hepatocyte nuclear factor 1 β inactivation The authors thank Leigh Pascoe for critical reading of the manuscript, Hne Blanch and Hung Bui of the CEPH/Fondation Jean Dausset for technical help in sequencing, and Drs. A. Saillant, E. Akodjenou, and E. Urvoas (Pediatric and Radiology Units, Hpitaux de Chartres, France) for referring patient B1 to E.J. and for performing liver ultrasound screening in family B.. <i>Gastroenterology</i> , 2003, 125, 1470-1475.	0.6	169

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55	Immune Contexture, Immunoscore, and Malignant Cell Molecular Subgroups for Prognostic and Theranostic Classifications of Cancers. <i>Advances in Immunology</i> , 2016, 130, 95-190.	1.1	160
56	Macrotrabecularâ€massive hepatocellular carcinoma: A distinctive histological subtype with clinical relevance. <i>Hepatology</i> , 2018, 68, 103-112.	3.6	159
57	The role of telomeres and telomerase in cirrhosis and liver cancer. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2019, 16, 544-558.	8.2	154
58	Liver Cancer Initiation Requires p53 Inhibition by CD44-Enhanced Growth Factor Signaling. <i>Cancer Cell</i> , 2018, 33, 1061-1077.e6.	7.7	151
59	Beta-catenin mutations in hepatocellular carcinoma correlate with a low rate of loss of heterozygosity. <i>Oncogene</i> , 1999, 18, 4044-4046.	2.6	149
60	High resolution deletion analysis of constitutional DNA from neurofibromatosis type 2 (NF2) patients using microarray-CGH. <i>Human Molecular Genetics</i> , 2001, 10, 271-282.	1.4	147
61	Molecular characterization of hepatocellular adenomas developed in patients with glycogen storage disease type I. <i>Journal of Hepatology</i> , 2013, 58, 350-357.	1.8	146
62	Loss of Trim24 (Tif1±) gene function confers oncogenic activity to retinoic acid receptor alpha. <i>Nature Genetics</i> , 2007, 39, 1500-1506.	9.4	145
63	Genomic Medicine and Implications for Hepatocellular Carcinoma Prevention and Therapy. <i>Gastroenterology</i> , 2019, 156, 492-509.	0.6	145
64	Overâ€expression of glutamine synthetase in focal nodular hyperplasia: a novel easy diagnostic tool in surgical pathology. <i>Liver International</i> , 2009, 29, 459-465.	1.9	143
65	Subtype Classification of Hepatocellular Adenoma. <i>Digestive Surgery</i> , 2010, 27, 39-45.	0.6	143
66	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.	0.6	141
67	Single nucleotide polymorphisms and risk of hepatocellular carcinoma in cirrhosis. <i>Journal of Hepatology</i> , 2012, 57, 663-674.	1.8	140
68	Genetics of Hepatobiliary Carcinogenesis. <i>Seminars in Liver Disease</i> , 2011, 31, 173-187.	1.8	138
69	Mutational signature analysis identifies <i><sc>MUTYH</sc></i> deficiency in colorectal cancers and adrenocortical carcinomas. <i>Journal of Pathology</i> , 2017, 242, 10-15.	2.1	130
70	Clinical Impact of Genomic Diversity From Early to Advanced Hepatocellular Carcinoma. <i>Hepatology</i> , 2020, 71, 164-182.	3.6	129
71	Integration of tumour and viral genomic characterisations in HBV-related hepatocellular carcinomas. <i>Gut</i> , 2015, 64, 820-829.	6.1	127
72	Clinical and molecular analysis of combined hepatocellular-cholangiocarcinomas. <i>Journal of Hepatology</i> , 2004, 41, 292-298.	1.8	126

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73	HOX11L2 expression defines a clinical subtype of pediatric T-ALL associated with poor prognosis. <i>Blood</i> , 2002, 100, 991-997.	0.6	125
74	Dissecting heterogeneity in malignant pleural mesothelioma through histo-molecular gradients for clinical applications. <i>Nature Communications</i> , 2019, 10, 1333.	5.8	125
75	Hepatocyte Nuclear Factor-1 α Gene Inactivation: Cosegregation between Liver Adenomatosis and Diabetes Phenotypes in Two Maturity-Onset Diabetes of the Young (MODY)3 Families. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2004, 89, 1476-1480.	1.8	124
76	HNF1 α Inactivation Promotes Lipogenesis in Human Hepatocellular Adenoma Independently of SREBP-1 and Carbohydrate-response Element-binding Protein (ChREBP) Activation. <i>Journal of Biological Chemistry</i> , 2007, 282, 14437-14446.	1.6	123
77	Molecular Classification of Malignant Pleural Mesothelioma: Identification of a Poor Prognosis Subgroup Linked to the Epithelial-to-Mesenchymal Transition. <i>Clinical Cancer Research</i> , 2014, 20, 1323-1334.	3.2	121
78	Molecular classification of hepatocellular adenoma in clinical practice. <i>Journal of Hepatology</i> , 2017, 67, 1074-1083.	1.8	119
79	Immunohistochemical Markers on Needle Biopsies Are Helpful for the Diagnosis of Focal Nodular Hyperplasia and Hepatocellular Adenoma Subtypes. <i>American Journal of Surgical Pathology</i> , 2012, 36, 1691-1699.	2.1	118
80	Cyclin A2/E1 activation defines a hepatocellular carcinoma subclass with a rearrangement signature of replication stress. <i>Nature Communications</i> , 2018, 9, 5235.	5.8	118
81	Negative impact of bone metastasis on outcome in clear-cell renal cell carcinoma treated with sunitinib. <i>Annals of Oncology</i> , 2011, 22, 794-800.	0.6	116
82	RIPK1 Suppresses a TRAF2-Dependent Pathway to Liver Cancer. <i>Cancer Cell</i> , 2017, 31, 94-109.	7.7	115
83	Revisiting the Pathology of Resected Benign Hepatocellular Nodules Using New Immunohistochemical Markers. <i>Seminars in Liver Disease</i> , 2011, 31, 091-103.	1.8	112
84	Germline hepatocyte nuclear factor 1 α and 1 β mutations in renal cell carcinomas. <i>Human Molecular Genetics</i> , 2005, 14, 603-614.	1.4	109
85	Unique Genomic Profile of Fibrolamellar Hepatocellular Carcinoma. <i>Gastroenterology</i> , 2015, 148, 806-818.e10.	0.6	109
86	Hepatitis B virus integrations promote local and distant oncogenic driver alterations in hepatocellular carcinoma. <i>Gut</i> , 2022, 71, 616-626.	6.1	106
87	Genomic Profiling Reveals Alternative Genetic Pathways of Meningioma Malignant Progression Dependent on the Underlying NF2 Status. <i>Clinical Cancer Research</i> , 2010, 16, 4155-4164.	3.2	103
88	Trial Watch: Monoclonal antibodies in cancer therapy. <i>Onc Immunology</i> , 2012, 1, 28-37.	2.1	103
89	Chromosome translocation based on illegitimate recombination in human tumors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 11786-11791.	3.3	101
90	Overexpression and role of the ATPase and putative DNA helicase RuvB-like 2 in human hepatocellular carcinoma. <i>Hepatology</i> , 2007, 46, 1108-1118.	3.6	100

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91	Inactivation of the <i>APC</i> Gene Is Constant in Adrenocortical Tumors from Patients with Familial Adenomatous Polyposis but Not Frequent in Sporadic Adrenocortical Cancers. <i>Clinical Cancer Research</i> , 2010, 16, 5133-5141.	3.2	97
92	Genomic structure of the EWS gene and its relationship to EWSR1, a site of tumor-associated chromosome translocation. <i>Genomics</i> , 1993, 18, 609-615.	1.3	94
93	Modeling a human hepatocellular carcinoma subset in mice through coexpression of met and point mutant β -catenin. <i>Hepatology</i> , 2016, 64, 1587-1605.	3.6	92
94	Inhibiting Glutamine-Dependent mTORC1 Activation Ameliorates Liver Cancers Driven by β -Catenin Mutations. <i>Cell Metabolism</i> , 2019, 29, 1135-1150.e6.	7.2	92
95	Genotype phenotype classification of hepatocellular adenoma. <i>World Journal of Gastroenterology</i> , 2007, 13, 2649.	1.4	90
96	Single-nucleotide polymorphisms associated with outcome in metastatic renal cell carcinoma treated with sunitinib. <i>British Journal of Cancer</i> , 2013, 108, 887-900.	2.9	88
97	The β -catenin pathway is activated in focal nodular hyperplasia but not in cirrhotic FNH-like nodules. <i>Journal of Hepatology</i> , 2008, 49, 61-71.	1.8	87
98	microRNA 193a-5p Regulates Levels of Nucleolar- and Spindle-Associated Protein 1 to Suppress Hepatocarcinogenesis. <i>Gastroenterology</i> , 2018, 155, 1951-1966.e26.	0.6	86
99	Polyploidy spectrum: a new marker in HCC classification. <i>Gut</i> , 2020, 69, 355-364.	6.1	82
100	Compliance With Hepatocellular Carcinoma Surveillance Guidelines Associated With Increased Lead-Time Adjusted Survival of Patients With Compensated Viral Cirrhosis: A Multi-Center Cohort Study. <i>Gastroenterology</i> , 2018, 155, 431-442.e10.	0.6	81
101	Dual Targeting of Histone Methyltransferase G9a and DNA Methyltransferase 1 for the Treatment of Experimental Hepatocellular Carcinoma. <i>Hepatology</i> , 2019, 69, 587-603.	3.6	81
102	Hepatocellular adenoma: what is new in 2008. <i>Hepatology International</i> , 2008, 2, 316-321.	1.9	78
103	TERT promoter mutations in primary liver tumors. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2016, 40, 9-14.	0.7	78
104	AXIN deficiency in human and mouse hepatocytes induces hepatocellular carcinoma in the absence of β -catenin activation. <i>Journal of Hepatology</i> , 2018, 68, 1203-1213.	1.8	78
105	Adeno-associated virus in the liver: natural history and consequences in tumour development. <i>Gut</i> , 2020, 69, 737-747.	6.1	78
106	KIF20A mRNA and Its Product MKlp2 Are Increased During Hepatocyte Proliferation and Hepatocarcinogenesis. <i>American Journal of Pathology</i> , 2012, 180, 131-140.	1.9	76
107	A functional screening identifies five micrnas controlling glypican-3: role of mir-1271 down-regulation in hepatocellular carcinoma. <i>Hepatology</i> , 2013, 57, 195-204.	3.6	76
108	Identification of Novel Oncogenes and Tumor Suppressors in Hepatocellular Carcinoma. <i>Seminars in Liver Disease</i> , 2010, 30, 075-086.	1.8	75

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109	A 17 α -Hydroxysteroid Dehydrogenase 13 Variant Protects From Hepatocellular Carcinoma Development in Alcoholic Liver Disease. <i>Hepatology</i> , 2019, 70, 231-240.	3.6	75
110	From the Editor's desk..... <i>Journal of Hepatology</i> , 2018, 69, 1-4.	1.8	74
111	Genetic alterations of malignant pleural mesothelioma: association with tumor heterogeneity and overall survival. <i>Molecular Oncology</i> , 2020, 14, 1207-1223.	2.1	74
112	Molecular classification of hepatocellular carcinoma. <i>Digestive and Liver Disease</i> , 2010, 42, S235-S241.	0.4	73
113	Molecular Subtypes of Clear-cell Renal Cell Carcinoma are Prognostic for Outcome After Complete Metastectomy. <i>European Urology</i> , 2018, 74, 474-480.	0.9	72
114	PNPLA3 and TM6SF2 variants as risk factors of hepatocellular carcinoma across various etiologies and severity of underlying liver diseases. <i>International Journal of Cancer</i> , 2019, 144, 533-544.	2.3	72
115	Recurrent inactivating mutations of <i>ARID2</i> in non-small cell lung carcinoma. <i>International Journal of Cancer</i> , 2013, 132, 2217-2221.	2.3	70
116	Cloning of a balanced translocation breakpoint in the DiGeorge syndrome critical region and isolation of a novel potential adhesion receptor gene in its vicinity. <i>Human Molecular Genetics</i> , 1995, 4, 551-558.	1.4	69
117	NF2 gene in neurofibromatosis type 2 patients. <i>Human Molecular Genetics</i> , 1998, 7, 2095-2101.	1.4	69
118	CXCR7 is up-regulated in human and murine hepatocellular carcinoma and is specifically expressed by endothelial cells. <i>European Journal of Cancer</i> , 2012, 48, 138-148.	1.3	68
119	Prognostic impact of baseline serum C-reactive protein in patients with metastatic renal cell carcinoma (RCC) treated with sunitinib. <i>BJU International</i> , 2014, 114, 81-89.	1.3	68
120	Overexpression and promoter mutation of the TERT gene in malignant pleural mesothelioma. <i>Oncogene</i> , 2014, 33, 3748-3752.	2.6	68
121	Interphase molecular cytogenetics of Ewing's sarcoma and peripheral neuroepithelioma t(11;22) with flanking and overlapping cosmid probes. <i>Cancer Genetics and Cytogenetics</i> , 1994, 74, 13-18.	1.0	67
122	Co-occurring Mutations of Tumor Suppressor Genes, <i>LATS2</i> and <i>NF2</i> , in Malignant Pleural Mesothelioma. <i>Clinical Cancer Research</i> , 2017, 23, 3191-3202.	3.2	67
123	Loss of hepatocyte nuclear factor 1 α function in human hepatocellular adenomas leads to aberrant activation of signaling pathways involved in tumorigenesis. <i>Hepatology</i> , 2010, 51, 557-566.	3.6	66
124	A Novel Epigenetic Phenotype Associated With the Most Aggressive Pathway of Bladder Tumor Progression. <i>Journal of the National Cancer Institute</i> , 2011, 103, 47-60.	3.0	66
125	Genetics of Hepatocellular Carcinoma: Approaches to Explore Molecular Diversity. <i>Hepatology</i> , 2021, 73, 14-26.	3.6	66
126	Nivolumab, nivolumab+ipilimumab, and VEGFR-tyrosine kinase inhibitors as first-line treatment for metastatic clear-cell renal cell carcinoma (BIONIKK): a biomarker-driven, open-label, non-comparative, randomised, phase 2 trial. <i>Lancet Oncology</i> , The, 2022, 23, 612-624.	5.1	66

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127	Specific association between alcohol intake, high grade of differentiation and 4q34-q35 deletions in hepatocellular carcinomas identified by high resolution allelotyping. <i>Oncogene</i> , 2002, 21, 1225-1232.	2.6	65
128	ESM1 as a Marker of Macrotrabecular-Massive Hepatocellular Carcinoma. <i>Clinical Cancer Research</i> , 2019, 25, 5859-5865.	3.2	64
129	Association of CYP1B1 Germ Line Mutations with Hepatocyte Nuclear Factor 1 α Mutated Hepatocellular Adenoma. <i>Cancer Research</i> , 2007, 67, 2611-2616.	0.4	62
130	Detection of plasma tumor DNA in head and neck squamous cell carcinoma by microsatellite typing and p53 mutation analysis. <i>Cancer Research</i> , 2000, 60, 707-11.	0.4	62
131	Mapping of human chromosome 22 with a panel of somatic cell hybrids. <i>Genomics</i> , 1991, 9, 721-727.	1.3	60
132	The <i>HOX</i> gene network in hepatocellular carcinoma. <i>International Journal of Cancer</i> , 2011, 129, 2577-2587.	2.3	60
133	Genetics of hepatocellular carcinoma: The next generation. <i>Journal of Hepatology</i> , 2014, 60, 224-226.	1.8	59
134	Role of Contrast-Enhanced Sonography in Differentiation of Subtypes of Hepatocellular Adenoma: Correlation with MRI Findings. <i>American Journal of Roentgenology</i> , 2012, 199, 341-348.	1.0	58
135	Hepatocyte nuclear factor 1 α suppresses steatosis-associated liver cancer by inhibiting PPAR γ transcription. <i>Journal of Clinical Investigation</i> , 2017, 127, 1873-1888.	3.9	58
136	Spectrum of <i>HNF1A</i> Somatic Mutations in Hepatocellular Adenoma Differs From That in Patients With MODY3 and Suggests Genotoxic Damage. <i>Diabetes</i> , 2010, 59, 1836-1844.	0.3	57
137	Proliferation Markers Are Associated with MET Expression in Hepatocellular Carcinoma and Predict Tivantinib Sensitivity <i>In Vitro</i> . <i>Clinical Cancer Research</i> , 2017, 23, 4364-4375.	3.2	57
138	Adenosine triphosphatase pontin is overexpressed in hepatocellular carcinoma and coregulated with reptin through a new posttranslational mechanism. <i>Hepatology</i> , 2009, 50, 1871-1883.	3.6	54
139	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.	1.8	54
140	Identification of molecular pathways involved in oxaliplatin-associated sinusoidal dilatation. <i>Journal of Hepatology</i> , 2012, 56, 869-876.	1.8	53
141	Rnd3/RhoE Is down-regulated in hepatocellular carcinoma and controls cellular invasion. <i>Hepatology</i> , 2012, 55, 1766-1775.	3.6	53
142	Palimpsest: an R package for studying mutational and structural variant signatures along clonal evolution in cancer. <i>Bioinformatics</i> , 2018, 34, 3380-3381.	1.8	53
143	Focal Nodular Hyperplasia and Hepatocellular Adenoma around the World Viewed through the Scope of the Immunopathological Classification. <i>International Journal of Hepatology</i> , 2013, 2013, 1-12.	0.4	52
144	Interethnic polymorphism of EWS intron 6: genome plasticity mediated by Alu retroposition and recombination. <i>Human Genetics</i> , 1997, 99, 357-363.	1.8	51

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145	Prognostic and theranostic impact of molecular subtypes and immune classifications in renal cell cancer (RCC) and colorectal cancer (CRC). <i>OncImmunology</i> , 2015, 4, e1049804.	2.1	51
146	Identification of New Members of the Gas2 and Ras Families in the 22q12 Chromosome Region. <i>Genomics</i> , 1996, 38, 247-254.	1.3	50
147	p16INK4A inactivation mechanisms in non-small-cell lung cancer patients occupationally exposed to asbestos. <i>Lung Cancer</i> , 2010, 67, 23-30.	0.9	50
148	STAT3 mutations identified in human hematologic neoplasms induce myeloid malignancies in a mouse bone marrow transplantation model. <i>Haematologica</i> , 2013, 98, 1748-1752.	1.7	50
149	p53 mutations in human tumors with chimericEWS/FLI/1 genes. <i>International Journal of Cancer</i> , 1994, 57, 336-340.	2.3	49
150	Unicolor and bicolor in situ hybridization in the diagnosis of peripheral neuroepithelioma and related tumors. <i>Genes Chromosomes and Cancer</i> , 1992, 5, 30-34.	1.5	48
151	Syntenic Relationships between Genomic Profiles of Fiber-Induced Murine and Human Malignant Mesothelioma. <i>American Journal of Pathology</i> , 2011, 178, 881-894.	1.9	48
152	The liver-specific microRNA miR-122*, the complementary strand of microRNA miR-122, acts as a tumor suppressor by modulating the p53/mouse double minute 2 homolog circuitry. <i>Hepatology</i> , 2016, 64, 1623-1636.	3.6	48
153	Molecular Profiling of Liver Tumors: Classification and Clinical Translation for Decision Making. <i>Seminars in Liver Disease</i> , 2014, 34, 363-375.	1.8	47
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