

# Eric Delwart

## List of Publications by Year in descending order

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Version: 2024-02-01

346  
papers

25,147  
citations

5558

82  
h-index

9073

144  
g-index

356  
all docs

356  
docs citations

356  
times ranked

18541  
citing authors

#	ARTICLE	IF	CITATIONS
1	A novel parvovirus (family Parvoviridae) in a freshwater fish, zander ( <i>Sander lucioperca</i> ). <i>Archives of Virology</i> , 2022, 167, 1163-1167.	0.9	4
2	Enteric Viruses Nucleic Acids Distribution along the Digestive Tract of Rhesus Macaques with Idiopathic Chronic Diarrhea. <i>Viruses</i> , 2022, 14, 638.	1.5	0
3	Human-stool-associated tusavirus (Parvoviridae) in domestic goats and sheep. <i>Archives of Virology</i> , 2022, 167, 1307-1310.	0.9	5
4	Virome in the cloaca of wild and breeding birds revealed a diversity of significant viruses. <i>Microbiome</i> , 2022, 10, 60.	4.9	32
5	Chikungunya Virus Asian Lineage Infection in the Amazon Region Is Maintained by Asiatic and Caribbean-Introduced Variants. <i>Viruses</i> , 2022, 14, 1445.	1.5	3
6	A New Circular Single-Stranded DNA Virus Related with Howler Monkey Associated Porprismacovirus 1 Detected in Children with Acute Gastroenteritis. <i>Viruses</i> , 2022, 14, 1472.	1.5	1
7	Detection and genetic characterization of a novel parvovirus (family Parvoviridae) in barn owls ( <i>Tyto Tj ETQq1 1 0.784314 rgBT /Over</i>	0.9	7
8	Viral Metagenomic Analysis of Cerebrospinal Fluid from Patients with Acute Central Nervous System Infections of Unknown Origin, Vietnam. <i>Emerging Infectious Diseases</i> , 2021, 27, 205-213.	2.0	11
9	Virome of Bat Guano from Nine Northern California Roosts. <i>Journal of Virology</i> , 2021, 95, .	1.5	24
10	Norovirus strains in patients with acute gastroenteritis in rural and low-income urban areas in northern Brazil. <i>Archives of Virology</i> , 2021, 166, 905-913.	0.9	4
11	CrAssphage and its bacterial host in cat feces. <i>Scientific Reports</i> , 2021, 11, 815.	1.6	14
12	Astrovirus Outbreak in an Animal Shelter Associated With Feline Vomiting. <i>Frontiers in Veterinary Science</i> , 2021, 8, 628082.	0.9	5
13	Guapiaçu virus, a new insect-specific flavivirus isolated from two species of <i>Aedes</i> mosquitoes from Brazil. <i>Scientific Reports</i> , 2021, 11, 4674.	1.6	11
14	Redondoviridae: High Prevalence and Possibly Chronic Shedding in Human Respiratory Tract, But No Zoonotic Transmission. <i>Viruses</i> , 2021, 13, 533.	1.5	4
15	Multiple clades of Husavirus in South America revealed by next generation sequencing. <i>PLoS ONE</i> , 2021, 16, e0248486.	1.1	4
16	ContigExtender: a new approach to improving de novo sequence assembly for viral metagenomics data. <i>BMC Bioinformatics</i> , 2021, 22, 119.	1.2	10
17	High Heterogeneity of Echoviruses in Brazilian Children with Acute Gastroenteritis. <i>Viruses</i> , 2021, 13, 595.	1.5	5
18	Adaptive Evolution of New Variants of Dengue Virus Serotype 1 Genotype V Circulating in the Brazilian Amazon. <i>Viruses</i> , 2021, 13, 689.	1.5	5

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19	Circovirus in Blood of a Febrile Horse with Hepatitis. <i>Viruses</i> , 2021, 13, 944.	1.5	6
20	New Variants of Squash Mosaic Viruses Detected in Human Fecal Samples. <i>Microorganisms</i> , 2021, 9, 1349.	1.6	2
21	Novel picornavirus (family Picornaviridae) from freshwater fishes ( <i>Perca fluviatilis</i> , Sander) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.9	6
22	New Parvoviruses and Picornavirus in Tissues and Feces of Foals with Interstitial Pneumonia. <i>Viruses</i> , 2021, 13, 1612.	1.5	6
23	Human astrovirus types 1, 4 and 5 circulating among children with acute gastroenteritis in a rural Brazilian state, 2010-2016. <i>Archives of Virology</i> , 2021, 166, 3165-3172.	0.9	4
24	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.	1.0	2
25	Investigation of Three Newly Identified Equine Parvoviruses in Blood and Nasal Fluid Samples of Clinically Healthy Horses and Horses with Acute Onset of Respiratory Disease. <i>Animals</i> , 2021, 11, 3006.	1.0	5
26	Composition of Eukaryotic Viruses and Bacteriophages in Individuals with Acute Gastroenteritis. <i>Viruses</i> , 2021, 13, 2365.	1.5	6
27	Idiopathic Chronic Diarrhea in Rhesus Macaques Is Not Associated with Enteric Viral Infections. <i>Viruses</i> , 2021, 13, 2503.	1.5	0
28	Detection and characterization of Ilheus and Iguape virus genomes in historical mosquito samples from Southern Brazil. <i>Acta Tropica</i> , 2020, 205, 105401.	0.9	12
29	<i>Aedes aegypti</i> from Amazon Basin Harbor High Diversity of Novel Viral Species. <i>Viruses</i> , 2020, 12, 866.	1.5	12
30	The Virome of Acute Respiratory Diseases in Individuals at Risk of Zoonotic Infections. <i>Viruses</i> , 2020, 12, 960.	1.5	17
31	Complex Virome in a Mesenteric Lymph Node from a Californian Sea Lion ( <i>Zalophus californianus</i> ) with Polyserositis and Steatitis. <i>Viruses</i> , 2020, 12, 793.	1.5	13
32	Virome of a Feline Outbreak of Diarrhea and Vomiting Includes Bocaviruses and a Novel Chapparvovirus. <i>Viruses</i> , 2020, 12, 506.	1.5	42
33	Genomic Analyses of Potential Novel Recombinant Human Adenovirus C in Brazil. <i>Viruses</i> , 2020, 12, 508.	1.5	6
34	Plasma virome of 781 Brazilians with unexplained symptoms of arbovirus infection include a novel parvovirus and densovirus. <i>PLoS ONE</i> , 2020, 15, e0229993.	1.1	19
35	Performance of Metagenomic Next-Generation Sequencing for the Diagnosis of Viral Meningoencephalitis in a Resource-Limited Setting. <i>Open Forum Infectious Diseases</i> , 2020, 7, ofaa046.	0.4	26
36	Genotyping Porcine Circovirus 3 (PCV-3) Nowadays: Does It Make Sense?. <i>Viruses</i> , 2020, 12, 265.	1.5	47

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37	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.	1.0	2
38	Report of the 2019 NIST-FDA workshop on standards for next generation sequencing detection of viral adventitious agents in biologics and biomanufacturing. <i>Biologicals</i> , 2020, 64, 76-82.	0.5	11
39	<i>Cressdnaviricota</i> : a Virus Phylum Unifying Seven Families of Rep-Encoding Viruses with Single-Stranded, Circular DNA Genomes. <i>Journal of Virology</i> , 2020, 94, .	1.5	118
40	Identification of a novel bovine copiparvovirus in pooled fetal bovine serum. <i>Virus Genes</i> , 2020, 56, 522-526.	0.7	4
41	Semen virome of men with HIV on or off antiretroviral treatment. <i>Aids</i> , 2020, 34, 827-832.	1.0	13
42	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
43	Viral gastroenteritis in Tocantins, Brazil: characterizing the diversity of human adenovirus F through next-generation sequencing and bioinformatics. <i>Journal of General Virology</i> , 2020, 101, 1280-1288.	1.3	17
44	Viruses in unexplained encephalitis cases in American black bears ( <i>Ursus americanus</i> ). <i>PLoS ONE</i> , 2020, 15, e0244056.	1.1	11
45	Genomic constellation of human Rotavirus A strains identified in Northern Brazil: a 6-year follow-up (2010-2016). <i>Revista Do Instituto De Medicina Tropical De Sao Paulo</i> , 2020, 62, e98.	0.5	9
46	Serendipitous Discovery of a Novel Murine Astrovirus Contaminating a Murine Helper T-cell Line and Incapable of Infecting Highly Immunodeficient Mice. <i>Comparative Medicine</i> , 2020, 70, 359-369.	0.4	5
47	Viruses in unexplained encephalitis cases in American black bears ( <i>Ursus americanus</i> ). , 2020, 15, e0244056.		0
48	Viruses in unexplained encephalitis cases in American black bears ( <i>Ursus americanus</i> ). , 2020, 15, e0244056.		0
49	Viruses in unexplained encephalitis cases in American black bears ( <i>Ursus americanus</i> ). , 2020, 15, e0244056.		0
50	Viruses in unexplained encephalitis cases in American black bears ( <i>Ursus americanus</i> ). , 2020, 15, e0244056.		0
51	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.	1.0	6
52	The rare enterovirus c99 and echovirus 29 strains in Brazil: potential risks associated to silent circulation. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2019, 114, e190160.	0.8	12
53	Lyon-IARC Polyomavirus DNA in Feces of Diarrheic Cats. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	15
54	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.	0.9	3

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55	Viruses in Horses with Neurologic and Respiratory Diseases. <i>Viruses</i> , 2019, 11, 942.	1.5	28
56	High Diversity and Novel Enteric Viruses in Fecal Viromes of Healthy Wild and Captive Thai Cynomolgus Macaques ( <i>Macaca fascicularis</i> ). <i>Viruses</i> , 2019, 11, 971.	1.5	18
57	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
58	One-step pentaplex real-time polymerase chain reaction assay for detection of zika, dengue, chikungunya, West Nile viruses and a human housekeeping gene. <i>Journal of Clinical Virology</i> , 2019, 120, 44-50.	1.6	15
59	Characterization of a peribunyavirus isolated from largemouth bass ( <i>Micropterus salmoides</i> ). <i>Virus Research</i> , 2019, 273, 197761.	1.1	9
60	Viral species richness and composition in young children with loose or watery stool in Ethiopia. <i>BMC Infectious Diseases</i> , 2019, 19, 53.	1.3	18
61	Genomic sequencing of a virus representing a novel type within the species <i>Dyopipapillomavirus 1</i> in an Indian River Lagoon bottlenose dolphin. <i>Archives of Virology</i> , 2019, 164, 767-774.	0.9	5
62	Gut virome of mammals and birds reveals high genetic diversity of the family Microviridae. <i>Virus Evolution</i> , 2019, 5, vez013.	2.2	37
63	Viruses in Vietnamese Patients Presenting with Community-Acquired Sepsis of Unknown Cause. <i>Journal of Clinical Microbiology</i> , 2019, 57, .	1.8	27
64	Recombinant Strains of Human Parechovirus in Rural Areas in the North of Brazil. <i>Viruses</i> , 2019, 11, 488.	1.5	10
65	Effect of Geographic Isolation on the Nasal Virome of Indigenous Children. <i>Journal of Virology</i> , 2019, 93, .	1.5	10
66	The first reptilian circovirus identified infects gut and liver tissues of black-headed pythons. <i>Veterinary Research</i> , 2019, 50, 35.	1.1	14
67	Chapparvovirus DNA Found in 4% of Dogs with Diarrhea. <i>Viruses</i> , 2019, 11, 398.	1.5	49
68	Composite Analysis of the Virome and Bacteriome of HIV/HPV Co-Infected Women Reveals Proxies for Immunodeficiency. <i>Viruses</i> , 2019, 11, 422.	1.5	30
69	Canine bufavirus in faeces and plasma of dogs with diarrhoea, China. <i>Emerging Microbes and Infections</i> , 2019, 8, 245-247.	3.0	21
70	Discovery of Cucumis melo endornavirus by deep sequencing of human stool samples in Brazil. <i>Virus Genes</i> , 2019, 55, 332-338.	0.7	13
71	Human sapovirus GI.2 and GI.3 from children with acute gastroenteritis in northern Brazil. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2019, 114, e180574.	0.8	9
72	A novel phage from periodontal pockets associated with chronic periodontitis. <i>Virus Genes</i> , 2019, 55, 381-393.	0.7	19

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73	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
74	Detection of RNA-Dependent RNA Polymerase of Hubei Reo-Like Virus 7 by Next-Generation Sequencing in <i>Aedes aegypti</i> and <i>Culex quinquefasciatus</i> Mosquitoes from Brazil. <i>Viruses</i> , 2019, 11, 147.	1.5	8
75	Viral metagenomics reveals significant viruses in the genital tract of apparently healthy dairy cows. <i>Archives of Virology</i> , 2019, 164, 1059-1067.	0.9	18
76	First identification of mammalian orthoreovirus type 3 by gut virome analysis in diarrheic child in Brazil. <i>Scientific Reports</i> , 2019, 9, 18599.	1.6	22
77	The fecal virome of red-crowned cranes. <i>Archives of Virology</i> , 2019, 164, 3-16.	0.9	59
78	Enteric Virome and Bacterial Microbiota in Children With Ulcerative Colitis and Crohn Disease. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2019, 68, 30-36.	0.9	89
79	Asian black bear ( <i>Ursus thibetanus</i> ) picornavirus related to seal aquamavirus A. <i>Archives of Virology</i> , 2019, 164, 653-656.	0.9	3
80	Detection and Characterization of Enterovirus B73 from a Child in Brazil. <i>Viruses</i> , 2019, 11, 16.	1.5	12
81	Nasal virome of dogs with respiratory infection signs include novel taupapillomaviruses. <i>Virus Genes</i> , 2019, 55, 191-197.	0.7	14
82	Evidence for an unknown agent antigenically related to the hepatitis E virus in dairy cows in the United States. <i>Journal of Medical Virology</i> , 2019, 91, 677-686.	2.5	23
83	Next-generation sequencing of dsRNA is greatly improved by treatment with the inexpensive denaturing reagent DMSO. <i>Microbial Genomics</i> , 2019, 5, .	1.0	8
84	Amdoparvovirus Infection in Red Pandas ( <i>Ailurus fulgens</i> ). <i>Veterinary Pathology</i> , 2018, 55, 552-561.	0.8	24
85	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
86	Taxonomy of the family Arenaviridae and the order Bunyavirales: update 2018. <i>Archives of Virology</i> , 2018, 163, 2295-2310.	0.9	157
87	Genomic analysis of a novel picornavirus from a migratory waterfowl, greater white-fronted goose ( <i>Anser albifrons</i> ). <i>Archives of Virology</i> , 2018, 163, 1087-1090.	0.9	7
88	Complete Genome Sequencing of a Novel Type of Omikronpapillomavirus 1 in Indian River Lagoon Bottlenose Dolphins ( <i>Tursiops truncatus</i> ). <i>Genome Announcements</i> , 2018, 6, .	0.8	3
89	Complete Genome Sequences of Six Human Bocavirus Strains from Patients with Acute Gastroenteritis in the North Region of Brazil. <i>Genome Announcements</i> , 2018, 6, .	0.8	8
90	Wuhan large pig roundworm virus identified in human feces in Brazil. <i>Virus Genes</i> , 2018, 54, 470-473.	0.7	9

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91	Sera of Peruvians with fever of unknown origins include viral nucleic acids from non-vertebrate hosts. <i>Virus Genes</i> , 2018, 54, 33-40.	0.7	19
92	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
93	A Novel Highly Divergent Strain of Cell Fusing Agent Virus (CFAV) in Mosquitoes from the Brazilian Amazon Region. <i>Viruses</i> , 2018, 10, 666.	1.5	11
94	Recombination Located over 2A-2B Junction Ribosome Frameshifting Region of Saffold Cardiovirus. <i>Viruses</i> , 2018, 10, 520.	1.5	9
95	Complex virome in feces from Amerindian children in isolated Amazonian villages. <i>Nature Communications</i> , 2018, 9, 4270.	5.8	51
96	Molecular characterization of a novel picobirnavirus in a chicken. <i>Archives of Virology</i> , 2018, 163, 3455-3458.	0.9	7
97	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
98	Full-length and defective enterovirus G genomes with distinct torovirus protease insertions are highly prevalent on a Chinese pig farm. <i>Archives of Virology</i> , 2018, 163, 2471-2476.	0.9	19
99	Global Distribution of Human Protoparvoviruses. <i>Emerging Infectious Diseases</i> , 2018, 24, 1292-1299.	2.0	21
100	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
101	Metagenomics analysis of the virome of 300 concentrates from a Swiss platelet bank. <i>Vox Sanguinis</i> , 2018, 113, 601-604.	0.7	9
102	Plasma virome of cattle from forest region revealed diverse small circular ssDNA viral genomes. <i>Virology Journal</i> , 2018, 15, 11.	1.4	27
103	Enteric virome of Ethiopian children participating in a clean water intervention trial. <i>PLoS ONE</i> , 2018, 13, e0202054.	1.1	29
104	Near full length genome of a recombinant (E/D) cosavirus strain from a rural area in the central region of Brazil. <i>Scientific Reports</i> , 2018, 8, 12304.	1.6	17
105	Virome of ~12 thousand <i>Culex</i> mosquitoes from throughout California. <i>Virology</i> , 2018, 523, 74-88.	1.1	88
106	Virome of US bovine calf serum. <i>Biologicals</i> , 2017, 46, 64-67.	0.5	39
107	A novel pulmonary polyomavirus in alpacas ( <i>Vicugna pacos</i> ). <i>Veterinary Microbiology</i> , 2017, 201, 49-55.	0.8	6
108	Rotavirus I in feces of a cat with diarrhea. <i>Virus Genes</i> , 2017, 53, 487-490.	0.7	19

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109	ENDEMIC INFECTION OF STRANDED SOUTHERN SEA OTTERS ( <i>ENHYDRA LUTRIS NEREIS</i> ) WITH NOVEL PARVOVIRUS, POLYOMAVIRUS, AND ADENOVIRUS. <i>Journal of Wildlife Diseases</i> , 2017, 53, 532-542.	0.3	25
110	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.	0.9	7
111	Densoviruses in oyster <i>Crassostrea ariakensis</i> . <i>Archives of Virology</i> , 2017, 162, 2153-2157.	0.9	11
112	Revisiting the taxonomy of the family Circoviridae: establishment of the genus Cyclovirus and removal of the genus Gyrovirus. <i>Archives of Virology</i> , 2017, 162, 1447-1463.	0.9	285
113	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
114	Virus taxonomy in the age of metagenomics. <i>Nature Reviews Microbiology</i> , 2017, 15, 161-168.	13.6	590
115	In utero cytomegalovirus infection and development of childhood acute lymphoblastic leukemia. <i>Blood</i> , 2017, 129, 1680-1684.	0.6	55
116	Small Circular Rep-Encoding Single-Stranded DNA Genomes in Peruvian Diarrhea Virome. <i>Genome Announcements</i> , 2017, 5, .	0.8	15
117	Genomes of viral isolates derived from different mosquitos species. <i>Virus Research</i> , 2017, 242, 49-57.	1.1	40
118	Genome Sequence of Canine Polyomavirus in Respiratory Secretions of Dogs with Pneumonia of Unknown Etiology. <i>Genome Announcements</i> , 2017, 5, .	0.8	3
119	A Naturally Transmitted Epitheliotropic Polyomavirus Pathogenic in Immunodeficient Rats: Characterization, Transmission, and Preliminary Epidemiologic Studies. <i>Toxicologic Pathology</i> , 2017, 45, 593-603.	0.9	10
120	Genomic and epidemiological characterisation of a dengue virus outbreak among blood donors in Brazil. <i>Scientific Reports</i> , 2017, 7, 15216.	1.6	40
121	Case-Control Comparison of Enteric Viromes in Captive Rhesus Macaques with Acute or Idiopathic Chronic Diarrhea. <i>Journal of Virology</i> , 2017, 91, .	1.5	46
122	Virome comparisons in wild-diseased and healthy captive giant pandas. <i>Microbiome</i> , 2017, 5, 90.	4.9	113
123	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
124	Spread of Chikungunya Virus East/Central/South African Genotype in Northeast Brazil. <i>Emerging Infectious Diseases</i> , 2017, 23, 1742-1744.	2.0	69
125	A novel polyomavirus from the nasal cavity of a giant panda ( <i>Ailuropoda melanoleuca</i> ). <i>Virology Journal</i> , 2017, 14, 207.	1.4	6
126	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

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127	Detection of a new species of torque teno mini virus from the gingival epithelium of patients with periodontitis. <i>Virus Genes</i> , 2017, 53, 823-830.	0.7	17
128	ICTV Virus Taxonomy Profile: Circoviridae. <i>Journal of General Virology</i> , 2017, 98, 1997-1998.	1.3	147
129	Zika Virus Tissue and Blood Compartmentalization in Acute Infection of Rhesus Macaques. <i>PLoS ONE</i> , 2017, 12, e0171148.	1.1	102
130	The blood DNA virome in 8,000 humans. <i>PLoS Pathogens</i> , 2017, 13, e1006292.	2.1	259
131	Two new species of betatorqueviruses identified in a human melanoma that metastasized to the brain. <i>Oncotarget</i> , 2017, 8, 105800-105808.	0.8	27
132	Genetic Characterization and Classification of Human and Animal Sapoviruses. <i>PLoS ONE</i> , 2016, 11, e0156373.	1.1	71
133	The Ancient Evolutionary History of Polyomaviruses. <i>PLoS Pathogens</i> , 2016, 12, e1005574.	2.1	190
134	Viral nucleic acids in human plasma pools. <i>Transfusion</i> , 2016, 56, 2248-2255.	0.8	77
135	A new protoparvovirus in human fecal samples and cutaneous T cell lymphomas (mycosis fungoides). <i>Virology</i> , 2016, 496, 299-305.	1.1	49
136	Epidemiology of two human protoparvoviruses, bufavirus and tusavirus. <i>Scientific Reports</i> , 2016, 6, 39267.	1.6	28
137	A Common Parvovirus in Deer from California, USA. <i>Journal of Wildlife Diseases</i> , 2016, 52, 962-964.	0.3	1
138	A new densovirus in cerebrospinal fluid from a case of anti-NMDA-receptor encephalitis. <i>Archives of Virology</i> , 2016, 161, 3231-3235.	0.9	15
139	A novel bocavirus from domestic mink, China. <i>Virus Genes</i> , 2016, 52, 887-890.	0.7	23
140	Identification of a novel human papillomavirus by metagenomic analysis of vaginal swab samples from pregnant women. <i>Virology Journal</i> , 2016, 13, 122.	1.4	28
141	Detection of a novel circovirus PCV3 in pigs with cardiac and multi-systemic inflammation. <i>Virology Journal</i> , 2016, 13, 184.	1.4	309
142	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
143	A novel species of torque teno mini virus (TTMV) in gingival tissue from chronic periodontitis patients. <i>Scientific Reports</i> , 2016, 6, 26739.	1.6	35
144	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

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145	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
146	The fecal virome of South and Central American children with diarrhea includes small circular DNA viral genomes of unknown origin. <i>Archives of Virology</i> , 2016, 161, 959-966.	0.9	36
147	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
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