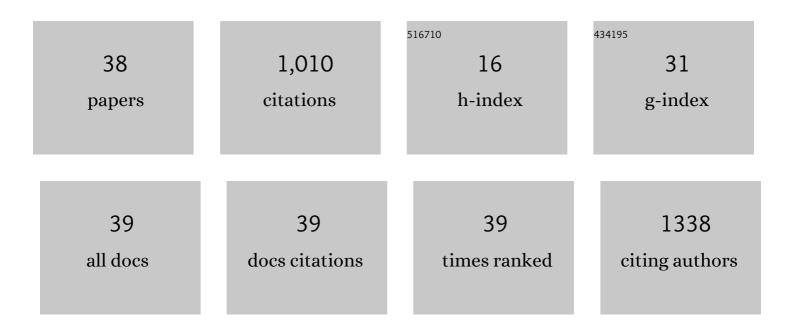
Akos Boros

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A novel parvovirus (family Parvoviridae) in a freshwater fish, zander (Sander lucioperca). Archives of Virology, 2022, 167, 1163-1167.	2.1	4
2	Development and Large-Scale Testing of a Novel One-Step Triplex RT-qPCR Assay for Simultaneous Detection of "Neurotropic―Porcine Sapeloviruses, Teschoviruses (Picornaviridae) and Type 3 Porcine Astroviruses (Astroviridae) in Various Samples including Nasal Swabs. Viruses, 2022, 14, 513.	3.3	0
3	Human-stool-associated tusavirus (Parvoviridae) in domestic goats and sheep. Archives of Virology, 2022, 167, 1307-1310.	2.1	5

Detection and genetic characterization of a novel parvovirus (family Parvoviridae) in barn owls (Tyto) Tj ETQq0 0 0 rgBT /OverJock 10 Tf

4		2.1	
5	Multiple Types of Novel Enteric Bopiviruses (Picornaviridae) with the Possibility of Interspecies Transmission Identified from Cloven-Hoofed Domestic Livestock (Ovine, Caprine and Bovine) in Hungary. Viruses, 2021, 13, 66.	3.3	9
6	Injury-Induced Innate Immune Response During Segment Regeneration of the Earthworm, Eisenia andrei. International Journal of Molecular Sciences, 2021, 22, 2363.	4.1	19
7	Novel picornavirus (family Picornaviridae) from freshwater fishes (Perca fluviatilis, Sander) Tj ETQq1 1 0.784314	rgBT /Ovei 2.1	rloçk 10 Tf S
8	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.	2.3	2
9	Review of Hepatitis E Virus in Rats: Evident Risk of Species Orthohepevirus C to Human Zoonotic Infection and Disease. Viruses, 2020, 12, 1148.	3.3	44
10	Detection and complete genome characterization of a novel RNA virus related to members of the Hepe-Virga clade in bird species, hoopoe (Upupa epops). Infection, Genetics and Evolution, 2020, 81, 104236.	2.3	2
11	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. Journal of Clinical Virology, 2020, 124, 104284.	3.1	11
12	High prevalence, genetic diversity and a potentially novel genotype of Sapelovirus A (Picornaviridae) in enteric and respiratory samples in Hungarian swine farms. Journal of General Virology, 2020, 101, 609-621.	2.9	13
13	Diverse picornaviruses are prevalent among free-living and laboratory rats (Rattus norvegicus) in Hungary and can cause disseminated infections. Infection, Genetics and Evolution, 2019, 75, 103988.	2.3	6
14	Analysis of a novel RNA virus in a wild northern white-breasted hedgehog (Erinaceus roumanicus). Archives of Virology, 2019, 164, 3065-3071.	2.1	3
15	Characterization of an integrated, endogenous mouse mammary tumor virus-like (MMTV) betaretrovirus genome in a black Syrian hamster (Mesocricetus auratus). Infection, Genetics and Evolution, 2019, 75, 103995.	2.3	1
16	Genetically highly divergent RNA virus with astrovirus-like (5′-end) and hepevirus-like (3′-end) genome organization in carnivorous birds, European roller (Coracias garrulus). Infection, Genetics and Evolution, 2019, 71, 215-223.	2.3	1
17	Identification of novel lumbricin homologues in Eisenia andrei earthworms. Developmental and Comparative Immunology, 2019, 90, 41-46.	2.3	16
18	Multiple divergent picobirnaviruses with functional prokaryotic Shine-Dalgarno ribosome binding sites present in cloacal sample of a diarrheic chicken. Virology, 2018, 525, 62-72.	2.4	26

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19	Detection of a novel RNA virus with hepatitis E virus-like non-structural genome organization in amphibian, agile frog (Rana dalmatina) tadpoles. Infection, Genetics and Evolution, 2018, 65, 112-116.	2.3	12
20	Genome characterization of a novel megrivirus-related avian picornavirus from a carnivorous wild bird, western marsh harrier (Circus aeruginosus). Archives of Virology, 2017, 162, 2781-2789.	2.1	7
21	Ljungan/Sebokele-like picornavirus in birds of prey, common kestrel (Falco tinnunculus) and red-footed falcon (F. vespertinus). Infection, Genetics and Evolution, 2017, 55, 14-19.	2.3	10
22	A novel avian-like hepatitis E virus in wild aquatic bird, little egret (Egretta garzetta), in Hungary. Infection, Genetics and Evolution, 2016, 46, 74-77.	2.3	21
23	Divergent hepatitis E virus in birds of prey, common kestrel (Falco tinnunculus) and red-footed falcon (F. vespertinus), Hungary. Infection, Genetics and Evolution, 2016, 43, 343-346.	2.3	52
24	A diarrheic chicken simultaneously co-infected with multiple picornaviruses: Complete genome analysis of avian picornaviruses representing up to six genera. Virology, 2016, 489, 63-74.	2.4	36
25	Detection and genome analysis of a novel (dima)rhabdovirus (Riverside virus) from Ochlerotatus sp. mosquitoes in Central Europe. Infection, Genetics and Evolution, 2016, 39, 336-341.	2.3	17
26	A novel posavirus-related single-stranded RNA virus from fish (Cyprinus carpio). Archives of Virology, 2015, 160, 565-568.	2.1	28
27	Novel dicistrovirus from bat guano. Archives of Virology, 2014, 159, 3453-3456.	2.1	22
28	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. Infection, Genetics and Evolution, 2014, 28, 333-338.	2.3	6
29	Avian picornaviruses: Molecular evolution, genome diversity and unusual genome features of a rapidly expanding group of viruses in birds. Infection, Genetics and Evolution, 2014, 28, 151-166.	2.3	37
30	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. Journal of Virology, 2014, 88, 6434-6443.	3.4	26
31	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. Journal of General Virology, 2013, 94, 1496-1509.	2.9	34
32	The Viruses of Wild Pigeon Droppings. PLoS ONE, 2013, 8, e72787.	2.5	108
33	Natural interspecies recombinant bovine/porcine enterovirus in sheep. Journal of General Virology, 2012, 93, 1941-1951.	2.9	69
34	Human enterovirus 109 (EV109) in acute paediatric respiratory disease in Hungary. Acta Microbiologica Et Immunologica Hungarica, 2012, 59, 285-290.	0.8	15
35	Identification and complete genome characterization of a novel picornavirus in turkey (Meleagris) Tj ETQq1 1	0.784314 rg 2.9	gBT_/Overlock
36	Novel Positive-Sense, Single-Stranded RNA (+ssRNA) Virus with Di-Cistronic Genome from Intestinal Content of Freshwater Carp (Cyprinus carpio). PLoS ONE, 2011, 6, e29145.	2,5	53

#	Article	IF	CITATIONS
37	Kobuviruses – a comprehensive review. Reviews in Medical Virology, 2011, 21, 32-41.	8.3	155
38	Kobuvirus in Domestic Sheep, Hungary. Emerging Infectious Diseases, 2010, 16, 869-870.	4.3	72