

Itzel E Calleja-Macias

List of Publications by Year in descending order

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16
papers

1,176
citations

567144

15
h-index

940416

16
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all docs

16
docs citations

16
times ranked

1087
citing authors

#	ARTICLE	IF	CITATIONS
1	Prevalence and 3-year persistence of human papillomavirus serotypes in asymptomatic patients in Northern Mexico. <i>International Journal of Gynecology and Obstetrics</i> , 2017, 136, 40-46.	1.0	5
2	Methylation of human papillomavirus 16, 18, 31, and 45 L2 and L1 genes and the cellular DAPK gene: Considerations for use as biomarkers of the progression of cervical neoplasia. <i>Virology</i> , 2014, 448, 314-321.	1.1	53
3	Muscarinic cholinergic signaling in cervical cancer cells affects cell motility via ERK1/2 signaling. <i>Life Sciences</i> , 2012, 91, 1093-1098.	2.0	19
4	Laser capture microdissection of cervical human papillomavirus infections: Copy number of the virus in cancerous and normal tissue and heterogeneous DNA methylation. <i>Virology</i> , 2009, 390, 261-267.	1.1	54
5	Cholinergic signaling through nicotinic acetylcholine receptors stimulates the proliferation of cervical cancer cells: An explanation for the molecular role of tobacco smoking in cervical carcinogenesis?. <i>International Journal of Cancer</i> , 2009, 124, 1090-1096.	2.3	29
6	Human papillomavirus-16 and -18 in penile carcinomas: DNA methylation, chromosomal recombination and genomic variation. <i>International Journal of Cancer</i> , 2008, 123, 1832-1840.	2.3	59
7	Effects of cellular differentiation, chromosomal integration and 5-aza-2'-deoxycytidine treatment on human papillomavirus-16 DNA methylation in cultured cell lines. <i>Virology</i> , 2008, 374, 292-303.	1.1	50
8	Human papillomavirus-16 DNA methylation patterns support a causal association of the virus with oral squamous cell carcinomas. <i>International Journal of Cancer</i> , 2007, 120, 2165-2169.	2.3	44
9	Methylation of the human papillomavirus-18 L1 gene: A biomarker of neoplastic progression?. <i>Virology</i> , 2006, 349, 175-183.	1.1	83
10	Genome variation of human papillomavirus types: Phylogenetic and medical implications. <i>International Journal of Cancer</i> , 2006, 118, 1071-1076.	2.3	257
11	Worldwide genomic diversity of the human papillomaviruses-53, 56, and 66, a group of high-risk HPVs unrelated to HPV-16 and HPV-18. <i>Virology</i> , 2005, 340, 95-104.	1.1	55
12	Worldwide Genomic Diversity of the High-Risk Human Papillomavirus Types 31, 35, 52, and 58, Four Close Relatives of Human Papillomavirus Type 16. <i>Journal of Virology</i> , 2005, 79, 13630-13640.	1.5	95
13	Papillomavirus Subtypes Are Natural and Old Taxa: Phylogeny of Human Papillomavirus Types 44 and 55 and 68a and -b. <i>Journal of Virology</i> , 2005, 79, 6565-6569.	1.5	39
14	Conserved Methylation Patterns of Human Papillomavirus Type 16 DNA in Asymptomatic Infection and Cervical Neoplasia. <i>Journal of Virology</i> , 2004, 78, 12762-12772.	1.5	165
15	Genomic diversity of human papillomavirus-16, 18, 31, and 35 isolates in a Mexican population and relationship to European, African, and Native American variants. <i>Virology</i> , 2004, 319, 315-323.	1.1	81
16	The human papillomavirus-18 genome is efficiently targeted by cellular DNA methylation. <i>Virology</i> , 2004, 324, 483-492.	1.1	88