## Koenraad Van Doorslaer

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Discovery of novel fish papillomaviruses: From the Antarctic to the commercial fish market. Virology, 2022, 565, 65-72.	2.4	10
2	Differentiating between viruses and virus species by writing their names correctly. Archives of Virology, 2022, 167, 1231-1234.	2.1	33
3	Coevolutionary Analysis Implicates Toll-Like Receptor 9 in Papillomavirus Restriction. MBio, 2022, 13, e0005422.	4.1	5
4	Mechanisms of DNA Virus Evolution. , 2021, , 71-78.		6
5	New World Cactaceae Plants Harbor Diverse Geminiviruses. Viruses, 2021, 13, 694.	3.3	8
6	HPV32â€related Heck's disease in a chronic graftâ€versusâ€host disease patient with longâ€ŧerm successful KTP laser treatment: A rare case report. Clinical Case Reports (discontinued), 2021, 9, e04253.	0.5	3
7	Changes to virus taxonomy and to the International Code of Virus Classification and Nomenclature ratified by the International Committee on Taxonomy of Viruses (2021). Archives of Virology, 2021, 166, 2633-2648.	2.1	219
8	Regulation of Human Papillomavirus 18 Genome Replication, Establishment, and Persistence by Sequences in the Viral Upstream Regulatory Region. Journal of Virology, 2021, 95, e0068621.	3.4	7
9	Complex evolutionary history of felid anelloviruses. Virology, 2021, 562, 176-189.	2.4	13
10	A novel lineage of polyomaviruses identified in bark scorpions. Virology, 2021, 563, 58-63.	2.4	9
11	Insertional oncogenesis by HPV70 revealed by multiple genomic analyses in a clinically HPVâ€negative cervical cancer. Genes Chromosomes and Cancer, 2020, 59, 84-95.	2.8	5
12	PuMA: A papillomavirus genome annotation tool. Virus Evolution, 2020, 6, veaa068.	4.9	12
13	Novel Circoviruses Detected in Feces of Sonoran Felids. Viruses, 2020, 12, 1027.	3.3	13
14	A Novel Divergent Geminivirus Identified in Asymptomatic New World Cactaceae Plants. Viruses, 2020, 12, 398.	3.3	10
15	3D Oral and Cervical Tissue Models for Studying Papillomavirus Hostâ€Pathogen Interactions. Current Protocols in Microbiology, 2020, 59, e129.	6.5	16
16	Vesicular trafficking permits evasion of cGAS/STING surveillance during initial human papillomavirus infection. PLoS Pathogens, 2020, 16, e1009028.	4.7	32
17	The Launch of an International Animal Papillomavirus Reference Center. Viruses, 2019, 11, 55.	3.3	10
18	Neisseria gonorrhoeae evades autophagic killing by downregulating CD46-cyt1 and remodeling lysosomes. PLoS Pathogens, 2019, 15, e1007495.	4.7	23

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19	Metagenomic Discovery of 83 New Human Papillomavirus Types in Patients with Immunodeficiency. MSphere, 2018, 3, .	2.9	75
20	ICTV Virus Taxonomy Profile: Papillomaviridae. Journal of General Virology, 2018, 99, 989-990.	2.9	140
21	Building (Viral) Phylogenetic Trees Using a Maximum Likelihood Approach. Current Protocols in Microbiology, 2018, 51, e63.	6.5	5
22	Fish polyomaviruses belong to two distinct evolutionary lineages. Journal of General Virology, 2018, 99, 567-573.	2.9	19
23	The Papillomavirus Episteme: a major update to the papillomavirus sequence database. Nucleic Acids Research, 2017, 45, D499-D506.	14.5	298
24	Persistence of an Oncogenic Papillomavirus Genome Requires <i>cis</i> Elements from the Viral Transcriptional Enhancer. MBio, 2017, 8, .	4.1	13
25	Roles of APOBEC3A and APOBEC3B in Human Papillomavirus Infection and Disease Progression. Viruses, 2017, 9, 233.	3.3	79
26	Unique genome organization of non-mammalian papillomaviruses provides insights into the evolution of viral early proteins. Virus Evolution, 2017, 3, vex027.	4.9	51
27	The Ancient Evolutionary History of Polyomaviruses. PLoS Pathogens, 2016, 12, e1005574.	4.7	190
28	Novel recombinant papillomavirus genomes expressing selectable genes. Scientific Reports, 2016, 6, 37782.	3.3	13
29	Molecular archeological evidence in support of the repeated loss of a papillomavirus gene. Scientific Reports, 2016, 6, 33028.	3.3	36
30	Role of the host restriction factor APOBEC3 on papillomavirus evolution. Virus Evolution, 2015, 1, vev015.	4.9	57
31	Degradation of Human PDZ-Proteins by Human Alphapapillomaviruses Represents an Evolutionary Adaptation to a Novel Cellular Niche. PLoS Pathogens, 2015, 11, e1004980.	4.7	20
32	Evolution of the Papillomaviridae. Virology, 2013, 445, 11-20.	2.4	176
33	The Papillomavirus Episteme: a central resource for papillomavirus sequence data and analysis. Nucleic Acids Research, 2012, 41, D571-D578.	14.5	188
34	Association between hTERT activation by HPV E6 proteins and oncogenic risk. Virology, 2012, 433, 216-219.	2.4	54
35	Papillomaviruses: evolution, Linnaean taxonomy and current nomenclature. Trends in Microbiology, 2011, 19, 49-50.	7.7	35
36	Sequence Imputation of HPV16 Genomes for Genetic Association Studies. PLoS ONE, 2011, 6, e21375.	2.5	70

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37	Classification of papillomaviruses (PVs) based on 189 PV types and proposal of taxonomic amendments. Virology, 2010, 401, 70-79.	2.4	1,377
38	Serological response to an HPV16 E7 based therapeutic vaccine in women with high-grade cervical dysplasia. Gynecologic Oncology, 2010, 116, 208-212.	1.4	30
39	Degradation of p53 by Human Alphapapillomavirus E6 Proteins Shows a Stronger Correlation with Phylogeny than Oncogenicity. PLoS ONE, 2010, 5, e12816.	2.5	53
40	Evolution of Human Papillomavirus Carcinogenicity. Advances in Virus Research, 2010, 77, 41-62.	2.1	29
41	Human Papillomaviruses: Genetic Basis of Carcinogenicity. Public Health Genomics, 2009, 12, 281-290.	1.0	113
42	Identification of Unusual E6 and E7 Proteins within Avian Papillomaviruses: Cellular Localization, Biophysical Characterization, and Phylogenetic Analysis. Journal of Virology, 2009, 83, 8759-8770.	3.4	33
43	Genomic characterization of two novel reptilian papillomaviruses, Chelonia mydas papillomavirus 1 and Caretta caretta papillomavirus 1. Virology, 2009, 383, 131-135.	2.4	67
44	Complete genomic characterization of a murine papillomavirus isolated from papillomatous lesions of a European harvest mouse (Micromys minutus). Journal of General Virology, 2007, 88, 1484-1488.	2.9	31
45	Ancient papillomavirus-host co-speciation in Felidae. Genome Biology, 2007, 8, R57.	9.6	152
46	Genetic characterization of the Capra hircus papillomavirus: A novel close-to-root artiodactyl papillomavirus. Virus Research, 2006, 118, 164-169.	2.2	26
47	Genetic characterization of the first chiropteran papillomavirus, isolated from a basosquamous carcinoma in an Egyptian fruit bat: The Rousettus aegyptiacus papillomavirus type 1. Veterinary Microbiology, 2006, 117, 267-275.	1.9	27
48	Isolation and cloning of a papillomavirus from a North American porcupine by using multiply primed rolling-circle amplification: the Erethizon dorsatum papillomavirus type 1. Virology, 2005, 331, 449-456.	2.4	45
49	Isolation and cloning of the raccoon (Procyon lotor) papillomavirus type 1 by using degenerate papillomavirus-specific primers. Journal of General Virology, 2005, 86, 2029-2033.	2.9	46