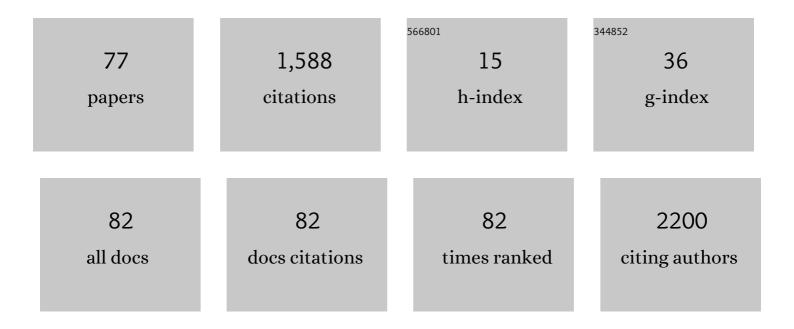
Kuttichantran Subramaniam

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Viable SARS-CoV-2 in the air of a hospital room with COVID-19 patients. International Journal of Infectious Diseases, 2020, 100, 476-482.	1.5	531
2	ICTV Virus Taxonomy Profile: Iridoviridae. Journal of General Virology, 2017, 98, 890-891.	1.3	162
3	Ranaviruses and other members of the family Iridoviridae: Their place in the virosphere. Virology, 2017, 511, 259-271.	1.1	75
4	<i>Megalocytivirus</i> infection in fish. Reviews in Aquaculture, 2012, 4, 221-233.	4.6	63
5	Megalocytivirus infection in cultured Nile tilapia Oreochromis niloticus. Diseases of Aquatic Organisms, 2016, 119, 253-258.	0.5	62
6	A New Family of DNA Viruses Causing Disease in Crustaceans from Diverse Aquatic Biomes. MBio, 2020, 11, .	1.8	62
7	Ranavirus phylogenomics: Signatures of recombination and inversions among bullfrog ranaculture isolates. Virology, 2017, 511, 330-343.	1.1	50
8	Detection of infectious spleen and kidney necrosis virus (ISKNV) and turbot reddish body iridovirus (TRBIV) from archival ornamental fish samples. Diseases of Aquatic Organisms, 2016, 122, 105-123.	0.5	32
9	Phylogenomic characterization of two novel members of the genus Megalocytivirus from archived ornamental fish samples. Diseases of Aquatic Organisms, 2018, 130, 11-24.	0.5	25
10	Detection and molecular characterization of infectious spleen and kidney necrosis virus from major ornamental fish breeding states in <scp>P</scp> eninsular <scp>M</scp> alaysia. Journal of Fish Diseases, 2014, 37, 609-618.	0.9	24
11	Genomic characterization of a novel calicivirus, FHMCV-2012, from baitfish in the USA. Archives of Virology, 2017, 162, 3619-3627.	0.9	24
12	Genomic Characterization of Tilapia Lake Virus Isolates Recovered from Moribund Nile Tilapia () Tj ETQq0 0 0 rgBT Announcements, 2020, 9, .	/Overlock 0.3	10 Tf 50 30 23
13	Complete Genome Sequence of a Tilapia Lake Virus Isolate Obtained from Nile Tilapia (Oreochromis) Tj ETQq1 1 0	.784314 r 0.8	gBT /Overlo
14	Molecular characterization of a novel reassortment Mammalian orthoreovirus type 2 isolated from a Florida white-tailed deer fawn. Virus Research, 2019, 270, 197642.	1.1	19
15	Genome Sequence of Tilapia Lake Virus Associated with Syncytial Hepatitis of Tilapia in an Ecuadorian Aquaculture Facility. Microbiology Resource Announcements, 2019, 8, .	0.3	17
16	Genomic and developmental characterisation of a novel bunyavirus infecting the crustacean Carcinus maenas. Scientific Reports, 2019, 9, 12957.	1.6	16
17	Three New Orbivirus Species Isolated from Farmed White-Tailed Deer (Odocoileus virginianus) in the United States. Viruses, 2020, 12, 13.	1.5	15
18	Characterization of a Novel Megalocytivirus Isolated from European Chub (Squalius cephalus). Viruses, 2019, 11, 440.	1.5	13

#	Article	IF	CITATIONS
19	Red seabream iridovirus associated with cultured Florida pompano Trachinotus carolinus mortality in Central America. Diseases of Aquatic Organisms, 2018, 130, 109-115.	0.5	13

20 Complete Genome Sequence of a <i>Bohle iridovirus </i> Isolate from Ornate Burrowing Frogs () Tj ETQq0 0 0 rgBT $_{0.8}^{/0}$ Verlock 10 Tf 50 7 $_{0.8}^{/0}$

21	Genome characterization of cetaceanpox virus from a managed Indo-Pacific bottlenose dolphin (Tursiops aduncus). Virus Research, 2020, 278, 197861.	1.1	12
22	Genome Sequence of a Ranavirus Isolated from Pike-Perch <i>Sander lucioperca</i> . Genome Announcements, 2016, 4, .	0.8	11
23	Genomic Sequence of a Ranavirus Isolated from Short-Finned Eel (<i>Anguilla australis</i>). Genome Announcements, 2016, 4, .	0.8	10
24	Novel cetacean morbillivirus in a rare Fraser's dolphin (Lagenodelphis hosei) stranding from Maui, Hawaiâ€ĩi. Scientific Reports, 2021, 11, 15986.	1.6	10
25	Earliest detection to date of SARS-CoV-2 in Florida: Identification together with influenza virus on the main entry door of a university building, February 2020. PLoS ONE, 2021, 16, e0245352.	1.1	10
26	Characterization of a novel picornavirus isolated from moribund aquacultured clownfish. Journal of General Virology, 2020, 101, 735-745.	1.3	10
27	Characterization of a peribunyavirus isolated from largemouth bass (Micropterus salmoides). Virus Research, 2019, 273, 197761.	1.1	9
28	Complete Genome Sequence of Mobuck Virus Isolated from a Florida White-Tailed Deer (Odocoileus) Tj ETQq0 0	0 rgBT /O	verlock 10 T
29	Complete Genome Sequence of an Ictalurid Herpesvirus 1 Strain Isolated from Blue Catfish (Ictalurus) Tj ETQq1 1	0,784314	4 rgBT /Over
30	Phylogenomic characterization of red seabream iridovirus from Florida pompano Trachinotus carolinus maricultured in the Caribbean Sea. Archives of Virology, 2019, 164, 1209-1212.	0.9	9
31	Genomic characterization of a novel pegivirus species from free-ranging bottlenose dolphins (Tursiops truncatus) in the Indian River Lagoon, Florida. Virus Research, 2019, 263, 98-101.	1.1	9
32	Genomic characterization of a novel circovirus from a stranded Longman's beaked whale (Indopacetus pacificus). Virus Research, 2020, 277, 197826.	1.1	9
33	Cytochrome oxidase gene sequencing reveals channel catfish ovary cell line is contaminated with brown bullhead cells. Journal of Fish Diseases, 2021, 44, 119-122.	0.9	9
34	Genome Sequence of a Gammaherpesvirus from a Common Bottlenose Dolphin (<i>Tursiops) Tj ETQq0 0 0 rgBT</i>	Overlock	10 Tf 50 14

35	Characterization of a novel rhabdovirus isolated from a stranded harbour porpoise (Phocoena) Tj ETQq1 1 0.784314 rgBT /Overlock 1 1.1	

Longitudinal analysis of pinnipeds in the northwest Atlantic provides insights on endemic circulation of phocine distemper virus. Proceedings of the Royal Society B: Biological Sciences, 2021, 288, 20211841.

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#	Article	IF	CITATIONS
37	A novel group of negative-sense RNA viruses associated with epizootics in managed and free-ranging freshwater turtles in Florida, USA. PLoS Pathogens, 2022, 18, e1010258.	2.1	8
38	Complete Genome Sequence of <i>Epizootic hemorrhagic disease virus</i> Serotype 6, Isolated from Florida White-Tailed Deer (Odocoileus virginianus). Genome Announcements, 2018, 6, .	0.8	7
39	In-Frame 12-Nucleotide Deletion within Open Reading Frame 3a in a SARS-CoV-2 Strain Isolated from a Patient Hospitalized with COVID-19. Microbiology Resource Announcements, 2021, 10, .	0.3	7
40	Genomic Sequencing of Ranaviruses Isolated from Turbot (Scophthalmus maximus) and Atlantic Cod () Tj ETQq0	0 0 rgBT / 0.8	Overlock 10
41	Genome Sequence of an Alphaherpesvirus from a Beluga Whale (Delphinapterus leucas). Genome Announcements, 2017, 5, .	0.8	6
42	Genomic Sequences of Epizootic Hemorrhagic Disease Viruses Isolated from Florida White-Tailed Deer. Genome Announcements, 2017, 5, .	0.8	6
43	Use of Acridine Orange to Visually Improve the Loop-mediated Isothermal Amplification for Detection of Infectious Spleen and Kidney Necrosis Virus. Fish Pathology, 2014, 49, 173-180.	0.4	5
44	Complete Genome Sequence of Human Coronavirus Strain 229E Isolated from Plasma Collected from a Haitian Child in 2016. Genome Announcements, 2017, 5, .	0.8	5
45	Complete genome sequence of a novel sea otterpox virus. Virus Genes, 2018, 54, 756-767.	0.7	5
46	Genomic sequencing of a virus representing a novel type within the species Dyopipapillomavirus 1 in an Indian River Lagoon bottlenose dolphin. Archives of Virology, 2019, 164, 767-774.	0.9	5
47	Characterization of ranaviruses isolated from lumpfish L. in the North Atlantic area: proposal for a new ranavirus species (European North Atlantic Ranavirus). Journal of General Virology, 2020, 101, 198-207.	1.3	5
48	Phylogenomic characterization of ranaviruses isolated from cultured fish and amphibians in Thailand. Facets, 2020, 5, 963-979.	1.1	5
49	Genetic characterization of esocid herpesvirus 1 (EsHV1). Diseases of Aquatic Organisms, 2016, 122, 1-11.	0.5	5
50	An alloherpesvirus infection of European perch Perca fluviatilis in Finland. Diseases of Aquatic Organisms, 2018, 128, 175-185.	0.5	5
51	Genomic Sequencing of Ranaviruses Isolated from Edible Frogs (Pelophylax esculentus). Genome Announcements, 2017, 5, .	0.8	4
52	Genome Sequence of a Novel Alloherpesvirus Isolated from Glass Catfish (Kryptopterus bicirrhis). Genome Announcements, 2018, 6, .	0.8	4
53	Genome Sequences of a Novel Strain of Big Cypress Orbivirus Isolated from a Dead Florida White-Tailed Deer (Odocoileus virginianus). Microbiology Resource Announcements, 2019, 8, .	0.3	4

54 Genome Sequence of a Yunnan Orbivirus Isolated from a Dead Florida White-Tailed Deer (Odocoileus) Tj ETQq0 0 0.78 BT /Overlock 10 Tf

#	Article	IF	CITATIONS
55	Genomic sequencing of a frog virus 3 strain from cultured American bullfrogs (Lithobates) Tj ETQq1 1 0.784314	rgBT/Ove	rloçk 10 Tf 50
56	Complete genome sequences of infectious spleen and kidney necrosis virus isolated from farmed albino rainbow sharks Epalzeorhynchos frenatum in the United States. Virus Genes, 2021, 57, 448-452.	0.7	4
57	Genome Sequence of a Ranavirus Isolated from a Red-Eared Slider (Trachemys scripta elegans) in Poland. Microbiology Resource Announcements, 2020, 9, .	0.3	4
58	Tilapia Lake Virus (TiLV): a Globally Emerging Threat to Tilapia Aquaculture. Edis, 2019, 2019, .	0.0	4
59	Genomic Characterization of Picornaviruses Isolated From Ribbon (Histriophoca fasciata) and Harbor (Phoca vitulina) Seals. Frontiers in Veterinary Science, 2020, 7, 554716.	0.9	4
60	Genomic characterization of psittacine adenovirus 2, a siadenovirus identified in a moribund African grey parrot (Psittacus erithacus). Archives of Virology, 2022, 167, 911-916.	0.9	4
61	Genome Sequence of Rotavirus A from a Florida Racing Pigeon (<i>Columba livia domestica</i>). Microbiology Resource Announcements, 2022, 11, e0114921.	0.3	4
62	Characterization of a Novel Reassortant Epizootic Hemorrhagic Disease Virus Serotype 6 Strain Isolated from Diseased White-Tailed Deer (Odocoileus virginianus) on a Florida Farm. Viruses, 2022, 14, 1012.	1.5	4
63	Complete Genome Sequencing of a Novel Type of <i>Omikronpapillomavirus 1</i> in Indian River Lagoon Bottlenose Dolphins (Tursiops truncatus). Genome Announcements, 2018, 6, .	0.8	3
64	Genomic sequence of a Bohle iridovirus strain isolated from a diseased boreal toad (Anaxyrus boreas) Tj ETQq0 0	0 rgBT /C	veglock 10 Tf
65	Characterization of mule deerpox virus in Florida white-tailed deer fawns expands the known host and geographic range of this emerging pathogen. Archives of Virology, 2019, 164, 51-61.	0.9	3
66	Genome Sequence of a CHeRI Orbivirus 3 Strain Isolated from a Dead White-Tailed Deer (Odocoileus) Tj ETQq0 0	0rgBT /C	Oveglock 10 Tf
67	Detection and Preliminary Characterization of Phocine Distemper Virus in a Stranded Harp Seal (Pagophilus groenlandicus) from the Gulf of St. Lawrence, Canada. Journal of Wildlife Diseases, 2020, 56, 646.	0.3	3
68	Characterization of an alphavirus isolated from a stranded harbor porpoise (Phocoena phocoena) from Alaska. Virus Research, 2021, 291, 198187.	1.1	3
69	Diagnosis of Fish Diseases. , 2019, , 46-88.		3
70	FATAL RANAVIRUS INFECTION IN A GROUP OF ZOO-HOUSED MELLER'S CHAMELEONS (TRIOCEROS MELLERI). Journal of Zoo and Wildlife Medicine, 2019, 50, 696.	0.3	3
71	Genomic Sequencing of Ranavirus Isolates from a Three-Spined Stickleback (<i>Gasterosteus) Tj ETQq1 1 0.784 2021, 10, e0090221.</i>	314 rgBT , 0.3	/Overlock 10 3
72	Molecular epidemiology of Epizootic haematopoietic necrosis virus (EHNV). Virology, 2017, 511, 320-329.	1.1	2

#	Article	IF	CITATIONS
73	Molecular Confirmation of Ranavirus Infection in Amphibians From Chad, Africa. Frontiers in Veterinary Science, 2021, 8, 733939.	0.9	2
74	PSEUDORABIES (AUJESZKY'S DISEASE) IS AN UNDERDIAGNOSED CAUSE OF DEATH IN THE FLORIDA PANTHER (PUMA CONCOLOR CORYI). Journal of Wildlife Diseases, 2021, 57, 784-798.	0.3	2
75	Isolation, Identification and Characterization of a Novel Megalocytivirus from Cultured Tilapia (Oreochromis spp.) from Southern California, USA. Animals, 2021, 11, 3524.	1.0	2
76	The complete mitochondrial genome of the Jaguar Loach (Yasuhikotakia splendida). Mitochondrial DNA Part B: Resources, 2021, 6, 1737-1739.	0.2	0
77	A Novel Orthoreovirus Isolated From Dead Stranded Harbor Seals From Puget Sound, Washington State, United States. Frontiers in Marine Science, 2021, 8, .	1.2	0