

CITATION REPORT

List of articles citing

Comparison of papillomavirus and immunodeficiency virus evolutionary patterns in the context of a papillomavirus vaccine

DOI: 10.1016/s1386-6532(00)00127-x
Journal of Clinical Virology, 2000, 19, 43-56.

Source: <https://exaly.com/paper-pdf/32062054/citation-report.pdf>

Version: 2024-04-28

This report has been generated based on the citations recorded by exaly.com for the above article. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

#	Paper	IF	Citations
28	Primary prevention of uterine cervix cancer: focus on vaccine history and current strategy. <i>Obstetrics and Gynecology Clinics of North America</i> , 2002 , 29, 843-68, ix	3.3	5
27	Mucosal human papillomaviruses encode four different E5 proteins whose chemistry and phylogeny correlate with malignant or benign growth. <i>Journal of Virology</i> , 2004 , 78, 13613-26	6.6	93
26	Classification of papillomaviruses. <i>Virology</i> , 2004 , 324, 17-27	3.6	2227
25	Overcoming antigenic diversity and improving vaccines using DNA shuffling and screening technologies. <i>Expert Opinion on Biological Therapy</i> , 2004 , 4, 589-97	5.4	7
24	Patterns of nucleotide difference in overlapping and non-overlapping reading frames of papillomavirus genomes. <i>Virus Research</i> , 2005 , 113, 81-8	6.4	48
23	Papillomaviruses: different genes have different histories. <i>Trends in Microbiology</i> , 2005 , 13, 514-21	12.4	114
22	Evidence of ancient papillomavirus recombination. <i>Journal of General Virology</i> , 2006 , 87, 2527-2531	4.9	56
21	Multiple evolutionary mechanisms drive papillomavirus diversification. <i>Molecular Biology and Evolution</i> , 2007 , 24, 1242-58	8.3	91
20	Evidence of recombination within human alpha-papillomavirus. <i>Virology Journal</i> , 2007 , 4, 33	6.1	33
19	Phylogenetic analysis of beta-papillomaviruses as inferred from nucleotide and amino acid sequence data. <i>Molecular Phylogenetics and Evolution</i> , 2007 , 42, 213-22	4.1	34
18	Novel papillomavirus isolates from <i>Erinaceus europaeus</i> (Erinaceidae, Insectivora) and the Cervidae (Artiodactyla), <i>Cervus timorensis</i> and <i>Pudu puda</i> , and phylogenetic analysis of partial sequence data. <i>Virus Genes</i> , 2008 , 36, 281-7	2.3	11
17	Detecting recombination and diversifying selection in human alpha-papillomavirus. <i>Infection, Genetics and Evolution</i> , 2008 , 8, 689-92	4.5	21
16	Vaccination and the evolutionary ecology of human papillomavirus. <i>Vaccine</i> , 2008 , 26 Suppl 3, C25-30	4.1	15
15	Isolation and genomic characterization of the first Norway rat (<i>Rattus norvegicus</i>) papillomavirus and its phylogenetic position within Pipapillomavirus, primarily infecting rodents. <i>Journal of General Virology</i> , 2009 , 90, 2609-2614	4.9	21
14	Modular organizations of novel cetacean papillomaviruses. <i>Molecular Phylogenetics and Evolution</i> , 2011 , 59, 34-42	4.1	41
13	Quantifying the phylodynamic forces driving papillomavirus evolution. <i>Molecular Biology and Evolution</i> , 2011 , 28, 2101-13	8.3	99
12	Molecular and evolutionary analysis of HPV16 E6 and E7 genes in Greek women. <i>Journal of Medical Microbiology</i> , 2013 , 62, 1688-1696	3.2	10

11	Measuring effectiveness of the cervical cancer vaccine in an Australian setting (the VACCINE study). <i>BMC Cancer</i> , 2013 , 13, 296	4.8	18
10	Identification of novel E6-E7 sequence variants of human papillomavirus 16. <i>Archives of Virology</i> , 2013 , 158, 821-8	2.6	5
9	Novel papillomaviruses in free-ranging Iberian bats: no virus-host co-evolution, no strict host specificity, and hints for recombination. <i>Genome Biology and Evolution</i> , 2014 , 6, 94-104	3.9	47
8	Analysis of the long control region of bovine papillomavirus type 1 associated with sarcoids in equine hosts indicates multiple cross-species transmission events and phylogeographical structure. <i>Journal of General Virology</i> , 2014 , 95, 2748-2756	4.9	28
7	Human papillomavirus detection in Corrientes, Argentina: High prevalence of type 58 and its phylogenetics. <i>Revista Argentina De Microbiologia</i> , 2015 , 47, 302-11	1.8	2
6	In vitro assessment of the effect of vaccine-targeted human papillomavirus (HPV) depletion on detection of non-vaccine HPV types: implications for post-vaccine surveillance studies. <i>Journal of Virological Methods</i> , 2015 , 214, 10-4	2.6	10
5	Could the human papillomavirus vaccines drive virulence evolution?. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015 , 282, 20141069	4.4	24
4	Genetic characterization and clinical implications of human papillomavirus type 16 (HPV16) variants from northeastern Argentina. <i>Infection, Genetics and Evolution</i> , 2015 , 29, 103-9	4.5	5
3	The Role of aDNA in Understanding the Coevolutionary Patterns of Human Sexually Transmitted Infections. <i>Genes</i> , 2018 , 9,	4.2	8
2	Papillomavirus vaccination for prevention and treatment of cervical carcinoma. 2001 , 3, 231-240		
1	The T350G Variation of Human Papillomavirus 16 E6 Gene Prevails in Oropharyngeal Cancer from a Small Cohort of Greek Patients. 2022 , 14, 1724		1