## **Akos Boros**

## List of Publications by Citations

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37 761 15 27 g-index

39 899 4.5 avg, IF L-index

#	Paper	IF	Citations
37	Kobuviruses - a comprehensive review. <i>Reviews in Medical Virology</i> , <b>2011</b> , 21, 32-41	11.7	134
36	The viruses of wild pigeon droppings. <i>PLoS ONE</i> , <b>2013</b> , 8, e72787	3.7	85
35	Kobuvirus in domestic sheep, Hungary. <i>Emerging Infectious Diseases</i> , <b>2010</b> , 16, 869-70	10.2	66
34	Natural interspecies recombinant bovine/porcine enterovirus in sheep. <i>Journal of General Virology</i> , <b>2012</b> , 93, 1941-1951	4.9	57
33	Novel positive-sense, single-stranded RNA (+ssRNA) virus with di-cistronic genome from intestinal content of freshwater carp (Cyprinus carpio). <i>PLoS ONE</i> , <b>2011</b> , 6, e29145	3.7	47
32	Identification and complete genome characterization of a novel picornavirus in turkey (Meleagris gallopavo). <i>Journal of General Virology</i> , <b>2012</b> , 93, 2171-2182	4.9	45
31	Divergent hepatitis E virus in birds of prey, common kestrel (Falco tinnunculus) and red-footed falcon (F. vespertinus), Hungary. <i>Infection, Genetics and Evolution</i> , <b>2016</b> , 43, 343-6	4.5	42
30	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> , <b>2014</b> , 28, 151-66	4.5	31
29	Genetic characterization of a novel picornavirus in turkeys (Meleagris gallopavo) distinct from turkey galliviruses and megriviruses and distantly related to the members of the genus Avihepatovirus. <i>Journal of General Virology</i> , <b>2013</b> , 94, 1496-1509	4.9	30
28	A diarrheic chicken simultaneously co-infected with multiple picornaviruses: Complete genome analysis of avian picornaviruses representing up to six genera. <i>Virology</i> , <b>2016</b> , 489, 63-74	3.6	27
27	A novel posavirus-related single-stranded RNA virus from fish (Cyprinus carpio). <i>Archives of Virology</i> , <b>2015</b> , 160, 565-8	2.6	23
26	Comparative complete genome analysis of chicken and Turkey megriviruses (family picornaviridae): long 3Umtranslated regions with a potential second open reading frame and evidence for possible recombination. <i>Journal of Virology</i> , <b>2014</b> , 88, 6434-43	6.6	21
25	Novel dicistrovirus from bat guano. <i>Archives of Virology</i> , <b>2014</b> , 159, 3453-6	2.6	18
24	A novel avian-like hepatitis E virus in wild aquatic bird, little egret (Egretta garzetta), in Hungary. <i>Infection, Genetics and Evolution</i> , <b>2016</b> , 46, 74-77	4.5	17
23	Multiple divergent picobirnaviruses with functional prokaryotic Shine-Dalgarno ribosome binding sites present in cloacal sample of a diarrheic chicken. <i>Virology</i> , <b>2018</b> , 525, 62-72	3.6	15
22	Human enterovirus 109 (EV109) in acute paediatric respiratory disease in Hungary. <i>Acta Microbiologica Et Immunologica Hungarica</i> , <b>2012</b> , 59, 285-90	1.8	13
21	Review of Hepatitis E Virus in Rats: Evident Risk of Species to Human Zoonotic Infection and Disease. <i>Viruses</i> , <b>2020</b> , 12,	6.2	13

20	Detection and genome analysis of a novel (dima)rhabdovirus (Riverside virus) from Ochlerotatus sp. mosquitoes in Central Europe. <i>Infection, Genetics and Evolution</i> , <b>2016</b> , 39, 336-341	4.5	11
19	Identification of novel lumbricin homologues in Eisenia andrei earthworms. <i>Developmental and Comparative Immunology</i> , <b>2019</b> , 90, 41-46	3.2	10
18	Detection of a novel RNA virus with hepatitis E virus-like non-structural genome organization in amphibian, agile frog (Rana dalmatina) tadpoles. <i>Infection, Genetics and Evolution</i> , <b>2018</b> , 65, 112-116	4.5	9
17	Genome characterization of a novel chicken picornavirus distantly related to the members of genus Avihepatovirus with a single 2A protein and a megrivirus-like 3WTR. <i>Infection, Genetics and Evolution</i> , <b>2014</b> , 28, 333-8	4.5	6
16	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> , <b>2020</b> , 124, 104284	14.5	6
15	Diverse picornaviruses are prevalent among free-living and laboratory rats (Rattus norvegicus) in Hungary and can cause disseminated infections. <i>Infection, Genetics and Evolution</i> , <b>2019</b> , 75, 103988	4.5	5
14	Ljungan/Sebokele-like picornavirus in birds of prey, common kestrel (Falco tinnunculus) and red-footed falcon (F. vespertinus). <i>Infection, Genetics and Evolution</i> , <b>2017</b> , 55, 14-19	4.5	5
13	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> , <b>2020</b> , 101, 609-621	4.9	5
12	Injury-Induced Innate Immune Response During Segment Regeneration of the Earthworm,. <i>International Journal of Molecular Sciences</i> , <b>2021</b> , 22,	6.3	5
11	Genome characterization of a novel megrivirus-related avian picornavirus from a carnivorous wild bird, western marsh harrier (Circus aeruginosus). <i>Archives of Virology</i> , <b>2017</b> , 162, 2781-2789	2.6	3
10	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> , <b>2021</b> , 13,	6.2	3
9	Detection and genetic characterization of a novel parvovirus (family Parvoviridae) in barn owls (Tyto alba) in Hungary. <i>Archives of Virology</i> , <b>2021</b> , 166, 231-236	2.6	2
8	Genetically highly divergent RNA virus with astrovirus-like (5 end) and hepevirus-like (3 end) genome organization in carnivorous birds, European roller (Coracias garrulus). <i>Infection, Genetics and Evolution</i> , 2019, 71, 215-223	4.5	1
7	Detection and complete genome characterization of a novel RNA virus related to members of the Hepe-Virga clade in bird species, hoopoe (Upupa epops). <i>Infection, Genetics and Evolution</i> , <b>2020</b> , 81, 104	12 <sup>1</sup> 38	1
6	Novel picornavirus (family Picornaviridae) from freshwater fishes (Perca fluviatilis, Sander lucioperca, and Ameiurus melas) in Hungary. <i>Archives of Virology</i> , <b>2021</b> , 166, 2627-2632	2.6	1
5	A novel parvovirus (family Parvoviridae) in a freshwater fish, zander (Sander lucioperca) <i>Archives of Virology</i> , <b>2022</b> , 167, 1163	2.6	1
4	Characterization of an integrated, endogenous mouse mammary tumor virus-like (MMTV) betaretrovirus genome in a black Syrian hamster (Mesocricetus auratus). <i>Infection, Genetics and Evolution</i> , <b>2019</b> , 75, 103995	4.5	0
3	Analysis of a novel RNA virus in a wild northern white-breasted hedgehog (Erinaceus roumanicus). <i>Archives of Virology</i> , <b>2019</b> , 164, 3065-3071	2.6	O

Genome characterization, prevalence and tissue distribution of astrovirus, hepevirus and norovirus among wild and laboratory rats (Rattus norvegicus) and mice (Mus musculus) in Hungary. *Infection, Genetics and Evolution,* **2021**, 93, 104942

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Human-stool-associated tusavirus (Parvoviridae) in domestic goats and sheep.. *Archives of Virology*, **2022**, 1

2.6