

Akos Boros

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

1,010
citations

516710

16
h-index

434195

31
g-index

39
all docs

39
docs citations

39
times ranked

1338
citing authors

#	ARTICLE	IF	CITATIONS
1	Kobuviruses – a comprehensive review. <i>Reviews in Medical Virology</i> , 2011, 21, 32-41.	8.3	155
2	The Viruses of Wild Pigeon Droppings. <i>PLoS ONE</i> , 2013, 8, e72787.	2.5	108
3	Kobuvirus in Domestic Sheep, Hungary. <i>Emerging Infectious Diseases</i> , 2010, 16, 869-870.	4.3	72
4	Natural interspecies recombinant bovine/porcine enterovirus in sheep. <i>Journal of General Virology</i> , 2012, 93, 1941-1951.	2.9	69
5	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.	2.5	53
6	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, 343-346.	2.3	52
7	Identification and complete genome characterization of a novel picornavirus in turkey (<i>Meleagris</i>) Tj ETQq1 1 0.784314 rgBT /Overl... 2.9 51	2.9	51
8	Review of Hepatitis E Virus in Rats: Evident Risk of Species Orthohepevirus C to Human Zoonotic Infection and Disease. <i>Viruses</i> , 2020, 12, 1148.	3.3	44
9	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, 151-166.	2.3	37
10	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.	2.4	36
11	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 <i>Avihepatovirus</i> . <i>Journal of General Virology</i> , 2013, 94, 1496-1509.	2.9	34
12	A novel posavirus-related single-stranded RNA virus from fish (<i>Cyprinus carpio</i>). <i>Archives of Virology</i> , 2015, 160, 565-568.	2.1	28
13	Comparative Complete Genome Analysis of Chicken and Turkey Megriviruses (Family Picornaviridae): Long 3' UTR Untranslated Regions with a Potential Second Open Reading Frame and Evidence for Possible Recombination. <i>Journal of Virology</i> , 2014, 88, 6434-6443.	3.4	26
14	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.	2.4	26
15	Novel dicistrovirus from bat guano. <i>Archives of Virology</i> , 2014, 159, 3453-3456.	2.1	22
16	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.	2.3	21
17	Injury-Induced Innate Immune Response During Segment Regeneration of the Earthworm, <i>Eisenia andrei</i> . <i>International Journal of Molecular Sciences</i> , 2021, 22, 2363.	4.1	19
18	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.	2.3	17

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19	Identification of novel lumbricin homologues in <i>Eisenia andrei</i> earthworms. <i>Developmental and Comparative Immunology</i> , 2019, 90, 41-46.	2.3	16
20	Human enterovirus 109 (EV109) in acute paediatric respiratory disease in Hungary. <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2012, 59, 285-290.	0.8	15
21	High prevalence, genetic diversity and a potentially novel genotype of Sapelovirus A (Picornaviridae) in enteric and respiratory samples in Hungarian swine farms. <i>Journal of General Virology</i> , 2020, 101, 609-621.	2.9	13
22	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.	2.3	12
23	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.	3.1	11
24	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.	2.3	10
25	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. <i>Viruses</i> , 2021, 13, 66.	3.3	9
26	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.	2.1	7
27	Detection and genetic characterization of a novel parvovirus (family Parvoviridae) in barn owls (<i>Tyto</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 302 Td	2.1	7
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. <i>Infection, Genetics and Evolution</i> , 2014, 28, 333-338.	2.3	6
29	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.	2.3	6
30	Novel picornavirus (family Picornaviridae) from freshwater fishes (<i>Perca fluviatilis</i> , Sander) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 302 Td	2.1	6
31	Human-stool-associated tusavirus (Parvoviridae) in domestic goats and sheep. <i>Archives of Virology</i> , 2022, 167, 1307-1310.	2.1	5
32	A novel parvovirus (family Parvoviridae) in a freshwater fish, zander (<i>Sander lucioperca</i>). <i>Archives of Virology</i> , 2022, 167, 1163-1167.	2.1	4
33	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.	2.1	3
34	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.	2.3	2
35	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.	2.3	2
36	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.	2.3	1

#	ARTICLE	IF	CITATIONS
37	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.	2.3	1
38	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. <i>Viruses</i> , 2022, 14, 513.	3.3	0