

# Koenraad Van Doorslaer

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3824939/publications.pdf>

Version: 2024-02-01

49  
papers

3,989  
citations

201674

27  
h-index

206112

48  
g-index

57  
all docs

57  
docs citations

57  
times ranked

4017  
citing authors

#	ARTICLE	IF	CITATIONS
1	Classification of papillomaviruses (PVs) based on 189 PV types and proposal of taxonomic amendments. <i>Virology</i> , 2010, 401, 70-79.	2.4	1,377
2	The Papillomavirus Episteme: a major update to the papillomavirus sequence database. <i>Nucleic Acids Research</i> , 2017, 45, D499-D506.	14.5	298
3	Changes to virus taxonomy and to the International Code of Virus Classification and Nomenclature ratified by the International Committee on Taxonomy of Viruses (2021). <i>Archives of Virology</i> , 2021, 166, 2633-2648.	2.1	219
4	The Ancient Evolutionary History of Polyomaviruses. <i>PLoS Pathogens</i> , 2016, 12, e1005574.	4.7	190
5	The Papillomavirus Episteme: a central resource for papillomavirus sequence data and analysis. <i>Nucleic Acids Research</i> , 2012, 41, D571-D578.	14.5	188
6	Evolution of the Papillomaviridae. <i>Virology</i> , 2013, 445, 11-20.	2.4	176
7	Ancient papillomavirus-host co-speciation in Felidae. <i>Genome Biology</i> , 2007, 8, R57.	9.6	152
8	ICTV Virus Taxonomy Profile: Papillomaviridae. <i>Journal of General Virology</i> , 2018, 99, 989-990.	2.9	140
9	Human Papillomaviruses: Genetic Basis of Carcinogenicity. <i>Public Health Genomics</i> , 2009, 12, 281-290.	1.0	113
10	Roles of APOBEC3A and APOBEC3B in Human Papillomavirus Infection and Disease Progression. <i>Viruses</i> , 2017, 9, 233.	3.3	79
11	Metagenomic Discovery of 83 New Human Papillomavirus Types in Patients with Immunodeficiency. <i>MSphere</i> , 2018, 3, .	2.9	75
12	Sequence Imputation of HPV16 Genomes for Genetic Association Studies. <i>PLoS ONE</i> , 2011, 6, e21375.	2.5	70
13	Genomic characterization of two novel reptilian papillomaviruses, <i>Chelonia mydas</i> papillomavirus 1 and <i>Caretta caretta</i> papillomavirus 1. <i>Virology</i> , 2009, 383, 131-135.	2.4	67
14	Role of the host restriction factor APOBEC3 on papillomavirus evolution. <i>Virus Evolution</i> , 2015, 1, vev015.	4.9	57
15	Association between hTERT activation by HPV E6 proteins and oncogenic risk. <i>Virology</i> , 2012, 433, 216-219.	2.4	54
16	Degradation of p53 by Human Alphapapillomavirus E6 Proteins Shows a Stronger Correlation with Phylogeny than Oncogenicity. <i>PLoS ONE</i> , 2010, 5, e12816.	2.5	53
17	Unique genome organization of non-mammalian papillomaviruses provides insights into the evolution of viral early proteins. <i>Virus Evolution</i> , 2017, 3, vex027.	4.9	51
18	Isolation and cloning of the raccoon ( <i>Procyon lotor</i> ) papillomavirus type 1 by using degenerate papillomavirus-specific primers. <i>Journal of General Virology</i> , 2005, 86, 2029-2033.	2.9	46

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19	Isolation and cloning of a papillomavirus from a North American porcupine by using multiply primed rolling-circle amplification: the Erethizon dorsatum papillomavirus type 1. <i>Virology</i> , 2005, 331, 449-456.	2.4	45
20	Molecular archeological evidence in support of the repeated loss of a papillomavirus gene. <i>Scientific Reports</i> , 2016, 6, 33028.	3.3	36
21	Papillomaviruses: evolution, Linnaean taxonomy and current nomenclature. <i>Trends in Microbiology</i> , 2011, 19, 49-50.	7.7	35
22	Identification of Unusual E6 and E7 Proteins within Avian Papillomaviruses: Cellular Localization, Biophysical Characterization, and Phylogenetic Analysis. <i>Journal of Virology</i> , 2009, 83, 8759-8770.	3.4	33
23	Differentiating between viruses and virus species by writing their names correctly. <i>Archives of Virology</i> , 2022, 167, 1231-1234.	2.1	33
24	Vesicular trafficking permits evasion of cGAS/STING surveillance during initial human papillomavirus infection. <i>PLoS Pathogens</i> , 2020, 16, e1009028.	4.7	32
25	Complete genomic characterization of a murine papillomavirus isolated from papillomatous lesions of a European harvest mouse ( <i>Micromys minutus</i> ). <i>Journal of General Virology</i> , 2007, 88, 1484-1488.	2.9	31
26	Serological response to an HPV16 E7 based therapeutic vaccine in women with high-grade cervical dysplasia. <i>Gynecologic Oncology</i> , 2010, 116, 208-212.	1.4	30
27	Evolution of Human Papillomavirus Carcinogenicity. <i>Advances in Virus Research</i> , 2010, 77, 41-62.	2.1	29
28	Genetic characterization of the first chiropteran papillomavirus, isolated from a basosquamous carcinoma in an Egyptian fruit bat: The <i>Rousettus aegyptiacus</i> papillomavirus type 1. <i>Veterinary Microbiology</i> , 2006, 117, 267-275.	1.9	27
29	Genetic characterization of the <i>Capra hircus</i> papillomavirus: A novel close-to-root artiodactyl papillomavirus. <i>Virus Research</i> , 2006, 118, 164-169.	2.2	26
30	<i>Neisseria gonorrhoeae</i> evades autophagic killing by downregulating CD46-cyt1 and remodeling lysosomes. <i>PLoS Pathogens</i> , 2019, 15, e1007495.	4.7	23
31	Degradation of Human PDZ-Proteins by Human Alphapapillomaviruses Represents an Evolutionary Adaptation to a Novel Cellular Niche. <i>PLoS Pathogens</i> , 2015, 11, e1004980.	4.7	20
32	Fish polyomaviruses belong to two distinct evolutionary lineages. <i>Journal of General Virology</i> , 2018, 99, 567-573.	2.9	19
33	3D Oral and Cervical Tissue Models for Studying Papillomavirus Host-Pathogen Interactions. <i>Current Protocols in Microbiology</i> , 2020, 59, e129.	6.5	16
34	Novel recombinant papillomavirus genomes expressing selectable genes. <i>Scientific Reports</i> , 2016, 6, 37782.	3.3	13
35	Persistence of an Oncogenic Papillomavirus Genome Requires <i>cis</i> Elements from the Viral Transcriptional Enhancer. <i>MBio</i> , 2017, 8, .	4.1	13
36	Novel Circoviruses Detected in Feces of Sonoran Felids. <i>Viruses</i> , 2020, 12, 1027.	3.3	13

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37	Complex evolutionary history of felid anelloviruses. <i>Virology</i> , 2021, 562, 176-189.	2.4	13
38	PuMA: A papillomavirus genome annotation tool. <i>Virus Evolution</i> , 2020, 6, veaa068.	4.9	12
39	The Launch of an International Animal Papillomavirus Reference Center. <i>Viruses</i> , 2019, 11, 55.	3.3	10
40	A Novel Divergent Geminivirus Identified in Asymptomatic New World Cactaceae Plants. <i>Viruses</i> , 2020, 12, 398.	3.3	10
41	Discovery of novel fish papillomaviruses: From the Antarctic to the commercial fish market. <i>Virology</i> , 2022, 565, 65-72.	2.4	10
42	A novel lineage of polyomaviruses identified in bark scorpions. <i>Virology</i> , 2021, 563, 58-63.	2.4	9
43	New World Cactaceae Plants Harbor Diverse Geminiviruses. <i>Viruses</i> , 2021, 13, 694.	3.3	8
44	Regulation of Human Papillomavirus 18 Genome Replication, Establishment, and Persistence by Sequences in the Viral Upstream Regulatory Region. <i>Journal of Virology</i> , 2021, 95, e0068621.	3.4	7
45	Mechanisms of DNA Virus Evolution. , 2021, , 71-78.		6
46	Building (Viral) Phylogenetic Trees Using a Maximum Likelihood Approach. <i>Current Protocols in Microbiology</i> , 2018, 51, e63.	6.5	5
47	Insertional oncogenesis by HPV70 revealed by multiple genomic analyses in a clinically HPV-negative cervical cancer. <i>Genes Chromosomes and Cancer</i> , 2020, 59, 84-95.	2.8	5
48	Coevolutionary Analysis Implicates Toll-Like Receptor 9 in Papillomavirus Restriction. <i>MBio</i> , 2022, 13, e0005422.	4.1	5
49	HPV32-related Heck's disease in a chronic graft-versus-host disease patient with long-term successful KTP laser treatment: A rare case report. <i>Clinical Case Reports (discontinued)</i> , 2021, 9, e04253.	0.5	3