

# Qian Tao

## List of Publications by Year in descending order

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182  
papers

10,376  
citations

22099

59  
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45213

90  
g-index

186  
all docs

186  
docs citations

186  
times ranked

12534  
citing authors

#	ARTICLE	IF	CITATIONS
1	Nasopharyngeal carcinoma: molecular pathogenesis and therapeutic developments. <i>Expert Reviews in Molecular Medicine</i> , 2007, 9, 1-24.	1.6	266
2	Functional epigenetics identifies a protocadherin PCDH10 as a candidate tumor suppressor for nasopharyngeal, esophageal and multiple other carcinomas with frequent methylation. <i>Oncogene</i> , 2006, 25, 1070-1080.	2.6	247
3	The Stress-Responsive Gene GADD45G Is a Functional Tumor Suppressor, with Its Response to Environmental Stresses Frequently Disrupted Epigenetically in Multiple Tumors. <i>Clinical Cancer Research</i> , 2005, 11, 6442-6449.	3.2	220
4	DNA methyltransferase 3B (DNMT3B) mutations in ICF syndrome lead to altered epigenetic modifications and aberrant expression of genes regulating development, neurogenesis and immune function. <i>Human Molecular Genetics</i> , 2008, 17, 690-709.	1.4	216
5	Nasopharyngeal carcinoma: an evolving paradigm. <i>Nature Reviews Clinical Oncology</i> , 2021, 18, 679-695.	12.5	207
6	Methylation of Protocadherin 10, a Novel Tumor Suppressor, Is Associated With Poor Prognosis in Patients With Gastric Cancer. <i>Gastroenterology</i> , 2009, 136, 640-651.e1.	0.6	190
7	<i>WNT5A</i> Exhibits Tumor-Suppressive Activity through Antagonizing the Wnt/ $\beta$ -Catenin Signaling, and Is Frequently Methylated in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2008, 14, 55-61.	3.2	181
8	Epstein-Barr Virus (EBV) in Endemic Burkitt's Lymphoma: Molecular Analysis of Primary Tumor Tissue. <i>Blood</i> , 1998, 91, 1373-1381.	0.6	169
9	Inactivation of Wnt inhibitory factor-1 (WIF1) expression by epigenetic silencing is a common event in breast cancer. <i>Carcinogenesis</i> , 2006, 27, 1341-1348.	1.3	169
10	Characterization of Epstein-Barr virus-infected B cells in patients with posttransplantation lymphoproliferative disease: disappearance after rituximab therapy does not predict clinical response. <i>Blood</i> , 2000, 96, 4055-4063.	0.6	167
11	Epigenetic Therapy Using Belinostat for Patients With Unresectable Hepatocellular Carcinoma: A Multicenter Phase I/II Study With Biomarker and Pharmacokinetic Analysis of Tumors From Patients in the Mayo Phase II Consortium and the Cancer Therapeutics Research Group. <i>Journal of Clinical Oncology</i> , 2012, 30, 3361-3367.	0.8	167
12	Epigenetic disruption of the WNT/ $\beta$ -catenin signaling pathway in human cancers. <i>Epigenetics</i> , 2009, 4, 307-312.	1.3	159
13	Nasal NK- and T-cell lymphomas share the same type of Epstein-Barr virus latency as nasopharyngeal carcinoma and Hodgkin's disease. , 1996, 68, 285-290.		143
14	The Tumor Suppressor UCHL1 Forms a Complex with p53/MDM2/ARF to Promote p53 Signaling and Is Frequently Silenced in Nasopharyngeal Carcinoma. <i>Clinical Cancer Research</i> , 2010, 16, 2949-2958.	3.2	136
15	Epigenetic identification of ubiquitin carboxyl-terminal hydrolase L1 as a functional tumor suppressor and biomarker for hepatocellular carcinoma and other digestive tumors. <i>Hepatology</i> , 2008, 48, 508-518.	3.6	134
16	Defective de novo methylation of viral and cellular DNA sequences in ICF syndrome cells. <i>Human Molecular Genetics</i> , 2002, 11, 2091-2102.	1.4	131
17	The candidate tumor suppressor gene BLU, located at the commonly deleted region 3p21.3, is an E2F-regulated, stress-responsive gene and inactivated by both epigenetic and genetic mechanisms in nasopharyngeal carcinoma. <i>Oncogene</i> , 2004, 23, 4793-4806.	2.6	130
18	Azacitidine Induces Demethylation of the Epstein-Barr Virus Genome in Tumors. <i>Journal of Clinical Oncology</i> , 2004, 22, 1373-1381.	0.8	129

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19	Cellular uptake, evolution, and excretion of silica nanoparticles in human cells. <i>Nanoscale</i> , 2011, 3, 3291.	2.8	121
20	The major 8p22 tumor suppressor DLC1 is frequently silenced by methylation in both endemic and sporadic nasopharyngeal, esophageal, and cervical carcinomas, and inhibits tumor cell colony formation. <i>Oncogene</i> , 2007, 26, 934-944.	2.6	119
21	Epigenetic silencing of a Ca <sup>2+</sup> -regulated Ras GTPase-activating protein RASAL defines a new mechanism of Ras activation in human cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 12353-12358.	3.3	118
22	Epstein-Barr virus (EBV) and its associated human cancers - Genetics, epigenetics, pathobiology and novel therapeutics. <i>Frontiers in Bioscience - Landmark</i> , 2006, 11, 2672.	3.0	116
23	KRAB Zinc Finger Protein ZNF382 Is a Proapoptotic Tumor Suppressor That Represses Multiple Oncogenes and Is Commonly Silenced in Multiple Carcinomas. <i>Cancer Research</i> , 2010, 70, 6516-6526.	0.4	116
24	The Ubiquitin Peptidase UCHL1 Induces G0/G1 Cell Cycle Arrest and Apoptosis Through Stabilizing p53 and Is Frequently Silenced in Breast Cancer. <i>PLoS ONE</i> , 2012, 7, e29783.	1.1	116
25	Promoter methylation of the Wnt/ $\beta$ -catenin signaling antagonist <i>Dkk3</i> is associated with poor survival in gastric cancer. <i>Cancer</i> , 2009, 115, 49-60.	2.0	115
26	Frequent epigenetic inactivation of secreted frizzled-related protein 2 (SFRP2) by promoter methylation in human gastric cancer. <i>British Journal of Cancer</i> , 2007, 97, 895-901.	2.9	112
27	ADAMTS9 is a functional tumor suppressor through inhibiting AKT/mTOR pathway and associated with poor survival in gastric cancer. <i>Oncogene</i> , 2013, 32, 3319-3328.	2.6	108
28	Receptor-type tyrosine-protein phosphatase $\hat{p}$ directly targets STAT3 activation for tumor suppression in nasal NK/T-cell lymphoma. <i>Blood</i> , 2015, 125, 1589-1600.	0.6	108
29	Epigenetic identification of ADAMTS18 as a novel 16q23.1 tumor suppressor frequently silenced in esophageal, nasopharyngeal and multiple other carcinomas. <i>Oncogene</i> , 2007, 26, 7490-7498.	2.6	106
30	Methylation Status of the Epstein-Barr Virus Major Latent Promoter C in Iatrogenic B Cell Lymphoproliferative Disease. <i>American Journal of Pathology</i> , 1999, 155, 619-625.	1.9	100
31	Protocadherin 17 acts as a tumour suppressor inducing tumour cell apoptosis and autophagy, and is frequently methylated in gastric and colorectal cancers. <i>Journal of Pathology</i> , 2013, 229, 62-73.	2.1	98
32	<i>CMTM3</i> , Located at the Critical Tumor Suppressor Locus 16q22.1, Is Silenced by CpG Methylation in Carcinomas and Inhibits Tumor Cell Growth through Inducing Apoptosis. <i>Cancer Research</i> , 2009, 69, 5194-5201.	0.4	95
33	The tumor suppressor Wnt inhibitory factor 1 is frequently methylated in nasopharyngeal and esophageal carcinomas. <i>Laboratory Investigation</i> , 2007, 87, 644-650.	1.7	93
34	OPCML Is a Broad Tumor Suppressor for Multiple Carcinomas and Lymphomas with Frequently Epigenetic Inactivation. <i>PLoS ONE</i> , 2008, 3, e2990.	1.1	92
35	The human cadherin 11 is a pro-apoptotic tumor suppressor modulating cell stemness through Wnt/ $\beta$ -catenin signaling and silenced in common carcinomas. <i>Oncogene</i> , 2012, 31, 3901-3912.	2.6	92
36	Evidence for lytic infection by Epstein-Barr virus in mucosal lymphocytes instead of nasopharyngeal epithelial cells in normal individuals.. <i>Journal of Medical Virology</i> , 1995, 45, 71-77.	2.5	91

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37	Epigenetic inactivation of the CpG demethylase TET1 as a DNA methylation feedback loop in human cancers. <i>Scientific Reports</i> , 2016, 6, 26591.	1.6	90
38	Epstein-Barr virus is localized in the tumour cells of nasal lymphomas of NK, T or B cell type. <i>International Journal of Cancer</i> , 1995, 60, 315-320.	2.3	89
39	Authentication of nasopharyngeal carcinoma tumor lines. <i>International Journal of Cancer</i> , 2008, 122, 2169-2171.	2.3	88
40	Stealth technology: how Epstein-Barr virus utilizes DNA methylation to cloak itself from immune detection. <i>Clinical Immunology</i> , 2003, 109, 53-63.	1.4	84
41	DACT1, an antagonist to Wnt/ $\beta$ -catenin signaling, suppresses tumor cell growth and is frequently silenced in breast cancer. <i>Breast Cancer Research</i> , 2013, 15, R23.	2.2	83
42	DNA methylation and the Epstein-Barr virus. <i>Seminars in Cancer Biology</i> , 1999, 9, 369-375.	4.3	82
43	Epigenetic disruption of the WNT/ $\beta$ -catenin signaling pathway in human cancers. <i>Epigenetics</i> , 2009, 4, 307-12.	1.3	82
44	Zinc-finger protein 331, a novel putative tumor suppressor, suppresses growth and invasiveness of gastric cancer. <i>Oncogene</i> , 2013, 32, 307-317.	2.6	76
45	Epigenetic disruption of two proapoptotic genes MAPK10/JNK3 and PTPN13/FAP-1 in multiple lymphomas and carcinomas through hypermethylation of a common bidirectional promoter. <i>Leukemia</i> , 2006, 20, 1173-1175.	3.3	75
46	Phospholipase C delta 1 is a novel 3p22.3 tumor suppressor involved in cytoskeleton organization, with its epigenetic silencing correlated with high-stage gastric cancer. <i>Oncogene</i> , 2009, 28, 2466-2475.	2.6	72
47	Epigenetic disruption of interferon- $\beta$ response through silencing the tumor suppressor interferon regulatory factor 8 in nasopharyngeal, esophageal and multiple other carcinomas. <i>Oncogene</i> , 2008, 27, 5267-5276.	2.6	71
48	WNT5A antagonizes WNT/ $\beta$ -catenin signaling and is frequently silenced by promoter CpG methylation in esophageal squamous cell carcinoma. <i>Cancer Biology and Therapy</i> , 2010, 10, 617-624.	1.5	71
49	<i>CMTM5</i> Exhibits Tumor Suppressor Activities and Is Frequently Silenced by Methylation in Carcinoma Cell Lines. <i>Clinical Cancer Research</i> , 2007, 13, 5756-5762.	3.2	69
50	Epigenetic inactivation of BCL6B, a novel functional tumour suppressor for gastric cancer, is associated with poor survival. <i>Gut</i> , 2012, 61, 977-985.	6.1	69
51	The Epstein-Barr Virus Major Latent Promoter Qp Is Constitutively Active, Hypomethylated, and Methylation Sensitive. <i>Journal of Virology</i> , 1998, 72, 7075-7083.	1.5	69
52	Frequent Hypermethylation of RASSF1A, TSLC1, High Viral Load of Epstein-Barr Virus DNA in Nasopharyngeal Carcinoma, Matched Tumor-Adjacent Tissues. <i>Neoplasia</i> , 2005, 7, 809-815.	2.3	68
53	STAT3 activation contributes directly to Epstein-Barr virus-mediated invasiveness of nasopharyngeal cancer cells <i>in vitro</i> . <i>International Journal of Cancer</i> , 2009, 125, 1884-1893.	2.3	67
54	DLC1-dependent parathyroid hormone-like hormone inhibition suppresses breast cancer bone metastasis. <i>Journal of Clinical Investigation</i> , 2014, 124, 1646-1659.	3.9	67

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55	Double restriction-enzyme digestion improves the coverage and accuracy of genome-wide CpG methylation profiling by reduced representation bisulfite sequencing. <i>BMC Genomics</i> , 2013, 14, 11.	1.2	64
56	A novel 3p22.3 gene CMTM7 represses oncogenic EGFR signaling and inhibits cancer cell growth. <i>Oncogene</i> , 2014, 33, 3109-3118.	2.6	64
57	The Epigenetic Modifier PRDM5 Functions as a Tumor Suppressor through Modulating WNT/ $\beta$ -Catenin Signaling and Is Frequently Silenced in Multiple Tumors. <i>PLoS ONE</i> , 2011, 6, e27346.	1.1	64
58	Frequent epigenetic inactivation of the RASSF1A tumor suppressor gene in Hodgkin's lymphoma. <i>Oncogene</i> , 2004, 23, 1326-1331.	2.6	63
59	High BCL6 expression predicts better prognosis, independent of BCL6 translocation status, translocation partner, or BCL6-deregulating mutations, in gastric lymphoma. <i>Blood</i> , 2006, 108, 2373-2383.	0.6	63
60	Paired box gene 5 is a novel tumor suppressor in hepatocellular carcinoma through interaction with p53 signaling pathway. <i>Hepatology</i> , 2011, 53, 843-853.	3.6	63
61	Epigenetic inactivation of paired box gene 5, a novel tumor suppressor gene, through direct upregulation of p53 is associated with prognosis in gastric cancer patients. <i>Oncogene</i> , 2012, 31, 3419-3430.	2.6	62
62	Patterned CpG Methylation of Silenced B Cell Gene Promoters in Classical Hodgkin Lymphoma-derived and Primary Effusion Lymphoma Cell Lines. <i>Journal of Molecular Biology</i> , 2005, 350, 631-640.	2.0	61
63	Protocadherin PCDH10, involved in tumor progression, is a frequent and early target of promoter hypermethylation in cervical cancer. <i>Genes Chromosomes and Cancer</i> , 2009, 48, 983-992.	1.5	61
64	DLEC1 is a functional 3p22.3 tumour suppressor silenced by promoter CpG methylation in colon and gastric cancers. <i>British Journal of Cancer</i> , 2009, 100, 663-669.	2.9	60
65	Epigenetic silencing of the WNT antagonist Dickkopf 3 disrupts normal Wnt/ $\beta$ -catenin signalling and apoptosis regulation in breast cancer cells. <i>Journal of Cellular and Molecular Medicine</i> , 2013, 17, 1236-1246.	1.6	60
66	CaMKII-mediated Beclin 1 phosphorylation regulates autophagy that promotes degradation of Id and neuroblastoma cell differentiation. <i>Nature Communications</i> , 2017, 8, 1159.	5.8	60
67	Epigenetic disruption of cell signaling in nasopharyngeal carcinoma. <i>Chinese Journal of Cancer</i> , 2011, 30, 231-239.	4.9	59
68	A survey of methylated candidate tumor suppressor genes in nasopharyngeal carcinoma. <i>International Journal of Cancer</i> , 2011, 128, 1393-1403.	2.3	59
69	The Metalloprotease ADAMTS8 Displays Antitumor Properties through Antagonizing EGFR-MEK-ERK Signaling and Is Silenced in Carcinomas by CpG Methylation. <i>Molecular Cancer Research</i> , 2014, 12, 228-238.	1.5	58
70	Recurrent ECSIT mutation encoding V140A triggers hyperinflammation and promotes hemophagocytic syndrome in extranodal NK/T cell lymphoma. <i>Nature Medicine</i> , 2018, 24, 154-164.	15.2	58
71	The preclinical activity of the histone deacetylase inhibitor PXD101 (belinostat) in hepatocellular carcinoma cell lines. <i>Investigational New Drugs</i> , 2010, 28, 107-114.	1.2	56
72	SOX10, a novel HMG-box-containing tumor suppressor, inhibits growth and metastasis of digestive cancers by suppressing the Wnt/ $\beta$ -catenin pathway. <i>Oncotarget</i> , 2014, 5, 10571-10583.	0.8	56

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73	Lipidâ€“Polymer Nanoparticles Encapsulating Doxorubicin and 2â€“Deoxy-5-azacytidine Enhance the Sensitivity of Cancer Cells to Chemical Therapeutics. <i>Molecular Pharmaceutics</i> , 2013, 10, 1901-1909.	2.3	53
74	Frequent epigenetic silencing of protocadherin 10 by methylation in multiple haematologic malignancies. <i>British Journal of Haematology</i> , 2007, 136, 829-832.	1.2	52
75	PLCD1 is a functional tumor suppressor inducing G <sub>2</sub> /M arrest and frequently methylated in breast cancer. <i>Cancer Biology and Therapy</i> , 2010, 10, 520-527.	1.5	52
76	Characterization of the nasopharyngeal carcinoma methylome identifies aberrant disruption of key signaling pathways and methylated tumor suppressor genes. <i>Epigenomics</i> , 2015, 7, 155-173.	1.0	52
77	The tumor suppressor gene DLEC1 is frequently silenced by DNA methylation in hepatocellular carcinoma and induces G1 arrest in cell cycle. <i>Journal of Hepatology</i> , 2008, 48, 433-441.	1.8	51
78	Coinfection of multiple strains of Epstein-Barr virus in immunocompetent normal individuals: reassessment of the viral carrier state. <i>Blood</i> , 2000, 95, 2443-2445.	0.6	51
79	WNT5A is epigenetically silenced in hematologic malignancies and inhibits leukemia cell growth as a tumor suppressor. <i>Blood</i> , 2007, 110, 4130-4131.	0.6	50
80	Genome-wide expression analysis using microarray identified complex signaling pathways modulated by hypoxia in nasopharyngeal carcinoma. <i>Cancer Letters</i> , 2007, 253, 74-88.	3.2	50
81	OVOL2 links stemness and metastasis via fine-tuning epithelial-mesenchymal transition in nasopharyngeal carcinoma. <i>Theranostics</i> , 2018, 8, 2202-2216.	4.6	50
82	DNA methylation downregulated ZDHHC1 suppresses tumor growth by altering cellular metabolism and inducing oxidative/ER stress-mediated apoptosis and pyroptosis. <i>Theranostics</i> , 2020, 10, 9495-9511.	4.6	50
83	Viral oncoprotein LMP1 disrupts p53-induced cell cycle arrest and apoptosis through modulating K63-linked ubiquitination of p53. <i>Cell Cycle</i> , 2012, 11, 2327-2336.	1.3	49
84	USP3 promotes breast cancer cell proliferation by deubiquitinating KLF5. <i>Journal of Biological Chemistry</i> , 2019, 294, 17837-17847.	1.6	49
85	Overexpression of Cyclooxygenase-2 in Nasopharyngeal Carcinoma and Association With Epidermal Growth Factor Receptor Expression. <i>JAMA Otolaryngology</i> , 2005, 131, 147.	1.5	48
86	Down-regulation of tyrosine aminotransferase at a frequently deleted region 16q22 contributes to the pathogenesis of hepatocellular carcinoma. <i>Hepatology</i> , 2010, 51, 1624-1634.	3.6	48
87	A novel isoform of the 8p22 tumor suppressor gene DLC1 suppresses tumor growth and is frequently silenced in multiple common tumors. <i>Oncogene</i> , 2011, 30, 1923-1935.	2.6	48
88	Characterization of naturally Epsteinâ€“Barr virus-infected gastric carcinoma cell line YCCEL1. <i>Journal of General Virology</i> , 2013, 94, 497-506.	1.3	47
89	BS69, a Specific Adaptor in the Latent Membrane Protein 1-Mediated c-Jun N-Terminal Kinase Pathway. <i>Molecular and Cellular Biology</i> , 2006, 26, 448-456.	1.1	46
90	Aberrant Promoter Methylation of <i>DLEC1</i> , a Critical 3p22 Tumor Suppressor for Renal Cell Carcinoma, is Associated With More Advanced Tumor Stage. <i>Journal of Urology</i> , 2010, 184, 731-737.	0.2	46

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91	Zinc-finger protein 545 is a novel tumour suppressor that acts by inhibiting ribosomal RNA transcription in gastric cancer. <i>Gut</i> , 2013, 62, 833-841.	6.1	46
92	Protocadherin 17 functions as a tumor suppressor suppressing Wnt/ $\beta$ 2-catenin signaling and cell metastasis and is frequently methylated in breast cancer. <i>Oncotarget</i> , 2016, 7, 51720-51732.	0.8	46
93	Oncogenic HOXB8 is driven by MYC-regulated super-enhancer and potentiates colorectal cancer invasiveness via BACH1. <i>Oncogene</i> , 2020, 39, 1004-1017.	2.6	45
94	Lack of somatic mutations in EGFR tyrosine kinase domain in hepatocellular and nasopharyngeal carcinoma. <i>Pharmacogenetics and Genomics</i> , 2006, 16, 73-74.	0.7	45
95	A Novel 19q13 Nucleolar Zinc Finger Protein Suppresses Tumor Cell Growth through Inhibiting Ribosome Biogenesis and Inducing Apoptosis but Is Frequently Silenced in Multiple Carcinomas. <i>Molecular Cancer Research</i> , 2012, 10, 925-936.	1.5	44
96	FEZF2, a novel 3p14 tumor suppressor gene, represses oncogene EZH2 and MDM2 expression and is frequently methylated in nasopharyngeal carcinoma. <i>Carcinogenesis</i> , 2013, 34, 1984-1993.	1.3	44
97	Aberrant promoter CpG methylation and its translational applications in breast cancer. <i>Chinese Journal of Cancer</i> , 2013, 32, 12-20.	4.9	44
98	The ECM protein LTBP2 is a suppressor of esophageal squamous cell carcinoma tumor formation but higher tumor expression associates with poor patient outcome. <i>International Journal of Cancer</i> , 2011, 129, 565-573.	2.3	43
99	Epigenetic identification of receptor tyrosine kinase-like orphan receptor 2 as a functional tumor suppressor inhibiting $\beta$ 2-catenin and AKT signaling but frequently methylated in common carcinomas. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 2179-2192.	2.4	43
100	Preclinical activity of gefitinib in non-keratinizing nasopharyngeal carcinoma cell lines and biomarkers of response. <i>Investigational New Drugs</i> , 2010, 28, 326-333.	1.2	40
101	Aberrant methylation of the 8p22 tumor suppressor gene DLC1 in renal cell carcinoma. <i>Cancer Letters</i> , 2007, 249, 220-226.	3.2	38
102	BCLB, methylated in hepatocellular carcinoma, is a starvation stress sensor that induces apoptosis and autophagy through the AMPK-mTOR signaling cascade. <i>Cancer Letters</i> , 2017, 395, 63-71.	3.2	38
103	Sox2 promotes tumor aggressiveness and epithelial-mesenchymal transition in tongue squamous cell carcinoma. <i>International Journal of Molecular Medicine</i> , 2018, 42, 1418-1426.	1.8	38
104	Promoter methylation of tumor suppressor genes in esophageal squamous cell carcinoma. <i>Chinese Journal of Cancer</i> , 2013, 32, 3-11.	4.9	36
105	The new 6q27 tumor suppressor DACT2, frequently silenced by CpG methylation, sensitizes nasopharyngeal cancer cells to paclitaxel and 5-FU toxicity via $\beta$ 2-catenin/Cdc25c signaling and G2/M arrest. <i>Clinical Epigenetics</i> , 2018, 10, 26.	1.8	34
106	Promoter hypermethylation of the cyclin-dependent kinase inhibitor (CDKI) gene p21WAF1/CIP1/SDI1 is rare in various lymphomas and carcinomas. <i>Blood</i> , 2004, 103, 743-746.	0.6	33
107	Epigenomic characterization of a p53-regulated 3p22.2 tumor suppressor that inhibits STAT3 phosphorylation via protein docking and is frequently methylated in esophageal and other carcinomas. <i>Theranostics</i> , 2018, 8, 61-77.	4.6	33
108	Methylation profiling of Epstein-Barr virus immediate-early gene promoters, BZLF1 and BRLF1 in tumors of epithelial, NK- and B-cell origins. <i>BMC Cancer</i> , 2012, 12, 125.	1.1	32

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109	Interferon regulatory factor 8 functions as a tumor suppressor in renal cell carcinoma and its promoter methylation is associated with patient poor prognosis. <i>Cancer Letters</i> , 2014, 354, 227-234.	3.2	32
110	<i>NGALR</i> Is Overexpressed and Regulated by Hypomethylation in Esophageal Squamous Cell Carcinoma. <i>Clinical Cancer Research</i> , 2008, 14, 7674-7681.	3.2	31
111	19q13 KRAB zinc-finger protein ZNF471 activates MAPK10/JNK3 signaling but is frequently silenced by promoter CpG methylation in esophageal cancer. <i>Theranostics</i> , 2020, 10, 2243-2259.	4.6	31
112	Dickkopf-related protein 2 induces G0/G1 arrest and apoptosis through suppressing Wnt/ $\beta$ -catenin signaling and is frequently methylated in breast cancer. <i>Oncotarget</i> , 2017, 8, 39443-39459.	0.8	31
113	Epigenetic silencing of BCL6B inactivates p53 signaling and causes human hepatocellular carcinoma cell resist to 5-FU. <i>Oncotarget</i> , 2015, 6, 11547-11560.	0.8	31
114	Dapper Homolog 1 Is a Novel Tumor Suppressor in Gastric Cancer through Inhibiting the Nuclear Factor- $\kappa$ B Signaling Pathway. <i>Molecular Medicine</i> , 2012, 18, 1402-1411.	1.9	30
115	The epigenetic modifier PBRM1 restricts the basal activity of the innate immune system by repressing retinoic acid-inducible gene-like receptor signalling and is a potential prognostic biomarker for colon cancer. <i>Journal of Pathology</i> , 2018, 244, 36-48.	2.1	30
116	Epstein-Barr-virus-infected nasopharyngeal intraepithelial lymphocytes. <i>Lancet, The</i> , 1995, 345, 1309-1310.	6.3	29
117	BTB/POZ zinc finger protein ZBTB16 inhibits breast cancer proliferation and metastasis through upregulating ZBTB28 and antagonizing BCL6/ZBTB27. <i>Clinical Epigenetics</i> , 2020, 12, 82.	1.8	29
118	Genome-Wide Screening for Genetic Alterations in Esophageal Cancer by aCGH Identifies 11q13 Amplification Oncogenes Associated with Nodal Metastasis. <i>PLoS ONE</i> , 2012, 7, e39797.	1.1	29
119	LTBP-2 confers pleiotropic suppression and promotes dormancy in a growth factor permissive microenvironment in nasopharyngeal carcinoma. <i>Cancer Letters</i> , 2012, 325, 89-98.	3.2	28
120	A single nucleotide polymorphism in the Epstein-Barr virus genome is strongly associated with a high risk of nasopharyngeal carcinoma. <i>Chinese Journal of Cancer</i> , 2015, 34, 563-72.	4.9	28
121	Tumor-Specific Methylation of the 8p22 Tumor Suppressor Gene DLC1 is an Epigenetic Biomarker for Hodgkin, Nasal NK/T-Cell and Other Types of Lymphomas. <i>Epigenetics</i> , 2007, 2, 15-21.	1.3	27
122	Zinc-Finger Protein 545 Inhibits Cell Proliferation as a Tumor Suppressor through Inducing Apoptosis and is Disrupted by Promoter Methylation in Breast Cancer. <i>PLoS ONE</i> , 2014, 9, e110990.	1.1	27
123	Apolipoprotein M Gene (APOM) Polymorphism Modifies Metabolic and Disease Traits in Type 2 Diabetes. <i>PLoS ONE</i> , 2011, 6, e17324.	1.1	27
124	The tumor suppressor interferon regulatory factor 8 inhibits $\beta$ -catenin signaling in breast cancers, but is frequently silenced by promoter methylation. <i>Oncotarget</i> , 2017, 8, 48875-48888.	0.8	27
125	Epigenetic silencing of the 3p22 tumor suppressor DLEC1 by promoter CpG methylation in non-Hodgkin and Hodgkin lymphomas. <i>Journal of Translational Medicine</i> , 2012, 10, 209.	1.8	26
126	The novel 19q13 KRAB zinc-finger tumour suppressor ZNF382 is frequently methylated in oesophageal squamous cell carcinoma and antagonises Wnt/ $\beta$ -catenin signalling. <i>Cell Death and Disease</i> , 2018, 9, 573.	2.7	26



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127	The epigenetic modifier CHD5 functions as a novel tumor suppressor for renal cell carcinoma and is predominantly inactivated by promoter CpG methylation. <i>Oncotarget</i> , 2016, 7, 21618-21630.	0.8	26
128	Oncogenic induction of cellular high CpG methylation by Epstein-Barr virus in malignant epithelial cells. <i>Chinese Journal of Cancer</i> , 2014, 33, 604-8.	4.9	25
129	Celecoxib reduces microvessel density in patients treated with nasopharyngeal carcinoma and induces changes in gene expression. <i>Annals of Oncology</i> , 2006, 17, 1625-1630.	0.6	24
130	STK31 Maintains the Undifferentiated State of Colon Cancer Cells. <i>Carcinogenesis</i> , 2012, 33, 2044-2053.	1.3	24
131	Epigenomic analysis of lung adenocarcinoma reveals novel DNA methylation patterns associated with smoking. <i>OncoTargets and Therapy</i> , 2013, 6, 1471.	1.0	24
132	DACT2 silencing by promoter CpG methylation disrupts its regulation of epithelial-to-mesenchymal transition and cytoskeleton reorganization in breast cancer cells. <i>Oncotarget</i> , 2016, 7, 70924-70935.	0.8	24
133	Epigenetic silencing of WNT5A in Epstein-Barr virus-associated gastric carcinoma. <i>Archives of Virology</i> , 2013, 158, 123-132.	0.9	23
134	Tumor suppressive BTB/POZ zinc-finger protein ZBTB28 inhibits oncogenic BCL6/ZBTB27 signaling to maintain p53 transcription in multiple carcinogenesis. <i>Theranostics</i> , 2019, 9, 8182-8195.	4.6	23
135	Selective loss of Bâ€œcell phenotype in lymphocyte predominant Hodgkin lymphoma. <i>Journal of Pathology</i> , 2007, 213, 429-440.	2.1	22
136	Epigenetic Silencing of a Proapoptotic Cell Adhesion Molecule, the Immunoglobulin Superfamily Member IGSF4, by Promoter CpG Methylation Protects Hodgkin Lymphoma Cells from Apoptosis. <i>American Journal of Pathology</i> , 2010, 177, 1480-1490.	1.9	22
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