Susan L Payne

List of Publications by Year in descending order

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SUSAN L DAVNE

#	Article	IF	CITATIONS
1	ICTV Virus Taxonomy Profile: Bornaviridae. Journal of General Virology, 2021, 102, .	2.9	24
2	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2021, 166, 3513-3566.	2.1	62
3	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2020, 165, 3023-3072.	2.1	184
4	Taxonomy of the order Mononegavirales: second update 2018. Archives of Virology, 2019, 164, 1233-1244.	2.1	70
5	Taxonomy of the order Mononegavirales: update 2019. Archives of Virology, 2019, 164, 1967-1980.	2.1	224
6	Strengthening the Interaction of the Virology Community with the International Committee on Taxonomy of Viruses (ICTV) by Linking Virus Names and Their Abbreviations to Virus Species. Systematic Biology, 2019, 68, 828-839.	5.6	11
7	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	2.1	153
8	Avian Vaccination. Veterinary Clinics of North America - Exotic Animal Practice, 2018, 21, 379-397.	0.7	4
9	Studies on immunity and immunopathogenesis of parrot bornaviral disease in cockatiels. Virology, 2018, 515, 81-91.	2.4	23
10	Taxonomy of the order Mononegavirales: update 2017. Archives of Virology, 2017, 162, 2493-2504.	2.1	173
11	Aquatic Bird Bornavirus-Associated Disease in Free-Living Canada Geese (<i>Branta canadensis</i>) in the Northeastern USA. Journal of Wildlife Diseases, 2017, 53, 607-611.	0.8	7
12	Apparent resolution of parrot bornavirus infection in cockatiels (Nymphicus hollandicus). Veterinary Medicine: Research and Reports, 2017, Volume 8, 31-36.	0.6	2
13	The pathogenesis of proventricular dilatation disease. Animal Health Research Reviews, 2016, 17, 110-126.	3.1	20
14	Horses naturally infected with EIAV harbor 2 distinct SU populations but are monophyletic with respect to IN. Virus Genes, 2016, 52, 71-80.	1.6	0
15	Taxonomy of the order Mononegavirales: update 2016. Archives of Virology, 2016, 161, 2351-2360.	2.1	407
16	Possibility and Challenges of Conversion of Current Virus Species Names to Linnaean Binomials. Systematic Biology, 2016, 66, syw096.	5.6	17
17	The pathogenesis of bornaviral diseases in mammals. Animal Health Research Reviews, 2016, 17, 92-109.	3.1	44
18	Avian Bornaviruses in North American Gulls. Journal of Wildlife Diseases, 2015, 51, 754-758.	0.8	17

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19	Characterization of a new genotype of avian bornavirus from wild ducks. Virology Journal, 2014, 11, 197.	3.4	24
20	Avian Bornaviruses: Diagnosis, Isolation, and Genotyping. Current Protocols in Microbiology, 2014, 34, 15I.1.1-33.	6.5	22
21	Complete Genome Sequence of Avian Bornavirus Genotype 1 from a Macaw with Proventricular Dilatation Disease. Journal of Virology, 2012, 86, 7023-7023.	3.4	3
22	Birds and bornaviruses. Animal Health Research Reviews, 2012, 13, 145-156.	3.1	52
23	Widespread avian bornavirus infection in mute swans in the Northeast United States. Veterinary Medicine: Research and Reports, 2012, 3, 49.	0.6	19
24	Proventricular Dilatation Disease in Cockatiels (Nymphicus hollandicus) After Infection With a Genotype 2 Avian Bornavirus. , 2011, 25, 199-204.		42
25	Detection and Characterization of a Distinct Bornavirus Lineage from Healthy Canada Geese (Branta) Tj ETQq1 1	l 0.78431 3.4	4 rgBT /Oved
26	Virulence Determinants of Equine Infectious Anemia Virus. Current HIV Research, 2010, 8, 66-72.	0.5	9
27	The diagnosis of proventricular dilatation disease: Use of a Western blot assay to detect antibodies against avian Borna virus. Veterinary Microbiology, 2010, 143, 196-201.	1.9	43
28	EIAV S2 enhances pro-inflammatory cytokine and chemokine response in infected macrophages. Virology, 2010, 397, 217-223.	2.4	29
29	Use of Avian Bornavirus Isolates to Induce Proventricular Dilatation Disease in Conures. Emerging Infectious Diseases, 2010, 16, 473-479.	4.3	89
30	The Isolation, Pathogenesis, Diagnosis, Transmission, and Control of Avian Bornavirus and Proventricular Dilatation Disease. Veterinary Clinics of North America - Exotic Animal Practice, 2010, 13, 495-508.	0.7	52
31	Equine Infectious Anemia Virus as a Model for Lentiviral Pathogenesis. , 2006, , 365-390.		0
32	The S2 accessory gene of equine infectious anemia virus is essential for expression of disease in ponies. Virology, 2006, 349, 22-30.	2.4	20
33	Influence of Long Terminal Repeat and Env on the Virulence Phenotype of Equine Infectious Anemia Virus. Journal of Virology, 2004, 78, 2478-2485.	3.4	23
34	Disease Induction by Virus Derived from Molecular Clones of Equine Infectious Anemia Virus. Journal of Virology, 1998, 72, 483-487.	3.4	27
35	Localization of conserved and variable antigenic domains of equine infectious anemia virus envelope glycoproteins using recombinant env-encoded protein fragments produced in Escherichia coli. Virology, 1989, 172, 609-615.	2.4	52
36	Antigenic variation and lentivirus persistence: Variations in envelope gene sequences during EIAV infection resemble changes reported for sequential isolates of HIV. Virology, 1987, 161, 321-331.	2.4	138