Arvind Varsani

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.

56 98 12,052 329 h-index g-index citations papers 6.6 16,244 4.5 355 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
329	Differentiating between viruses and virus species by writing their names correctly <i>Archives of Virology</i> , 2022 , 167, 1231	2.6	5
328	Novel adenovirus associated with common tern (Sterna hirundo) chicks <i>Archives of Virology</i> , 2022 , 167, 659	2.6	1
327	Microvirus Genomes Identified in Fecal Samples from Yellow-Bellied Marmots <i>Microbiology Resource Announcements</i> , 2022 , e0121821	1.3	
326	Diverse single-stranded DNA viruses identified in New Zealand (Aotearoa) South Island robin (Petroica australis) fecal samples. <i>Virology</i> , 2022 , 565, 38-51	3.6	O
325	Discovery of novel fish papillomaviruses: From the Antarctic to the commercial fish market. <i>Virology</i> , 2022 , 565, 65-72	3.6	O
324	A PCR-Based Retrospective Study for Beak and Feather Disease Virus (BFDV) in Five Wild Populations of Parrots from Australia, Argentina and New Zealand. <i>Diversity</i> , 2022 , 14, 148	2.5	2
323	Genomes of Bacteriophages Belonging to the Orders and Identified in Fecal Samples from Pacific Flying Fox () from the Kingdom of Tonga <i>Microbiology Resource Announcements</i> , 2022 , 11, e0003822	1.3	O
322	Viral Aggregation: The Knowns and Unknowns Viruses, 2022, 14,	6.2	1
321	Coevolutionary Analysis Implicates Toll-Like Receptor 9 in Papillomavirus Restriction <i>MBio</i> , 2022 , e000	05/482	O
320	Identification of a Novel Myxoma Virus C7-Like Host Range Factor That Enabled a Species Leap from Rabbits to Hares <i>MBio</i> , 2022 , e0346121	7.8	1
319	RNA Virus Gene Signatures Detected in Patients With Cardiomyopathy After Chemotherapy; A Pilot Study <i>Frontiers in Cardiovascular Medicine</i> , 2022 , 9, 821162	5.4	O
318	Detection of Human, Porcine and Canine Picornaviruses in Municipal Sewage Sludge Using Pan-Enterovirus Amplicon-based Long-read Illumina Sequencing <i>Emerging Microbes and Infections</i> , 2022 , 1-25	18.9	0
317	ICTV Virus Taxonomy Profile: 2021 Journal of General Virology, 2021 , 102,	4.9	16
316	Predicting the zoonotic capacity of mammals to transmit SARS-CoV-2. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021 , 288, 20211651	4.4	15
315	Presence and Persistence of Putative Lytic and Temperate Bacteriophages in Vaginal Metagenomes from South African Adolescents <i>Viruses</i> , 2021 , 13,	6.2	2
314	Establishment of five new genera in the family Geminiviridae: Citlodavirus, Maldovirus, Mulcrilevirus, Opunvirus, and Topilevirus. <i>Archives of Virology</i> , 2021 , 1	2.6	7
313	Perspective on taxonomic classification of uncultivated viruses. <i>Current Opinion in Virology</i> , 2021 , 51, 207-215	7.5	5

(2021-2021)

312	An investigation of antibacterial activity of New Zealand seaweed-associated marine bacteria. <i>Future Microbiology</i> , 2021 , 16, 1167-1179	2.9	
311	Adaptation of the polony technique to quantify Gokushovirinae, a diverse group of single-stranded DNA phage. <i>Environmental Microbiology</i> , 2021 , 23, 6622-6636	5.2	1
310	RDP5: a computer program for analyzing recombination in, and removing signals of recombination from, nucleotide sequence datasets. <i>Virus Evolution</i> , 2021 , 7, veaa087	3.7	55
309	Genome Sequences of Microviruses Identified in Gila Monster Feces. <i>Microbiology Resource Announcements</i> , 2021 , 10,	1.3	3
308	Occurrence of lettuce chlorosis virus in Passiflora spp. in Brazil. <i>Journal of Plant Pathology</i> , 2021 , 103, 443-447	1	1
307	A rapid assessment of wastewater for genomic surveillance of SARS-CoV-2 variants at sewershed scale in Louisville, KY 2021 ,		7
306	Genome characterization of parsley severe stunt-associated virus in Iran. Virus Genes, 2021, 57, 293-301	2.3	1
305	New World Cactaceae Plants Harbor Diverse Geminiviruses. <i>Viruses</i> , 2021 , 13,	6.2	2
304	Molecular Epidemiology of Citrus Leprosis Virus C: A New Viral Lineage and Phylodynamic of the Main Viral Subpopulations in the Americas. <i>Frontiers in Microbiology</i> , 2021 , 12, 641252	5.7	7
303	A Pilot Study Investigating the Dynamics of Pigeon Circovirus Recombination in Domesticated Pigeons Housed in a Single Loft. <i>Viruses</i> , 2021 , 13,	6.2	2
302	Genome Sequences of Microviruses Identified in a Sample from a Sewage Treatment Oxidation Pond. <i>Microbiology Resource Announcements</i> , 2021 , 10,	1.3	2
301	First report of citrus tristeza virus in commercial citrus orchards in Tunisia. <i>Journal of Plant Pathology</i> , 2021 , 103, 1051-1052	1	1
300	Discovery of Known and Novel Viruses in Wild and Cultivated Blueberry in Florida through Viral Metagenomic Approaches. <i>Viruses</i> , 2021 , 13,	6.2	1
299	Circular DNA viruses identified in short-finned pilot whale and orca tissue samples. <i>Virology</i> , 2021 , 559, 156-164	3.6	1
298	crAssphage genomes identified in fecal samples of an adult and infants with evidence of positive genomic selective pressure within tail protein genes. <i>Virus Research</i> , 2021 , 292, 198219	6.4	2
297	Risk assessment of SARS-CoV-2 in Antarctic wildlife. <i>Science of the Total Environment</i> , 2021 , 755, 143357	210.2	10
296	Predicting the zoonotic capacity of mammals to transmit SARS-CoV-2 2021 ,		6
295	VirSorter2: a multi-classifier, expert-guided approach to detect diverse DNA and RNA viruses. <i>Microbiome</i> , 2021 , 9, 37	16.6	69

294	Diverse cressdnaviruses and an anellovirus identified in the fecal samples of yellow-bellied marmots. <i>Virology</i> , 2021 , 554, 89-96	3.6	5
293	Complete Genome Sequence of a Phapecoctavirus Isolated from a Pigeon Cloacal Swab Sample. <i>Microbiology Resource Announcements</i> , 2021 , 10,	1.3	1
292	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
291	Family Genomoviridae: 2021 taxonomy update. Archives of Virology, 2021, 166, 2911-2926	2.6	5
290	Genomic skimming and nanopore sequencing uncover cryptic hybridization in one of world's most threatened primates. <i>Scientific Reports</i> , 2021 , 11, 17279	4.9	3
289	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
288	Use of hemagglutinin and neuraminidase amplicon-based high-throughput sequencing with variant analysis to detect co-infection and resolve identical consensus sequences of seasonal influenza in a university setting. <i>BMC Infectious Diseases</i> , 2021 , 21, 810	4	О
287	Taxonomic updates for the genus Gyrovirus (family Anelloviridae): recognition of several new members and establishment of species demarcation criteria. <i>Archives of Virology</i> , 2021 , 166, 2937-2942	2.6	5
286	Taxonomic update for mammalian anelloviruses (family Anelloviridae). <i>Archives of Virology</i> , 2021 , 166, 2943-2953	2.6	6
285	High-throughput sequencing of SARS-CoV-2 in wastewater provides insights into circulating variants. <i>Water Research</i> , 2021 , 205, 117710	12.5	25
284	Wastewater Monitoring Raises Privacy and Ethical Considerations. <i>IEEE Transactions on Technology and Society</i> , 2021 , 2, 116-121	5.2	4
283	Integrating Viral Metagenomics into an Ecological Framework. <i>Annual Review of Virology</i> , 2021 , 8, 133-1	58 .6	10
282	Novel viruses belonging to the family Circoviridae identified in wild American wigeon samples. <i>Archives of Virology</i> , 2021 , 166, 3437-3441	2.6	
281	Wastewater-Based Epidemiology and Long-Read Sequencing to Identify Enterovirus Circulation in Three Municipalities in Maricopa County, Arizona, Southwest United States between June and October 2020. <i>Viruses</i> , 2021 , 13,	6.2	3
2 80	A 2021 taxonomy update for the family Smacoviridae. <i>Archives of Virology</i> , 2021 , 166, 3245-3253	2.6	2
279	Taxonomy update for the family Alphasatellitidae: new subfamily, genera, and species. <i>Archives of Virology</i> , 2021 , 166, 3503-3511	2.6	4
278	Identification of novel circovirus and anelloviruses from wolverines using a non-invasive faecal sampling approach. <i>Infection, Genetics and Evolution</i> , 2021 , 93, 104914	4.5	1
277	Complex evolutionary history of felid anelloviruses. <i>Virology</i> , 2021 , 562, 176-189	3.6	O

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276	Circoviruses and cycloviruses identified in Weddell seal fecal samples from McMurdo Sound, Antarctica. <i>Infection, Genetics and Evolution</i> , 2021 , 95, 105070	4.5	О
275	A novel lineage of polyomaviruses identified in bark scorpions. <i>Virology</i> , 2021 , 563, 58-63	3.6	2
274	Agricultural practices drive biological loads, seasonal patterns and potential pathogens in the aerobiome of a mixed-land-use dryland. <i>Science of the Total Environment</i> , 2021 , 798, 149239	10.2	3
273	How virulent are emerging maize-infecting mastreviruses?. <i>Archives of Virology</i> , 2021 , 166, 955-959	2.6	1
272	Pan-Enterovirus Amplicon-Based High-Throughput Sequencing Detects the Complete Capsid of a EVA71 Genotype C1 Variant via Wastewater-Based Epidemiology in Arizona. <i>Viruses</i> , 2021 , 13,	6.2	3
271	High-throughput sequencing of SARS-CoV-2 in wastewater provides insights into circulating variants 2021 ,		22
270	An 81-Nucleotide Deletion in SARS-CoV-2 ORF7a Identified from Sentinel Surveillance in Arizona (January to March 2020). <i>Journal of Virology</i> , 2020 , 94,	6.6	91
269	Sorghum mastrevirus-associated alphasatellites: new geminialphasatellites associated with an African streak mastrevirus infecting wild Poaceae plants on Reunion Island. <i>Archives of Virology</i> , 2020 , 165, 1925-1928	2.6	2
268	Viruses representing two new genomovirus species identified in citrus from Tunisia. <i>Archives of Virology</i> , 2020 , 165, 1225-1229	2.6	3
267	Global Organization and Proposed Megataxonomy of the Virus World. <i>Microbiology and Molecular Biology Reviews</i> , 2020 , 84,	13.2	178
267		13.2	178
Í	Biology Reviews, 2020, 84,		,
266	Biology Reviews, 2020, 84, Isolation and characterization of a novel geminivirus from parsley. Virus Research, 2020, 286, 198056 Frequent cross-species transmissions of foamy virus between domestic and wild felids. Virus	6.4	3
266 265	Biology Reviews, 2020, 84, Isolation and characterization of a novel geminivirus from parsley. Virus Research, 2020, 286, 198056 Frequent cross-species transmissions of foamy virus between domestic and wild felids. Virus Evolution, 2020, 6, vez058 Immune protection is dependent on the gut microbiome in a lethal mouse gammaherpesviral	6.4	3 7
266 265 264	Isolation and characterization of a novel geminivirus from parsley. <i>Virus Research</i> , 2020 , 286, 198056 Frequent cross-species transmissions of foamy virus between domestic and wild felids. <i>Virus Evolution</i> , 2020 , 6, vez058 Immune protection is dependent on the gut microbiome in a lethal mouse gammaherpesviral infection. <i>Scientific Reports</i> , 2020 , 10, 2371 Virus Discovery in Desert Tortoise Fecal Samples: Novel Circular Single-Stranded DNA Viruses.	6.4 3.7 4.9	3 7 10
266 265 264 263	Isolation and characterization of a novel geminivirus from parsley. <i>Virus Research</i> , 2020 , 286, 198056 Frequent cross-species transmissions of foamy virus between domestic and wild felids. <i>Virus Evolution</i> , 2020 , 6, vez058 Immune protection is dependent on the gut microbiome in a lethal mouse gammaherpesviral infection. <i>Scientific Reports</i> , 2020 , 10, 2371 Virus Discovery in Desert Tortoise Fecal Samples: Novel Circular Single-Stranded DNA Viruses. <i>Viruses</i> , 2020 , 12, Coinfections of Novel Polyomavirus, Anelloviruses and a Recombinant Strain of Myxoma	6.4 3.7 4.9	3 7 10 9
266 265 264 263	Isolation and characterization of a novel geminivirus from parsley. <i>Virus Research</i> , 2020, 286, 198056 Frequent cross-species transmissions of foamy virus between domestic and wild felids. <i>Virus Evolution</i> , 2020, 6, vez058 Immune protection is dependent on the gut microbiome in a lethal mouse gammaherpesviral infection. <i>Scientific Reports</i> , 2020, 10, 2371 Virus Discovery in Desert Tortoise Fecal Samples: Novel Circular Single-Stranded DNA Viruses. <i>Viruses</i> , 2020, 12, Coinfections of Novel Polyomavirus, Anelloviruses and a Recombinant Strain of Myxoma Virus-MYXV-Tol Identified in Iberian Hares. <i>Viruses</i> , 2020, 12, : a Virus Phylum Unifying Seven Families of Rep-Encoding Viruses with Single-Stranded, Circular	6.4 3.7 4.9 6.2	3 7 10 9

258	Author response: Discovery of several thousand highly diverse circular DNA viruses 2020,		5
257	Symptom evolution following the emergence of maize streak virus. <i>ELife</i> , 2020 , 9,	8.9	9
256	Genome characterization of cetaceanpox virus from a managed Indo-Pacific bottlenose dolphin (Tursiops aduncus). <i>Virus Research</i> , 2020 , 278, 197861	6.4	3
255	Novel nanovirus and associated alphasatellites identified in milk vetch plants with chlorotic dwarf disease in Iran. <i>Virus Research</i> , 2020 , 276, 197830	6.4	4
254	Binomial nomenclature for virus species: a consultation. <i>Archives of Virology</i> , 2020 , 165, 519-525	2.6	27
253	Transmission of the Bean-Associated Cytorhabdovirus by the Whitefly MEAM1. Viruses, 2020 , 12,	6.2	8
252	Identification and Distribution of Novel Cressdnaviruses and Circular molecules in Four Penguin Species in South Georgia and the Antarctic Peninsula. <i>Viruses</i> , 2020 , 12,	6.2	3
251	Genome Sequences of Novel Torque Teno Viruses Identified in Human Brain Tissue. <i>Microbiology Resource Announcements</i> , 2020 , 9,	1.3	1
250	Blueberry red ringspot virus genomes from Florida inferred through analysis of blueberry root transcriptomes. <i>Scientific Reports</i> , 2020 , 10, 12043	4.9	2
249	VIRIDIC-A Novel Tool to Calculate the Intergenomic Similarities of Prokaryote-Infecting Viruses. <i>Viruses</i> , 2020 , 12,	6.2	57
248	The Vaginal Virome-Balancing Female Genital Tract Bacteriome, Mucosal Immunity, and Sexual and Reproductive Health Outcomes?. <i>Viruses</i> , 2020 , 12,	6.2	3
247	Unveiling Crucivirus Diversity by Mining Metagenomic Data. <i>MBio</i> , 2020 , 11,	7.8	7
246	Novel Circoviruses Detected in Feces of Sonoran Felids. <i>Viruses</i> , 2020 , 12,	6.2	3
245	Diverse genomoviruses representing twenty-nine species identified associated with plants. <i>Archives of Virology</i> , 2020 , 165, 2891-2901	2.6	5
244	Identification of Circovirus Genome in a Chinstrap Penguin () and Adlie Penguin () on the Antarctic Peninsula. <i>Viruses</i> , 2020 , 12,	6.2	5
243	First report of barley yellow dwarf virus in maize in Tunisia. <i>Journal of Plant Pathology</i> , 2020 , 102, 227-2	27	O
242	A Novel Divergent Geminivirus Identified in Asymptomatic New World Cactaceae Plants. <i>Viruses</i> , 2020 , 12,	6.2	5
241	Exploring the diversity of Poaceae-infecting mastreviruses on Reunion Island using a viral metagenomics-based approach. <i>Scientific Reports</i> , 2019 , 9, 12716	4.9	5

240	Single Stranded DNA Viruses Associated with Capybara Faeces Sampled in Brazil. Viruses, 2019, 11,	6.2	19
239	Kombucha: a novel model system for cooperation and conflict in a complex multi-species microbial ecosystem. <i>PeerJ</i> , 2019 , 7, e7565	3.1	38
238	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
237	Genetic Characterization of a Recombinant Myxoma Virus in the Iberian Hare (). Viruses, 2019, 11,	6.2	20
236	Unravelling the Single-Stranded DNA Virome of the New Zealand Blackfly. Viruses, 2019, 11,	6.2	11
235	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
234	Genome Sequences of Microviruses Associated with. <i>Microbiology Resource Announcements</i> , 2019 , 8,	1.3	5
233	Evidence of age-related improvement in the foraging efficiency of Adlie penguins. <i>Scientific Reports</i> , 2019 , 9, 3375	4.9	11
232	Diverse single-stranded DNA viruses associated with honey bees (Apis mellifera). <i>Infection, Genetics and Evolution</i> , 2019 , 71, 179-188	4.5	16
231	Molecular and partial biological characterization of the coat protein sequences of Iranian alfalfa mosaic virus isolates. <i>Journal of Plant Pathology</i> , 2019 , 101, 735-742	1	6
230	Diverse genomoviruses representing eight new and one known species identified in feces and nests of house finches (Haemorhous mexicanus). <i>Archives of Virology</i> , 2019 , 164, 2345-2350	2.6	7
229	Multiple origins of prokaryotic and eukaryotic single-stranded DNA viruses from bacterial and archaeal plasmids. <i>Nature Communications</i> , 2019 , 10, 3425	17.4	55
228	Evolution and ecology of plant viruses. <i>Nature Reviews Microbiology</i> , 2019 , 17, 632-644	22.2	72
227	Novel smacoviruses identified in the faeces of two wild felids: North American bobcat and African lion. <i>Archives of Virology</i> , 2019 , 164, 2395-2399	2.6	3
226	Tracking harmful chemicals and pathogens using the Human Health Observatory at ASU. <i>Online Journal of Public Health Informatics</i> , 2019 , 11,	0.3	5
225	Single-Stranded DNA Viruses in Antarctic Cryoconite Holes. <i>Viruses</i> , 2019 , 11,	6.2	17
224	Identification of the wild and cultivated hosts of wheat dwarf virus and oat dwarf virus in Iran. <i>VirusDisease</i> , 2019 , 30, 545-550	3.4	0
223	Identification of a Novel Adlie Penguin Circovirus at Cape Crozier (Ross Island, Antarctica). <i>Viruses</i> , 2019 , 11,	6.2	10

222	Novel circular DNA viruses associated with Apiaceae and Poaceae from South Africa and New Zealand. <i>Archives of Virology</i> , 2019 , 164, 237-242	2.6	5
221	Minimum Information about an Uncultivated Virus Genome (MIUViG). <i>Nature Biotechnology</i> , 2019 , 37, 29-37	44.5	180
220	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. <i>Systematic Biology</i> , 2019 , 68, 828-839	8.4	5
219	Geometagenomics illuminates the impact of agriculture on the distribution and prevalence of plant viruses at the ecosystem scale. <i>ISME Journal</i> , 2018 , 12, 173-184	11.9	72
218	Smacoviridae: a new family of animal-associated single-stranded DNA viruses. <i>Archives of Virology</i> , 2018 , 163, 2005-2015	2.6	39
217	Genome Sequences of Two Single-Stranded DNA Viruses Identified in. <i>Genome Announcements</i> , 2018 , 6,		4
216	Viruses associated with Antarctic wildlife: From serology based detection to identification of genomes using high throughput sequencing. <i>Virus Research</i> , 2018 , 243, 91-105	6.4	41
215	Discovery of the first maize-infecting mastrevirus in the Americas using a vector-enabled metagenomics approach. <i>Archives of Virology</i> , 2018 , 163, 263-267	2.6	6
214	Genomoviruses associated with mountain and western pine beetles. Virus Research, 2018, 256, 17-20	6.4	8
213	Novel anelloviruses identified in buccal swabs of Antarctic fur seals. <i>Virus Genes</i> , 2018 , 54, 719-723	2.3	9
212	Recombinant Goose Circoviruses Circulating in Domesticated and Wild Geese in Poland. <i>Viruses</i> , 2018 , 10,	6.2	6
211	Pervasive Chimerism in the Replication-Associated Proteins of Uncultured Single-Stranded DNA Viruses. <i>Viruses</i> , 2018 , 10,	6.2	36
210	Alphasatellitidae: a new family with two subfamilies for the classification of geminivirus- and nanovirus-associated alphasatellites. <i>Archives of Virology</i> , 2018 , 163, 2587-2600	2.6	78
209	Identification of circular single-stranded DNA viruses in faecal samples of Canada lynx (Lynx canadensis), moose (Alces alces) and snowshoe hare (Lepus americanus) inhabiting the Colorado San Juan Mountains. <i>Infection, Genetics and Evolution</i> , 2018 , 64, 1-8	4.5	17
208	From Spatial Metagenomics to Molecular Characterization of Plant Viruses: A Geminivirus Case Study. <i>Advances in Virus Research</i> , 2018 , 101, 55-83	10.7	20
207	Virus discovery in all three major lineages of terrestrial arthropods highlights the diversity of single-stranded DNA viruses associated with invertebrates. <i>PeerJ</i> , 2018 , 6, e5761	3.1	30
206	Diverse papillomaviruses identified in Weddell seals. <i>Journal of General Virology</i> , 2018 , 99, 549-557	4.9	8
205	Fish polyomaviruses belong to two distinct evolutionary lineages. <i>Journal of General Virology</i> , 2018 , 99, 567-573	4.9	12

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204	Nanovirus-alphasatellite complex identified in Vicia cracca in the Rhtle delta region of France. <i>Archives of Virology</i> , 2018 , 163, 695-700	2.6	14
203	Molecular characterization of faba bean necrotic yellows viruses in Tunisia. <i>Archives of Virology</i> , 2018 , 163, 687-694	2.6	11
202	Identification of an anellovirus and genomoviruses in ixodid ticks. Virus Genes, 2018, 54, 155-159	2.3	14
201	Turnip leaf curl disease associated with two begomoviruses in south-eastern Iran. <i>Tropical Plant Pathology</i> , 2018 , 43, 165-169	2.5	5
200	Genome Sequence of a Gyrovirus Associated with Ashy Storm-Petrel. <i>Microbiology Resource Announcements</i> , 2018 , 7,	1.3	5
199	The Westward Journey of Alfalfa Leaf Curl Virus. <i>Viruses</i> , 2018 , 10,	6.2	6
198	Identification of a new turncurtovirus in the leafhopper Circulifer haematoceps and the host plant species Sesamum indicum. <i>Virus Genes</i> , 2018 , 54, 840-845	2.3	6
197	Notes on recombination and reassortment in multipartite/segmented viruses. <i>Current Opinion in Virology</i> , 2018 , 33, 156-166	7.5	26
196	First World Report of Cucurbit Aphid-Borne Yellows Virus Infecting Passionfruit. <i>Plant Disease</i> , 2018 , 102, 2665-2665	1.5	11
195	Genome Sequence of a Single-Stranded DNA Virus Identified in Gila Monster Feces. <i>Microbiology Resource Announcements</i> , 2018 , 7,	1.3	3
194	Passion Fruit Chlorotic Mottle Virus: Molecular Characterization of a New Divergent Geminivirus in Brazil. <i>Viruses</i> , 2018 , 10,	6.2	26
193	Changes to taxonomy and the International Code of Virus Classification and Nomenclature ratified by the International Committee on Taxonomy of Viruses (2018). <i>Archives of Virology</i> , 2018 , 163, 2601-26	5 3 1 ⁶	187
192	Identification of a polyomavirus in Weddell seal (Leptonychotes weddellii) from the Ross Sea (Antarctica). <i>Archives of Virology</i> , 2017 , 162, 1403-1407	2.6	29
191	New strains of chickpea chlorotic dwarf virus discovered on diseased papaya and tomato plants in Burkina Faso. <i>Archives of Virology</i> , 2017 , 162, 1791-1794	2.6	12
190	Capulavirus and Grablovirus: two new genera in the family Geminiviridae. <i>Archives of Virology</i> , 2017 , 162, 1819-1831	2.6	166
189	Evolutionary history of ssDNA bacilladnaviruses features horizontal acquisition of the capsid gene from ssRNA nodaviruses. <i>Virology</i> , 2017 , 504, 114-121	3.6	35
188	The role of Kenya in the trans-African spread of maize streak virus strain A. <i>Virus Research</i> , 2017 , 232, 69-76	6.4	9
187	Identification of a Nanovirus-Alphasatellite Complex in Sophora alopecuroides. <i>Virus Research</i> , 2017 , 235, 24-32	6.4	19

186	50 years of the International Committee on Taxonomy of Viruses: progress and prospects. <i>Archives of Virology</i> , 2017 , 162, 1441-1446	2.6	53
185	Genome sequences of a capulavirus infecting Plantago lanceolata in the land archipelago of Finland. <i>Archives of Virology</i> , 2017 , 162, 2041-2045	2.6	29
184	Genome Sequences of , , , and Identified in Leafhoppers. <i>Genome Announcements</i> , 2017 , 5,		11
183	Changes to taxonomy and the International Code of Virus Classification and Nomenclature ratified by the International Committee on Taxonomy of Viruses (2017). <i>Archives of Virology</i> , 2017 , 162, 2505-25	538	398
182	Comparative analysis of common regions found in babuviruses and alphasatellite molecules. <i>Archives of Virology</i> , 2017 , 162, 849-855	2.6	3
181	Genomovirus Genomes Recovered from Sampled in Florida, USA. <i>Genome Announcements</i> , 2017 , 5,		11
180	Tracking effluent discharges in undisturbed stony soil and alluvial gravel aquifer using synthetic DNA tracers. <i>Science of the Total Environment</i> , 2017 , 592, 144-152	10.2	18
179	Genome Sequence of Cauliflower Mosaic Virus Identified in Earwigs () through a Metagenomic Approach. <i>Genome Announcements</i> , 2017 , 5,		3
178	Novel Single-Stranded DNA Virus Genomes Recovered from Chimpanzee Feces Sampled from the Mambilla Plateau in Nigeria. <i>Genome Announcements</i> , 2017 , 5,		5
177	Revisiting the taxonomy of the family Circoviridae: establishment of the genus Cyclovirus and removal of the genus Gyrovirus. <i>Archives of Virology</i> , 2017 , 162, 1447-1463	2.6	182
176	Consensus statement: Virus taxonomy in the age of metagenomics. <i>Nature Reviews Microbiology</i> , 2017 , 15, 161-168	22.2	375
175	Unique genome organization of non-mammalian papillomaviruses provides insights into the evolution of viral early proteins. <i>Virus Evolution</i> , 2017 , 3, vex027	3.7	27
174	Occurrence of a novel mastrevirus in sugarcane germplasm collections in Florida, Guadeloupe and Rùnion. <i>Virology Journal</i> , 2017 , 14, 146	6.1	13
173	Sequence-based taxonomic framework for the classification of uncultured single-stranded DNA viruses of the family. <i>Virus Evolution</i> , 2017 , 3, vew037	3.7	67
172	Diverse and highly recombinant anelloviruses associated with Weddell seals in Antarctica. <i>Virus Evolution</i> , 2017 , 3, vex017	3.7	39
171	ICTV Virus Taxonomy Profile: Geminiviridae. <i>Journal of General Virology</i> , 2017 , 98, 131-133	4.9	400
170	Possibility and Challenges of Conversion of Current Virus Species Names to Linnaean Binomials. <i>Systematic Biology</i> , 2017 , 66, 463-473	8.4	12
169	A novel geminivirus identified in tomato and cleome plants sampled in Brazil. <i>Virus Research</i> , 2017 , 240, 175-179	6.4	16

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167	First Report of Maize yellow mosaic virus Infecting Maize in Brazil. <i>Plant Disease</i> , 2017 , 101, 2156	1.5	19
166	Molecular diversity, geographic distribution and host range of monocot-infecting mastreviruses in Africa and surrounding islands. <i>Virus Research</i> , 2017 , 238, 171-178	6.4	9
165	Novel mastreviruses identified in Australian wild rice. <i>Virus Research</i> , 2017 , 238, 193-197	6.4	10
164	ICTV Virus Taxonomy Profile: Circoviridae. <i>Journal of General Virology</i> , 2017 , 98, 1997-1998	4.9	84
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154	Comparison of Illumina de novo assembled and Sanger sequenced viral genomes: A case study for RNA viruses recovered from the plant pathogenic fungus Sclerotinia sclerotiorum. <i>Virus Research</i> , 2016 , 219, 51-57	6.4	16
153	Molecular diversity of turncurtoviruses in Iran. Archives of Virology, 2016, 161, 551-61	2.6	13
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151	Complete Genome Sequence of a Genomovirus Associated with Common Bean Plant Leaves in Brazil. <i>Genome Announcements</i> , 2016 , 4,		6

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148	Circular replication-associated protein encoding DNA viruses identified in the faecal matter of various animals in New Zealand. <i>Infection, Genetics and Evolution</i> , 2016 , 43, 151-64	4.5	56
147	Molecular characterization and prevalence of two capulaviruses: Alfalfa leaf curl virus from France and Euphorbia caput-medusae latent virus from South Africa. <i>Virology</i> , 2016 , 493, 142-53	3.6	33
146	Divergent evolutionary and epidemiological dynamics of cassava mosaic geminiviruses in Madagascar. <i>BMC Evolutionary Biology</i> , 2016 , 16, 182	3	15
145	Geometric morphometrics and molecular systematics of Xanthocnemis sobrina (McLachlan, 1873) (Odonata: Coenagrionidae) and comparison to its congeners. <i>Zootaxa</i> , 2016 , 4078, 84-120	0.5	1
144	Identification of novel Bromus- and Trifolium-associated circular DNA viruses. <i>Archives of Virology</i> , 2015 , 160, 1303-11	2.6	24
143	Size exclusion-based purification and PCR-based quantitation of MS2 bacteriophage particles for environmental applications. <i>Journal of Virological Methods</i> , 2015 , 213, 135-8	2.6	7
142	Diverse small circular DNA viruses circulating amongst estuarine molluscs. <i>Infection, Genetics and Evolution</i> , 2015 , 31, 284-95	4.5	44
141	Alfalfa Leaf Curl Virus: an Aphid-Transmitted Geminivirus. <i>Journal of Virology</i> , 2015 , 89, 9683-8	6.6	47
140	Discovery of a novel circular DNA virus in the Forbes sea star, Asterias forbesi. <i>Archives of Virology</i> , 2015 , 160, 2349-51	2.6	23
139	Identification of diverse circular single-stranded DNA viruses in adult dragonflies and damselflies (Insecta: Odonata) of Arizona and Oklahoma, USA. <i>Infection, Genetics and Evolution</i> , 2015 , 30, 278-287	4.5	54
138	Adsorption of Rotavirus, MS2 Bacteriophage and Surface-Modified Silica Nanoparticles to Hydrophobic Matter. <i>Food and Environmental Virology</i> , 2015 , 7, 261-8	4	20
137	Revision of Begomovirus taxonomy based on pairwise sequence comparisons. <i>Archives of Virology</i> , 2015 , 160, 1593-619	2.6	430
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134	Genome Sequence of a Diverse Goose Circovirus Recovered from Greylag Goose. <i>Genome Announcements</i> , 2015 , 3,		7
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130	Phylogenetic diversity and co-evolutionary signals among trophic levels change across a habitat edge. <i>Journal of Animal Ecology</i> , 2015 , 84, 364-72	4.7	19	
129	Avian Polyomavirus Genome Sequences Recovered from Parrots in Captive Breeding Facilities in Poland. <i>Genome Announcements</i> , 2015 , 3,		5	
128	Genome Sequences of Poaceae-Associated Gemycircularviruses from the Pacific Ocean Island of Tonga. <i>Genome Announcements</i> , 2015 , 3,		13	
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125	Genome Sequences of Beak and Feather Disease Virus in Urban Rainbow Lorikeets (Trichoglossus haematodus). <i>Genome Announcements</i> , 2015 , 3,		2	
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34	Complete sequences of tomato leaf curl Palampur virus isolates infecting cucurbits in Iran. <i>Archives of Virology</i> , 2009 , 154, 1015-8	2.6	24
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21	Six novel begomoviruses infecting tomato and associated weeds in Southeastern Brazil. <i>Archives of Virology</i> , 2008 , 153, 1985-9	2.6	73
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14	evolution through recombination. <i>Journal of General Virology</i> , 2007 , 88, 3458-3468 HIV type 1 subtype C gag and nef diversity in Southern Africa. <i>AIDS Research and Human Retroviruses</i> , 2007 , 23, 477-81	1.6	18
14	evolution through recombination. <i>Journal of General Virology</i> , 2007 , 88, 3458-3468 HIV type 1 subtype C gag and nef diversity in Southern Africa. <i>AIDS Research and Human Retroviruses</i> , 2007 , 23, 477-81 Evidence of ancient papillomavirus recombination. <i>Journal of General Virology</i> , 2006 , 87, 2527-2531 Plant-produced cottontail rabbit papillomavirus L1 protein protects against tumor challenge: a	1.6	18 56
14 13	evolution through recombination. <i>Journal of General Virology</i> , 2007 , 88, 3458-3468 HIV type 1 subtype C gag and nef diversity in Southern Africa. <i>AIDS Research and Human Retroviruses</i> , 2007 , 23, 477-81 Evidence of ancient papillomavirus recombination. <i>Journal of General Virology</i> , 2006 , 87, 2527-2531 Plant-produced cottontail rabbit papillomavirus L1 protein protects against tumor challenge: a proof-of-concept study. <i>Vaccine Journal</i> , 2006 , 13, 845-53 Recombination patterns in aphthoviruses mirror those found in other picornaviruses. <i>Journal of</i>	1.6	18 56 46
14 13 12	evolution through recombination. <i>Journal of General Virology</i> , 2007 , 88, 3458-3468 HIV type 1 subtype C gag and nef diversity in Southern Africa. <i>AIDS Research and Human Retroviruses</i> , 2007 , 23, 477-81 Evidence of ancient papillomavirus recombination. <i>Journal of General Virology</i> , 2006 , 87, 2527-2531 Plant-produced cottontail rabbit papillomavirus L1 protein protects against tumor challenge: a proof-of-concept study. <i>Vaccine Journal</i> , 2006 , 13, 845-53 Recombination patterns in aphthoviruses mirror those found in other picornaviruses. <i>Journal of Virology</i> , 2006 , 80, 11827-32 Restoration of native folding of single-stranded DNA sequences through reverse mutations: an	1.6 4.9 6.6	18 56 46 234
14 13 12 11	HIV type 1 subtype C gag and nef diversity in Southern Africa. AIDS Research and Human Retroviruses, 2007, 23, 477-81 Evidence of ancient papillomavirus recombination. Journal of General Virology, 2006, 87, 2527-2531 Plant-produced cottontail rabbit papillomavirus L1 protein protects against tumor challenge: a proof-of-concept study. Vaccine Journal, 2006, 13, 845-53 Recombination patterns in aphthoviruses mirror those found in other picornaviruses. Journal of Virology, 2006, 80, 11827-32 Restoration of native folding of single-stranded DNA sequences through reverse mutations: an indication of a new epigenetic mechanism. Archives of Biochemistry and Biophysics, 2006, 453, 108-22 Transient expression of Human papillomavirus type 16 L1 protein in Nicotiana benthamiana using	1.6 4.9 6.6 4.1	18 56 46 234 20

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3	VIRIDIC 🖪 novel tool to calculate the intergenomic similarities of prokaryote-infecting viruses		16
2	Discovery of several thousand highly diverse circular DNA viruses		5
1	crAssphage abundance and genomic selective pressure correlate with altered bacterial abundance in the fecal microbiota of South African mother-infant dyads		2