

# Lea HoÅ;njak

## List of Publications by Year in descending order

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35  
papers

395  
citations

759233

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h-index

839539

18  
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36  
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36  
docs citations

36  
times ranked

649  
citing authors

#	ARTICLE	IF	CITATIONS
1	Viral Metagenomic Data Analyses of Five New World Bat Species from Argentina: Identification of 35 Novel DNA Viruses. <i>Microorganisms</i> , 2022, 10, 266.	3.6	10
2	Risk factors for the development of high-grade dysplasia and carcinoma in patients with laryngeal squamous cell papillomas: Large retrospective cohort study. <i>Head and Neck</i> , 2021, 43, 956-966.	2.0	4
3	First Report of Phodopus sungorus Papillomavirus Type 1 Infection in Roborovski Hamsters (Phodopus) Tj ETQq1 1 0,784314 rgBT /Ov	3.3	1
4	Molecular and Phylogenetic Characterization of Novel Papillomaviruses Isolated from Oral and Anogenital Neoplasms of Japanese Macaques ( <i>Macaca fuscata</i> ). <i>Viruses</i> , 2021, 13, 630.	3.3	0
5	The genetic diversity of human papillomavirus types from the species Gammapapillomavirus 15: HPV135, HPV146, and HPV179. <i>PLoS ONE</i> , 2021, 16, e0249829.	2.5	0
6	Molecular Characterization of Human Papillomavirus Type 159 (HPV159). <i>Viruses</i> , 2021, 13, 1668.	3.3	1
7	Usefulness of high-risk human papillomavirus mRNA silver in situ hybridization diagnostic assay in oropharyngeal squamous cell carcinomas. <i>Pathology Research and Practice</i> , 2021, 226, 153585.	2.3	2
8	Determination of Causative Human Papillomavirus Type in Tissue Specimens of Common Warts Based on Estimated Viral Loads. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 4.	3.9	12
9	Assessing Gammapapillomavirus infections of mucosal epithelia with two broad-spectrum PCR protocols. <i>BMC Infectious Diseases</i> , 2020, 20, 274.	2.9	8
10	A Preliminary Study of the Virome of the South American Free-Tailed Bats ( <i>Tadarida brasiliensis</i> ) and Identification of Two Novel Mammalian Viruses. <i>Viruses</i> , 2020, 12, 422.	3.3	22
11	Dose-Modifying Factor of Radiation Therapy with Concurrent Cisplatin Treatment in HPV-Positive Squamous Cell Carcinoma: A Preclinical Study. <i>Radiation Research</i> , 2018, 189, 644.	1.5	11
12	New Insights into the Evolutionary and Genomic Landscape of Molluscum Contagiosum Virus (MCV) based on Nine MCV1 and Six MCV2 Complete Genome Sequences. <i>Viruses</i> , 2018, 10, 586.	3.3	13
13	Clinical, Histopathological, and Virological Evaluation of 203 Patients With a Clinical Diagnosis of Molluscum Contagiosum. <i>Open Forum Infectious Diseases</i> , 2018, 5, ofy298.	0.9	12
14	High prevalence of Gammapapillomaviruses (Gamma-PVs) in pre-malignant cutaneous lesions of immunocompetent individuals using a new broad-spectrum primer system, and identification of HPV210, a novel Gamma-PV type. <i>Virology</i> , 2018, 525, 182-191.	2.4	12
15	Detection of novel Betapapillomaviruses and Gammapapillomaviruses in eyebrow hair follicles using a single-tube "hanging droplet"™ PCR assay with modified pan-PV CODEHOP primers. <i>Journal of General Virology</i> , 2018, 99, 109-118.	2.9	10
16	A systematic literature review of studies reporting human papillomavirus (HPV) prevalence in esophageal carcinoma over 36 years (1982-2017). <i>Acta Dermatovenerologica Alpina, Panonica Et Adriatica</i> , 2018, 27, 127-136.	0.1	2
17	Anal verrucous carcinoma is not related to infection with human papillomaviruses and should be distinguished from giant condyloma (Buschke-Lowenstein tumour). <i>Histopathology</i> , 2017, 70, 938-945.	2.9	23
18	The effect of single nucleotide polymorphisms in G-rich regions of high-risk human papillomaviruses on structural diversity of DNA. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2017, 1861, 1229-1236.	2.4	21

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19	Molecular characterization, tissue tropism, and genetic variability of the novel Mupapillomavirus type HPV204 and phylogenetically related types HPV1 and HPV63. <i>PLoS ONE</i> , 2017, 12, e0175892.	2.5	4
20	Natural history of human papillomavirus infection of sun-exposed healthy skin of immunocompetent individuals over three climatic seasons and identification of HPV209, a novel betapapillomavirus. <i>Journal of General Virology</i> , 2017, 98, 1334-1348.	2.9	9
21	Molecular characterization, prevalence and clinical relevance of Phodopus sungorus papillomavirus type 1 (PsuPV1) naturally infecting Siberian hamsters ( <i>Phodopus sungorus</i> ). <i>Journal of General Virology</i> , 2017, 98, 2799-2809.	2.9	2
22	Global Genomic Diversity of Human Papillomavirus 11 Based on 433 Isolates and 78 Complete Genome Sequences. <i>Journal of Virology</i> , 2016, 90, 5503-5513.	3.4	20
23	Detection of alpha human papillomaviruses in archival formalin-fixed, paraffin-embedded (FFPE) tissue specimens. <i>Journal of Clinical Virology</i> , 2016, 76, S88-S97.	3.1	29
24	Development of a novel multiplex type-specific quantitative real-time PCR for detection and differentiation of infections with human papillomavirus types HPV2, HPV27, and HPV57. <i>Acta Dermatovenerologica Alpina, Panonica Et Adriatica</i> , 2016, 25, 65-71.	0.1	5
25	Hepatitis D virus infection in Slovenian patients with chronic hepatitis B virus infection: a national prevalence study and literature review. <i>Acta Dermatovenerologica Alpina, Panonica Et Adriatica</i> , 2016, 25, 49-53.	0.1	3
26	Identification of a Novel Human Papillomavirus, Type HPV199, Isolated from a Nasopharynx and Anal Canal, and Complete Genomic Characterization of Papillomavirus Species Gamma-12. <i>PLoS ONE</i> , 2015, 10, e0138628.	2.5	14
27	Morphological characteristics of conjunctival squamous papillomas in relation to human papillomavirus infection. <i>British Journal of Ophthalmology</i> , 2015, 99, 431-436.	3.9	28
28	Characterization of Two Novel Gammapapillomaviruses, HPV179 and HPV184, Isolated from Common Warts of a Renal-Transplant Recipient. <i>PLoS ONE</i> , 2015, 10, e0119154.	2.5	18
29	Commercially available kits for manual and automatic extraction of nucleic acids from formalin-fixed, paraffin-embedded (FFPE) tissues. <i>Acta Dermatovenerologica Alpina, Panonica Et Adriatica</i> , 2015, 24, 47-53.	0.1	19
30	Genome announcement: complete genome sequence of a novel Mupapillomavirus, HPV204. <i>Acta Dermatovenerologica Alpina, Panonica Et Adriatica</i> , 2015, 24, 21-3.	0.1	5
31	Betapapillomaviruses in the anal canal of HIV positive and HIV negative men who have sex with men. <i>Journal of Clinical Virology</i> , 2014, 61, 237-241.	3.1	16
32	Global Genomic Diversity of Human Papillomavirus 6 Based on 724 Isolates and 190 Complete Genome Sequences. <i>Journal of Virology</i> , 2014, 88, 7307-7316.	3.4	33
33	Role of human papillomaviruses in esophageal carcinoma: an updated systematic review from 1982 to 2013. <i>Future Virology</i> , 2014, 9, 69-86.	1.8	2
34	Rapid detection and typing of Molluscum contagiosum virus by FRET-based real-time PCR. <i>Journal of Virological Methods</i> , 2013, 187, 431-434.	2.1	15
35	Complete Genome Sequence of a Novel Human Betapapillomavirus, HPV-159. <i>Genome Announcements</i> , 2013, 1, .	0.8	8