## William C Wilson

List of Publications by Year in descending order

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132	4,255	35	57
papers	citations	h-index	g-index
137 all docs	137 docs citations	137 times ranked	3937 citing authors

#	Article	IF	CITATIONS
1	Infection and transmission of ancestral SARS-CoV-2 and its alpha variant in pregnant white-tailed deer. Emerging Microbes and Infections, 2022, 11, 95-112.	6.5	77
2	Detection of Rift Valley Fever Virus in Aedes (Aedimorphus) durbanensis, South Africa. Pathogens, 2022, 11, 125.	2.8	4
3	World Society for Virology first international conference: Tackling global virus epidemics. Virology, 2022, 566, 114-121.	2.4	2
4	Rift Valley fever virus Gn V5-epitope tagged virus enables identification of UBR4 as a Gn interacting protein that facilitates Rift Valley fever virus production. Virology, 2022, 567, 65-76.	2.4	3
5	Susceptibility of sheep to experimental co-infection with the ancestral lineage of SARS-CoV-2 and its alpha variant. Emerging Microbes and Infections, 2022, 11, 662-675.	6.5	21
6	Experimental re-infected cats do not transmit SARS-CoV-2. Emerging Microbes and Infections, 2021, 10, 638-650.	6.5	48
7	Susceptibility of Midge and Mosquito Vectors to SARS-CoV-2. Journal of Medical Entomology, 2021, 58, 1948-1951.	1.8	14
8	Mechanical transmission of SARS-CoV-2 by house flies. Parasites and Vectors, 2021, 14, 214.	2.5	30
9	Exposure of Culicoides sonorensis to Enzootic Strains of Bluetongue Virus Demonstrates Temperature- and Virus-Specific Effects on Virogenesis. Viruses, 2021, 13, 1016.	3.3	7
10	Preliminary Evaluation of a Recombinant Rift Valley Fever Virus Glycoprotein Subunit Vaccine Providing Full Protection against Heterologous Virulent Challenge in Cattle. Vaccines, 2021, 9, 748.	4.4	7
11	Large-Scale International Validation of an Indirect ELISA Based on Recombinant Nucleocapsid Protein of Rift Valley Fever Virus for the Detection of IgG Antibody in Domestic Ruminants. Viruses, 2021, 13, 1651.	3.3	1
12	High dose of vesicular stomatitis virus-vectored Ebola virus vaccine causes vesicular disease in swine without horizontal transmission. Emerging Microbes and Infections, 2021, 10, 651-663.	6.5	5
13	Perspectives on the Changing Landscape of Epizootic Hemorrhagic Disease Virus Control. Viruses, 2021, 13, 2268.	3.3	7
14	SARS-CoV-2 infection, disease and transmission in domestic cats. Emerging Microbes and Infections, 2020, 9, 2322-2332.	6.5	215
15	Editorial: Emerging Arboviruses. Frontiers in Veterinary Science, 2020, 7, 593872.	2.2	2
16	African Swine Fever Virus: An Emerging DNA Arbovirus. Frontiers in Veterinary Science, 2020, 7, 215.	2.2	211
17	Livestock Challenge Models of Rift Valley Fever for Agricultural Vaccine Testing. Frontiers in Veterinary Science, 2020, 7, 238.	2.2	7
18	Effect of Environmental Temperature on the Ability of <i>Culex tarsalis</i> and <i>Aedes taeniorhynchus</i> (Diptera: Culicidae) to Transmit Rift Valley Fever Virus. Vector-Borne and Zoonotic Diseases, 2020, 20, 454-460.	1.5	8

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19	Evaluation of A Baculovirus-Expressed VP2 Subunit Vaccine for the Protection of White-Tailed Deer (Odocoileus virginianus) from Epizootic Hemorrhagic Disease. Vaccines, 2020, 8, 59.	4.4	7
20	Evaluation of an Indirect Enzyme-Linked Immunosorbent Assay Based on Recombinant Baculovirus-Expressed Rift Valley Fever Virus Nucleoprotein as the Diagnostic Antigen. Journal of Clinical Microbiology, 2019, 57, .	3.9	8
21	Rift Valley Fever Virus: Propagation, Quantification, and Storage. Current Protocols in Microbiology, 2019, 55, e92.	6.5	17
22	Identification and evaluation of antivirals for Rift Valley fever virus. Veterinary Microbiology, 2019, 230, 110-116.	1.9	10
23	Interâ€serotype reassortment among epizootic haemorrhagic disease viruses in the United States. Transboundary and Emerging Diseases, 2019, 66, 1809-1820.	3.0	2
24	Complete Genome Sequence of a 2016 Bluetongue Virus Serotype 3 Isolate from Louisiana. Microbiology Resource Announcements, 2019, 8, .	0.6	0
25	EHDV-2 Infection Prevalence Varies in Culicoides sonorensis after Feeding on Infected White-Tailed Deer over the Course of Viremia. Viruses, 2019, 11, 371.	3.3	10
26	A multiplex fluorescence microsphere immunoassay for increased understanding of Rift Valley fever immune responses in ruminants in Kenya. Journal of Virological Methods, 2019, 269, 70-76.	2.1	5
27	Rift Valley Fever Viral RNA Detection by <i>In Situ</i> Hybridization in Formalin-Fixed, Paraffin-Embedded Tissues. Vector-Borne and Zoonotic Diseases, 2019, 19, 553-556.	1.5	10
28	Schmallenberg Diseaseâ€"A Newly Emerged Culicoides-Borne Viral Disease of Ruminants. Viruses, 2019, 11, 1065.	3.3	28
29	Molecular aspects of Rift Valley fever virus and the emergence of reassortants. Virus Genes, 2019, 55, 1-11.	1.6	40
30	Bluetongue and epizootic hemorrhagic disease viruses: recent developments with these globally re-emerging arboviral infections of ruminants. Current Opinion in Virology, 2019, 34, 56-62.	5.4	52
31	Evaluation of 2012 US EHDV-2 outbreak isolates for genetic determinants of cattle infection. Journal of General Virology, 2019, 100, 556-567.	2.9	4
32	Immunogenicity and efficacy of Schmallenberg virus envelope glycoprotein subunit vaccines. Journal of Veterinary Science, 2019, 20, e58.	1.3	5
33	Evaluation of Fluorescence Microsphere Immunoassay for Detection of Antibodies to Rift Valley Fever Virus Nucleocapsid Protein and Glycoproteins. Journal of Clinical Microbiology, 2018, 56, .	3.9	17
34	JARID2 Functions as a Tumor Suppressor in Myeloid Neoplasms by Repressing Self-Renewal in Hematopoietic Progenitor Cells. Cancer Cell, 2018, 34, 741-756.e8.	16.8	44
35	Virological and Serological Responses of Sheep and Cattle to Experimental Schmallenberg Virus Infection. Vector-Borne and Zoonotic Diseases, 2018, 18, 697-703.	1.5	4
36	Preliminary evaluation of diagnostic accuracy and precision of a competitive ELISA for detection of antibodies to Rift Valley fever virus in cattle and sheep sera. Journal of Virological Methods, 2018, 262, 6-11.	2.1	5

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37	Susceptibility of White-Tailed Deer to Rift Valley Fever Virus. Emerging Infectious Diseases, 2018, 24, 1717-1719.	4.3	31
38	Detection of Multiple Pathogens in Serum Using Silica-Encapsulated Nanotags in a Surface-Enhanced Raman Scattering-Based Immunoassay. Journal of Agricultural and Food Chemistry, 2018, 66, 5707-5712.	5.2	28
39	Recent US bluetongue virus serotype 3 isolates found outside of Florida indicate evidence of reassortment with co-circulating endemic serotypes. Journal of General Virology, 2018, 99, 157-168.	2.9	37
40	Comparison of two zoonotic viruses from the order Bunyavirales. Current Opinion in Virology, 2017, 27, 36-41.	5.4	10
41	Current Status of Rift Valley Fever Vaccine Development. Vaccines, 2017, 5, 29.	4.4	102
42	Distinct virulence of Rift Valley fever phlebovirus strains from different genetic lineages in a mouse model. PLoS ONE, 2017, 12, e0189250.	2.5	23
43	Experimental Infection of Calves by Two Genetically-Distinct Strains of Rift Valley Fever Virus. Viruses, 2016, 8, 145.	3.3	33
44	Multiplex Detection of IgG and IgM to Rift Valley Fever Virus Nucleoprotein, Nonstructural Proteins, and Glycoprotein in Ovine and Bovine. Vector-Borne and Zoonotic Diseases, 2016, 16, 550-557.	1.5	13
45	A Recombinant Rift Valley Fever Virus Glycoprotein Subunit Vaccine Confers Full Protection against Rift Valley Fever Challenge in Sheep. Scientific Reports, 2016, 6, 27719.	3.3	50
46	Collaborative Control of Cell Cycle Progression by the RNA Exonuclease Dis3 and Ras Is Conserved Across Species. Genetics, 2016, 203, 749-762.	2.9	19
47	Development of a sheep challenge model for Rift Valley fever. Virology, 2016, 489, 128-140.	2.4	38
48	Molecular evolution of epizootic hemorrhagic disease viruses in North America based on historical isolates using motif fingerprints. Virus Genes, 2016, 52, 495-508.	1.6	9
49	Mouse model for the Rift Valley fever virus MP12 strain infection. Veterinary Microbiology, 2016, 195, 70-77.	1.9	14
50	Complete Genome Sequence of Two Rift Valley Fever Virus Strains Isolated from Outbreaks in Saudi Arabia (2000) and Kenya (2006 to 2007). Genome Announcements, 2016, 4, .	0.8	7
51	Deletion of Rb1 induces both hyperproliferation and cell death in murine germinal center B cells. Experimental Hematology, 2016, 44, 161-165.e4.	0.4	9
52	Rift Valley Fever Virus., 2016,, 553-561.		0
53	Comparison of Rift Valley fever virus replication in North American livestock and wildlife cell lines. Frontiers in Microbiology, 2015, 6, 664.	3.5	30
54	Whole Genome Sequence of Multiple Myeloma-Prone C57BL/KaLwRij Mouse Strain Suggests the Origin of Disease Involves Multiple Cell Types. PLoS ONE, 2015, 10, e0127828.	2.5	26

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55	Whole genome sequence analysis of circulating <i>Bluetongue virus</i> serotype 11 strains from the United States including two domestic canine isolates. Journal of Veterinary Diagnostic Investigation, 2015, 27, 442-448.	1.1	10
56	Evaluation of the Efficacy, Potential for Vector Transmission, and Duration of Immunity of MP-12, an Attenuated Rift Valley Fever Virus Vaccine Candidate, in Sheep. Vaccine Journal, 2015, 22, 930-937.	3.1	27
57	Diagnostic Tools for Bluetongue and Epizootic Hemorrhagic Disease Viruses Applicable to North American Veterinary Diagnosticians. Vector-Borne and Zoonotic Diseases, 2015, 15, 364-373.	1.5	13
58	Orbiviruses: A North American Perspective. Vector-Borne and Zoonotic Diseases, 2015, 15, 335-338.	1.5	10
59	Genetic characterization of epizootic hemorrhagic disease virus strains isolated from cattle in Israel. Journal of General Virology, 2015, 96, 1400-1410.	2.9	9
60	Lesser-known bunyavirus infections. OIE Revue Scientifique Et Technique, 2015, 34, 419-429.	1.2	6
61	Whole Genome Sequence of Multiple Myelomaâ€Prone C57BL/KaLwRij Mouse Strain Suggests the Origin of Disease Involves Multiple Cell Types. FASEB Journal, 2015, 29, 926.9.	0.5	0
62	Molecular evolution of American field strains of bluetongue and epizootic haemorrhagic disease viruses. Veterinaria Italiana, 2015, 51, 269-73.	0.5	6
63	Rift Valley Fever Virus Incorporates the 78 kDa Glycoprotein into Virions Matured in Mosquito C6/36 Cells. PLoS ONE, 2014, 9, e87385.	2.5	50
64	A Glycoprotein Subunit Vaccine Elicits a Strong Rift Valley Fever Virus Neutralizing Antibody Response in Sheep. Vector-Borne and Zoonotic Diseases, 2014, 14, 746-756.	1.5	47
65	Development of a Rift Valley fever virus viremia challenge model in sheep and goats. Vaccine, 2014, 32, 2337-2344.	3.8	27
66	Evaluation of lamb and calf responses to Rift Valley fever MP-12 vaccination. Veterinary Microbiology, 2014, 172, 44-50.	1.9	28
67	Whole genome sequencing and phylogenetic analysis of <i>Bluetongue virus</i> serotype 2 strains isolated in the Americas including a novel strain from the western United States. Journal of Veterinary Diagnostic Investigation, 2014, 26, 553-557.	1.1	13
68	Efficacy of a recombinant Rift Valley fever virus MP-12 with NSm deletion as a vaccine candidate in sheep. Vaccine, 2014, 32, 2345-2349.	3.8	28
69	Development of a Rift Valley fever real-time RT-PCR assay that can detect all three genome segments. Journal of Virological Methods, 2013, 193, 426-431.	2.1	39
70	Experimental infection of white-tailed deer (Odocoileus virginianus) with Northern European bluetongue virus serotype 8. Veterinary Microbiology, 2013, 166, 347-355.	1.9	27
71	Rift Valley Fever Virus Structural and Nonstructural Proteins: Recombinant Protein Expression and Immunoreactivity Against Antisera from Sheep. Vector-Borne and Zoonotic Diseases, 2013, 13, 619-629.	1.5	33
72	Surface-enhanced Raman scattering (SERS) detection of multiple viral antigens using magnetic capture of SERS-active nanoparticles. Biosensors and Bioelectronics, 2013, 41, 316-321.	10.1	134

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73	Novel Serotype of Bluetongue Virus, Western North America. Emerging Infectious Diseases, 2013, 19, 665-6.	4.3	29
74	Rift Valley Fever Risk Map Model and Seroprevalence in Selected Wild Ungulates and Camels from Kenya. PLoS ONE, 2013, 8, e66626.	2.5	77
75	Diagnostic Approaches for Rift Valley Fever. Developments in Biologicals, 2013, 135, 73-78.	0.5	3
76	Surface-Enhanced Raman Scattering Detection of DNAs Derived from Virus Genomes Using Au-Coated Paramagnetic Nanoparticles. Langmuir, 2012, 28, 4030-4037.	3.5	65
77	Development and evaluation of one-step rRT-PCR and immunohistochemical methods for detection of Rift Valley fever virus in biosafety level 2 diagnostic laboratories. Journal of Virological Methods, 2012, 179, 373-382.	2.1	30
78	Surface-Enhanced Raman Scattering Detection of DNA Derived from the West Nile Virus Genome Using Magnetic Capture of Raman-Active Gold Nanoparticles. Analytical Chemistry, 2011, 83, 254-260.	6.5	119
79	Epizootic haemorragic disease. Research in Veterinary Science, 2011, 91, 1-17.	1.9	135
80	A versatile SERS-based immunoassay for immunoglobulin detection using antigen-coated gold nanoparticles and malachite green-conjugated protein A/G. Biosensors and Bioelectronics, 2010, 26, 1009-1015.	10.1	79
81	An improved Real-Time Polymerase Chain Reaction for the Simultaneous Detection of All Serotypes of <i>Epizootic Hemorrhagic Disease Virus</i> Journal of Veterinary Diagnostic Investigation, 2010, 22, 588-593.	1.1	30
82	Detection of a novel reassortant epizootic hemorrhagic disease virus (EHDV) in the USA containing RNA segments derived from both exotic (EHDV-6) and endemic (EHDV-2) serotypes. Journal of General Virology, 2010, 91, 430-439.	2.9	84
83	Potential for North American Mosquitoes (Diptera: Culicidae) to Transmit Rift Valley Fever Virus. Journal of Medical Entomology, 2010, 47, 884-889.	1.8	99
84	Investigation of a bluetongue disease epizootic caused by bluetongue virus serotype 17 in sheep in Wyoming. Journal of the American Veterinary Medical Association, 2010, 237, 955-959.	0.5	16
85	Potential for North American Mosquitoes (Diptera: Culicidae) to Transmit Rift Valley Fever Virus. Journal of Medical Entomology, 2010, 47, 884-889.	1.8	78
86	Field Evaluation of a Multiplex Real-Time Reverse Transcription Polymerase Chain Reaction Assay for Detection of <i>Vesicular Stomatitis Virus</i> Journal of Veterinary Diagnostic Investigation, 2009, 21, 179-186.	1.1	23
87	A Multiplex Real-Time Reverse Transcription Polymerase Chain Reaction Assay for Detection and Differentiation of <i>Bluetongue Virus </i> and <i>Epizootic Hemorrhagic Disease Virus </i> Journal of Veterinary Diagnostic Investigation, 2009, 21, 760-770.	1.1	34
88	Detection of All Eight Serotypes of <i>Epizootic Hemorrhagic Disease Virus </i> by Real-Time Reverse Transcription Polymerase Chain Reaction. Journal of Veterinary Diagnostic Investigation, 2009, 21, 220-225.	1.1	28
89	SERS detection of indirect viral DNA capture using colloidal gold and methylene blue as a Raman label. Biosensors and Bioelectronics, 2009, 25, 674-681.	10.1	62
90	Current status of bluetongue virus in the Americas. , 2009, , 197-221.		6

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91	The NS3 proteins of global strains of bluetongue virus evolve into regional topotypes through negative (purifying) selection. Veterinary Microbiology, 2008, 126, 91-100.	1.9	67
92	Bluetongue virus serotype 17 sequence variation associated with neutralization. DNA Sequence, 2008, 19, 237-240.	0.7	11
93	Developing a Research Agenda and a Comprehensive National Prevention and Response Plan for Rift Valley Fever in the United States. Emerging Infectious Diseases, 2007, 13, e1-e1.	4.3	12
94	A Rift Valley fever risk surveillance system for Africa using remotely sensed data: potential for use on other continents. Veterinaria Italiana, 2007, 43, 663-74.	0.5	25
95	Midgut and salivary gland transcriptomes of the arbovirus vector <i>Culicoides sonorensis</i> (Diptera: Ceratopogonidae). Insect Molecular Biology, 2005, 14, 121-136.	2.0	81
96	Studies on overwintering of bluetongue viruses in insects. Journal of General Virology, 2005, 86, 453-462.	2.9	69
97	Vector Competence of <i>Culicoides sonorensis</i> (Diptera: Ceratopogonidae) for Vesicular Stomatitis Virus. Journal of Medical Entomology, 2005, 42, 409-418.	1.8	42
98	Culture-Independent Analysis of Midgut Microbiota in the Arbovirus Vector <i>Culicoides sonorensis</i> (Diptera: Ceratopogonidae). Journal of Medical Entomology, 2004, 41, 340-348.	1.8	69
99	Antigen Capture Competitive Enzyme-Linked Immunosorbent Assays Using Baculovirus-Expressed Antigens for Diagnosis of Bluetongue Virus and Epizootic Hemorrhagic Disease Virus. Journal of Clinical Microbiology, 2004, 42, 518-523.	3.9	46
100	The S7 gene and VP7 protein are highly conserved among temporally and geographically distinct American isolates of epizootic hemorrhagic disease virus. Virus Research, 2003, 94, 129-133.	2.2	18
101	Grasshoppers (Orthoptera: Acrididae) Could Serve as Reservoirs and Vectors of Vesicular Stomatitis Virus. Journal of Medical Entomology, 2003, 40, 957-963.	1.8	33
102	Replication of bluetongue virus and epizootic hemorrhagic disease virus in pulmonary artery endothelial cells obtained from cattle, sheep, and deer. American Journal of Veterinary Research, 2003, 64, 860-865.	0.6	18
103	Differentially expressed midgut transcripts in Culicoides sonorensis (Diptera: Ceratopogonidae) following Orbivirus (Reoviridae) oral feeding. Insect Molecular Biology, 2002, 11, 595-604.	2.0	16
104	Verification of bluetongue virus S9 segment nucleotide sequences. Virus Research, 2001, 81, 93-101.	2.2	0
105	Validation of a reverse transcriptase multiplex PCR test for the serotype determination of U.S. isolates of bluetongue virus. Veterinary Microbiology, 2000, 76, 105-115.	1.9	26
106	Phylogenetic analysis of the S7 gene does not segregate Chinese strains of bluetongue virus into a single topotype. Archives of Virology, 2000, 145, 1163-1171.	2.1	36
107	Phylogenetic relationships of bluetongue viruses based on gene S7. Virus Research, 2000, 67, 141-151.	2.2	61
108	Preliminary Description of a Polymerase Chain Reaction Test for Bluetongue and Epizootic Hemorrhagic Disease Viral RNA in Bovine Semen. Journal of Veterinary Diagnostic Investigation, 1999, 11, 377-379.	1.1	6

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109	Activity Pattern Analysis by Means of Sequence-Alignment Methods. Environment and Planning A, 1998, 30, 1017-1038.	3.6	128
110	PCR Detection of North American and Central African Isolates of Epizootic Hemorrhagic Disease Virus (EHDV) Based on Genome Segment 10 of EHDV Serotype 1. Journal of Clinical Microbiology, 1998, 36, 2604-2608.	3.9	12
111	Epizootic Hemorrhagic Disease: Analysis of Tissues by Amplification and In Situ Hybridization Reveals Widespread Orbivirus Infection at Low Copy Numbers. Journal of Virology, 1998, 72, 3863-3871.	3.4	28
112	The Effects of Pharmacological and Lentivirus-Induced Immune Suppression on Orbivirus Pathogenesis: Assessment of Virus Burden in Blood Monocytes and Tissues by Reverse Transcription-In Situ PCR. Journal of Virology, 1998, 72, 5599-5609.	3.4	25
113	Bluetongue Virus Detection: A Safer Reverse-Transcriptase Polymerase Chain Reaction for Prediction of Viremia in Sheep. Journal of Veterinary Diagnostic Investigation, 1997, 9, 118-124.	1.1	26
114	VP7: an attachment protein of bluetongue virus for cellular receptors in Culicoides variipennis Journal of General Virology, 1997, 78, 1617-1623.	2.9	53
115	Sequence and Cognitive Analyses of Two Virulence-Associated Markers of Bluetongue Virus Serotype 17. Intervirology, 1997, 40, 226-231.	2.8	5
116	Molecular Characterization of the Segment 2 Gene of Epizootic Hemorrhagic Disease Virus Serotype 2: Gene Sequence and Genetic Diversity. Virology, 1996, 224, 555-560.	2.4	25
117	A model for the membrane topology of the NS3 protein as predicted from the sequence of segment 10 of epizootic haemorrhagic disease virus serotype 1. Archives of Virology, 1995, 140, 799-805.	2.1	24
118	Development of polymerase chain reaction for specific identification of epizootic hemorrhagic disease virus serotype 1. Archives of Virology, 1995, 140, 2273-2281.	2.1	18
119	Application of PCR for Specific Identification of Epizootic Hemorrhagic Disease Virus Serotype 2. Journal of Veterinary Diagnostic Investigation, 1995, 7, 388-392.	1.1	13
120	Complete nucleotide sequence of RNA segment 3 of bluetongue virus serotype 2 (Ona-A). Phylogenetic analyses reveal the probable origin and relationship with other orbiviruses. Virus Research, 1995, 35, 247-261.	2.2	52
121	Geographical genetic variation in the gene encoding VP3 from the Alberta isolate of epizootic hemorrhagic disease virus. Virus Research, 1995, 36, 279-286.	2.2	22
122	Development of a nested-PCR test based on sequence analysis of epizootic hemorrhagic disease viruses non-structural protein 1 (NS1). Virus Research, 1994, 31, 357-365.	2.2	37
123	Sequence analysis of the non-structural protein 2 from epizootic hemorrhagic disease viruses. Virus Research, 1994, 34, 63-68.	2.2	13
124	The smallest gene of the orbivirus, epizootic hemorrhagic disease, is expressed m virus-infected cells as two proteins and the expression differs from that of the cognate gene of bluetongue virus. Virus Research, 1994, 32, 353-364.	2.2	25
125	Nested and multiplex polymerase chain reactions for the identification of bluetongue virus infection in the biting midge, Culicoides variipennis. Journal of Virological Methods, 1993, 45, 39-47.	2.1	46

Bluetongue Virus in Sheep and Cattle and Culicoides variipennis and C. stellifer (Diptera:) Tj ETQq $0\ 0\ 0$  rgBT /Overlock  $10\ Tf\ 50\ 62\ Td$  (Control of the control o

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127	Molecular comparison of VP3 from bluetongue and epizootic hemorrhagic disease viruses. Virus Research, 1991, 21, 225-236.	2.2	21
128	A RNA virus in cells from Culicoides variipennis. Journal of Invertebrate Pathology, 1991, 57, 200-205.	3.2	30
129	Detection of Epizootic Hemorrhagic Disease Virus in Culicoides variipennis (Diptera:) Tj ETQq1 1 0.784314 rgBT	/Oyerlock	10 <sub>11</sub> f 50 662
130	Cytokine Modulation of the Interaction Between Bluetongue Virus and Endothelial Cells in vitro. Veterinary Pathology, 1991, 28, 524-532.	1.7	23
131	Limits of Detection of Bluetongue Virus with Different Assay Systems. Journal of Veterinary Diagnostic Investigation, 1990, 2, 103-106.	1.1	12
132	Development and optimization of a hybridization assay for epizootic hemorrhagic disease viruses. Journal of Virological Methods, 1990, 30, 173-181.	2.1	10