

UrÅjka Kuhar

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

214
citations

1163117

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27
all docs

27
docs citations

27
times ranked

360
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of novel reassortant mammalian orthoreoviruses from bats in Slovenia. BMC Veterinary Research, 2018, 14, 264.	1.9	27
2	Complete Genome Sequence of Lumpy Skin Disease Virus Isolate SERBIA/Bujanovac/2016, Detected during an Outbreak in the Balkan Area. Genome Announcements, 2017, 5, .	0.8	21
3	Development and validation of TaqMan probe based real time PCR assays for the specific detection of genotype A and B small ruminant lentivirus strains. BMC Veterinary Research, 2013, 9, 172.	1.9	20
4	Phylogenetic analysis of small ruminant lentiviruses detected in Slovenia. Veterinary Microbiology, 2013, 162, 201-206.	1.9	17
5	The first detection and whole genome characterization of the G6P[15] group A rotavirus strain from roe deer. Veterinary Microbiology, 2016, 191, 52-59.	1.9	17
6	Complete Genome Sequence of the Porcine Epidemic Diarrhea Virus Strain SLO/JH-11/2015. Genome Announcements, 2016, 4, .	0.8	13
7	First Complete Genome of Lake Sinai Virus Lineage 3 and Genetic Diversity of Lake Sinai Virus Strains From Honey Bees and Bumble Bees. Journal of Economic Entomology, 2020, 113, 1055-1061.	1.8	13
8	Whole genome sequence and a phylogenetic analysis of the G8P[14] group A rotavirus strain from roe deer. BMC Veterinary Research, 2017, 13, 353.	1.9	10
9	Screening of Mosquitoes for West Nile Virus and Usutu Virus in Croatia, 2015â€“2020. Tropical Medicine and Infectious Disease, 2021, 6, 45.	2.3	9
10	Complete Genome of Chronic Bee Paralysis Virus Strain SLO/M92/2010, Detected from Apis mellifera carnica. Genome Announcements, 2017, 5, .	0.8	6
11	Complete Genome Sequence of Roe Deer Picobirnavirus Strain PBV/roe_deer/SLO/D38-14/2014. Genome Announcements, 2017, 5, .	0.8	6
12	Effect of Composting Under Semipermeable Film on the Sewage Sludge Virome. Microbial Ecology, 2019, 78, 895-903.	2.8	6
13	Twenty Years of Passive Disease Surveillance of Roe Deer (Capreolus capreolus) in Slovenia. Animals, 2021, 11, 407.	2.3	6
14	Genome sequence of an aichivirus detected in a common pipistrelle bat (Pipistrellus pipistrellus). Archives of Virology, 2020, 165, 1019-1022.	2.1	5
15	Postâ€“epidemic investigation of Schmallenberg virus in wild ruminants in Slovenia. Transboundary and Emerging Diseases, 2020, 67, 1708-1715.	3.0	5
16	Complete Genome Sequence of a Bovine Viral Diarrhea Virus Subgenotype 1e Strain, SLO/2407/2006, Isolated in Slovenia. Genome Announcements, 2016, 4, .	0.8	4
17	Detection and Whole-Genome Analysis of a Zoonotic G8P[14] Rotavirus Strain Isolated from a Child with Diarrhea. Genome Announcements, 2017, 5, .	0.8	4
18	High detection rate and high genetic diversity of genogroup I Picobirnaviruses from roe deer. Infection, Genetics and Evolution, 2019, 73, 210-213.	2.3	4

#	ARTICLE	IF	CITATIONS
19	A molecular survey, whole genome sequencing and phylogenetic analysis of astroviruses from roe deer. BMC Veterinary Research, 2020, 16, 68.	1.9	4
20	Molecular and genetic characteristics of small ruminant lentiviruses in Slovenia. Acta Veterinaria Hungarica, 2013, 61, 135-146.	0.5	3
21	A Comprehensive Study of Cutaneous Fibropapillomatosis in Free-Ranging Roe Deer (<i>Capreolus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock of Papillomaviruses. Viruses, 2020, 12, 1001.	3.3	3
22	Detection of Herpesviruses in Wild Bird Casualties in Slovenia. Frontiers in Veterinary Science, 2022, 9, 822212.	2.2	3
23	Passive Disease Surveillance of Alpine Chamois (<i>Rupicapra r. rupicapra</i>) in Slovenia between 2000 and 2020. Animals, 2022, 12, 1119.	2.3	3
24	Herpesvirus Infection in a Breeding Population of Two Coexisting Strix Owls. Animals, 2021, 11, 2519.	2.3	2
25	Serological Survey of Small Ruminant Lentivirus Infections in Free-Ranging Mouflon and Chamois in Slovenia. Animals, 2022, 12, 1032.	2.3	2
26	Identification of a Novel Papillomavirus Type (MfoiPV1) Associated with Acrochordon in a Stone Marten (<i>Martes foina</i>). Pathogens, 2021, 10, 539.	2.8	1
27	Nearly Complete Genome Sequences of Two Bovine Viral Diarrhea Virus Isolates, Subtype 1f Strain SLO/1170/2000 and Subtype 1d Strain SLO/2416/2002. Microbiology Resource Announcements, 2019, 8, .	0.6	0