

Akos Boros

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

37
papers

761
citations

15
h-index

27
g-index

39
ext. papers

899
ext. citations

4.5
avg, IF

4.07
L-index

#	Paper	IF	Citations
37	A novel parvovirus (family Parvoviridae) in a freshwater fish, zander (<i>Sander lucioperca</i>).. <i>Archives of Virology</i> , 2022 , 167, 1163	2.6	1
36	Human-stool-associated tusavirus (Parvoviridae) in domestic goats and sheep.. <i>Archives of Virology</i> , 2022 , 1	2.6	0
35	Novel picornavirus (family Picornaviridae) from freshwater fishes (<i>Perca fluviatilis</i> , <i>Sander lucioperca</i> , and <i>Ameiurus melas</i>) in Hungary. <i>Archives of Virology</i> , 2021 , 166, 2627-2632	2.6	1
34	Detection and genetic characterization of a novel parvovirus (family Parvoviridae) in barn owls (<i>Tyto alba</i>) in Hungary. <i>Archives of Virology</i> , 2021 , 166, 231-236	2.6	2
33	Multiple Types of Novel Enteric Bopiviruses () with the Possibility of Interspecies Transmission Identified from Cloven-Hoofed Domestic Livestock (Ovine, Caprine and Bovine) in Hungary. <i>Viruses</i> , 2021 , 13,	6.2	3
32	Injury-Induced Innate Immune Response During Segment Regeneration of the Earthworm,. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	5
31	Genome characterization, prevalence and tissue distribution of astrovirus, hepevirus and norovirus among wild and laboratory rats (<i>Rattus norvegicus</i>) and mice (<i>Mus musculus</i>) in Hungary. <i>Infection, Genetics and Evolution</i> , 2021 , 93, 104942	4.5	0
30	Detection and complete genome characterization of a novel RNA virus related to members of the Hepe-Virga clade in bird species, hoopoe (<i>Upupa epops</i>). <i>Infection, Genetics and Evolution</i> , 2020 , 81, 104236	4.5	1
29	Four-year long (2014-2017) clinical and laboratory surveillance of hepatitis E virus infections using combined antibody, molecular, antigen and avidity detection methods: Increasing incidence and chronic HEV case in Hungary. <i>Journal of Clinical Virology</i> , 2020 , 124, 104284	14.5	6
28	High prevalence, genetic diversity and a potentially novel genotype of Sapelovirus A () in enteric and respiratory samples in Hungarian swine farms. <i>Journal of General Virology</i> , 2020 , 101, 609-621	4.9	5
27	Review of Hepatitis E Virus in Rats: Evident Risk of Species to Human Zoonotic Infection and Disease. <i>Viruses</i> , 2020 , 12,	6.2	13
26	Characterization of an integrated, endogenous mouse mammary tumor virus-like (MMTV) betaretrovirus genome in a black Syrian hamster (<i>Mesocricetus auratus</i>). <i>Infection, Genetics and Evolution</i> , 2019 , 75, 103995	4.5	0
25	Genetically highly divergent RNA virus with astrovirus-like (5'end) and hepevirus-like (3'end) genome organization in carnivorous birds, European roller (<i>Coracias garrulus</i>). <i>Infection, Genetics and Evolution</i> , 2019 , 71, 215-223	4.5	1
24	Diverse picornaviruses are prevalent among free-living and laboratory rats (<i>Rattus norvegicus</i>) in Hungary and can cause disseminated infections. <i>Infection, Genetics and Evolution</i> , 2019 , 75, 103988	4.5	5
23	Analysis of a novel RNA virus in a wild northern white-breasted hedgehog (<i>Erinaceus roumanicus</i>). <i>Archives of Virology</i> , 2019 , 164, 3065-3071	2.6	0
22	Identification of novel lumbricin homologues in <i>Eisenia andrei</i> earthworms. <i>Developmental and Comparative Immunology</i> , 2019 , 90, 41-46	3.2	10
21	Detection of a novel RNA virus with hepatitis E virus-like non-structural genome organization in amphibian, agile frog (<i>Rana dalmatina</i>) tadpoles. <i>Infection, Genetics and Evolution</i> , 2018 , 65, 112-116	4.5	9

20	Multiple divergent picobirnaviruses with functional prokaryotic Shine-Dalgarno ribosome binding sites present in cloacal sample of a diarrheic chicken. <i>Virology</i> , 2018 , 525, 62-72	3.6	15
19	Genome characterization of a novel megrivirus-related avian picornavirus from a carnivorous wild bird, western marsh harrier (<i>Circus aeruginosus</i>). <i>Archives of Virology</i> , 2017 , 162, 2781-2789	2.6	3
18	Ljungan/Sebokele-like picornavirus in birds of prey, common kestrel (<i>Falco tinnunculus</i>) and red-footed falcon (<i>F. vespertinus</i>). <i>Infection, Genetics and Evolution</i> , 2017 , 55, 14-19	4.5	5
17	A novel avian-like hepatitis E virus in wild aquatic bird, little egret (<i>Egretta garzetta</i>), in Hungary. <i>Infection, Genetics and Evolution</i> , 2016 , 46, 74-77	4.5	17
16	Divergent hepatitis E virus in birds of prey, common kestrel (<i>Falco tinnunculus</i>) and red-footed falcon (<i>F. vespertinus</i>), Hungary. <i>Infection, Genetics and Evolution</i> , 2016 , 43, 343-6	4.5	42
15	A diarrheic chicken simultaneously co-infected with multiple picornaviruses: Complete genome analysis of avian picornaviruses representing up to six genera. <i>Virology</i> , 2016 , 489, 63-74	3.6	27
14	Detection and genome analysis of a novel (dima)rhabdovirus (Riverside virus) from <i>Ochlerotatus</i> sp. mosquitoes in Central Europe. <i>Infection, Genetics and Evolution</i> , 2016 , 39, 336-341	4.5	11
13	A novel posavirus-related single-stranded RNA virus from fish (<i>Cyprinus carpio</i>). <i>Archives of Virology</i> , 2015 , 160, 565-8	2.6	23
12	Avian picornaviruses: molecular evolution, genome diversity and unusual genome features of a rapidly expanding group of viruses in birds. <i>Infection, Genetics and Evolution</i> , 2014 , 28, 151-66	4.5	31
11	Comparative complete genome analysis of chicken and Turkey megriviruses (family picornaviridae): long 3' untranslated regions with a potential second open reading frame and evidence for possible recombination. <i>Journal of Virology</i> , 2014 , 88, 6434-43	6.6	21
10	Novel dicistrovirus from bat guano. <i>Archives of Virology</i> , 2014 , 159, 3453-6	2.6	18
9	Genome characterization of a novel chicken picornavirus distantly related to the members of genus Avihepatovirus with a single 2A protein and a megrivirus-like 3'UTR. <i>Infection, Genetics and Evolution</i> , 2014 , 28, 333-8	4.5	6
8	Genetic characterization of a novel picornavirus in turkeys (<i>Meleagris gallopavo</i>) distinct from turkey galliviruses and megriviruses and distantly related to the members of the genus Avihepatovirus. <i>Journal of General Virology</i> , 2013 , 94, 1496-1509	4.9	30
7	The viruses of wild pigeon droppings. <i>PLoS ONE</i> , 2013 , 8, e72787	3.7	85
6	Identification and complete genome characterization of a novel picornavirus in turkey (<i>Meleagris gallopavo</i>). <i>Journal of General Virology</i> , 2012 , 93, 2171-2182	4.9	45
5	Natural interspecies recombinant bovine/porcine enterovirus in sheep. <i>Journal of General Virology</i> , 2012 , 93, 1941-1951	4.9	57
4	Human enterovirus 109 (EV109) in acute paediatric respiratory disease in Hungary. <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2012 , 59, 285-90	1.8	13
3	Novel positive-sense, single-stranded RNA (+ssRNA) virus with di-cistronic genome from intestinal content of freshwater carp (<i>Cyprinus carpio</i>). <i>PLoS ONE</i> , 2011 , 6, e29145	3.7	47

- 2 Kobuviruses - a comprehensive review. *Reviews in Medical Virology*, **2011**, 21, 32-41 11.7 134
- 1 Kobuvirus in domestic sheep, Hungary. *Emerging Infectious Diseases*, **2010**, 16, 869-70 10.2 66