

Complete sequence of Great Island virus and comparison of proteins of Kemerovo, Lipovnik and Tribec viruses (genus *Orthomyxovirus*)

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Serotype Specific Primers and Gel-Based RT-PCR Assays for "Typing" African Horse Sickness Virus: Identification of Strains from Africa. PLoS ONE, 2011, 6, e25686.	2.5	13
2	Complete Genome Characterisation of a Novel 26th Bluetongue Virus Serotype from Kuwait. PLoS ONE, 2011, 6, e26147.	2.5	151
3	RNA segment 9 exists as a duplex concatemer in an Australian strain of epizootic haemorrhagic disease virus (EHDV): Genetic analysis and evidence for the presence of concatemers as a normal feature of orbivirus replication. Virology, 2011, 420, 164-171.	2.4	15
4	Umatilla Virus Genome Sequencing and Phylogenetic Analysis: Identification of Stretch Lagoon Orbivirus as a New Member of the Umatilla virus Species. PLoS ONE, 2011, 6, e23605.	2.5	37
5	Molecular characterization of eight segments of Scylla serrata reovirus (SsRV) provides the complete genome sequence. Archives of Virology, 2012, 157, 1551-1557.	2.1	9
6	Confusions in orbivirus protein classification. Virology Journal, 2012, 9, 166.	3.4	8
7	Full Genome Sequencing and Genetic Characterization of Eubenangee Viruses Identify Pata Virus as a Distinct Species within the Genus Orbivirus. PLoS ONE, 2012, 7, e31911.	2.5	31
8	Tick-borne viruses in Europe. Parasitology Research, 2012, 111, 9-36.	1.6	204
9	Genetic characterization of Tribevirus and Kemerovo virus, two tick-transmitted human-pathogenic Orbiviruses. Virology, 2012, 423, 68-76.	2.4	44
10	A novel mosquito-borne Orbivirus species found in South-east Asia. Journal of General Virology, 2013, 94, 1051-1057.	2.9	24
12	Endogenous tick viruses and modulation of tick-borne pathogen growth. Frontiers in Cellular and Infection Microbiology, 2013, 3, 25.	3.9	29
13	Tibet Orbivirus, a Novel Orbivirus Species Isolated from Anopheles maculatus Mosquitoes in Tibet, China. PLoS ONE, 2014, 9, e88738.	2.5	26
14	Genetic and biological characterization of selected Changuinola viruses (Reoviridae, Orbivirus) from Brazil. Journal of General Virology, 2014, 95, 2251-2259.	2.9	10
15	Characterization of Two Strains of Tribevirus Isolated in Ukraine. Vector-Borne and Zoonotic Diseases, 2014, 14, 808-816.	1.5	8
16	The Characterization of RNA Viruses in Tropical Seawater Using Targeted PCR and Metagenomics. MBio, 2014, 5, e01210-14.	4.1	69
17	Prevalence of Kemerovo virus in ixodid ticks from the Russian Federation. Ticks and Tick-borne Diseases, 2014, 5, 651-655.	2.7	23
18	First isolation and characterization of a mosquito-borne orbivirus belonging to the species Umatilla virus in East Asia. Archives of Virology, 2014, 159, 2675-2685.	2.1	14
19	Mobuck virus genome sequence and phylogenetic analysis: identification of a novel Orbivirus isolated from a white-tailed deer in Missouri, USA. Journal of General Virology, 2014, 95, 110-116.	2.9	17

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20	First detection of Kemerovo virus in <i>Ixodes pavlovskyi</i> and <i>Ixodes persulcatus</i> ticks collected in Novosibirsk region, Russia. <i>Ticks and Tick-borne Diseases</i> , 2014, 5, 494-496.	2.7	17
21	Immunisation with bacterial expressed VP2 and VP5 of bluetongue virus (BTV) protect IFNAR ² knock-out (IFNAR ² /Δ ²) mice from homologous lethal challenge. <i>Vaccine</i> , 2014, 32, 4059-4067.	3.8	26
22	Characterising Non-Structural Protein NS4 of African Horse Sickness Virus. <i>PLoS ONE</i> , 2015, 10, e0124281.	2.5	29
23	Double-Stranded RNA Viruses. , 2015, , 113-133.		0
24	A Review of Knowledge Gaps and Tools for Orbivirus Research. <i>Vector-Borne and Zoonotic Diseases</i> , 2015, 15, 339-347.	1.5	25
25	Genetic Characterization of the Tick-Borne Orbiviruses. <i>Viruses</i> , 2015, 7, 2185-2209.	3.3	36
26	Genetic and biological characterization of Muko virus, a new distinct member of the species Great Island virus (genus Orbivirus, family Reoviridae), isolated from ixodid ticks in Japan. <i>Archives of Virology</i> , 2015, 160, 2965-2977.	2.1	17
27	Colorado Tick Fever and Other Arthropod Borne Reoviridae. , 2016, , 841-852.		1
28	Article Commentary: Virus Discovery Using Tick Cell Lines. <i>Evolutionary Bioinformatics</i> , 2016, 12s2, EBO.S39675.	1.2	16
29	Identification of the Genome Segments of Bluetongue Virus Serotype 26 (Isolate KUW2010/02) that Restrict Replication in a <i>Culicoides sonorensis</i> Cell Line (KC Cells). <i>PLoS ONE</i> , 2016, 11, e0149709.	2.5	22
30	The evolution of transmission mode. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160083.	4.0	80
31	Occurrence and genetic variability of Kemerovo virus in <i>Ixodes</i> ticks from different regions of Western Siberia, Russia and Kazakhstan. <i>Infection, Genetics and Evolution</i> , 2017, 47, 56-63.	2.3	20
32	Viral population analysis of the taiga tick, <i>Ixodes persulcatus</i> , by using Batch Learning Self-Organizing Maps and BLAST search. <i>Journal of Veterinary Medical Science</i> , 2019, 81, 401-410.	0.9	30
33	Discovery and Characterization of Bukakata orbivirus (Reoviridae:Orbivirus), a Novel Virus from a Ugandan Bat. <i>Viruses</i> , 2019, 11, 209.	3.3	17
34	Failure to Remove Bluetongue Serotype 8 Virus (BTV-8) From in vitro Produced and in vivo Derived Bovine Embryos and Subsequent Transmission of BTV-8 to Recipient Cows After Embryo Transfer. <i>Frontiers in Veterinary Science</i> , 2019, 6, 432.	2.2	7
35	Genetic diversity of Kemerovo virus and phylogenetic relationships within the Great Island virus genetic group. <i>Ticks and Tick-borne Diseases</i> , 2020, 11, 101333.	2.7	4
36	Isolation and characterization of Wad Medani virus obtained in the tuva Republic of Russia. <i>Ticks and Tick-borne Diseases</i> , 2021, 12, 101612.	2.7	3
37	African horse sickness virus NS4 protein is an important virulence factor and interferes with JAK-STAT signaling during viral infection. <i>Virus Research</i> , 2021, 298, 198407.	2.2	5

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38	Whole Genome Sequencing Strategies and Development of Orbivirus Sequence Database: Implications for Novel dsRNA Virus Detection. , 2013, , 237-255.		1
39	Detection of a Fourth Orbivirus Non-Structural Protein. PLoS ONE, 2011, 6, e25697.	2.5	174
40	Full Genome Sequencing of Corriparta Virus, Identifies California Mosquito Pool Virus as a Member of the Corriparta virus Species. PLoS ONE, 2013, 8, e70779.	2.5	12
41	Full-Genome Characterisation of Orungo, Lebombo and Changuinola Viruses Provides Evidence for Co-Evolution of Orbiviruses with Their Arthropod Vectors. PLoS ONE, 2014, 9, e86392.	2.5	33
42	Full Genome Characterization of the Culicoides-Borne Marsupial Orbiviruses: Wallal Virus, Mudjinbarry Virus and Warrego Viruses. PLoS ONE, 2014, 9, e108379.	2.5	9
43	Orbiviruses. Livestock Diseases and Management, 2020, , 161-214.	0.5	1
45	Extended characterisation of five archival tick-borne viruses provides insights for virus discovery in Australian ticks. Parasites and Vectors, 2022, 15, 59.	2.5	2
46	Genome-scale molecular and phylogenetic characterization of Middle Point orbiviruses from Australia. Journal of General Virology, 2021, 102, .	2.9	0
47	Full-genome characterisation of a putative novel serotype of Yonaguni orbivirus isolated from cattle in Yunnan province, China. Virus Genes, 0, , .	1.6	1
48	Genetic Characterization of DH13M98, <i>Umatilla Virus</i> , Isolated from <i>Culex tritaeniorhynchus</i> Giles in Yunnan Province, China. Vector-Borne and Zoonotic Diseases, 2023, 23, 35-43.	1.5	1
49	The role of sand flies as vectors of viruses other than phleboviruses. Journal of General Virology, 2023, 104, .	2.9	4
50	Characterization of a Novel Orbivirus from Cattle Reveals Active Circulation of a Previously Unknown and Pathogenic Orbivirus in Ruminants in Kenya. MSphere, 2023, 8, .	2.9	1
51	Reoviruses: Colorado Tick Fever Virus and Other Vector-Borne Reoviruses. , 2023, , 1-26.		0
52	Orbivirus NS4 Proteins Play Multiple Roles to Dampen Cellular Responses. Viruses, 2023, 15, 1908.	3.3	0
53	Isolation of Epizootic Hemorrhagic Disease Virus Serotype 10 from <i>Culicoides tainanus</i> and Associated Infections in Livestock in Yunnan, China. Viruses, 2024, 16, 175.	3.3	0
54	Age- and Sex-Associated Pathogenesis of Cell Culture-Passaged Kemerovo Virus in IFNAR ^{−/−} Mice. International Journal of Molecular Sciences, 2024, 25, 3177.	4.1	0