

Lise M A De Strooper

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

730
citations

687363

13
h-index

996975

15
g-index

15
all docs

15
docs citations

15
times ranked

648
citing authors

#	ARTICLE	IF	CITATIONS
1	Methylation Analysis of the <i>FAM19A4</i> Gene in Cervical Scrapes Is Highly Efficient in Detecting Cervical Carcinomas and Advanced CIN2/3 Lesions. <i>Cancer Prevention Research</i> , 2014, 7, 1251-1257.	1.5	97
2	<i>CADM1</i> , <i>MAL</i> and <i>miR124-2</i> methylation analysis in cervical scrapes to detect cervical and endometrial cancer. <i>Journal of Clinical Pathology</i> , 2014, 67, 1067-1071.	2.0	82
3	Validation of the <i>FAM19A4</i> / <i>mir124-2</i> DNA methylation test for both lavage- and brush-based self-samples to detect cervical (pre)cancer in HPV-positive women. <i>Gynecologic Oncology</i> , 2016, 141, 341-347.	1.4	80
4	Genome-wide DNA Methylation Profiling Reveals Methylation Markers Associated with 3q Gain for Detection of Cervical Precancer and Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 3813-3822.	7.0	68
5	Cervical cancer risk in HPV-positive women after a negative <i>FAM19A4/mir124-2</i> methylation test: A post hoc analysis in the POBASCAM trial with 14 year follow-up. <i>International Journal of Cancer</i> , 2018, 143, 1541-1548.	5.1	63
6	Comparing the performance of <i>FAM19A4</i> methylation analysis, cytology and HPV16/18 genotyping for the detection of cervical (pre)cancer in high-risk HPV-positive women of a gynecologic outpatient population (COMETH study). <i>International Journal of Cancer</i> , 2016, 138, 992-1002.	5.1	60
7	<i>FAM19A4</i> methylation analysis in self-samples compared with cervical scrapes for detecting cervical (pre)cancer in HPV-positive women. <i>British Journal of Cancer</i> , 2016, 115, 579-587.	6.4	55
8	Development of a multiplex methylation-specific PCR as candidate triage test for women with an HPV-positive cervical scrape. <i>BMC Cancer</i> , 2012, 12, 551.	2.6	54
9	Management of high-risk HPV-positive women for detection of cervical (pre)cancer. <i>Expert Review of Molecular Diagnostics</i> , 2016, 16, 961-974.	3.1	45
10	Combined <i>CADM1</i> / <i>MAL</i> Methylation and Cytology Testing for Colposcopy Triage of High-Risk HPV-Positive Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1933-1937.	2.5	39
11	Long-term CIN3+ risk of HPV positive women after triage with <i>FAM19A4/mir124-2</i> methylation analysis. <i>Gynecologic Oncology</i> , 2019, 154, 368-373.	1.4	32
12	Classification of high-grade cervical intraepithelial neoplasia by p16 ^{ink4a} , Ki67, HPV E4 _{scp} and <i>FAM19A4</i> / <i>miR124-2</i> methylation status demonstrates considerable heterogeneity with potential consequences for management. <i>International Journal of Cancer</i> , 2021, 149, 707-716.	5.1	26
13	Evaluation of six methylation markers derived from genome-wide screens for detection of cervical precancer and cancer. <i>Epigenomics</i> , 2020, 12, 1569-1578.	2.1	15
14	Defining hrHPV genotypes in cervical intraepithelial neoplasia by laser capture microdissection supports reflex triage of self-samples using HPV16/18 and <i>FAM19A4/mir124-2</i> methylation. <i>Gynecologic Oncology</i> , 2018, 151, 311-318.	1.4	7
15	Complementarity between miRNA expression analysis and DNA methylation analysis in hrHPV-positive cervical scrapes for the detection of cervical disease. <i>Epigenetics</i> , 2019, 14, 558-567.	2.7	7