

Peter J Walker

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

133
papers

5,534
citations

42
h-index

69
g-index

139
ext. papers

6,720
ext. citations

4.1
avg, IF

5.4
L-index

#	Paper	IF	Citations
133	Differentiating between viruses and virus species by writing their names correctly.. <i>Archives of Virology</i> , 2022 , 167, 1231	2.6	5
132	Revised Taxonomy of Rhabdoviruses Infecting Fish and Marine Mammals. <i>Animals</i> , 2022 , 12, 1363	3.1	0
131	ICTV Virus Taxonomy Profile: Rhabdoviridae 2022. <i>Journal of General Virology</i> , 2022 , 103,	4.9	8
130	Distribution of Culicoides biting midges (Diptera: Ceratopogonidae) in southern Australia and insight into the Culicoides victoriae morpho-variants. <i>Austral Entomology</i> , 2021 , 60, 525-534	1.1	
129	Vesicular Stomatitis Virus and Bovine Ephemeral Fever Virus (Rhabdoviridae) 2021 , 875-883		
128	Changes to virus taxonomy and to the International Code of Virus Classification and Nomenclature ratified by the International Committee on Taxonomy of Viruses (2021). <i>Archives of Virology</i> , 2021 , 166, 2633-2648	2.6	52
127	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2021 , 166, 3513-3566	2.6	10
126	ICTV Virus Taxonomy Profile:. <i>Journal of General Virology</i> , 2021 , 102,	4.9	1
125	Experimental bluetongue virus infection of Culicoides austropalpalis, collected from a farm environment in Victoria, Australia.. <i>Veterinaria Italiana</i> , 2021 , 57, 341-345	1	
124	Hayes Yard virus: a novel ephemerovirus isolated from a bull with severe clinical signs of bovine ephemeral fever is most closely related to Puchong virus. <i>Veterinary Research</i> , 2020 , 51, 58	3.8	3
123	Binomial nomenclature for virus species: a consultation. <i>Archives of Virology</i> , 2020 , 165, 519-525	2.6	27
122	Characterization of Port Bolivar Virus, a Novel Entomobirnavirus) Isolated from Mosquitoes Collected in East Texas, USA. <i>Viruses</i> , 2020 , 12,	6.2	4
121	Taxonomy of the order Mononegavirales: second update 2018. <i>Archives of Virology</i> , 2019 , 164, 1233-1244.6	4.6	50
120	Additional changes to taxonomy ratified in a special vote by the International Committee on Taxonomy of Viruses (October 2018). <i>Archives of Virology</i> , 2019 , 164, 943-946	2.6	66
119	Exploiting the Legacy of the Arbovirus Hunters. <i>Viruses</i> , 2019 , 11,	6.2	7
118	Changes to virus taxonomy and the International Code of Virus Classification and Nomenclature ratified by the International Committee on Taxonomy of Viruses (2019). <i>Archives of Virology</i> , 2019 , 164, 2417-2429	2.6	171
117	Taxonomy of the order Mononegavirales: update 2019. <i>Archives of Virology</i> , 2019 , 164, 1967-1980	2.6	133

116	Evolutionary history of Simbu serogroup orthobunyaviruses in the Australian episystem. <i>Virology</i> , 2019 , 535, 32-44	3.6	7
115	Characterization of Three Novel Viruses from the Families , and , Isolated from Dead Birds Collected during West Nile Virus Surveillance in Harris County, Texas. <i>Viruses</i> , 2019 , 11,	6.2	4
114	Genomic characterisation of Cuiaba and Charleville viruses: arboviruses (family Rhabdoviridae, genus Sripuvirus) infecting reptiles and amphibians. <i>Virus Genes</i> , 2019 , 55, 87-94	2.3	7
113	Taxonomy of the Family Arenaviridae and the order Bunyvirales: update 2018. <i>Archives of Virology</i> , 2018 , 163, 2295-2310	2.6	108
112	Taxonomy of the order Mononegavirales: update 2018. <i>Archives of Virology</i> , 2018 , 163, 2283-2294	2.6	111
111	ICTV Virus Taxonomy Profile: Rhabdoviridae. <i>Journal of General Virology</i> , 2018 , 99, 447-448	4.9	110
110	Taxonomy of the order Mononegavirales: update 2017. <i>Archives of Virology</i> , 2017 , 162, 2493-2504	2.6	137
109	A large-scale serological survey of Akabane virus infection in cattle, yak, sheep and goats in China. <i>Veterinary Microbiology</i> , 2017 , 207, 7-12	3.3	9
108	Genomic analysis of bluetongue virus episystems in Australia and Indonesia. <i>Veterinary Research</i> , 2017 , 48, 82	3.8	11
107	Possibility and Challenges of Conversion of Current Virus Species Names to Linnaean Binomials. <i>Systematic Biology</i> , 2017 , 66, 463-473	8.4	12
106	Beatrice Hill Virus Represents a Novel Species in the Genus Tibrovirus (Mononegavirales: Rhabdoviridae). <i>Genome Announcements</i> , 2017 , 5,		3
105	Almendravirus: A Proposed New Genus of Rhabdoviruses Isolated from Mosquitoes in Tropical Regions of the Americas. <i>American Journal of Tropical Medicine and Hygiene</i> , 2017 , 96, 100-109	3.2	20
104	Detection of Elizabethkingia spp. in Culicoides Biting Midges, Australia. <i>Emerging Infectious Diseases</i> , 2017 , 23, 1409-1410	10.2	4
103	A Global Genomic Characterization of Nairoviruses Identifies Nine Discrete Genogroups with Distinctive Structural Characteristics and Host-Vector Associations. <i>American Journal of Tropical Medicine and Hygiene</i> , 2016 , 94, 1107-1122	3.2	32
102	Genotyping of whole genome amplified reduced representation libraries reveals a cryptic population of Culicoides brevitarsis in the Northern Territory, Australia. <i>BMC Genomics</i> , 2016 , 17, 769	4.5	9
101	Taxonomy of the order Mononegavirales: update 2016. <i>Archives of Virology</i> , 2016 , 161, 2351-60	2.6	324
100	Discovery of novel rhabdoviruses in the blood of healthy individuals from West Africa. <i>PLoS Neglected Tropical Diseases</i> , 2015 , 9, e0003631	4.8	36
99	Evolution of genome size and complexity in the rhabdoviridae. <i>PLoS Pathogens</i> , 2015 , 11, e1004664	7.6	103

98	Detection of Low-Level Cardinium and Wolbachia Infections in Culicoides. <i>Applied and Environmental Microbiology</i> , 2015 , 81, 6177-88	4.8	42
97	Genomic Characterization of Yogue, Kasokero, Issyk-Kul, Keterah, Gossas, and Thiafora Viruses: Nairoviruses Naturally Infecting Bats, Shrews, and Ticks. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015 , 93, 1041-51	3.2	28
96	Ledantevirus: a proposed new genus in the Rhabdoviridae has a strong ecological association with bats. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015 , 92, 405-10	3.2	23
95	Delineation of the population genetic structure of <i>Culicoides imicola</i> in East and South Africa. <i>Parasites and Vectors</i> , 2015 , 8, 660	4	8
94	Epidemiology and control of bovine ephemeral fever. <i>Veterinary Research</i> , 2015 , 46, 124	3.8	50
93	Assessment of population genetic structure in the arbovirus vector midge, <i>Culicoides brevitarsis</i> (Diptera: Ceratopogonidae), using multi-locus DNA microsatellites. <i>Veterinary Research</i> , 2015 , 46, 108	3.8	10
92	Wongabel rhabdovirus accessory protein U3 targets the SWI/SNF chromatin remodeling complex. <i>Journal of Virology</i> , 2015 , 89, 1377-88	6.6	1
91	Cullin4 Is Pro-Viral during West Nile Virus Infection of <i>Culex</i> Mosquitoes. <i>PLoS Pathogens</i> , 2015 , 11, e1005643	5.43	23
90	Evolution of bovine ephemeral fever virus in the Australian epizootic system. <i>Journal of Virology</i> , 2014 , 88, 1525-35	3.5	35
89	Bovine ephemeral fever rhabdovirus σ protein has viroporin-like properties and binds importin β and importin γ . <i>Journal of Virology</i> , 2014 , 88, 1591-603	6.6	32
88	Evolution of bluetongue virus serotype 1 in northern Australia over 30 years. <i>Journal of Virology</i> , 2014 , 88, 13981-9	6.6	22
87	Stability of the WSSV ORF94 VNTR genotype marker during passage in marine shrimp, freshwater crayfish and freshwater prawns. <i>Diseases of Aquatic Organisms</i> , 2014 , 111, 249-57	1.7	2
86	Long-distance aerial dispersal modelling of <i>Culicoides</i> biting midges: case studies of incursions into Australia. <i>BMC Veterinary Research</i> , 2014 , 10, 135	2.7	32
85	Arboretum and Puerto Almendras viruses: two novel rhabdoviruses isolated from mosquitoes in Peru. <i>Journal of General Virology</i> , 2014 , 95, 787-792	4.9	34
84	Mesoniviruses are mosquito-specific viruses with extensive geographic distribution and host range. <i>Virology Journal</i> , 2014 , 11, 97	6.1	52
83	Characteristics of cyprinid herpesvirus 3 in different phases of infection: implications for disease transmission and control. <i>Virus Research</i> , 2014 , 188, 45-53	6.4	24
82	Low frequency of paleoviral infiltration across the avian phylogeny. <i>Genome Biology</i> , 2014 , 15, 539	18.3	43
81	Dicer-2-dependent activation of <i>Culex Vago</i> occurs via the TRAF-Rel2 signaling pathway. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e2823	4.8	75

80	Seasonal drivers of the epidemiology of arthropod-borne viruses in Australia. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e3325	4.8	26
79	Koolpinyah and Yata viruses: two newly recognised ephemeroviruses from tropical regions of Australia and Africa. <i>Veterinary Microbiology</i> , 2014 , 174, 547-553	3.3	7
78	Characterization of Farmington virus, a novel virus from birds that is distantly related to members of the family Rhabdoviridae. <i>Virology Journal</i> , 2013 , 10, 219	6.1	11
77	Malpais spring virus is a new species in the genus vesiculovirus. <i>Virology Journal</i> , 2013 , 10, 69	6.1	10
76	Niakha virus: a novel member of the family Rhabdoviridae isolated from phlebotomine sandflies in Senegal. <i>Virology</i> , 2013 , 444, 80-9	3.6	20
75	Kolente virus, a rhabdovirus species isolated from ticks and bats in the Republic of Guinea. <i>Journal of General Virology</i> , 2013 , 94, 2609-2615	4.9	23
74	Modelling spatio-temporal patterns of long-distance Culicoides dispersal into northern Australia. <i>Preventive Veterinary Medicine</i> , 2013 , 110, 312-22	3.1	37
73	A reverse-transcription PCR method for detecting all known ephemeroviruses in clinical samples. <i>Journal of Virological Methods</i> , 2013 , 191, 128-35	2.6	8
72	Kotonkan and Obodhiang viruses: African ephemeroviruses with large and complex genomes. <i>Virology</i> , 2012 , 425, 143-53	3.6	21
71	Evaluation of long-distance dispersal of Culicoides midges into northern Australia using a migration model. <i>Medical and Veterinary Entomology</i> , 2012 , 26, 334-40	2.4	30
70	Secreted Vago restricts West Nile virus infection in Culex mosquito cells by activating the Jak-STAT pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 18915-20	11.5	206
69	Koi herpesvirus encodes and expresses a functional interleukin-10. <i>Journal of Virology</i> , 2012 , 86, 11512-20	206	29
68	Malakal virus from Africa and Kimberley virus from Australia are geographic variants of a widely distributed ephemerovirus. <i>Virology</i> , 2012 , 433, 236-44	3.6	14
67	Genomic sequences of Australian bluetongue virus prototype serotypes reveal global relationships and possible routes of entry into Australia. <i>Journal of Virology</i> , 2012 , 86, 6724-31	6.6	48
66	Longitudinal disease studies in small-holder black tiger shrimp (<i>Penaeus monodon</i>) farms in Andhra Pradesh, India. I. High prevalence of WSSV infection and low incidence of disease outbreaks in BMP ponds. <i>Aquaculture</i> , 2011 , 318, 277-282	4.4	8
65	Longitudinal disease studies in small-holder black tiger shrimp (<i>Penaeus monodon</i>) ponds in Andhra Pradesh, India. II. Multiple WSSV genotypes associated with disease outbreaks in ponds seeded with uninfected postlarvae. <i>Aquaculture</i> , 2011 , 319, 18-24	4.4	19
64	Longitudinal disease studies in small-holder black tiger shrimp (<i>Penaeus monodon</i>) farms in Andhra Pradesh, India. III. A complex dynamic of WSSV infection and WSSV genotype distribution in farmed shrimp and wild crustaceans. <i>Aquaculture</i> , 2011 , 319, 319-327	4.4	11
63	Rhabdovirus accessory genes. <i>Virus Research</i> , 2011 , 162, 110-25	6.4	118

62	Mass extinctions, biodiversity and mitochondrial function: are bats special reservoirs for emerging viruses?. <i>Current Opinion in Virology</i> , 2011 , 1, 649-57	7.5	120
61	Isolation and characterization of koi herpesvirus (KHV) from Indonesia: identification of a new genetic lineage. <i>Journal of Fish Diseases</i> , 2011 , 34, 87-101	2.6	45
60	RNA-binding domain in the nucleocapsid protein of gill-associated nidovirus of penaeid shrimp. <i>PLoS ONE</i> , 2011 , 6, e22156	3.7	6
59	Pathogenicity of gill-associated virus and Mourilyan virus during mixed infections of black tiger shrimp (<i>Penaeus monodon</i>). <i>Journal of General Virology</i> , 2011 , 92, 893-901	4.9	10
58	Glycosylation of gp116 and gp64 envelope proteins of yellow head virus of <i>Penaeus monodon</i> shrimp. <i>Journal of General Virology</i> , 2010 , 91, 2463-73	4.9	11
57	Emerging viral diseases of fish and shrimp. <i>Veterinary Research</i> , 2010 , 41, 51	3.8	248
56	A proposal to change existing virus species names to non-Latinized binomials. <i>Archives of Virology</i> , 2010 , 155, 1909-19	2.6	24
55	A consensus real-time RT-PCR for detection of all genotypic variants of yellow head virus of penaeid shrimp. <i>Journal of Virological Methods</i> , 2010 , 167, 5-9	2.6	5
54	Ngaingan virus, a macropod-associated rhabdovirus, contains a second glycoprotein gene and seven novel open reading frames. <i>Virology</i> , 2010 , 399, 98-108	3.6	27
53	A virulent isolate of yellow head nidovirus contains a deformed envelope glycoprotein gp116. <i>Virology</i> , 2009 , 384, 192-200	3.6	24
52	Homologous genetic recombination in the yellow head complex of nidoviruses infecting <i>Penaeus monodon</i> shrimp. <i>Virology</i> , 2009 , 390, 79-88	3.6	20
51	Viral disease emergence in shrimp aquaculture: origins, impact and the effectiveness of health management strategies. <i>Reviews in Aquaculture</i> , 2009 , 1, 125-154	8.9	141
50	Detection of Laem-Singh virus in cultured <i>Penaeus monodon</i> shrimp from several sites in the Indo-Pacific region. <i>Diseases of Aquatic Organisms</i> , 2009 , 84, 195-200	1.7	12
49	Genetic diversity in the yellow head nidovirus complex. <i>Virology</i> , 2008 , 380, 213-25	3.6	58
48	Consensus RT-nested PCR detection of yellow head complex genotypes in penaeid shrimp. <i>Journal of Virological Methods</i> , 2008 , 153, 168-75	2.6	14
47	RNA transcription analysis and completion of the genome sequence of yellow head nidovirus. <i>Virus Research</i> , 2008 , 136, 157-65	6.4	47
46	A key gene of the RNA interference pathway in the black tiger shrimp, <i>Penaeus monodon</i> : identification and functional characterisation of Dicer-1. <i>Fish and Shellfish Immunology</i> , 2008 , 24, 223-33	4.3	76
45	Antigenic characterization of bovine ephemeral fever rhabdovirus G and GNS glycoproteins expressed from recombinant baculoviruses. <i>Archives of Virology</i> , 2008 , 153, 1657-65	2.6	14

44	A TaqMan real-time RT-PCR for quantifying Mourilyan virus infection levels in penaeid shrimp tissues. <i>Journal of Virological Methods</i> , 2006 , 137, 265-71	2.6	14
43	Identification of the nucleocapsid, tegument, and envelope proteins of the shrimp white spot syndrome virus virion. <i>Journal of Virology</i> , 2006 , 80, 3021-9	6.6	155
42	Structural and antigenic analysis of the yellow head virus nucleocapsid protein p20. <i>Virus Research</i> , 2006 , 116, 21-9	6.4	21
41	Phylogenetic relationships among rhabdoviruses inferred using the L polymerase gene. <i>Journal of General Virology</i> , 2005 , 86, 2849-2858	4.9	117
40	RT-nested PCR detection of Mourilyan virus in Australian <i>Penaeus monodon</i> and its tissue distribution in healthy and moribund prawns. <i>Diseases of Aquatic Organisms</i> , 2005 , 66, 91-104	1.7	20
39	Genomic organization, biology, and diagnosis of Taura syndrome virus and yellowhead virus of penaeid shrimp. <i>Advances in Virus Research</i> , 2004 , 63, 353-421	10.7	33
38	The gene encoding the nucleocapsid protein of Gill-associated nidovirus of <i>Penaeus monodon</i> prawns is located upstream of the glycoprotein gene. <i>Journal of Virology</i> , 2004 , 78, 8935-41	6.6	25
37	Multiplex RT-nested PCR differentiation of gill-associated virus (Australia) from yellow head virus (Thailand) of <i>Penaeus monodon</i> . <i>Journal of Virological Methods</i> , 2004 , 117, 49-59	2.6	39
36	Detection and differentiation of yellow head complex viruses using monoclonal antibodies. <i>Diseases of Aquatic Organisms</i> , 2003 , 57, 193-200	1.7	43
35	Identification and analysis of gp116 and gp64 structural glycoproteins of yellow head nidovirus of <i>Penaeus monodon</i> shrimp. <i>Journal of General Virology</i> , 2003 , 84, 863-873	4.9	51
34	High variation in repetitive DNA fragment length for white spot syndrome virus (WSSV) isolates in Thailand. <i>Diseases of Aquatic Organisms</i> , 2003 , 54, 253-7	1.7	45
33	Detection of gill-associated virus (GAV) by in situ hybridization during acute and chronic infections of <i>Penaeus monodon</i> and <i>P. esculentus</i> . <i>Diseases of Aquatic Organisms</i> , 2003 , 56, 1-10	1.7	25
32	The complete genome sequence of gill-associated virus of <i>Penaeus monodon</i> prawns indicates a gene organisation unique among nidoviruses. <i>Archives of Virology</i> , 2002 , 147, 1977-87	2.6	64
31	Complete ORF1b-gene sequence indicates yellow head virus is an invertebrate nidovirus. <i>Diseases of Aquatic Organisms</i> , 2002 , 50, 87-93	1.7	50
30	Vertical transmission of gill-associated virus (GAV) in the black tiger prawn <i>Penaeus monodon</i> . <i>Diseases of Aquatic Organisms</i> , 2002 , 50, 95-104	1.7	40
29	Gill-associated nidovirus of <i>Penaeus monodon</i> prawns transcribes 3'coterminal subgenomic mRNAs that do not possess 5'leader sequences. <i>Journal of General Virology</i> , 2002 , 83, 927-935	4.9	53
28	RNA polymerase (L) gene and genome terminal sequences of ephemeroviruses bovine ephemeral fever virus and Adelaide River virus indicate a close relationship to vesiculoviruses. <i>Virus Research</i> , 2000 , 70, 87-95	6.4	25
27	Detection of Australian gill-associated virus (GAV) and lymphoid organ virus (LOV) of <i>Penaeus monodon</i> by RT-nested PCR. <i>Diseases of Aquatic Organisms</i> , 2000 , 39, 159-67	1.7	42

26	Differences in the susceptibility of some penaeid prawn species to gill-associated virus (GAV) infection. <i>Diseases of Aquatic Organisms</i> , 2000 , 42, 221-5	1.7	28
25	Gill-associated virus of <i>Penaeus monodon</i> prawns: an invertebrate virus with ORF1a and ORF1b genes related to arteri- and coronaviruses. <i>Journal of General Virology</i> , 2000 , 81, 1473-84	4.9	114
24	Yellow head virus from Thailand and gill-associated virus from Australia are closely related but distinct prawn viruses. <i>Diseases of Aquatic Organisms</i> , 1999 , 36, 153-7	1.7	54
23	A yellow-head like virus from <i>Penaeus monodon</i> cultured in Australia. <i>Diseases of Aquatic Organisms</i> , 1997 , 31, 169-179	1.7	75
22	Vaccinia virus-expressed bovine ephemeral fever virus G but not G(NS) glycoprotein induces neutralizing antibodies and protects against experimental infection. <i>Journal of General Virology</i> , 1996 , 77 (Pt 4), 631-40	4.9	29
21	Adelaide River virus nucleoprotein gene: analysis of phylogenetic relationships of ephemeroviruses and other rhabdoviruses. <i>Journal of General Virology</i> , 1995 , 76 (Pt 4), 995-9	4.9	15
20	Structural and antigenic analysis of the nucleoprotein of bovine ephemeral fever rhabdovirus. <i>Journal of General Virology</i> , 1994 , 75 (Pt 8), 1889-99	4.9	36
19	Complex genome organization in the GNS-L intergenic region of Adelaide River rhabdovirus. <i>Virology</i> , 1994 , 203, 63-72	3.6	31
18	Effective vaccination of cattle using the virion G protein of bovine ephemeral fever virus as an antigen. <i>Vaccine</i> , 1994 , 12, 845-50	4.1	24
17	Adelaide river rhabdovirus expresses consecutive glycoprotein genes as polycistronic mRNAs: new evidence of gene duplication as an evolutionary process. <i>Virology</i> , 1993 , 195, 719-31	3.6	31
16	Infectivity of bovine leukaemia virus infected cattle: an ELISA for detecting antigens expressed in vitro cultured lymphocytes. <i>Veterinary Microbiology</i> , 1992 , 30, 137-50	3.3	16
15	A blocking ELISA for the detection of specific antibodies to bovine ephemeral fever virus. <i>Journal of Immunological Methods</i> , 1992 , 151, 289-97	2.5	14
14	The genome of bovine ephemeral fever rhabdovirus contains two related glycoprotein genes. <i>Virology</i> , 1992 , 191, 49-61	3.6	63
13	Proteins of bovine ephemeral fever virus. <i>Journal of General Virology</i> , 1991 , 72 (Pt 1), 67-74	4.9	53
12	Mapping of antigenic sites on the bovine ephemeral fever virus glycoprotein using monoclonal antibodies. <i>Journal of General Virology</i> , 1990 , 71 (Pt 9), 2065-72	4.9	39
11	An enzyme-linked immunosorbent assay for detection of bovine leukaemia virus p24 antibody in cattle. <i>Journal of Virological Methods</i> , 1990 , 28, 47-57	2.6	15
10	Bovine ephemeral fever and rhabdoviruses endemic to Australia. <i>Australian Veterinary Journal</i> , 1989 , 66, 398-400	1.2	9
9	Detection of reassortant orbiviruses (Wallal serogroup) in a prototype strain isolated from a pool of biting midges (<i>Culicoides dycei</i>). <i>Journal of General Virology</i> , 1989 , 70 (Pt 4), 1011-6	4.9	1

8	Variation in dengue type 2 viruses isolated in Bangkok during 1980. <i>Journal of General Virology</i> , 1988 , 69 (Pt 3), 591-602	4.9	23
7	A protein immunoblot test for detection of bovine leukemia virus p24 antibody in cattle and experimentally infected sheep. <i>Journal of Virological Methods</i> , 1987 , 15, 201-11	2.6	37
6	Orbiviruses 1983 , 287-357		27
5	Comparison of bluetongue type 20 with certain viruses of the bluetongue and Eubenangee serological groups of orbiviruses. <i>Journal of General Virology</i> , 1981 , 57, 251-61	4.9	40
4	Genetic Analysis of Orbiviruses by Using RNase T(1) Oligonucleotide Fingerprints. <i>Journal of Virology</i> , 1980 , 34, 583-91	6.6	23
3	The isolation of recombinants between related orbiviruses. <i>Journal of General Virology</i> , 1978 , 41, 333-424.9	4.9	36
2	Electrophoretic separation of double-stranded RNA genome segments from Warrego and Mitchell River viruses. <i>Archives of Virology</i> , 1977 , 54, 153-8	2.6	12
1	Molecular Biology and Pathogenesis of Roniviruses361-377		3