Renske D M Steenbergen

List of Publications by Year
in descending order

146 papers	6,345 citations	57758 44 h-index	85541 71 g-index
148 all docs	148 docs citations	148 times ranked	6505 citing authors

#	Article	IF	Citations
1	FAM19A4/miR124-2 Methylation Testing and Human Papillomavirus (HPV) 16/18 Genotyping in HPV-Positive Women Under the Age of 30 Years. Clinical Infectious Diseases, 2023, 76, e827-e834.	5.8	4
2	Performance of <scp>DNA</scp> methylation analysis of <scp>ASCL1</scp>, <scp>LHX8</scp>, <scp>ST6GALNAC5</scp>, <scp>GHSR</scp>, <scp>ZIC1</scp> and <scp>SST</scp> for the triage of <scp>HPV</scp>â€positive women: Results from a Dutch primary <scp>HPV</scp>â€based screening cohort. International Journal of Cancer, 2022, 150, 440-449.	5.1	17
3	Dynamics of methylated cell-free DNA in the urine of non-small cell lung cancer patients. Epigenetics, 2022, 17, 1057-1069.	2.7	5
4	Direct bisulphite conversion of cervical samples for DNA methylation analysis. Epigenetics, 2022, 17, 1173-1179.	2.7	6
5	The association between viral load and concurrent human papillomavirus infection at the genital and anal sites of young women and the impact of vaccination. Tumour Virus Research, 2022, 13, 200233.	3.8	2
6	Bladder cancer detection in urine using DNA methylation markers: a technical and prospective preclinical validation. Clinical Epigenetics, 2022, 14, 19.	4.1	16
7	Post-treatment monitoring by ASCL1/LHX8 methylation analysis in women with HIV treated for cervical intraepithelial neoplasia grade 2/3. Aids, 2022, Publish Ahead of Print, .	2.2	1
8	HPV and DNA Methylation Testing in Urine for Cervical Intraepithelial Neoplasia and Cervical Cancer Detection. Clinical Cancer Research, 2022, 28, 2061-2068.	7.0	24
9	Functional Screen for microRNAs Suppressing Anchorage-Independent Growth in Human Cervical Cancer Cells. International Journal of Molecular Sciences, 2022, 23, 4791.	4.1	3
10	Clinical Regression of High-Grade Cervical Intraepithelial Neoplasia Is Associated With Absence of FAM19A4/miR124-2 DNA Methylation (CONCERVE Study). Journal of Clinical Oncology, 2022, 40, 3037-3046.	1.6	25
11	Lyon IARC Polyomavirus Displays Transforming Activities in Primary Human Cells. Journal of Virology, 2022, 96, .	3.4	2
12	Cancer Risk Stratification of Anal Intraepithelial Neoplasia in Human Immunodeficiency Virusâ€"Positive Men by Validated Methylation Markers Associated With Progression to Cancer. Clinical Infectious Diseases, 2021, 72, 2154-2163.	5.8	36
13	A systematic review on mutation markers for bladder cancer diagnosis in urine. BJU International, 2021, 127, 12-27.	2.5	14
14	Vulvar intraepithelial neoplasia: Incidence and longâ€term risk of vulvar squamous cell carcinoma. International Journal of Cancer, 2021, 148, 90-98.	5.1	49
15	Delta-Like Ligandâ€"Notch1 Signaling Is Selectively Modulated by HPV16 E6 to Promote Squamous Cell Proliferation and Correlates with Cervical Cancer Prognosis. Cancer Research, 2021, 81, 1909-1921.	0.9	16
16	The Origin of Tumor DNA in Urine of Urogenital Cancer Patients: Local Shedding and Transrenal Excretion. Cancers, 2021, 13, 535.	3.7	9
17	<scp>DNA</scp> methylation markers for cancer risk prediction of vulvar intraepithelial neoplasia. International Journal of Cancer, 2021, 148, 2481-2488.	5.1	17
18	Reply to Fang and Buchwald. Journal of Infectious Diseases, 2021, 224, 1271-1272.	4.0	1

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19	DNA methylation markers have universal prognostic value for anal cancer risk in HIVâ€negative and HIVâ€positive individuals. Molecular Oncology, 2021, 15, 3024-3036.	4.6	13
20	Impact of Collection Volume and DNA Extraction Method on the Detection of Biomarkers and HPV DNA in First-Void Urine. Molecules, 2021, 26, 1989.	3.8	10
21	Triage of human papillomavirus infected women by methylation analysis in first-void urine. Scientific Reports, 2021, 11, 7862.	3.3	15
22	Classification of highâ€grade cervical intraepithelial neoplasia by p16^{ink4a}, Kiâ€67, <scp>HPV E4</scp> and <scp>FAM19A4</scp>/<scp>miR124</scp>â€2 methylation status demonstrates considerable heterogeneity with potential consequences for management. International Journal of Cancer, 2021, 149, 707-716.	5.1	26
23	Characterisation of anal intraepithelial neoplasia and anal cancer in <scp>HIV</scp>â€positive men by immunohistochemical markers p16, Kiâ€67, <scp>HPVâ€E4</scp> and <scp>DNA</scp> methylation markers. International Journal of Cancer, 2021, 149, 1833-1844.	5.1	6
24	Oncogenic HPV promotes the expression of the long noncoding RNA lnc-FANCI-2 through E7 and YY1. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	31
25	Biomarker Expression in Multifocal Vulvar High-Grade Squamous Intraepithelial Lesions. Cancers, 2021, 13, 5646.	3.7	1
26	FAM19A4/miR124â€2 methylation in invasive cervical cancer: A retrospective crossâ€sectional worldwide study. International Journal of Cancer, 2020, 147, 1215-1221.	5.1	40
27	miR-9-5p Exerts a Dual Role in Cervical Cancer and Targets Transcription Factor TWIST1. Cells, 2020, 9, 65.	4.1	25
28	DPHL: A DIA Pan-human Protein Mass Spectrometry Library for Robust Biomarker Discovery. Genomics, Proteomics and Bioinformatics, 2020, 18, 104-119.	6.9	51
29	Evaluation of six methylation markers derived from genome-wide screens for detection of cervical precancer and cancer. Epigenomics, 2020, 12, 1569-1578.	2.1	15
30	DNA methylation markers for endometrial cancer detection in minimally invasive samples: a systematic review. Epigenomics, 2020, 12, 1661-1672.	2.1	7
31	Non-invasive detection of endometrial cancer by DNA methylation analysis in urine. Clinical Epigenetics, 2020, 12, 165.	4.1	22
32	Methylation analysis in urine fractions for optimal CIN3 and cervical cancer detection. Papillomavirus Research (Amsterdam, Netherlands), 2020, 9, 100193.	4.5	15
33	Identification of Deregulated Pathways, Key Regulators, and Novel miRNA-mRNA Interactions in HPV-Mediated Transformation. Cancers, 2020, 12, 700.	3.7	20
34	Comparative Analysis of Urine Fractions for Optimal Bladder Cancer Detection Using DNA Methylation Markers. Cancers, 2020, 12, 859.	3.7	31
35	Host Cell Deoxyribonucleic Acid Methylation Markers for the Detection of High-grade Anal Intraepithelial Neoplasia and Anal Cancer. Clinical Infectious Diseases, 2019, 68, 1110-1117.	5.8	25
36	Altered microRNA processing proteins in HPV-induced cancers. Current Opinion in Virology, 2019, 39, 23-32.	5.4	15

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37	A two-gene methylation signature for the diagnosis of bladder cancer in urine. Epigenomics, 2019, 11, 337-347.	2.1	23
38	Circulating Tumor DNA Analysis: Clinical Implications for Colorectal Cancer Patients. A Systematic Review. JNCI Cancer Spectrum, 2019, 3, pkz042.	2.9	22
39	Long-term CIN3+ risk of HPV positive women after triage with FAM19A4/miR124-2 methylation analysis. Gynecologic Oncology, 2019, 154, 368-373.	1.4	32
40	HPV16 variant analysis in primary and recurrent CIN2/3 lesions demonstrates presence of the same consensus variant. Papillomavirus Research (Amsterdam, Netherlands), 2019, 7, 168-172.	4.5	6
41	Complementarity between miRNA expression analysis and DNA methylation analysis in hrHPV-positive cervical scrapes for the detection of cervical disease. Epigenetics, 2019, 14, 558-567.	2.7	7
42	Cervical cancer detection by DNA methylation analysis in urine. Scientific Reports, 2019, 9, 3088.	3.3	35
43	The use of molecular markers for cervical screening of women living with HIV in South Africa. Aids, 2019, 33, 2035-2042.	2.2	20
44	Genomeâ€wide microRNA analysis of HPVâ€positive selfâ€samples yields novel triage markers for early detection of cervical cancer. International Journal of Cancer, 2019, 144, 372-379.	5.1	29
45	The diagnostic accuracy of methylation markers in urine for the detection of bladder cancer: a systematic review. Epigenomics, 2018, 10, 673-687.	2.1	24
46	Cervical cancer risk in HPVâ€positive women after a negative FAM19A4/mir124â€2 methylation test: A post hoc analysis in the POBASCAM trial with 14 year followâ€up. International Journal of Cancer, 2018, 143, 1541-1548.	5.1	63
47	Identification and Validation of a 3-Gene Methylation Classifier for HPV-Based Cervical Screening on Self-Samples. Clinical Cancer Research, 2018, 24, 3456-3464.	7.0	55
48	The effect of ART on cervical cancer precursor lesions. Lancet HIV,the, 2018, 5, e6-e8.	4.7	6
49	HPV16-Related Cervical Cancers and Precancers Have Increased Levels of Host Cell DNA Methylation in Women Living with HIV. International Journal of Molecular Sciences, 2018, 19, 3297.	4.1	7
50	Detection of hypermethylated genes as markers for cervical screening in women living with HIV. Journal of the International AIDS Society, 2018, 21, e25165.	3.0	18
51	Host-cell DNA methylation patterns during high-risk HPV-induced carcinogenesis reveal a heterogeneous nature of cervical pre-cancer. Epigenetics, 2018, 13, 769-778.	2.7	43
52	Three-tiered score for Ki-67 and p16^{ink4a} improves accuracy and reproducibility of grading CIN lesions. Journal of Clinical Pathology, 2018, 71, 981-988.	2.0	33
53	Triage of high-risk HPV-positive women in population-based screening by miRNA expression analysis in cervical scrapes; a feasibility study. Clinical Epigenetics, 2018, 10, 76.	4.1	18
54	Molecular heterogeneity in human papillomavirusâ€dependent and â€independent vulvar carcinogenesis. Cancer Medicine, 2018, 7, 4542-4553.	2.8	21

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55	HPV E4 expression and DNA hypermethylation of CADM1, MAL, and miR124-2 genes in cervical cancer and precursor lesions. Modern Pathology, 2018, 31, 1842-1850.	5.5	37
56	Genome-wide DNA Methylation Profiling Reveals Methylation Markers Associated with 3q Gain for Detection of Cervical Precancer and Cancer. Clinical Cancer Research, 2017, 23, 3813-3822.	7.0	68
57	Development of a replicationâ€deficient adenoviral vectorâ€based vaccine candidate for the interception of HPV16â€and HPV18â€induced infections and disease. International Journal of Cancer, 2017, 141, 393-404.	5.1	19
58	Novel molecular subtypes of cervical cancer â€" potential clinical consequences. Nature Reviews Clinical Oncology, 2017, 14, 397-398.	27.6	18
59	DNA hypermethylation analysis in sputum of asymptomatic subjects at risk for lung cancer participating in the NELSON trial: argument for maximum screening interval of 2â€…years. Journal of Clinical Pathology, 2017, 70, 250-254.	2.0	21
60	A Strategy to Find Suitable Reference Genes for miRNA Quantitative PCR Analysis and Its Application to Cervical Specimens. Journal of Molecular Diagnostics, 2017, 19, 625-637.	2.8	21
61	Selection of women at risk for cervical cancer in an HIV-infected South African population. Aids, 2017, 31, 1945-1953.	2.2	17
62	Good performance of p16/kiâ€67 dualâ€stained cytology for surveillance of women treated for highâ€grade CIN. International Journal of Cancer, 2017, 140, 423-430.	5.1	16
63	High Levels of EBV-Encoded RNA 1 (EBER1) Trigger Interferon and Inflammation-Related Genes in Keratinocytes Expressing HPV16 E6/E7. PLoS ONE, 2017, 12, e0169290.	2.5	14
64	Symptomatic HPV-related neovaginal lesions in transgender women: case series and review of literature. Sexually Transmitted Infections, 2016, 92, 499-501.	1.9	24
65	Validation of the FAM19A4 / mir124-2 DNA methylation test for both lavage- and brush-based self-samples to detect cervical (pre)cancer in HPV-positive women. Gynecologic Oncology, 2016, 141, 341-347.	1.4	80
66	Molecular events leading to HPV-induced high grade neoplasia. Papillomavirus Research (Amsterdam,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	4.5	40
67	Management of high-risk HPV-positive women for detection of cervical (pre)cancer. Expert Review of Molecular Diagnostics, 2016, 16, 961-974.	3.1	45
68	FAM19A4 methylation analysis in self-samples compared with cervical scrapes for detecting cervical (pre)cancer in HPV-positive women. British Journal of Cancer, 2016, 115, 579-587.	6.4	55
69	Performance of CADM1/MAL-methylation analysis for monitoring of women treated for high-grade CIN. Gynecologic Oncology, 2016, 143, 135-142.	1.4	18
70	Prevalence of Neovaginal High-Risk Human Papillomavirus Among Transgender Women in The Netherlands. Sexually Transmitted Diseases, 2016, 43, 503-505.	1.7	29
71	Comparing the performance of FAM19A4 methylation analysis, cytology and HPV16/18 genotyping for the detection of cervical (pre)cancer in highâ€risk HPVâ€positive women of a gynecologic outpatient population (COMETH study). International Journal of Cancer, 2016, 138, 992-1002.	5.1	60
72	Genome-wide methylome analysis using MethylCap-seq uncovers 4 hypermethylated markers with high sensitivity for both adeno- and squamous-cell cervical carcinoma. Oncotarget, 2016, 7, 80735-80750.	1.8	15

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73	Immortalization capacity of HPV types is inversely related to chromosomal instability. Oncotarget, 2016, 7, 37608-37621.	1.8	25
74	Aberrant methylation-mediated silencing of microRNAs contributes to HPV-induced anchorage independence. Oncotarget, 2016, 7, 43805-43819.	1.8	22
75	Association Between Type-specific HPV Infections and hTERT DNA Methylation in Patients with Invasive Cervical Cancer. Cancer Genomics and Proteomics, 2016, 13, 483-492.	2.0	10
76	Oncogenic Role of miR-15a-3p in 13q Amplicon-Driven Colorectal Adenoma-to-Carcinoma Progression. PLoS ONE, 2015, 10, e0132495.	2.5	22
77	Somatic mutation in PIK3CA is a late event in cervical carcinogenesis. Journal of Pathology: Clinical Research, 2015, 1, 207-211.	3.0	24
78	Longitudinal assessment of DNA methylation changes during HPVE6E7-induced immortalization of primary keratinocytes. Epigenetics, 2015, 10, 73-81.	2.7	29
79	Mining for viral fragments in methylation enriched sequencing data. Frontiers in Genetics, 2015, 6, 16.	2.3	5
80	Follow-up of high-risk HPV positive women by combined cytology and bi-marker CADM1/MAL methylation analysis on cervical scrapes. Gynecologic Oncology, 2015, 137, 55-59.	1.4	22
81	Interplay between promoter methylation and chromosomal loss in gene silencing at 3p11-p14 in cervical cancer. Epigenetics, 2015, 10, 970-980.	2.7	47
82	Combined CADM1/MAL Methylation and Cytology Testing for Colposcopy Triage of High-Risk HPV-Positive Women. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1933-1937.	2.5	39
83	Methylation-mediated repression of PRDM14 contributes to apoptosis evasion in HPV-positive cancers. Carcinogenesis, 2014, 35, 2611-2618.	2.8	35
84	Combined sputum hypermethylation and eNose analysis for lung cancer diagnosis. Journal of Clinical Pathology, 2014, 67, 707-711.	2.0	56
85	tigaR: integrative significance analysis of temporal differential gene expression induced by genomic abnormalities. BMC Bioinformatics, 2014, 15, 327.	2.6	2
86	CADM1, MAL and miR124-2 methylation analysis in cervical scrapes to detect cervical and endometrial cancer. Journal of Clinical Pathology, 2014, 67, 1067-1071.	2.0	82
87	Geneâ€dosage dependent overexpression at the 13q amplicon identifies DIS3 as candidate oncogene in colorectal cancer progression. Genes Chromosomes and Cancer, 2014, 53, 339-348.	2.8	31
88	Lactate transporters and vascular factors in HPV-induced squamous cell carcinoma of the uterine cervix. BMC Cancer, 2014, 14, 751.	2.6	23
89	Differential In Vitro Immortalization Capacity of Eleven, Probable High-Risk Human Papillomavirus Types. Journal of Virology, 2014, 88, 1714-1724.	3.4	27
90	Clinical implications of (epi)genetic changes in HPV-induced cervical precancerous lesions. Nature Reviews Cancer, 2014, 14, 395-405.	28.4	295

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91	Methylation Analysis of the FAM19A4 Gene in Cervical Scrapes Is Highly Efficient in Detecting Cervical Carcinomas and Advanced CIN2/3 Lesions. Cancer Prevention Research, 2014, 7, 1251-1257.	1.5	97
92	Focal chromosomal copy number aberrations in cancerâ€"Needles in a genome haystack. Biochimica Et Biophysica Acta - Molecular Cell Research, 2014, 1843, 2698-2704.	4.1	55
93	Methylation marker analysis and HPV16/18 genotyping in high-risk HPV positive self-sampled specimens to identify women with high grade CIN or cervical cancer. Gynecologic Oncology, 2014, 135, 58-63.	1.4	45
94	Chromosomal gains and losses in human papillomavirus-associated neoplasia of the lower genital tract â€" A systematic review and meta-analysis. European Journal of Cancer, 2014, 50, 85-98.	2.8	70
95	Triage by methylation-marker testing versus cytology in women who test HPV-positive on self-collected cervicovaginal specimens (PROHTECT-3): a randomised controlled non-inferiority trial. Lancet Oncology, The, 2014, 15, 315-322.	10.7	147
96	Detection limits of DNA copy number alterations in heterogeneous cell populations. Cellular Oncology (Dordrecht), 2013, 36, 27-36.	4.4	18
97	Promoter methylation of Wnt-antagonists in polypoid and nonpolypoid colorectal adenomas. BMC Cancer, 2013, 13, 603.	2.6	23
98	Methylation-specific digital karyotyping of HPV16E6E7-expressing human keratinocytes identifies novel methylation events in cervical carcinogenesis. Journal of Pathology, 2013, 231, 53-62.	4.5	48
99	Identification of eight candidate target genes of the recurrent 3p12â€"p14 loss in cervical cancer by integrative genomic profiling. Journal of Pathology, 2013, 230, 59-69.	4.5	37
100	CADM1andMALpromoter methylation levels in hrHPV-positive cervical scrapes increase proportional to degree and duration of underlying cervical disease. International Journal of Cancer, 2013, 133, 1293-1299.	5.1	100
101	Focal aberrations indicate EYA2 and hsaâ€miRâ€375 as oncogene and tumor suppressor in cervical carcinogenesis. Genes Chromosomes and Cancer, 2013, 52, 56-68.	2.8	76
102	Methylation-mediated transcriptional repression of microRNAs during cervical carcinogenesis. Epigenetics, 2013, 8, 220-228.	2.7	67
103	High-Risk Human Papillomavirusâ€"Positive Lung Cancer: Molecular Evidence for a Pattern of Pulmonary Metastasis. Journal of Thoracic Oncology, 2013, 8, 711-718.	1.1	39
104	Total RNA Isolation after Laser-capture Microdissection of Human Cervical Squamous Epithelial Cells from Fresh Frozen Tissue. Bio-protocol, 2013, 3, .	0.4	0
105	Comprehensive analysis of human papillomavirus prevalence and the potential role of low-risk types in verrucous carcinoma. Modern Pathology, 2012, 25, 1354-1363.	5.5	66
106	Development of a multiplex methylation-specific PCR as candidate triage test for women with an HPV-positive cervical scrape. BMC Cancer, 2012, 12, 551.	2.6	54
107	Chromosomal profiles of highâ€grade cervical intraepithelial neoplasia relate to duration of preceding highâ€risk human papillomavirus infection. International Journal of Cancer, 2012, 131, E579-85.	5.1	37
108	The functional role of Notch signaling in HPV-mediated transformation is dose-dependent and linked to AP-1 alterations. Cellular Oncology (Dordrecht), 2012, 35, 77-84.	4.4	13

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109	Methylation status of the E2 binding sites of HPV16 in cervical lesions determined with the LuminexÂ® xMAPâ„¢ system. Virology, 2012, 422, 357-365.	2.4	30
110	HPV type-related chromosomal profiles in high-grade cervical intraepithelial neoplasia. BMC Cancer, 2012, 12, 36.	2.6	10
111	CGH arrays compared for DNA isolated from formalinâ€fixed, paraffinâ€embedded material. Genes Chromosomes and Cancer, 2012, 51, 344-352.	2.8	33
112	Comprehensive CADM1 promoter methylation analysis in NSCLC and normal lung specimens. Lung Cancer, 2011, 72, 316-321.	2.0	6
113	Epigenetic markers for early detection of nasopharyngeal carcinoma in a high risk population. Molecular Cancer, 2011, 10, 48.	19.2	68
114	PIK3CA-mediated PI3-kinase signalling is essential for HPV-induced transformation in vitro. Molecular Cancer, 2011, 10, 71.	19.2	47
115	Immortalization of oral keratinocytes by functional inactivation of the p53 and pRb pathways. International Journal of Cancer, 2011, 128, 1596-1605.	5.1	84
116	Combined CADM1 and MAL promoter methylation analysis to detect (preâ€)malignant cervical lesions in highâ€risk HPVâ€positive women. International Journal of Cancer, 2011, 129, 2218-2225.	5.1	87
117	Combined Promoter Methylation Analysis of CADM1 and MAL: An Objective Triage Tool for High-Risk Human Papillomavirus DNAâ€"Positive Women. Clinical Cancer Research, 2011, 17, 2459-2465.	7.0	119
118	hTERT promoter activity and CpG methylation in HPV-induced carcinogenesis. BMC Cancer, 2010, 10, 271.	2.6	53
119	Methylation-mediated silencing and tumour suppressive function of hsa-miR-124 in cervical cancer. Molecular Cancer, 2010, 9, 167.	19.2	217
120	The dynamic DNA methylomes of double-stranded DNA viruses associated with human cancer. Genome Research, 2009, 19, 438-451.	5.5	218
121	Chromosomal Signatures of a Subset of High-Grade Premalignant Cervical Lesions Closely Resemble Invasive Carcinomas. Cancer Research, 2009, 69, 647-655.	0.9	53
122	Repression of MAL tumour suppressor activity by promoter methylation during cervical carcinogenesis. Journal of Pathology, 2009, 219, 327-336.	4.5	95
123	Genomic profiling identifies common HPV-associated chromosomal alterations in squamous cell carcinomas of cervix and head and neck. BMC Medical Genomics, 2009, 2, 32.	1.5	56
124	Association between dense CADM1 promoter methylation and reduced protein expression in highâ€grade CIN and cervical SCC. Journal of Pathology, 2008, 215, 388-397.	4.5	92
125	Gene expression profiling to identify markers associated with deregulated hTERT in HPVâ€transformed keratinocytes and cervical cancer. International Journal of Cancer, 2008, 122, 877-888.	5.1	26
126	Integrated genomic and transcriptional profiling identifies chromosomal loci with altered gene expression in cervical cancer. Genes Chromosomes and Cancer, 2008, 47, 890-905.	2.8	59

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127	Specific betapapillomaviruses associated with squamous cell carcinoma of the skin inhibit UVB-induced apoptosis of primary human keratinocytes. Journal of General Virology, 2008, 89, 2303-2314.	2.9	59
128	Alterations in AP-1 and AP-1 Regulatory Genes during HPV-Induced Carcinogenesis. Analytical Cellular Pathology, 2008, 30, 77-87.	1.4	25
129	A Role for EZH2 in Silencing of IFN-Î³ Inducible MHC2TA Transcription in Uveal Melanoma. Journal of Immunology, 2007, 179, 5317-5325.	0.8	51
130	Molecular Markers for Cervical Cancer. , 2006, , 73-81.		0
131	HPVâ€mediated cervical carcinogenesis: concepts and clinical implications. Journal of Pathology, 2006, 208, 152-164.	4.5	360
132	Increased gene copy numbers at chromosome 20q are frequent in both squamous cell carcinomas and adenocarcinomas of the cervix. Journal of Pathology, 2006, 209, 220-230.	4.5	96
133	Oncolytic Adenovirus Expressing a p53 Variant Resistant to Degradation by HPV E6 Protein Exhibits Potent and Selective Replication in Cervical Cancer. Molecular Therapy, 2005, 12, 1083-1090.	8.2	25
134	HPV-mediated transformation of the anogenital tract. Journal of Clinical Virology, 2005, 32, 25-33.	3.1	130
135	TSLC1 Gene Silencing in Cervical Cancer Cell Lines and Cervical Neoplasia. Journal of the National Cancer Institute, 2004, 96, 294-305.	6.3	194
136	Clonal Selection for Transcriptionally Active Viral Oncogenes during Progression to Cancer. Journal of Virology, 2004, 78, 11172-11186.	3.4	116
137	Elevated hTERT mRNA levels: A potential determinant of bronchial squamous cell carcinoma (in) Tj ETQq1 1 0.784314 rgBT /Overlock	5.1	21
138	Assessment of TGF-Î²1-mediated growth inhibition of HPV-16- and HPV-18-transfected foreskin keratinocytes during and following immortalization. Archives of Dermatological Research, 2003, 295, 297-304.	1.9	3
139	Telomerase activity in high-grade cervical lesions is associated with allelic imbalance at 6Q14-22. International Journal of Cancer, 2003, 105, 577-582.	5.1	16
140	Down-Regulation of GATA-3 Expression during Human Papillomavirus-Mediated Immortalization and Cervical Carcinogenesis. American Journal of Pathology, 2002, 160, 1945-1951.	3.8	39
141	Quantitative reverse transcription-polymerase chain reaction measurement of HASH1 (ASCL1), a marker for small cell lung carcinomas with neuroendocrine features. Clinical Cancer Research, 2002, 8, 1082-6.	7.0	35
142	Telomerase Suppression by Chromosome 6 in a Human Papillomavirus Type 16-Immortalized Keratinocyte Cell Line and in a Cervical Cancer Cell Line. Journal of the National Cancer Institute, 2001, 93, 865-872.	6.3	67
143	Non-random allelic losses at 3p, 11p and 13q during HPV-mediated immortalization and concomitant loss of terminal differentiation of human keratinocytes. , 1998, 76, 412-417.		31
144	Hemidesmosome Formation Is Initiated by the Î²4 Integrin Subunit, Requires Complex Formation of Î²4 and HD1/Plectin, and Involves a Direct Interaction between Î²4 and the Bullous Pemphigoid Antigen 180. Journal of Cell Biology, 1998, 142, 271-284.	5.2	171

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145	Viral E6-E7 Transcription in the Basal Layer of Organotypic Cultures without Apparent p21cip1 Protein Precedes Immortalization of Human Papillomavirus Type 16- and 18-Transfected Human Keratinocytes. Journal of Virology, 1998, 72, 749-757.	3.4	47
146	Three Sensitive Methods for the Detection of Cytomegalovirus in Lung Tissue of Patients with Interstitial Pneumonitis. American Journal of Clinical Pathology, 1990, 93, 491-494.	0.7	41