

# Szilvia L Farkas

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1713052/publications.pdf>

Version: 2024-02-01

39  
papers

1,436  
citations

430874

18  
h-index

330143

37  
g-index

39  
all docs

39  
docs citations

39  
times ranked

2191  
citing authors

#	ARTICLE	IF	CITATIONS
1	Taxonomy of the order Mononegavirales: update 2016. <i>Archives of Virology</i> , 2016, 161, 2351-2360.	2.1	407
2	Candidate New Rotavirus Species in Sheltered Dogs, Hungary. <i>Emerging Infectious Diseases</i> , 2015, 21, 660-663.	4.3	170
3	Zoonotic transmission of rotavirus: surveillance and control. <i>Expert Review of Anti-Infective Therapy</i> , 2015, 13, 1337-1350.	4.4	95
4	First detection and analysis of a fish circovirus. <i>Journal of General Virology</i> , 2011, 92, 1817-1821.	2.9	67
5	A parvovirus isolated from royal python ( <i>Python regius</i> ) is a member of the genus Dependovirus. <i>Journal of General Virology</i> , 2004, 85, 555-561.	2.9	62
6	Genomic and phylogenetic analyses of an adenovirus isolated from a corn snake ( <i>Elaphe guttata</i> ) imply a common origin with members of the proposed new genus Atadenovirus. <i>Journal of General Virology</i> , 2002, 83, 2403-2410.	2.9	62
7	Adenovirus and mycoplasma infection in an ornate box turtle ( <i>Terrapene ornata ornata</i> ) in Hungary. <i>Veterinary Microbiology</i> , 2009, 138, 169-173.	1.9	48
8	Completion of the genome analysis of snake adenovirus type 1, a representative of the reptilian lineage within the novel genus Atadenovirus. <i>Virus Research</i> , 2008, 132, 132-139.	2.2	43
9	Unique genomic organization of a novel Avipoxvirus detected in turkey ( <i>Meleagris gallopavo</i> ). <i>Infection, Genetics and Evolution</i> , 2015, 35, 221-229.	2.3	39
10	Detection and characterization of a divergent avian reovirus strain from a broiler chicken with central nervous system disease. <i>Archives of Virology</i> , 2013, 158, 2583-2588.	2.1	34
11	Novel NSP4 genotype in a camel G10P[15] rotavirus strain. <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2012, 59, 411-421.	0.8	33
12	Full genome characterization of human Rotavirus A strains isolated in Cameroon, 2010–2011: Diverse combinations of the G and P genes and lack of reassortment of the backbone genes. <i>Infection, Genetics and Evolution</i> , 2014, 28, 537-560.	2.3	31
13	Rotavirus strains in neglected animal species including lambs, goats and camelids. <i>VirusDisease</i> , 2014, 25, 215-222.	2.0	25
14	Detection and sequencing of West Nile virus RNA from human urine and serum samples during the 2014 seasonal period. <i>Archives of Virology</i> , 2016, 161, 1797-1806.	2.1	25
15	Whole-genome sequencing of a green bush viper reovirus reveals a shared evolutionary history between reptilian and unusual mammalian orthoreoviruses. <i>Archives of Virology</i> , 2014, 159, 153-158.	2.1	24
16	Lineage diversification, homo- and heterologous reassortment and recombination shape the evolution of chicken orthoreoviruses. <i>Scientific Reports</i> , 2016, 6, 36960.	3.3	23
17	Complete genome analysis identifies TvÅrminne avian virus as a candidate new species within the genus Orthoreovirus. <i>Journal of General Virology</i> , 2014, 95, 898-904.	2.9	19
18	Sequencing and phylogenetic analysis identifies candidate members of a new picornavirus genus in terrestrial tortoise species. <i>Archives of Virology</i> , 2015, 160, 811-816.	2.1	18

#	ARTICLE	IF	CITATIONS
19	Isolation and complete genome characterization of novel reassortant orthoreovirus from common vole ( <i>Microtus arvalis</i> ). <i>Virus Genes</i> , 2017, 53, 307-311.	1.6	17
20	Genomic sequence and phylogenetic analyses of two novel orthoreovirus strains isolated from Pekin ducks in 2014 in Germany. <i>Virus Research</i> , 2018, 257, 57-62.	2.2	17
21	Detection of shared genes among Asian and European waterfowl reoviruses in the whole genome constellations. <i>Infection, Genetics and Evolution</i> , 2014, 28, 55-57.	2.3	16
22	Neuroinvasive influenza virus A(H5N8) in fattening ducks, Hungary, 2015. <i>Infection, Genetics and Evolution</i> , 2016, 43, 418-423.	2.3	15
23	Novel mastadenovirus infection and clinical disease in a pygmy marmoset ( <i>Callithrix [Cebuella] Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 462</i> )	1.9	14
24	Integrated circoviral rep-like sequences in the genome of cyprinid fish. <i>Virus Genes</i> , 2013, 47, 374-377.	1.6	13
25	Ubiquiter circovirus sequences raise challenges in laboratory diagnosis: The case of honey bee and bee mite, reptiles, and free living amoebae. <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2015, 62, 57-73.	0.8	13
26	Phylogenetic analysis of a novel reassortant orthoreovirus strain detected in partridge ( <i>Perdix Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462</i> )	2.2	13
27	Whole genome sequencing of a rare rotavirus from archived stool sample demonstrates independent zoonotic origin of human G8P[14] strains in Hungary. <i>Virus Research</i> , 2017, 227, 96-103.	2.2	13
28	The fecal virome of domesticated animals. <i>VirusDisease</i> , 2014, 25, 150-157.	2.0	11
29	The complete genome sequence of a European goose reovirus strain. <i>Archives of Virology</i> , 2014, 159, 2165-2169.	2.1	10
30	Reassortant human group C rotaviruses in Hungary. <i>Infection, Genetics and Evolution</i> , 2015, 34, 410-414.	2.3	10
31	Whole genome characterization of a chelonian orthoreovirus strain identifies significant genetic diversity and may classify reptile orthoreoviruses into distinct species. <i>Virus Research</i> , 2016, 215, 94-98.	2.2	10
32	Development of a consensus reverse transcription PCR assay for the specific detection of tortoise picornaviruses. <i>Journal of Veterinary Diagnostic Investigation</i> , 2016, 28, 309-314.	1.1	7
33	Coding-complete sequencing classifies parrot bornavirus 5 into a novel virus species. <i>Archives of Virology</i> , 2015, 160, 2763-2768.	2.1	6
34	Complete Genome Sequence of a Genotype G23P[37] Pheasant Rotavirus Strain Identified in Hungary. <i>Genome Announcements</i> , 2016, 4, .	0.8	6
35	Whole genome sequencing and phylogenetic characterization of brown bullhead ( <i>Ameiurus Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 462</i> ) Evolution, 2016, 45, 402-407.	2.3	5
36	Novel adenovirus detected in kowari ( <i>Dasyuroides byrnei</i> ) with pneumonia. <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2017, 64, 81-90.	0.8	5

#	ARTICLE	IF	CITATIONS
37	Classification and characterization of a laboratory chicken rotavirus strain carrying G7P[35] neutralization antigens on the genotype 4 backbone gene configuration. <i>Biologicals</i> , 2014, 42, 299-304.	1.4	4
38	Genome Sequences of Three Turkey Orthoreovirus Strains Isolated in Hungary. <i>Genome Announcements</i> , 2015, 3, .	0.8	3
39	Genomic characterization of avian and neoavian orthoreoviruses detected in pheasants. <i>Virus Research</i> , 2021, 297, 198349.	2.2	3