

# Manon van Engeland

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6201139/publications.pdf>

Version: 2024-02-01

128  
papers

11,802  
citations

41344

49  
h-index

27406

106  
g-index

129  
all docs

129  
docs citations

129  
times ranked

17113  
citing authors

#	ARTICLE	IF	CITATIONS
1	Differential Orthopedia Homeobox expression in pulmonary carcinoids is associated with changes in <scp>DNA</scp> methylation. <i>International Journal of Cancer</i> , 2022, 150, 1987-1997.	5.1	4
2	Molecular pathways in post-colonoscopy versus detected colorectal cancers: results from a nested caseâ€“control study. <i>British Journal of Cancer</i> , 2022, 126, 865-873.	6.4	6
3	Evaluation of a seven gene mutational profile as a prognostic factor in a population-based study of clear cell renal cell carcinoma. <i>Scientific Reports</i> , 2022, 12, 6478.	3.3	1
4	Technical considerations in PCR-based assay design for diagnostic DNA methylation cancer biomarkers. <i>Clinical Epigenetics</i> , 2022, 14, 56.	4.1	5
5	Genetic Profiling of Colorectal Carcinomas of Patients with Primary Sclerosing Cholangitis and Inflammatory Bowel Disease. <i>Inflammatory Bowel Diseases</i> , 2022, , .	1.9	2
6	Lessons from a systematic literature search on diagnostic DNA methylation biomarkers for colorectal cancer: how to increase research value and decrease research waste. <i>Clinical and Translational Gastroenterology</i> , 2022, Publish Ahead of Print, .	2.5	1
7	Diagnostic DNA Methylation Biomarkers for Renal Cell Carcinoma: A Systematic Review. <i>European Urology Oncology</i> , 2021, 4, 215-226.	5.4	12
8	Molecular profiles of response to neoadjuvant chemoradiotherapy in oesophageal cancers to develop personalized treatment strategies. <i>Molecular Oncology</i> , 2021, 15, 901-914.	4.6	7
9	Identification of DNA methylation markers for early detection of CRC indicates a role for nervous system-related genes in CRC. <i>Clinical Epigenetics</i> , 2021, 13, 80.	4.1	22
10	SOX17 expression and its downâ€“regulation by promoter methylation in cervical adenocarcinoma in situ and adenocarcinoma. <i>Histopathology</i> , 2020, 76, 383-393.	2.9	15
11	Past, Present and Future of Epigenetics in Adrenocortical Carcinoma. <i>Cancers</i> , 2020, 12, 1218.	3.7	21
12	Switches of SOX17 and SOX2 expression in the development of squamous metaplasia and squamous intraepithelial lesions of the uterine cervix. <i>Cancer Medicine</i> , 2020, 9, 6330-6343.	2.8	8
13	Germline polymorphisms in the Von Hippel-Lindau and Hypoxia-inducible factor 1-alpha genes, gene-environment and gene-gene interactions and renal cell cancer. <i>Scientific Reports</i> , 2020, 10, 137.	3.3	5
14	Prognostic DNA methylation markers for hormone receptor breast cancer: a systematic review. <i>Breast Cancer Research</i> , 2020, 22, 13.	5.0	29
15	The trans-DATA study: aims and design of a translational breast cancer prognostic marker identification study. <i>Diagnostic and Prognostic Research</i> , 2019, 3, 20.	1.8	1
16	Gene Promoter Methylation in Endometrial Carcinogenesis. <i>Pathology and Oncology Research</i> , 2019, 25, 659-667.	1.9	8
17	Analysis of DNA methylation in cancer: location revisited. <i>Nature Reviews Clinical Oncology</i> , 2018, 15, 459-466.	27.6	486
18	Prognostic DNA methylation markers for sporadic colorectal cancer: a systematic review. <i>Clinical Epigenetics</i> , 2018, 10, 35.	4.1	38

#	ARTICLE	IF	CITATIONS
19	Cost-effectiveness of High-performance Biomarker Tests vs Fecal Immunochemical Test for Noninvasive Colorectal Cancer Screening. <i>Clinical Gastroenterology and Hepatology</i> , 2018, 16, 504-512.e11.	4.4	36
20	A combined literature and in silico analysis enlightens the role of the NDRG family in the gut. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2018, 1862, 2140-2151.	2.4	11
21	Epigenetics in renal cell cancer: mechanisms and clinical applications. <i>Nature Reviews Urology</i> , 2018, 15, 430-451.	3.8	115
22	Promoter CpG island methylation in ion transport mechanisms and associated dietary intakes jointly influence the risk of clear-cell renal cell cancer. <i>International Journal of Epidemiology</i> , 2017, 46, dyw266.	1.9	18
23	Details matter: the role of genomic location and assay standardization in DNA methylation analyses. <i>Epigenomics</i> , 2017, 9, 933-935.	2.1	5
24	Assessing opportunities for coordinated R&D in early cancer detection and management in Europe. <i>International Journal of Cancer</i> , 2017, 140, 1700-1701.	5.1	1
25	Prognostic DNA methylation markers for renal cell carcinoma: a systematic review. <i>Epigenomics</i> , 2017, 9, 1243-1257.	2.1	44
26	Molecular stool testing as an alternative for surveillance colonoscopy: a cross-sectional cohort study. <i>BMC Cancer</i> , 2017, 17, 116.	2.6	37
27	A Four-Gene Promoter Methylation Marker Panel Consisting of <i>GREM1</i> , <i>NEURL</i> , <i>LAD1</i> , and <i>NEFH</i> Predicts Survival of Clear Cell Renal Cell Cancer Patients. <i>Clinical Cancer Research</i> , 2017, 23, 2006-2018.	7.0	51
28	Energy restriction at young age, genetic variants in the insulin-like growth factor pathway and colorectal cancer risk in the Netherlands Cohort Study. <i>International Journal of Cancer</i> , 2017, 140, 272-284.	5.1	5
29	Lifestyle, Diet, and Colorectal Cancer Risk According to (Epi)genetic Instability: Current Evidence and Future Directions of Molecular Pathological Epidemiology. <i>Current Colorectal Cancer Reports</i> , 2017, 13, 455-469.	0.5	91
30	Decoy receptor 1 (DCR1) promoter hypermethylation and response to irinotecan in metastatic colorectal cancer. <i>Oncotarget</i> , 2017, 8, 63140-63154.	1.8	19
31	The emerging role of GATA transcription factors in development and disease. <i>Expert Reviews in Molecular Medicine</i> , 2016, 18, e3.	3.9	172
32	Alcohol and Dietary Folate Intake and Promoter CpG Island Methylation in Clear-Cell Renal Cell Cancer. <i>Nutrition and Cancer</i> , 2016, 68, 1097-1107.	2.0	9
33	Potential role of gene-environment interactions in ion transport mechanisms in the etiology of renal cell cancer. <i>Scientific Reports</i> , 2016, 6, 34262.	3.3	7
34	Analysis of RET promoter CpG island methylation using methylation-specific PCR (MSP), pyrosequencing, and methylation-sensitive high-resolution melting (MS-HRM): impact on stage II colon cancer patient outcome. <i>Clinical Epigenetics</i> , 2016, 8, 44.	4.1	18
35	Genetic Variants in the Insulin-like Growth Factor Pathway and Colorectal Cancer Risk in the Netherlands Cohort Study. <i>Scientific Reports</i> , 2015, 5, 14126.	3.3	16
36	<i>Spectrin Repeat Containing Nuclear Envelope 1</i> and <i>Forkhead Box Protein E1</i> Are Promising Markers for the Detection of Colorectal Cancer in Blood. <i>Cancer Prevention Research</i> , 2015, 8, 157-164.	1.5	29

#	ARTICLE	IF	CITATIONS
37	Polymorphisms in genes of the renin-angiotensin-aldosterone system and renal cell cancer risk: Interplay with hypertension and intakes of sodium, potassium and fluid. <i>International Journal of Cancer</i> , 2015, 136, 1104-1116.	5.1	44
38	Promoter Methylation of <i>CDO1</i> Identifies Clear-Cell Renal Cell Cancer Patients with Poor Survival Outcome. <i>Clinical Cancer Research</i> , 2015, 21, 3492-3500.	7.0	50
39	Formalin-fixed, paraffin-embedded (FFPE) tissue epigenomics using Infinium HumanMethylation450 BeadChip assays. <i>Laboratory Investigation</i> , 2015, 95, 833-842.	3.7	40
40	Mitochondrial DNA copy number in colorectal cancer: between tissue comparisons, clinicopathological characteristics and survival. <i>Carcinogenesis</i> , 2015, 36, bgv151.	2.8	36
41	Prognostic Significance of Promoter Hypermethylation and Diminished Gene Expression of <i>SYNPO2</i> in Melanoma. <i>Journal of Investigative Dermatology</i> , 2015, 135, 2328-2331.	0.7	13
42	Methylation of <i>RASSF10</i> promotes cell proliferation and serves as a docetaxel resistant marker in human breast cancer. <i>Discovery Medicine</i> , 2015, 20, 261-71.	0.5	3
43	<i>CHFR</i> Promoter Methylation Indicates Poor Prognosis in Stage II Microsatellite Stable Colorectal Cancer. <i>Clinical Cancer Research</i> , 2014, 20, 3261-3271.	7.0	29
44	Promoter CpG island methylation of <i>RET</i> predicts poor prognosis in stage II colorectal cancer patients. <i>Molecular Oncology</i> , 2014, 8, 679-688.	4.6	33
45	<i>MEN1</i> Gene Mutation and Reduced Expression Are Associated With Poor Prognosis in Pulmonary Carcinoids. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2014, 99, E374-E378.	3.6	62
46	Dietary acrylamide intake and the risk of colorectal cancer with specific mutations in <i>KRAS</i> and <i>APC</i> . <i>Carcinogenesis</i> , 2014, 35, 1032-1038.	2.8	31
47	Promoter CpG Island Hypermethylation in Dysplastic Nevus and Melanoma: <i>CLDN11</i> as an Epigenetic Biomarker for Malignancy. <i>Journal of Investigative Dermatology</i> , 2014, 134, 2957-2966.	0.7	38
48	Body Size, Physical Activity, Early-Life Energy Restriction, and Associations with Methylated Insulin-like Growth Factor-1 Binding Protein Genes in Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1852-1862.	2.5	22
49	Chordoma: the entity. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2014, 1846, 655-669.	7.4	47
50	Epigenetics in radiotherapy: Where are we heading?. <i>Radiotherapy and Oncology</i> , 2014, 111, 168-177.	0.6	43
51	The mTOR Pathway and the Role of Energy Balance Throughout Life in Colorectal Cancer Etiology and Prognosis: Unravelling Mechanisms Through a Multidimensional Molecular Epidemiologic Approach. <i>Current Nutrition Reports</i> , 2013, 2, 19-26.	4.3	19
52	The CpG Island Methylator Phenotype: What's in a Name?. <i>Cancer Research</i> , 2013, 73, 5858-5868.	0.9	154
53	Promoter methylation of <i>Wnt</i> -antagonists in polypoid and nonpolypoid colorectal adenomas. <i>BMC Cancer</i> , 2013, 13, 603.	2.6	23
54	Emerging evidence for <i>CHFR</i> as a cancer biomarker: from tumor biology to precision medicine. <i>Cancer and Metastasis Reviews</i> , 2013, 33, 161-71.	5.9	30

#	ARTICLE	IF	CITATIONS
55	Frequent Inactivation of Cysteine Dioxygenase Type 1 Contributes to Survival of Breast Cancer Cells and Resistance to Anthracyclines. <i>Clinical Cancer Research</i> , 2013, 19, 3201-3211.	7.0	77
56	An exploration of pathways involved in lung carcinoid progression using gene expression profiling. <i>Carcinogenesis</i> , 2013, 34, 2726-2737.	2.8	49
57	Tracking the Molecular Features of Nonpolypoid Colorectal Neoplasms: A Systematic Review and Meta-Analysis. <i>American Journal of Gastroenterology</i> , 2013, 108, 1042-1056.	0.4	45
58	Dietary heme iron and the risk of colorectal cancer with specific mutations in KRAS and APC. <i>Carcinogenesis</i> , 2013, 34, 2757-2766.	2.8	57
59	Molecular markers and the future of colorectal cancer screening. <i>Colorectal Cancer</i> , 2013, 2, 95-97.	0.8	0
60	Physical Activity, Occupational Sitting Time, and Colorectal Cancer Risk in the Netherlands Cohort Study. <i>American Journal of Epidemiology</i> , 2013, 177, 514-530.	3.4	60
61	CD44 and OTP Are Strong Prognostic Markers for Pulmonary Carcinoids. <i>Clinical Cancer Research</i> , 2013, 19, 2197-2207.	7.0	77
62	Chromosome 5q Loss in Colorectal Flat Adenomas. <i>Clinical Cancer Research</i> , 2012, 18, 4560-4569.	7.0	30
63	Body size and risk for colorectal cancers showing BRAF mutations or microsatellite instability: a pooled analysis. <i>International Journal of Epidemiology</i> , 2012, 41, 1060-1072.	1.9	65
64	DNA Methylation of Phosphatase and Actin Regulator 3 Detects Colorectal Cancer in Stool and Complements FIT. <i>Cancer Prevention Research</i> , 2012, 5, 464-472.	1.5	46
65	Promoter CpG island methylation markers in colorectal cancer: the road ahead. <i>Epigenomics</i> , 2012, 4, 179-194.	2.1	38
66	Analytical sensitivity and stability of DNA methylation testing in stool samples for colorectal cancer detection. <i>Cellular Oncology (Dordrecht)</i> , 2012, 35, 309-315.	4.4	25
67	Comprehensive Mutation Analysis in Colorectal Flat Adenomas. <i>PLoS ONE</i> , 2012, 7, e41963.	2.5	20
68	Epigenetic Changes in Basal Cell Carcinoma Affect SHH and WNT Signaling Components. <i>PLoS ONE</i> , 2012, 7, e51710.	2.5	38
69	The adenoma hunt in colorectal cancer screening: defining the target. <i>Journal of Pathology</i> , 2012, 226, 1-6.	4.5	30
70	The CpG island methylator phenotype in colorectal cancer: Progress and problems. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012, 1825, 77-85.	7.4	89
71	Taxane resistance in breast cancer: A closed HER2 circuit?. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012, 1825, 197-206.	7.4	22
72	Genetics and epigenetics of cutaneous malignant melanoma: A concert out of tune. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012, 1826, 89-102.	7.4	46

#	ARTICLE	IF	CITATIONS
73	Colorectal Cancer Epigenetics: Complex Simplicity. <i>Journal of Clinical Oncology</i> , 2011, 29, 1382-1391.	1.6	180
74	A <i>Let-7</i> MicroRNA SNP in the <i>KRAS</i> 3'UTR Is Prognostic in Early-Stage Colorectal Cancer. <i>Clinical Cancer Research</i> , 2011, 17, 7723-7731.	7.0	106
75	Body Size, Physical Activity and Risk of Colorectal Cancer with or without the CpG Island Methylator Phenotype (CIMP). <i>PLoS ONE</i> , 2011, 6, e18571.	2.5	64
76	Molecular Tests for Colorectal Cancer Screening. <i>Clinical Colorectal Cancer</i> , 2011, 10, 8-23.	2.3	90
77	Body Size and Colorectal Cancer Risk After 16.3 Years of Follow-up: An Analysis From the Netherlands Cohort Study. <i>American Journal of Epidemiology</i> , 2011, 174, 1127-1139.	3.4	43
78	Analysis of Promoter CpG Island Hypermethylation in Cancer: Location, Location, Location!. <i>Clinical Cancer Research</i> , 2011, 17, 4225-4231.	7.0	121
79	Dietary methyl donors, methyl metabolizing enzymes, and epigenetic regulators: diet-gene interactions and promoter CpG island hypermethylation in colorectal cancer. <i>Cancer Causes and Control</i> , 2011, 22, 1-12.	1.8	37
80	Characteristics of triple-negative breast cancer. <i>Journal of Cancer Research and Clinical Oncology</i> , 2011, 137, 183-192.	2.5	225
81	Alcohol consumption, alcohol dehydrogenase 1C (ADH1C) genotype, and risk of colorectal cancer in the Netherlands Cohort Study on diet and cancer. <i>Alcohol</i> , 2011, 45, 217-225.	1.7	14
82	Genomic and Epigenomic Integration Identifies a Prognostic Signature in Colon Cancer. <i>Clinical Cancer Research</i> , 2011, 17, 1535-1545.	7.0	136
83	Methylation-associated dysregulation of the suppressor of cytokine signaling-3 gene in multiple myeloma. <i>Epigenetics</i> , 2011, 6, 1047-1052.	2.7	23
84	Loss of SerpinA5 protein expression is associated with advanced-stage serous ovarian tumors. <i>Modern Pathology</i> , 2011, 24, 463-470.	5.5	29
85	Test Performance of Immunologic Fecal Occult Blood Testing and Sigmoidoscopy Compared with Primary Colonoscopy Screening for Colorectal Advanced Adenomas. <i>Cancer Prevention Research</i> , 2011, 4, 1563-1571.	1.5	40
86	Early onset MSI-H colon cancer with MLH1 promoter methylation, is there a genetic predisposition?. <i>BMC Cancer</i> , 2010, 10, 180.	2.6	45
87	VHL and HIF signalling in renal cell carcinogenesis. <i>Journal of Pathology</i> , 2010, 221, 125-138.	4.5	258
88	The <i>N-cyc</i> downstream regulated gene (NDRG) family: diverse functions, multiple applications. <i>FASEB Journal</i> , 2010, 24, 4153-4166.	0.5	249
89	Viewing the Epigenetics of Colorectal Cancer through the Window of Folic Acid Effects. <i>Cancer Prevention Research</i> , 2010, 3, 1509-1512.	1.5	10
90	Childhood and adolescent energy restriction and subsequent colorectal cancer risk: results from the Netherlands Cohort Study. <i>International Journal of Epidemiology</i> , 2010, 39, 1333-1344.	1.9	51

#	ARTICLE	IF	CITATIONS
91	Prognostic Significance of Gremlin1 (GREM1) Promoter CpG Island Hypermethylation in Clear Cell Renal Cell Carcinoma. <i>American Journal of Pathology</i> , 2010, 176, 575-584.	3.8	66
92	Early Life Exposure to Famine and Colorectal Cancer Risk: A Role for Epigenetic Mechanisms. <i>PLoS ONE</i> , 2009, 4, e7951.	2.5	104
93	GATA4 and GATA5 are Potential Tumor Suppressors and Biomarkers in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2009, 15, 3990-3997.	7.0	166
94	Prostate Cancer Detected by Methylated Gene Markers in Histopathologically Cancer-Negative Tissues from Men with Subsequent Positive Biopsies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2717-2722.	2.5	42
95	N-Myc Downstream-Regulated Gene 4 ( NDRG4 ): A Candidate Tumor Suppressor Gene and Potential Biomarker for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2009, 101, 916-927.	6.3	180
96	Genetic Variants of Methyl Metabolizing Enzymes and Epigenetic Regulators: Associations with Promoter CpG Island Hypermethylation in Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 3086-3096.	2.5	78
97	Promoter CpG island hypermethylation- and H3K9me3 and H3K27me3-mediated epigenetic silencing targets the deleted in colon cancer (DCC) gene in colorectal carcinogenesis without affecting neighboring genes on chromosomal region 18q21. <i>Carcinogenesis</i> , 2009, 30, 1041-1048.	2.8	46
98	Methylation of <i>TFPI2</i> in Stool DNA: A Potential Novel Biomarker for the Detection of Colorectal Cancer. <i>Cancer Research</i> , 2009, 69, 4691-4699.	0.9	204
99	Different angiogenic potential in low and high grade sporadic clear cell renal cell carcinoma is not related to alterations in the von Hippel-Lindau gene. <i>Cellular Oncology</i> , 2009, 31, 371-82.	1.9	12
100	Genetics and epigenetics of renal cell cancer. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2008, 1785, 133-155.	7.4	110
101	An Inactivating Mutation in HDAC2 Leads to Dysregulation of Apoptosis Mediated by APAF1. <i>Gastroenterology</i> , 2008, 135, 1654-1664.e2.	1.3	50
102	Associations of dietary methyl donor intake with MLH1 promoter hypermethylation and related molecular phenotypes in sporadic colorectal cancer. <i>Carcinogenesis</i> , 2008, 29, 1765-1773.	2.8	89
103	Integrated analysis of chromosomal, microsatellite and epigenetic instability in colorectal cancer identifies specific associations between promoter methylation of pivotal tumour suppressor and DNA repair genes and specific chromosomal alterations. <i>Carcinogenesis</i> , 2008, 29, 434-439.	2.8	59
104	Genetic and Epigenetic Alterations in the von Hippel-Lindau Gene: the Influence on Renal Cancer Prognosis. <i>Clinical Cancer Research</i> , 2008, 14, 782-787.	7.0	65
105	Alcohol Consumption and Mutations or Promoter Hypermethylation of the von Hippel-Lindau Gene in Renal Cell Carcinoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 3543-3550.	2.5	9
106	Dietary Folate, Methionine, Riboflavin, and Vitamin B-6 and Risk of Sporadic Colorectal Cancer. <i>Journal of Nutrition</i> , 2008, 138, 2372-2378.	2.9	80
107	Pharmacoepigenomics in colorectal cancer: a step forward in predicting prognosis and treatment response. <i>Pharmacogenomics</i> , 2008, 9, 1903-1916.	1.3	23
108	Lamin A/C Is a Risk Biomarker in Colorectal Cancer. <i>PLoS ONE</i> , 2008, 3, e2988.	2.5	186

#	ARTICLE	IF	CITATIONS
109	Identification of Epigenetically Silenced Genes in Tumor Endothelial Cells. <i>Cancer Research</i> , 2007, 67, 4138-4148.	0.9	126
110	Comparing the DNA Hypermethylome with Gene Mutations in Human Colorectal Cancer. <i>PLoS Genetics</i> , 2007, 3, e157.	3.5	307
111	Dietary Folate Intake in Combination with MTHFR C677T Genotype and Promoter Methylation of Tumor Suppressor and DNA Repair Genes in Sporadic Colorectal Adenomas. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 327-333.	2.5	43
112	Dual targeting of epigenetic therapy in cancer. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2007, 1775, 76-91.	7.4	85
113	Dietary Folate and APC Mutations in Sporadic Colorectal Cancer. <i>Journal of Nutrition</i> , 2006, 136, 3015-3021.	2.9	22
114	TP53 overexpression in recurrent endometrial carcinoma. <i>Gynecologic Oncology</i> , 2006, 100, 397-404.	1.4	28
115	Angiostatic activity of DNA methyltransferase inhibitors. <i>Molecular Cancer Therapeutics</i> , 2006, 5, 467-475.	4.1	93
116	Epigenetic Regulation of Tumor Endothelial Cell Anergy: Silencing of Intercellular Adhesion Molecule-1 by Histone Modifications. <i>Cancer Research</i> , 2006, 66, 10770-10777.	0.9	139
117	Dietary folate intake and k-ras mutations in sporadic colon and rectal cancer in the Netherlands Cohort Study. <i>International Journal of Cancer</i> , 2005, 114, 824-830.	5.1	23
118	Differential Gene Expression in Ovarian Tumors Reveals Dusp 4 and Serpina 5 As Key Regulators for Benign Behavior of Serous Borderline Tumors. <i>Journal of Clinical Oncology</i> , 2005, 23, 7257-7264.	1.6	82
119	CHFR promoter hypermethylation in colon cancer correlates with the microsatellite instability phenotype. <i>Carcinogenesis</i> , 2005, 26, 1152-1156.	2.8	81
120	APC mutations in sporadic colorectal carcinomas from The Netherlands Cohort Study. <i>Carcinogenesis</i> , 2004, 25, 1219-1226.	2.8	73
121	Epigenetic inactivation of SFRP genes allows constitutive WNT signaling in colorectal cancer. <i>Nature Genetics</i> , 2004, 36, 417-422.	21.4	976
122	GATA-4 and GATA-5 Transcription Factor Genes and Potential Downstream Antitumor Target Genes Are Epigenetically Silenced in Colorectal and Gastric Cancer. <i>Molecular and Cellular Biology</i> , 2003, 23, 8429-8439.	2.3	234
123	K-ras mutations and RASSF1A promoter methylation in colorectal cancer. <i>Oncogene</i> , 2002, 21, 3792-3795.	5.9	168
124	A genomic screen for genes upregulated by demethylation and histone deacetylase inhibition in human colorectal cancer. <i>Nature Genetics</i> , 2002, 31, 141-149.	21.4	820
125	Annexin V-Affinity assay: A review on an apoptosis detection system based on phosphatidylserine exposure. <i>Cytometry</i> , 1998, 31, 1-9.	1.8	1,567
126	The Effect of the Cyclin-Dependent Kinase Inhibitor Olomoucine on Cell Cycle Kinetics. <i>Experimental Cell Research</i> , 1997, 236, 4-15.	2.6	103

#	ARTICLE	IF	CITATIONS
127	Plasma Membrane Alterations and Cytoskeletal Changes in Apoptosis. <i>Experimental Cell Research</i> , 1997, 235, 421-430.	2.6	176
128	A novel assay to measure loss of plasma membrane asymmetry during apoptosis of adherent cells in culture. <i>Cytometry</i> , 1996, 24, 131-139.	1.8	451