

Gábor Reuter

List of Publications by Year in descending order

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113
papers

5,809
citations

94269

37
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79541

73
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128
all docs

128
docs citations

128
times ranked

4707
citing authors

#	ARTICLE	IF	CITATIONS
1	Norovirus Illness Is a Global Problem: Emergence and Spread of Norovirus GII.4 Variants, 2001–2007. <i>Journal of Infectious Diseases</i> , 2009, 200, 802-812.	1.9	596
2	Increase in viral gastroenteritis outbreaks in Europe and epidemic spread of new norovirus variant. <i>Lancet</i> , The, 2004, 363, 682-688.	6.3	458
3	ICTV Virus Taxonomy Profile: Picornaviridae. <i>Journal of General Virology</i> , 2017, 98, 2421-2422.	1.3	374
4	Update: proposed reference sequences for subtypes of hepatitis E virus (species Orthohepevirus A). <i>Journal of General Virology</i> , 2020, 101, 692-698.	1.3	221
5	Emergence of a novel GII.17 norovirus – End of the GII.4 era?. <i>Eurosurveillance</i> , 2015, 20, .	3.9	204
6	Analysis of Integrated Virological and Epidemiological Reports of Norovirus Outbreaks Collected within the Foodborne Viruses in Europe Network from 1 July 2001 to 30 June 2006. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2959-2965.	1.8	193
7	Molecular surveillance of norovirus, 2005–16: an epidemiological analysis of data collected from the NoroNet network. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 545-553.	4.6	193
8	Kobuviruses – a comprehensive review. <i>Reviews in Medical Virology</i> , 2011, 21, 32-41.	3.9	155
9	Serologic Assays Specific to Immunoglobulin M Antibodies against Hepatitis E Virus: Pangenotypic Evaluation of Performances. <i>Clinical Infectious Diseases</i> , 2010, 51, e24-e27.	2.9	152
10	Complete nucleotide and amino acid sequences and genetic organization of porcine kobuvirus, a member of a new species in the genus Kobuvirus, family Picornaviridae. <i>Archives of Virology</i> , 2009, 154, 101-108.	0.9	134
11	Characterization and zoonotic potential of endemic hepatitis E virus (HEV) strains in humans and animals in Hungary. <i>Journal of Clinical Virology</i> , 2009, 44, 277-281.	1.6	118
12	Candidate New Species of Kobuvirus in Porcine Hosts. <i>Emerging Infectious Diseases</i> , 2008, 14, 1968-1970.	2.0	115
13	The Viruses of Wild Pigeon Droppings. <i>PLoS ONE</i> , 2013, 8, e72787.	1.1	108
14	Emergence of New Norovirus Variants on Spring Cruise Ships and Prediction of Winter Epidemics. <i>Emerging Infectious Diseases</i> , 2008, 14, 238-243.	2.0	102
15	Recommendations for the nomenclature of enteroviruses and rhinoviruses. <i>Archives of Virology</i> , 2020, 165, 793-797.	0.9	93
16	Use of Norovirus Genotype Profiles to Differentiate Origins of Foodborne Outbreaks. <i>Emerging Infectious Diseases</i> , 2010, 16, 617-624.	2.0	87
17	Feline fecal virome reveals novel and prevalent enteric viruses. <i>Veterinary Microbiology</i> , 2014, 171, 102-111.	0.8	83
18	Sequence heterogeneity among human picobirnaviruses detected in a gastroenteritis outbreak. <i>Archives of Virology</i> , 2003, 148, 2281-2291.	0.9	76

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19	Detection of Hepatitis E virus in samples of animal origin collected in Hungary. <i>Veterinary Microbiology</i> , 2010, 143, 106-116.	0.8	75
20	Evidence of the etiological predominance of norovirus in gastroenteritis outbreaks – emerging new-variant and recombinant noroviruses in Hungary. <i>Journal of Medical Virology</i> , 2005, 76, 598-607.	2.5	73
21	Kobuvirus in Domestic Sheep, Hungary. <i>Emerging Infectious Diseases</i> , 2010, 16, 869-870.	2.0	72
22	Natural interspecies recombinant bovine/porcine enterovirus in sheep. <i>Journal of General Virology</i> , 2012, 93, 1941-1951.	1.3	69
23	Novel circular single-stranded DNA virus from turkey faeces. <i>Archives of Virology</i> , 2014, 159, 2161-2164.	0.9	67
24	Outbreaks of Neuroinvasive Astrovirus Associated with Encephalomyelitis, Weakness, and Paralysis among Weaned Pigs, Hungary. <i>Emerging Infectious Diseases</i> , 2017, 23, 1982-1993.	2.0	66
25	Reovirus identified as cause of disease in young geese. <i>Avian Pathology</i> , 2003, 32, 129-138.	0.8	65
26	Identification of a novel astrovirus in a domestic pig in Hungary. <i>Archives of Virology</i> , 2011, 156, 125-128.	0.9	64
27	Detection of Aichi virus shedding in a child with enteric and extraintestinal symptoms in Hungary. <i>Archives of Virology</i> , 2009, 154, 1529-1532.	0.9	55
28	Incidence, Diversity, and Molecular Epidemiology of Sapoviruses in Swine across Europe. <i>Journal of Clinical Microbiology</i> , 2010, 48, 363-368.	1.8	55
29	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.	1.1	53
30	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.	1.0	52
31	Epidemic spread of recombinant noroviruses with four capsid types in Hungary. <i>Journal of Clinical Virology</i> , 2006, 35, 84-88.	1.6	51
32	Data quality of 5 years of central norovirus outbreak reporting in the European Network for food-borne viruses. <i>Journal of Public Health</i> , 2008, 30, 82-90.	1.0	51
33	Identification and complete genome characterization of a novel picornavirus in turkey (<i>Meleagris</i>) Tj ETQq1 1 0.784314 rgBT /Overl... 1.3 51	1.3	51
34	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.	1.5	44
35	Enteric caliciviruses in domestic pigs in Hungary. <i>Archives of Virology</i> , 2007, 152, 611-614.	0.9	43
36	Bovine Kobuvirus in Europe. <i>Emerging Infectious Diseases</i> , 2009, 15, 822-823.	2.0	42

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37	Evolution of Porcine Kobuvirus Infection, Hungary. <i>Emerging Infectious Diseases</i> , 2010, 16, 696-698.	2.0	42
38	Nonsuppurative (Aseptic) Meningoencephalomyelitis Associated with Neurovirulent Astrovirus Infections in Humans and Animals. <i>Clinical Microbiology Reviews</i> , 2018, 31, .	5.7	39
39	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.	1.0	37
40	Two Closely Related Novel Picornaviruses in Cattle and Sheep in Hungary from 2008 to 2009, Proposed as Members of a New Genus in the Family Picornaviridae. <i>Journal of Virology</i> , 2012, 86, 13295-13302.	1.5	36
41	Non-primate hepacivirus infection with apparent hepatitis in a horse – Short communication. <i>Acta Veterinaria Hungarica</i> , 2014, 62, 422-427.	0.2	36
42	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.	1.1	36
43	Porcine teschovirus in wild boars in Hungary. <i>Archives of Virology</i> , 2012, 157, 1573-1578.	0.9	34
44	Astrovirus in wild boars (<i>Sus scrofa</i>) in Hungary. <i>Archives of Virology</i> , 2012, 157, 1143-1147.	0.9	34
45	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.	1.3	34
46	Characterization of a novel porcine enterovirus in domestic pig in Hungary. <i>Infection, Genetics and Evolution</i> , 2011, 11, 1096-1102.	1.0	33
47	Identification of a novel astrovirus in domestic sheep in Hungary. <i>Archives of Virology</i> , 2012, 157, 323-327.	0.9	33
48	Novel picornavirus in domesticated common quail (<i>Coturnix coturnix</i>) in Hungary. <i>Archives of Virology</i> , 2012, 157, 525-530.	0.9	30
49	Porcine kobuvirus in wild boars (<i>Sus scrofa</i>). <i>Archives of Virology</i> , 2013, 158, 281-282.	0.9	30
50	A Highly Divergent Picornavirus Infecting the Gut Epithelia of Zebrafish (<i>Danio rerio</i>) in Research Institutions Worldwide. <i>Zebrafish</i> , 2019, 16, 291-299.	0.5	30
51	Molecular epidemiology of human calicivirus gastroenteritis outbreaks in Hungary, 1998 to 2000. <i>Journal of Medical Virology</i> , 2002, 68, 390-398.	2.5	28
52	Characterization of a novel porcine enterovirus in wild boars in Hungary. <i>Archives of Virology</i> , 2012, 157, 981-986.	0.9	28
53	Novel seadornavirus (family Reoviridae) related to Banna virus in Europe. <i>Archives of Virology</i> , 2013, 158, 2163-2167.	0.9	28
54	A novel posavirus-related single-stranded RNA virus from fish (<i>Cyprinus carpio</i>). <i>Archives of Virology</i> , 2015, 160, 565-568.	0.9	28

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55	Detection of a mammalian-like astrovirus in bird, European roller (<i>Coracias garrulus</i>). <i>Infection, Genetics and Evolution</i> , 2015, 34, 114-121.	1.0	27
56	Detection and genetic characterization of a novel parvovirus distantly related to human bufavirus in domestic pigs. <i>Archives of Virology</i> , 2016, 161, 1033-1037.	0.9	27
57	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.	1.5	26
58	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.	1.1	26
59	Identification and complete genome analysis of kobuvirus in faecal samples of European roller (<i>Coracias garrulus</i>): for the first time in a bird. <i>Archives of Virology</i> , 2015, 160, 345-351.	0.9	23
60	Novel dicistrovirus from bat guano. <i>Archives of Virology</i> , 2014, 159, 3453-3456.	0.9	22
61	Genetic characterization of a novel picornavirus distantly related to the marine mammal-infecting aquamaviruses in a long-distance migrant bird species, European roller (<i>Coracias garrulus</i>). <i>Journal of General Virology</i> , 2013, 94, 2029-2035.	1.3	21
62	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.	1.0	21
63	Novel picornavirus in domestic rabbits (<i>Oryctolagus cuniculus</i> var. <i>domestica</i>). <i>Infection, Genetics and Evolution</i> , 2016, 37, 117-122.	1.0	20
64	Detection and follow-up of torque teno midi virus (a small anellovirus) in nasopharyngeal aspirates and three other human body fluids in children. <i>Archives of Virology</i> , 2011, 156, 1537-1541.	0.9	19
65	A highly divergent picornavirus in an amphibian, the smooth newt (<i>Lissotriton vulgaris</i>). <i>Journal of General Virology</i> , 2015, 96, 2607-2613.	1.3	19
66	Genetic drift of norovirus genotype GII-4 in seven consecutive epidemic seasons in Hungary. <i>Journal of Clinical Virology</i> , 2008, 42, 135-140.	1.6	18
67	A tortoise-infecting picornavirus expands the host range of the family Picornaviridae. <i>Archives of Virology</i> , 2015, 160, 1319-1323.	0.9	18
68	Genome analysis of a novel, highly divergent picornavirus from common kestrel (<i>Falco tinnunculus</i>): The first non-enteroviral picornavirus with type-I-like IRES. <i>Infection, Genetics and Evolution</i> , 2015, 32, 425-431.	1.0	18
69	Identification of a novel variant of human hepatitis E virus in Hungary. <i>Journal of Clinical Virology</i> , 2006, 36, 100-102.	1.6	17
70	Seroepidemiology of hepatitis E virus in patients with non-A, non-B, non-C hepatitis in Hungary. <i>Journal of Medical Virology</i> , 2007, 79, 927-930.	2.5	17
71	Frequency and phylogeny of norovirus in diarrheic children in Istanbul, Turkey. <i>Journal of Clinical Virology</i> , 2011, 51, 160-164.	1.6	17
72	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.	1.0	17

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73	Saliviruses-the first knowledge about a newly discovered human picornavirus. <i>Reviews in Medical Virology</i> , 2017, 27, e1904.	3.9	17
74	Molecular detection and sequence analysis of human caliciviruses from acute gastroenteritis outbreaks in Hungary. <i>Journal of Medical Virology</i> , 2002, 67, 567-573.	2.5	16
75	Dicpivirus (family Picornaviridae) in wild Northern white-breasted hedgehog (<i>Erinaceus roumanicus</i>). <i>Archives of Virology</i> , 2018, 163, 175-181.	0.9	16
76	Human enterovirus 109 (EV109) in acute paediatric respiratory disease in Hungary. <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2012, 59, 285-290.	0.4	15
77	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.	1.3	13
78	Genetic characterization of a second novel picornavirus from an amphibian host, smooth newt (<i>Lissotriton vulgaris</i>). <i>Archives of Virology</i> , 2017, 162, 1043-1050.	0.9	12
79	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.	1.0	12
80	Co-circulation of genotype IA and new variant IB hepatitis A virus in outbreaks of acute hepatitis in Hungaryâ€”2003/2004. <i>Journal of Medical Virology</i> , 2006, 78, 1392-1397.	2.5	11
81	Complete genome characterization of mosavirus (family Picornaviridae) identified in droppings of a European roller (<i>Coracias garrulus</i>) in Hungary. <i>Archives of Virology</i> , 2014, 159, 2723-2729.	0.9	11
82	A cluster of salivirus A1 (Picornaviridae) infections in newborn babies with acute gastroenteritis in a neonatal hospital unit in Hungary. <i>Archives of Virology</i> , 2016, 161, 1671-1677.	0.9	11
83	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.	1.6	11
84	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.	1.0	10
85	Intra-host analysis of hepaciviral glycoprotein evolution reveals signatures associated with viral persistence and clearance. <i>Virus Evolution</i> , 2022, 8, veac007.	2.2	10
86	Detection and characterization of human parechoviruses in archived cell cultures, in Hungary. <i>Journal of Clinical Virology</i> , 2010, 47, 379-381.	1.6	9
87	Cytotoxic T lymphocytes mediate neuronal injury in patients with X-linked agammaglobulinemia and progressive neurodegenerative disease. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2011, 66, 1617-1618.	2.7	9
88	Novel 5â€™-3â€™RACE Method for Amplification and Determination of Single-Stranded RNAs Through Double-Stranded RNA (dsRNA) Intermediates. <i>Molecular Biotechnology</i> , 2015, 57, 974-981.	1.3	9
89	A novel passerivirus (family Picornaviridae) in an outbreak of enteritis with high mortality in estrildid finches (<i>Uraeginthus</i> sp.). <i>Archives of Virology</i> , 2018, 163, 1063-1071.	0.9	9
90	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.	1.5	9

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91	Detection of Aichi virus in South Korea. Archives of Virology, 2014, 159, 1835-1839.	0.9	8
92	Rabovirus: a proposed new picornavirus genus that is phylogenetically basal to enteroviruses and sapeloviruses. Archives of Virology, 2015, 160, 2569-2575.	0.9	8
93	Genome characterization of a novel megrivirus-related avian picornavirus from a carnivorous wild bird, western marsh harrier (<i>Circus aeruginosus</i>). Archives of Virology, 2017, 162, 2781-2789.	0.9	7
94	Genomic analysis of a novel picornavirus from a migratory waterfowl, greater white-fronted goose (<i>Anser albifrons</i>). Archives of Virology, 2018, 163, 1087-1090.	0.9	7
95	Molecular characterization of a novel picobirnavirus in a chicken. Archives of Virology, 2018, 163, 3455-3458.	0.9	7
96	Detection and genetic characterization of a novel parvovirus (family Parvoviridae) in barn owls (<i>Tyto</i>). Tj ETQq0 0 0 ggBT /Overlock 10 Tf 5	0.9	6
97	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.	1.0	6
98	Diverse picornaviruses are prevalent among free-living and laboratory rats (<i>Rattus norvegicus</i>) in Hungary and can cause disseminated infections. Infection, Genetics and Evolution, 2019, 75, 103988.	1.0	6
99	Novel picornavirus (family Picornaviridae) from freshwater fishes (<i>Perca fluviatilis</i> , Sander). Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5	0.9	6
100	Human-stool-associated tusavirus (Parvoviridae) in domestic goats and sheep. Archives of Virology, 2022, 167, 1307-1310.	0.9	5
101	An archived serum sample as a clue for identifying the primary source of a nosocomial hepatitis C virus outbreak in a haemodialysis unit. Archives of Virology, 2014, 159, 2207-2212.	0.9	4
102	A novel parvovirus (family Parvoviridae) in a freshwater fish, zander (<i>Sander lucioperca</i>). Archives of Virology, 2022, 167, 1163-1167.	0.9	4
103	Secondary structure analysis of swine pasivirus (family Picornaviridae) RNA reveals a type-IV IRES and a parechovirus-like 3' UTR organization. Archives of Virology, 2015, 160, 1363-1366.	0.9	3
104	Analysis of a novel RNA virus in a wild northern white-breasted hedgehog (<i>Erinaceus roumanicus</i>). Archives of Virology, 2019, 164, 3065-3071.	0.9	3
105	Performance of serologic assays specific to IgM antibodies against hepatitis E virus: pangentypic evaluation. International Journal of Infectious Diseases, 2010, 14, e239.	1.5	2
106	Co-infection with coxsackievirus A5 and norovirus GII.4 could have been the trigger of the first episode of severe acute encephalopathy in a six-year-old child with the intermittent form of maple syrup urine disease (MSUD). Archives of Virology, 2017, 162, 1757-1763.	0.9	2
107	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>). Infection, Genetics and Evolution, 2020, 81, 104236.	1.0	2
108	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. Infection, Genetics and Evolution, 2021, 93, 104942.	1.0	2

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109	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.	1.0	1
110	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.	1.0	1
111	KOBUVIRUS DETECTION IN THE CRITICALLY ENDANGERED PYGMY HOG (<i>PORCULA SALVANIA</i>), INDIA. <i>Journal of Zoo and Wildlife Medicine</i> , 2021, 52, 343-347.	0.3	1
112	Epidemiological, Clinicopathological and Virological Features of Merkel Cell Carcinomas in Medical Center of University of Pács, Hungary (2007–2012). <i>Pathology and Oncology Research</i> , 2016, 22, 71-77.	0.9	0
113	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.	1.5	0