

# Qian Tao

## List of Publications by Year in descending order

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182  
papers

10,376  
citations

22099

59  
h-index

45213

90  
g-index

186  
all docs

186  
docs citations

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times ranked

12534  
citing authors

#	ARTICLE	IF	CITATIONS
1	Diverse Ras-related GTPase DIRAS2, downregulated by PSMD2 in a proteasome-mediated way, inhibits colorectal cancer proliferation by blocking NF- $\kappa$ B signaling. <i>International Journal of Biological Sciences</i> , 2022, 18, 1039-1050.	2.6	4
2	Zinc-finger protein 382 antagonises CDC25A and ZEB1 signaling pathway in breast cancer. <i>Genes and Diseases</i> , 2022, , .	1.5	0
3	Zinc finger protein 280C contributes to colorectal tumorigenesis by maintaining epigenetic repression at H3K27me3-marked loci. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	5
4	Low-dose selenite synergizes with KRAS inhibitor as a dual apoptotic and ferroptotic agent in lung adenocarcinoma.. <i>Journal of Clinical Oncology</i> , 2022, 40, e21039-e21039.	0.8	1
5	Nasopharyngeal carcinoma: an evolving paradigm. <i>Nature Reviews Clinical Oncology</i> , 2021, 18, 679-695.	12.5	207
6	Cancer cells escape p53 $\Delta$ ™s tumor suppression through ablation of ZDHHC1-mediated p53 palmitoylation. <i>Oncogene</i> , 2021, 40, 5416-5426.	2.6	16
7	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
8	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
9	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
10	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
11	Monoamine oxidase A is down-regulated in EBV-associated nasopharyngeal carcinoma. <i>Scientific Reports</i> , 2020, 10, 6115.	1.6	10
12	Targeting the polycomb repressive complex-2 related proteins with novel combinational strategies for nasopharyngeal carcinoma. <i>American Journal of Cancer Research</i> , 2020, 10, 3267-3284.	1.4	4
13	Classic SRY $\Delta$ box protein SOX7 functions as a tumor suppressor regulating WNT signaling and is methylated in renal cell carcinoma. <i>FASEB Journal</i> , 2019, 33, 254-263.	0.2	11
14	USP3 promotes breast cancer cell proliferation by deubiquitinating KLF5. <i>Journal of Biological Chemistry</i> , 2019, 294, 17837-17847.	1.6	49
15	Tumor Suppression of Ras GTPase-Activating Protein RASA5 through Antagonizing Ras Signaling Perturbation in Carcinomas. <i>IScience</i> , 2019, 21, 1-18.	1.9	12
16	Upregulation of interleukin-8 and activin A induces osteoclastogenesis in ameloblastoma. <i>International Journal of Molecular Medicine</i> , 2019, 43, 2329-2340.	1.8	11
17	Low Expression and Promoter Hypermethylation of the Tumour Suppressor SLIT2, are Associated with Adverse Patient Outcomes in Diffuse Large B Cell Lymphoma. <i>Pathology and Oncology Research</i> , 2019, 25, 1223-1231.	0.9	8
18	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

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19	The 19q13 KRAB Zinc-finger protein <i>ZFP82</i> suppresses the growth and invasion of esophageal carcinoma cells through inhibiting <i>NF-<math>\kappa</math>B</i> transcription and inducing apoptosis. <i>Epigenomics</i> , 2019, 11, 65-80.	1.0	12
20	The 3p14.2 tumour suppressor <i>ADAMTS9</i> is inactivated by promoter CpG methylation and inhibits tumour cell growth in breast cancer. <i>Journal of Cellular and Molecular Medicine</i> , 2018, 22, 1257-1271.	1.6	22
21	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
22	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
23	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
24	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
25	OVOL2 links stemness and metastasis via fine-tuning epithelial-mesenchymal transition in nasopharyngeal carcinoma. <i>Theranostics</i> , 2018, 8, 2202-2216.	4.6	50
26	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
27	Epstein-Barr Virus-Induced Epigenetic Pathogenesis of Viral-Associated Lymphoepithelioma-Like Carcinomas and Natural Killer/T-Cell Lymphomas. <i>Pathogens</i> , 2018, 7, 63.	1.2	16
28	The new 6q27 tumor suppressor DACT2, frequently silenced by CpG methylation, sensitizes nasopharyngeal cancer cells to paclitaxel and 5-FU toxicity via $\beta$ -catenin/Cdc25c signaling and G2/M arrest. <i>Clinical Epigenetics</i> , 2018, 10, 26.	1.8	34
29	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
30	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
31	Epigenomic and Functional Characterization of Junctophilin 3 (JPH3) as a Novel Tumor Suppressor Being Frequently Inactivated by Promoter CpG Methylation in Digestive Cancers. <i>Theranostics</i> , 2017, 7, 2150-2163.	4.6	15
32	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
33	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
34	Paired box 5 is a frequently methylated lung cancer tumour suppressor gene interfering $\beta$ -catenin signalling and <i>GADD45G</i> expression. <i>Journal of Cellular and Molecular Medicine</i> , 2016, 20, 842-854.	1.6	21
35	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
36	Protocadherin 17 functions as a tumor suppressor suppressing Wnt/ $\beta$ -catenin signaling and cell metastasis and is frequently methylated in breast cancer. <i>Oncotarget</i> , 2016, 7, 51720-51732.	0.8	46

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37	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
38	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
39	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
40	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
41	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
42	DLEC1, a 3p tumor suppressor, represses NF- $\hat{p}$ B signaling and is methylated in prostate cancer. <i>Journal of Molecular Medicine</i> , 2015, 93, 691-701.	1.7	21
43	Hypermethylation of ZNF545 is associated with poor prognosis in patients with early-stage hepatocellular carcinoma after thermal ablation: Table A1. <i>Gut</i> , 2015, 64, 1836-1837.	6.1	9
44	Abstract 2950: Epigenetic silencing BCL6B inactivates p53 signaling and causes human hepatocellular carcinoma cell resist to 5-FU. , 2015, , .		1
45	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
46	OPCML is frequently methylated in human colorectal cancer and its restored expression reverses EMT via downregulation of smad signaling. <i>American Journal of Cancer Research</i> , 2015, 5, 1635-48.	1.4	19
47	Preclinical evaluation of afatinib (BIBW2992) in esophageal squamous cell carcinoma (ESCC). <i>American Journal of Cancer Research</i> , 2015, 5, 3588-99.	1.4	11
48	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
49	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
50	Epigenetic identification of receptor tyrosine kinase-like orphan receptor 2 as a functional tumor suppressor inhibiting $\hat{I}^2$ -catenin and AKT signaling but frequently methylated in common carcinomas. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 2179-2192.	2.4	43
51	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
52	The Metalloprotease ADAMTS8 Displays Antitumor Properties through Antagonizing EGFR $\hat{a}$ MEK $\hat{a}$ ERK Signaling and Is Silenced in Carcinomas by CpG Methylation. <i>Molecular Cancer Research</i> , 2014, 12, 228-238.	1.5	58
53	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
54	DLC1-dependent parathyroid hormone $\hat{a}$ 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	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
56	Silencing of hypoxia-inducible tumor suppressor lysyl oxidase gene by promoter methylation activates carbonic anhydrase IX in nasopharyngeal carcinoma. <i>American Journal of Cancer Research</i> , 2014, 4, 789-800.	1.4	11
57	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
58	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
59	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
60	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
61	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
62	Epigenetic silencing of WNT5A in Epstein-Barr virus-associated gastric carcinoma. <i>Archives of Virology</i> , 2013, 158, 123-132.	0.9	23
63	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
64	Lipid-Polymer Nanoparticles Encapsulating Doxorubicin and 2-Deoxy-5-azacytidine Enhance the Sensitivity of Cancer Cells to Chemical Therapeutics. <i>Molecular Pharmaceutics</i> , 2013, 10, 1901-1909.	2.3	53
65	Characterization of naturally Epstein-Barr virus-infected gastric carcinoma cell line YCCEL1. <i>Journal of General Virology</i> , 2013, 94, 497-506.	1.3	47
66	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
67	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
68	Epigenomic analysis of lung adenocarcinoma reveals novel DNA methylation patterns associated with smoking. <i>OncoTargets and Therapy</i> , 2013, 6, 1471.	1.0	24
69	A phase IB trial of 5-azacitidine (5AC) and suberoylanilide hydroxamic acid (SAHA) in patients with metastatic or locally recurrent nasopharyngeal carcinoma (NPC) and NK-T cell lymphoma. <i>Journal of Clinical Oncology</i> , 2013, 31, e17017-e17017.	0.8	1
70	Aberrant promoter CpG methylation and its translational applications in breast cancer. <i>Chinese Journal of Cancer</i> , 2013, 32, 12-20.	4.9	44
71	Promoter methylation of tumor suppressor genes in esophageal squamous cell carcinoma. <i>Chinese Journal of Cancer</i> , 2013, 32, 3-11.	4.9	36
72	Cancer research in an era when epigenetics is no longer "epi" - challenges and opportunities. <i>Chinese Journal of Cancer</i> , 2013, 32, 1-2.	4.9	5

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73	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
74	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
75	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
76	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
77	STK31 Maintains the Undifferentiated State of Colon Cancer Cells. <i>Carcinogenesis</i> , 2012, 33, 2044-2053.	1.3	24
78	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
79	Physiological pathway of human cell damage induced by genotoxic crystalline silica nanoparticles. <i>Biomaterials</i> , 2012, 33, 7540-7546.	5.7	21
80	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
81	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
82	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
83	Chromatin regulators with tumor suppressor properties and their alterations in human cancers. <i>Epigenomics</i> , 2012, 4, 537-549.	1.0	19
84	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
85	Hypomethylation and Over-Expression of the Beta Isoform of BLIMP1 is Induced by Epstein-Barr Virus Infection of B Cells; Potential Implications for the Pathogenesis of EBV-Associated Lymphomas. <i>Pathogens</i> , 2012, 1, 83-101.	1.2	10
86	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
87	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
88	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
89	Cellular uptake, evolution, and excretion of silica nanoparticles in human cells. <i>Nanoscale</i> , 2011, 3, 3291.	2.8	121
90	Epigenetic disruption of cell signaling in nasopharyngeal carcinoma. <i>Chinese Journal of Cancer</i> , 2011, 30, 231-239.	4.9	59

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91	Nasopharyngeal carcinoma: molecular pathogenesis and therapeutic developments – CORRIGENDUM. Expert Reviews in Molecular Medicine, 2011, 13, .	1.6	0
92	A novel isoform of the 8p22 tumor suppressor gene DLC1 suppresses tumor growth and is frequently silenced in multiple common tumors. Oncogene, 2011, 30, 1923-1935.	2.6	48
93	Paired box gene 5 is a novel tumor suppressor in hepatocellular carcinoma through interaction with p53 signaling pathway. Hepatology, 2011, 53, 843-853.	3.6	63
94	A survey of methylated candidate tumor suppressor genes in nasopharyngeal carcinoma. International Journal of Cancer, 2011, 128, 1393-1403.	2.3	59
95	The ECM protein LTBP2 is a suppressor of esophageal squamous cell carcinoma tumor formation but higher tumor expression associates with poor patient outcome. International Journal of Cancer, 2011, 129, 565-573.	2.3	43
96	Apolipoprotein M Gene (APOM) Polymorphism Modifies Metabolic and Disease Traits in Type 2 Diabetes. PLoS ONE, 2011, 6, e17324.	1.1	27
97	The Epigenetic Modifier PRDM5 Functions as a Tumor Suppressor through Modulating WNT/β-Catenin Signaling and Is Frequently Silenced in Multiple Tumors. PLoS ONE, 2011, 6, e27346.	1.1	64
98	Epigenetic Silencing of the Receptor Tyrosine Phosphatase, PTPRK, Located At the Frequently Deleted 6q22.33-q23.2 Region, Leads to Tumor Growth Via the Constitutive Activation of STAT3 in Nasal-Type NK/T-Cell Lymphoma. Blood, 2011, 118, 1378-1378.	0.6	0
99	CD44 activation in mature B-cell malignancies by a novel recurrent IGH translocation. Blood, 2010, 115, 2458-2461.	0.6	13
100	The preclinical activity of the histone deacetylase inhibitor PXD101 (belinostat) in hepatocellular carcinoma cell lines. Investigational New Drugs, 2010, 28, 107-114.	1.2	56
101	Preclinical activity of gefitinib in non-keratinizing nasopharyngeal carcinoma cell lines and biomarkers of response. Investigational New Drugs, 2010, 28, 326-333.	1.2	40
102	Down-regulation of tyrosine aminotransferase at a frequently deleted region 16q22 contributes to the pathogenesis of hepatocellular carcinoma. Hepatology, 2010, 51, 1624-1634.	3.6	48
103	KRAB Zinc Finger Protein ZNF382 Is a Proapoptotic Tumor Suppressor That Represses Multiple Oncogenes and Is Commonly Silenced in Multiple Carcinomas. Cancer Research, 2010, 70, 6516-6526.	0.4	116
104	The Tumor Suppressor UCHL1 Forms a Complex with p53/MDM2/ARF to Promote p53 Signaling and Is Frequently Silenced in Nasopharyngeal Carcinoma. Clinical Cancer Research, 2010, 16, 2949-2958.	3.2	136
105	WNT5A antagonizes WNT/β-catenin signaling and is frequently silenced by promoter CpG methylation in esophageal squamous cell carcinoma. Cancer Biology and Therapy, 2010, 10, 617-624.	1.5	71
106	PLCD1 is a functional tumor suppressor inducing G <sub>2</sub> /M arrest and frequently methylated in breast cancer. Cancer Biology and Therapy, 2010, 10, 520-527.	1.5	52
107	Aberrant Promoter Methylation of DLEC1, a Critical 3p22 Tumor Suppressor for Renal Cell Carcinoma, is Associated With More Advanced Tumor Stage. Journal of Urology, 2010, 184, 731-737.	0.2	46
108	Epigenetic Silencing of a Proapoptotic Cell Adhesion Molecule, the Immunoglobulin Superfamily Member IGSF4, by Promoter CpG Methylation Protects Hodgkin Lymphoma Cells from Apoptosis. American Journal of Pathology, 2010, 177, 1480-1490.	1.9	22



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109	Epigenetic disruption of the WNT/ $\beta$ -catenin signaling pathway in human cancers. <i>Epigenetics</i> , 2009, 4, 307-312.	1.3	159
110	<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
111	Frequent concomitant epigenetic silencing of the stress-responsive tumor suppressor gene <i>CADM1</i> , and its interacting partner <i>DAL1</i> in nasal NK/T-cell lymphoma. <i>International Journal of Cancer</i> , 2009, 124, 1572-1578.	2.3	21
112	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
113	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
114	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
115	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
116	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
117	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
118	Epigenetic disruption of the WNT/ $\beta$ -catenin signaling pathway in human cancers. <i>Epigenetics</i> , 2009, 4, 307-12.	1.3	82
119	Integrative cancer epigenetics identifies novel tumor suppressor genes for common asian tumors. <i>Cell Biology International</i> , 2008, 32, S4-S4.	1.4	0
120	The KRAB domain-containing zinc finger protein ZNF382 is a potent tumor suppressor with frequent epigenetic inactivation in nasopharyngeal, esophageal and other carcinomas. <i>Cell Biology International</i> , 2008, 32, S5-S5.	1.4	0
121	Authentication of nasopharyngeal carcinoma tumor lines. <i>International Journal of Cancer</i> , 2008, 122, 2169-2171.	2.3	88
122	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
123	Epigenetic disruption of interferon- $\gamma$ 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
124	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
125	<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
126	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



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127	<i>WNT5A</i> Exhibits Tumor-Suppressive Activity through Antagonizing the Wnt/ $\beta^2$ -Catenin Signaling, and Is Frequently Methylated in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2008, 14, 55-61.	3.2	181
128	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
129	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
130	Tumor-Specific Methylation of the 8p22 Tumor Suppressor Gene <i>DLC1</i> 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
131	<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
132	Epigenetic silencing of a Ca <sup>2+</sup> -regulated Ras GTPase-activating protein <i>RASAL</i> 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
133	Nasopharyngeal carcinoma: molecular pathogenesis and therapeutic developments. <i>Expert Reviews in Molecular Medicine</i> , 2007, 9, 1-24.	1.6	266
134	<i>WNT5A</i> 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
135	Aberrant methylation of the 8p22 tumor suppressor gene <i>DLC1</i> in renal cell carcinoma. <i>Cancer Letters</i> , 2007, 249, 220-226.	3.2	38
136	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
137	Down-regulation of <i>ATM</i> protein in HRS cells of nodular sclerosis Hodgkin's lymphoma in children occurs in the absence of <i>ATM</i> gene inactivation. <i>Journal of Pathology</i> , 2007, 213, 329-336.	2.1	14
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