

# UrÅjka Kuhar

## List of Publications by Year in descending order

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Version: 2024-02-01

27  
papers

214  
citations

1163117

8  
h-index

1125743

13  
g-index

27  
all docs

27  
docs citations

27  
times ranked

360  
citing authors

#	ARTICLE	IF	CITATIONS
1	Detection of Herpesviruses in Wild Bird Casualties in Slovenia. <i>Frontiers in Veterinary Science</i> , 2022, 9, 822212.	2.2	3
2	Serological Survey of Small Ruminant Lentivirus Infections in Free-Ranging Mouflon and Chamois in Slovenia. <i>Animals</i> , 2022, 12, 1032.	2.3	2
3	Passive Disease Surveillance of Alpine Chamois ( <i>Rupicapra r. rupicapra</i> ) in Slovenia between 2000 and 2020. <i>Animals</i> , 2022, 12, 1119.	2.3	3
4	Twenty Years of Passive Disease Surveillance of Roe Deer ( <i>Capreolus capreolus</i> ) in Slovenia. <i>Animals</i> , 2021, 11, 407.	2.3	6
5	Screening of Mosquitoes for West Nile Virus and Usutu Virus in Croatia, 2015–2020. <i>Tropical Medicine and Infectious Disease</i> , 2021, 6, 45.	2.3	9
6	Identification of a Novel Papillomavirus Type (MfoiPV1) Associated with Acrochordon in a Stone Marten ( <i>Martes foina</i> ). <i>Pathogens</i> , 2021, 10, 539.	2.8	1
7	Herpesvirus Infection in a Breeding Population of Two Coexisting Strix Owls. <i>Animals</i> , 2021, 11, 2519.	2.3	2
8	A Comprehensive Study of Cutaneous Fibropapillomatosis in Free-Ranging Roe Deer ( <i>Capreolus</i> ) of Papillomaviruses. <i>Viruses</i> , 2020, 12, 1001.	3.3	3
9	First Complete Genome of Lake Sinai Virus Lineage 3 and Genetic Diversity of Lake Sinai Virus Strains From Honey Bees and Bumble Bees. <i>Journal of Economic Entomology</i> , 2020, 113, 1055-1061.	1.8	13
10	A molecular survey, whole genome sequencing and phylogenetic analysis of astroviruses from roe deer. <i>BMC Veterinary Research</i> , 2020, 16, 68.	1.9	4
11	Genome sequence of an aichivirus detected in a common pipistrelle bat ( <i>Pipistrellus pipistrellus</i> ). <i>Archives of Virology</i> , 2020, 165, 1019-1022.	2.1	5
12	Post-epidemic investigation of Schmallenberg virus in wild ruminants in Slovenia. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 1708-1715.	3.0	5
13	High detection rate and high genetic diversity of genogroup I Picobirnaviruses from roe deer. <i>Infection, Genetics and Evolution</i> , 2019, 73, 210-213.	2.3	4
14	Effect of Composting Under Semipermeable Film on the Sewage Sludge Virome. <i>Microbial Ecology</i> , 2019, 78, 895-903.	2.8	6
15	Nearly Complete Genome Sequences of Two Bovine Viral Diarrhea Virus Isolates, Subtype 1f Strain SLO/1170/2000 and Subtype 1d Strain SLO/2416/2002. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	0
16	Identification of novel reassortant mammalian orthoreoviruses from bats in Slovenia. <i>BMC Veterinary Research</i> , 2018, 14, 264.	1.9	27
17	Complete Genome Sequence of Lumpy Skin Disease Virus Isolate SERBIA/Bujanovac/2016, Detected during an Outbreak in the Balkan Area. <i>Genome Announcements</i> , 2017, 5, .	0.8	21
18	Complete Genome of Chronic Bee Paralysis Virus Strain SLO/M92/2010, Detected from <i>Apis mellifera carnica</i> . <i>Genome Announcements</i> , 2017, 5, .	0.8	6

#	ARTICLE	IF	CITATIONS
19	Whole genome sequence and a phylogenetic analysis of the G8P[14] group A rotavirus strain from roe deer. <i>BMC Veterinary Research</i> , 2017, 13, 353.	1.9	10
20	Complete Genome Sequence of Roe Deer Picobirnavirus Strain PBV/roe_deer/SLO/D38-14/2014. <i>Genome Announcements</i> , 2017, 5, .	0.8	6
21	Detection and Whole-Genome Analysis of a Zoonotic G8P[14] Rotavirus Strain Isolated from a Child with Diarrhea. <i>Genome Announcements</i> , 2017, 5, .	0.8	4
22	Complete Genome Sequence of the Porcine Epidemic Diarrhea Virus Strain SLO/JH-11/2015. <i>Genome Announcements</i> , 2016, 4, .	0.8	13
23	Complete Genome Sequence of a Bovine Viral Diarrhea Virus Subgenotype 1e Strain, SLO/2407/2006, Isolated in Slovenia. <i>Genome Announcements</i> , 2016, 4, .	0.8	4
24	The first detection and whole genome characterization of the G6P[15] group A rotavirus strain from roe deer. <i>Veterinary Microbiology</i> , 2016, 191, 52-59.	1.9	17
25	Development and validation of TaqMan probe based real time PCR assays for the specific detection of genotype A and B small ruminant lentivirus strains. <i>BMC Veterinary Research</i> , 2013, 9, 172.	1.9	20
26	Phylogenetic analysis of small ruminant lentiviruses detected in Slovenia. <i>Veterinary Microbiology</i> , 2013, 162, 201-206.	1.9	17
27	Molecular and genetic characteristics of small ruminant lentiviruses in Slovenia. <i>Acta Veterinaria Hungarica</i> , 2013, 61, 135-146.	0.5	3