

# UALCAN: A Portal for Facilitating Tumor Subgroup Gen

Neoplasia

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Prostate Cancer: An Update on Molecular Pathology with Clinical Implications. European Urology Supplements, 2017, 16, 253-271.	0.1	2
2	STRAP Promotes Stemness of Human Colorectal Cancer via Epigenetic Regulation of the NOTCH Pathway. Cancer Research, 2017, 77, 5464-5478.	0.9	69
3	Identification of key genes of papillary thyroid cancer using integrated bioinformatics analysis. Journal of Endocrinological Investigation, 2018, 41, 1237-1245.	3.3	31
4	Germline Mutations in the Mitochondrial 2-Oxoglutarate/Malate Carrier <i>SLC25A11</i> Gene Confer a Predisposition to Metastatic Paragangliomas. Cancer Research, 2018, 78, 1914-1922.	0.9	96
5	Making Use of Cancer Genomic Databases. Current Protocols in Molecular Biology, 2018, 121, 19.14.1-19.14.13.	2.9	13
6	MethSurv: a web tool to perform multivariable survival analysis using DNA methylation data. Epigenomics, 2018, 10, 277-288.	2.1	381
7	Gene mutation patterns of Chinese acute myeloid leukemia patients by targeted next-generation sequencing and bioinformatic analysis. Clinica Chimica Acta, 2018, 479, 25-37.	1.1	9
8	C-C motif chemokine 22 predicts postoperative prognosis and adjuvant chemotherapeutic benefits in patients with stage II/III gastric cancer. OncoImmunology, 2018, 7, e1433517.	4.6	16
9	miR-34a Regulates Expression of the Stathmin-1 Oncoprotein and Prostate Cancer Progression. Molecular Cancer Research, 2018, 16, 1125-1137.	3.4	51
10	Targeting CDC7 improves sensitivity to chemotherapy of esophageal squamous cell carcinoma. OncoTargets and Therapy, 2019, Volume 12, 63-74.	2.0	11
11	Ca <sup>2+</sup> -Dependent Transcriptional Repressors KCNIP and Regulation of Prognosis Genes in Glioblastoma. Frontiers in Molecular Neuroscience, 2018, 11, 472.	2.9	27
12	Centromere protein U expression promotes non-small-cell lung cancer cell proliferation through FOXM1 and predicts poor survival. Cancer Management and Research, 2018, Volume 10, 6971-6984.	1.9	23
13	Transcriptional expressions of Chromobox 1/2/3/6/8 as independent indicators for survivals in hepatocellular carcinoma patients. Aging, 2018, 10, 3450-3473.	3.1	43
14	Bayesian variable selection for parametric survival model with applications to cancer omics data. Human Genomics, 2018, 12, 49.	2.9	11
15	Prior Knowledge Driven Joint NMF Algorithm for ceRNA Co-Module Identification. International Journal of Biological Sciences, 2018, 14, 1822-1833.	6.4	17
16	Solute Carrier Family 27 Member 4 (SLC27A4) Enhances Cell Growth, Migration, and Invasion in Breast Cancer Cells. International Journal of Molecular Sciences, 2018, 19, 3434.	4.1	54
17	Telomere length, <i>ZNF208</i> genetic variants and risk of chronic obstructive pulmonary disease in the Hainan Li population. Journal of Gene Medicine, 2018, 20, e3061.	2.8	4
18	Interaction of polymorphisms in xeroderma pigmentosum group C with cigarette smoking and pancreatic cancer risk. Oncology Letters, 2018, 16, 5631-5638.	1.8	7

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19	MARCH5 overexpression contributes to tumor growth and metastasis and associates with poor survival in breast cancer. <i>Cancer Management and Research</i> , 2019, Volume 11, 201-215.	1.9	16
20	Integrated analysis of the impact of age on genetic and clinical aspects of hepatocellular carcinoma. <i>Aging</i> , 2018, 10, 2079-2097.	3.1	6
21	HDAC is indispensable for IFN- $\gamma$ -induced B7-H1 expression in gastric cancer. <i>Clinical Epigenetics</i> , 2018, 10, 153.	4.1	38
22	Long Non-coding RNA DLEU1 Promotes Proliferation and Invasion by Interacting With miR-381 and Enhancing HOXA13 Expression in Cervical Cancer. <i>Frontiers in Genetics</i> , 2018, 9, 629.	2.3	42
23	Upregulated FFAR4 correlates with the epithelial-mesenchymal transition and an unfavorable prognosis in human cholangiocarcinoma. <i>Cancer Biomarkers</i> , 2018, 23, 353-361.	1.7	5
24	Bioinformatics analyses of significant genes, related pathways and candidate prognostic biomarkers in glioblastoma. <i>Molecular Medicine Reports</i> , 2018, 18, 4185-4196.	2.4	39
25	Identifying novel candidate biomarkers of RCC based on WGCNA analysis. <i>Personalized Medicine</i> , 2018, 15, 381-394.	1.5	12
26	Identification of invasion-metastasis-associated microRNAs in hepatocellular carcinoma based on bioinformatic analysis and experimental validation. <i>Journal of Translational Medicine</i> , 2018, 16, 266.	4.4	79
27	MiPanda: A Resource for Analyzing and Visualizing Next-Generation Sequencing Transcriptomics Data. <i>Neoplasia</i> , 2018, 20, 1144-1149.	5.3	20
28	ANRIL promotes chemoresistance via disturbing expression of ABCC1 by regulating the expression of Let-7a in colorectal cancer. <i>Bioscience Reports</i> , 2018, 38, .	2.4	43
29	High Expression of Glycolytic Genes in Cirrhosis Correlates With the Risk of Developing Liver Cancer. <i>Frontiers in Cell and Developmental Biology</i> , 2018, 6, 138.	3.7	56
30	Bioinformatics identification of crucial genes and pathways associated with hepatocellular carcinoma. <i>Bioscience Reports</i> , 2018, 38, .	2.4	43
31	Regulation of the master regulator FOXM1 in cancer. <i>Cell Communication and Signaling</i> , 2018, 16, 57.	6.5	241
32	Exosomal zinc transporter ZIP4 promotes cancer growth and is a novel diagnostic biomarker for pancreatic cancer. <i>Cancer Science</i> , 2018, 109, 2946-2956.	3.9	116
33	Analysis of Transcription Factor-Related Regulatory Networks Based on Bioinformatics Analysis and Validation in Hepatocellular Carcinoma. <i>BioMed Research International</i> , 2018, 2018, 1-16.	1.9	42
34	FOXC1 induces cancer stem cell-like properties through upregulation of beta-catenin in NSCLC. <i>Journal of Experimental and Clinical Cancer Research</i> , 2018, 37, 220.	8.6	69
35	LncRNA CDKN2B-AS1 promotes tumor growth and metastasis of human hepatocellular carcinoma by targeting let-7c-5p/NAP1L1 axis. <i>Cancer Letters</i> , 2018, 437, 56-66.	7.2	131
36	Inhibition of Proteasomal Deubiquitinase by Silver Complex Induces Apoptosis in Non-Small Cell Lung Cancer Cells. <i>Cellular Physiology and Biochemistry</i> , 2018, 49, 780-797.	1.6	20

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37	Overexpression of signal sequence receptor $\beta^3$ predicts poor survival in patients with hepatocellular carcinoma. Human Pathology, 2018, 81, 47-54.	2.0	13
38	RNA-seq Reveals the Overexpression of IGSF9 in Endometrial Cancer. Journal of Oncology, 2018, 2018, 1-13.	1.3	12
39	miR-134 targets PDCD7 to reduce E-cadherin expression and enhance oral cancer progression. International Journal of Cancer, 2018, 143, 2892-2904.	5.1	58
40	MiR-490-3p Functions As a Tumor Suppressor by Inhibiting Oncogene VDAC1 Expression in Colorectal Cancer. Journal of Cancer, 2018, 9, 1218-1230.	2.5	50
41	Wnt receptor Frizzled 8 is a target of ERG in prostate cancer. Prostate, 2018, 78, 1311-1320.	2.3	25
42	ANLN functions as a key candidate gene in cervical cancer as determined by integrated bioinformatic analysis. Cancer Management and Research, 2018, Volume 10, 663-670.	1.9	74
43	Bioinformatic identification of key genes and analysis of prognostic values in clear cell renal cell carcinoma. Oncology Letters, 2018, 16, 1747-1757.	1.8	25
44	Identification of potential target genes in pancreatic ductal adenocarcinoma by bioinformatics analysis. Oncology Letters, 2018, 16, 2453-2461.	1.8	26
45	Prognostic Value of HMGA2 in Human Cancers: A Meta-Analysis Based on Literatures and TCGA Datasets. Frontiers in Physiology, 2018, 9, 776.	2.8	21
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47	The Effect of GPRC5a on the Proliferation, Migration Ability, Chemotherapy Resistance, and Phosphorylation of GSK-3 $\beta$ in Pancreatic Cancer. International Journal of Molecular Sciences, 2018, 19, 1870.	4.1	28
48	Dependence receptor UNC5A restricts luminal to basal breast cancer plasticity and metastasis. Breast Cancer Research, 2018, 20, 35.	5.0	14
49	Eukaryotic Translation Initiation Factor 4 Gamma 1 (eIF4G1) is upregulated during Prostate cancer progression and modulates cell growth and metastasis. Scientific Reports, 2018, 8, 7459.	3.3	31
50	A Role for De Novo Purine Metabolic Enzyme PAICS in Bladder Cancer Progression. Neoplasia, 2018, 20, 894-904.	5.3	50
51	Identification of biomarkers for Barcelona Clinic Liver Cancer staging and overall survival of patients with hepatocellular carcinoma. PLoS ONE, 2018, 13, e0202763.	2.5	40
52	Construction and analysis of mRNA, miRNA, lncRNA, and TF regulatory networks reveal the key genes associated with prostate cancer. PLoS ONE, 2018, 13, e0198055.	2.5	58
53	Aurovertin B sensitizes colorectal cancer cells to NK cell recognition and lysis. Biochemical and Biophysical Research Communications, 2018, 503, 3057-3063.	2.1	13
54	Biochemical and Epigenetic Insights into L-2-Hydroxyglutarate, a Potential Therapeutic Target in Renal Cancer. Clinical Cancer Research, 2018, 24, 6433-6446.	7.0	54

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55	Dowregulation of OTX1 attenuates gastric cancer cell proliferation, migration and invasion. <i>Oncology Reports</i> , 2018, 40, 1907-1916.	2.6	18
56	Competing endogenous RNA analysis reveals the regulatory potency of circRNA_036186 in HNSCC. <i>International Journal of Oncology</i> , 2018, 53, 1529-1543.	3.3	11
57	JMJD6 promotes hepatocellular carcinoma carcinogenesis by targeting CDK4. <i>International Journal of Cancer</i> , 2019, 144, 2489-2500.	5.1	52
58	Integrative Epigenetic and Gene Expression Analysis of Renal Tumor Progression to Metastasis. <i>Molecular Cancer Research</i> , 2019, 17, 84-96.	3.4	37
59	Dual regulatory role of CCNA2 in modulating CDK6 and METâ€mediated cellâ€cycle pathway and EMT progression is blocked by miRâ€381â€3p in bladder cancer. <i>FASEB Journal</i> , 2019, 33, 1374-1388.	0.5	60
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62	MKL1 overexpression predicts poor prognosis in patients with papillary thyroid cancer and promotes nodal metastasis. <i>Journal of Cell Science</i> , 2019, 132, .	2.0	9
63	Long noncoding RNA DIO3OS interacts with miR-122 to promote proliferation and invasion of pancreatic cancer cells through upregulating ALDOA. <i>Cancer Cell International</i> , 2019, 19, 202.	4.1	35
64	Expression and Role of Methylenetetrahydrofolate Dehydrogenase 1 Like (MTHFD1L) in Bladder Cancer. <i>Translational Oncology</i> , 2019, 12, 1416-1424.	3.7	21
65	Prognostic roles of the transcriptional expression of exportins in hepatocellular carcinoma. <i>Bioscience Reports</i> , 2019, 39, .	2.4	14
66	Overexpression of TWA1 predicts poor prognosis in patients with gastric cancer. <i>Pathology Research and Practice</i> , 2019, 215, 152594.	2.3	4
67	Comprehensive analysis of peroxiredoxins expression profiles and prognostic values in breast cancer. <i>Biomarker Research</i> , 2019, 7, 16.	6.8	22
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69	ENKUR acts as a tumor suppressor in lung adenocarcinoma cells through PI3K/Akt and MAPK/ERK signaling pathways. <i>Journal of Cancer</i> , 2019, 10, 3975-3984.	2.5	13
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73	Upregulated Expression of CUX1 Correlates with Poor Prognosis in Glioma Patients: a Bioinformatic Analysis. <i>Journal of Molecular Neuroscience</i> , 2019, 69, 527-537.	2.3	5
74	Identification of LCN1 as a Potential Biomarker for Breast Cancer by Bioinformatic Analysis. <i>DNA and Cell Biology</i> , 2019, 38, 1088-1099.	1.9	17
75	Identification of Potential Crucial Genes and Key Pathways in Breast Cancer Using Bioinformatic Analysis. <i>Frontiers in Genetics</i> , 2019, 10, 695.	2.3	161
76	Identification of CFTR as a novel key gene in chromophobe renal cell carcinoma through bioinformatics analysis. <i>Oncology Letters</i> , 2019, 18, 1767-1774.	1.8	5
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78	Mechanosensitive ion channel Piezo1 promotes prostate cancer development through the activation of the Akt/mTOR pathway and acceleration of cell cycle. <i>International Journal of Oncology</i> , 2019, 55, 629-644.	3.3	55
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83	Screening of key genes and prediction of therapeutic agents in Arsenic-induced lung carcinoma. <i>Cancer Biomarkers</i> , 2019, 25, 351-360.	1.7	7
84	UBL4A inhibits autophagy-mediated proliferation and metastasis of pancreatic ductal adenocarcinoma via targeting LAMP1. <i>Journal of Experimental and Clinical Cancer Research</i> , 2019, 38, 297.	8.6	40
85	CENPE promotes lung adenocarcinoma proliferation and is directly regulated by FOXM1. <i>International Journal of Oncology</i> , 2019, 55, 257-266.	3.3	39
86	The systemic activin response to pancreatic cancer: implications for effective cancer cachexia therapy. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2019, 10, 1083-1101.	7.3	46
87	High expression of CDH3 predicts a good prognosis for colon adenocarcinoma patients. <i>Experimental and Therapeutic Medicine</i> , 2019, 18, 841-847.	1.8	11
88	Characterizing methylation regulated miRNA in carcinoma of the human uterine cervix. <i>Life Sciences</i> , 2019, 232, 116668.	4.3	10
89	miR-125a-5p Functions as Tumor Suppressor microRNA And Is a Marker of Locoregional Recurrence And Poor prognosis in Head And Neck Cancer. <i>Neoplasia</i> , 2019, 21, 849-862.	5.3	39
90	EZH2 regulates PD-L1 expression via HIF-1 $\alpha$ in non-small cell lung cancer cells. <i>Biochemical and Biophysical Research Communications</i> , 2019, 517, 201-209.	2.1	51

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95	Relation of AURKB over-expression to low survival rate in BCRA and reversine-modulated aurora B kinase in breast cancer cell lines. Cancer Cell International, 2019, 19, 166.	4.1	37
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105	Knockdown of COPS3 inhibits the progress of prostate cancer through reducing phosphorylated p38 MAPK expression and impairs the epithelial-mesenchymal transition process. Prostate, 2019, 79, 1823-1831.	2.3	7
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109	<p>TCDD-Inducible Poly-ADP-Ribose Polymerase (TIPARP), A Novel Therapeutic Target Of Breast Cancer</p>. Cancer Management and Research, 2019, Volume 11, 8991-9004.	1.9	13
110	Identification of Prognostic and Metastatic Alternative Splicing Signatures in Kidney Renal Clear Cell Carcinoma. Frontiers in Bioengineering and Biotechnology, 2019, 7, 270.	4.1	55
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117	miR-451a inhibits cancer growth, epithelial-mesenchymal transition and induces apoptosis in papillary thyroid cancer by targeting PSMB8. Journal of Cellular and Molecular Medicine, 2019, 23, 8067-8075.	3.6	31
118	A novel BMI-1 inhibitor QW24 for the treatment of stem-like colorectal cancer. Journal of Experimental and Clinical Cancer Research, 2019, 38, 422.	8.6	23
119	NAMPT and NAPRT, Key Enzymes in NAD Salvage Synthesis Pathway, Are of Negative Prognostic Value in Colorectal Cancer. Frontiers in Oncology, 2019, 9, 736.	2.8	39
120	Comprehensive analysis of histone modification-associated genes on differential gene expression and prognosis in gastric cancer. Experimental and Therapeutic Medicine, 2019, 18, 2219-2230.	1.8	9
121	Elevated SPINK2 gene expression is a predictor of poor prognosis in acute myeloid leukemia. Oncology Letters, 2019, 18, 2877-2884.	1.8	9
122	Bioinformatics-based discovery of PYGM and TNNC2 as potential biomarkers of head and neck squamous cell carcinoma. Bioscience Reports, 2019, 39, .	2.4	20
123	LPP and RYR2 Gene Polymorphisms Correlate with the Risk and the Prognosis of Astrocytoma. Journal of Molecular Neuroscience, 2019, 69, 628-635.	2.3	8
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126	FABP5 is correlated with poor prognosis and promotes tumour cell growth and metastasis in clear cell renal cell carcinoma. European Journal of Pharmacology, 2019, 862, 172637.	3.5	34



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128	The Emerging Role of NANOG as an Early Cancer Risk Biomarker in Patients with Oral Potentially Malignant Disorders. Journal of Clinical Medicine, 2019, 8, 1376.	2.4	25
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130	Expression and clinical significance of CPS1 in glioblastoma multiforme. Current Research in Translational Medicine, 2019, 67, 123-128.	1.8	15
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132	WNT5B governs the phenotype of basal-like breast cancer by activating WNT signaling. Cell Communication and Signaling, 2019, 17, 109.	6.5	45
133	&lt;p&gt;Overexpression of KRT17 promotes proliferation and invasion of non-small cell lung cancer and indicates poor prognosis&lt;/p&gt;. Cancer Management and Research, 2019, Volume 11, 7485-7497.	1.9	53
134	Prexasertib treatment induces homologous recombination deficiency and synergizes with olaparib in triple-negative breast cancer cells. Breast Cancer Research, 2019, 21, 104.	5.0	45
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140	UFM1 suppresses invasive activities of gastric cancer cells by attenuating the expression of PDK1 through PI3K/AKT signaling. Journal of Experimental and Clinical Cancer Research, 2019, 38, 410.	8.6	42
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144	&lt;p&gt;Downregulated miR-585-3p promotes cell growth and proliferation in colon cancer by upregulating &lt;em&gt;PSME3&lt;/em&gt;&lt;/p&gt;. OncoTargets and Therapy, 2019, Volume 12, 6525-6534.	2.0	14

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146	Tribbles Homolog 3 Involved in Radiation Response of Triple Negative Breast Cancer Cells by Regulating Notch1 Activation. <i>Cancers</i> , 2019, 11, 127.	3.7	22
147	Lymphoid enhancer-binding factor-1 promotes stemness and poor differentiation of hepatocellular carcinoma by directly activating the NOTCH pathway. <i>Oncogene</i> , 2019, 38, 4061-4074.	5.9	31
148	Bladder cancer stage-associated hub genes revealed by WGCNA co-expression network analysis. <i>Hereditas</i> , 2019, 156, 7.	1.4	79
149	GALNT6 Promotes Tumorigenicity and Metastasis of Breast Cancer Cell via $\beta$ -catenin/MUC1-C Signaling Pathway. <i>International Journal of Biological Sciences</i> , 2019, 15, 169-182.	6.4	20
150	MiR-30a-5p frequently downregulated in prostate cancer inhibits cell proliferation via targeting PCLAF. <i>Artificial Cells, Nanomedicine and Biotechnology</i> , 2019, 47, 278-289.	2.8	30
151	Comprehensive bioinformation analysis of methylated and differentially expressed genes in esophageal squamous cell carcinoma. <i>Molecular Omics</i> , 2019, 15, 88-100.	2.8	9
152	Eleven genes associated with progression and prognosis of endometrial cancer (EC) identified by comprehensive bioinformatics analysis. <i>Cancer Cell International</i> , 2019, 19, 136.	4.1	61
153	The BET inhibitor JQ1 attenuates double-strand break repair and sensitizes models of pancreatic ductal adenocarcinoma to PARP inhibitors. <i>EBioMedicine</i> , 2019, 44, 419-430.	6.1	76
154	Parvifoline AA Promotes Susceptibility of Hepatocarcinoma to Natural Killer Cell-Mediated Cytolysis by Targeting Peroxiredoxin. <i>Cell Chemical Biology</i> , 2019, 26, 1122-1132.e6.	5.2	15
155	Knockdown of long noncoding RNA PVT1 suppresses cell proliferation and invasion of colorectal cancer via upregulation of microRNA-214-3p. <i>American Journal of Physiology - Renal Physiology</i> , 2019, 317, G222-G232.	3.4	57
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159	RECQL5 plays an essential role in maintaining genome stability and viability of triple-negative breast cancer cells. <i>Cancer Medicine</i> , 2019, 8, 4743-4752.	2.8	14
160	UDP-glucose accelerates SNAI1 mRNA decay and impairs lung cancer metastasis. <i>Nature</i> , 2019, 571, 127-131.	27.8	140
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1704	Identification and Validation of Afatinib Potential Drug Resistance Gene BIRC5 in Non-Small Cell Lung Cancer. <i>Frontiers in Oncology</i> , 2021, 11, 763035.	2.8	5
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1955	Identification of key miRNAs in prostate cancer progression based on miRNA-mRNA network construction. <i>Computational and Structural Biotechnology Journal</i> , 2022, 20, 864-873.	4.1	4
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1966	KLF16 Downregulates the Expression of Tumor Suppressor Gene TGFBR3 to Promote Bladder Cancer Proliferation and Migration. <i>Cancer Management and Research</i> , 2022, Volume 14, 465-477.	1.9	8
1967	Iron Activates cGAS-STING Signaling and Promotes Hepatic Inflammation. <i>Journal of Agricultural and Food Chemistry</i> , 2022, 70, 2211-2220.	5.2	14
1968	Ubiquitin-conjugating enzyme 2C (UBE2C) is a poor prognostic biomarker in invasive breast cancer. <i>Breast Cancer Research and Treatment</i> , 2022, 192, 529-539.	2.5	11
1969	Characterization of the prognostic and diagnostic values of ALKBH family members in non-small cell lung cancer. <i>Pathology Research and Practice</i> , 2022, 231, 153809.	2.3	6
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1976	Aberrant epigenetic and transcriptional events associated with breast cancer risk. <i>Clinical Epigenetics</i> , 2022, 14, 21.	4.1	14
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2026	FAM83A is a potential biomarker for breast cancer initiation. <i>Biomarker Research</i> , 2022, 10, 8.	6.8	9
2027	MYSM1 induces apoptosis and sensitizes TNBC cells to cisplatin via RSK3â€™phospho-BAD pathway. <i>Cell Death Discovery</i> , 2022, 8, 84.	4.7	5
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