

David Prangishvili

List of Publications by Year in descending order

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80
papers

4,841
citations

81839

39
h-index

102432

66
g-index

83
all docs

83
docs citations

83
times ranked

2872
citing authors

#	ARTICLE	IF	CITATIONS
1	Viruses of the Archaea: a unifying view. <i>Nature Reviews Microbiology</i> , 2006, 4, 837-848.	13.6	344
2	Genomics of Bacterial and Archaeal Viruses: Dynamics within the Prokaryotic Virosphere. <i>Microbiology and Molecular Biology Reviews</i> , 2011, 75, 610-635.	2.9	210
3	Viruses of archaea: Structural, functional, environmental and evolutionary genomics. <i>Virus Research</i> , 2018, 244, 181-193.	1.1	175
4	Taxonomy of prokaryotic viruses: 2017 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. <i>Archives of Virology</i> , 2018, 163, 1125-1129.	0.9	172
5	The enigmatic archaeal virosphere. <i>Nature Reviews Microbiology</i> , 2017, 15, 724-739.	13.6	169
6	Casposons: a new superfamily of self-synthesizing DNA transposons at the origin of prokaryotic CRISPR-Cas immunity. <i>BMC Biology</i> , 2014, 12, 36.	1.7	156
7	The archeoviruses. <i>FEMS Microbiology Reviews</i> , 2011, 35, 1035-1054.	3.9	150
8	Genetic elements in the extremely thermophilic archaeon <i>Sulfolobus</i> . <i>Extremophiles</i> , 1998, 2, 131-140.	0.9	148
9	The Wonderful World of Archaeal Viruses. <i>Annual Review of Microbiology</i> , 2013, 67, 565-585.	2.9	145
10	Taxonomy of prokaryotic viruses: 2018-2019 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. <i>Archives of Virology</i> , 2020, 165, 1253-1260.	0.9	144
11	Genome Analyses of Icelandic Strains of <i>Sulfolobus islandicus</i> , Model Organisms for Genetic and Virus-Host Interaction Studies. <i>Journal of Bacteriology</i> , 2011, 193, 1672-1680.	1.0	139
12	AFV1, a novel virus infecting hyperthermophilic archaea of the genus <i>acidianus</i> . <i>Virology</i> , 2003, 315, 68-79.	1.1	124
13	Sequences and Replication of Genomes of the Archaeal Rudiviruses SIRV1 and SIRV2: Relationships to the Archaeal Lipothrixvirus SIFV and Some Eukaryal Viruses. <i>Virology</i> , 2001, 291, 226-234.	1.1	112
14	Structural and Genomic Properties of the Hyperthermophilic Archaeal Virus ATV with an Extracellular Stage of the Reproductive Cycle. <i>Journal of Molecular Biology</i> , 2006, 359, 1203-1216.	2.0	110
15	Virus-mediated archaeal hecatomb in the deep seafloor. <i>Science Advances</i> , 2016, 2, e1600492.	4.7	107
16	A virus that infects a hyperthermophile encapsidates A-form DNA. <i>Science</i> , 2015, 348, 914-917.	6.0	98
17	Four newly isolated fuselloviruses from extreme geothermal environments reveal unusual morphologies and a possible interviral recombination mechanism. <i>Environmental Microbiology</i> , 2009, 11, 2849-2862.	1.8	85
18	Massive Activation of Archaeal Defense Genes during Viral Infection. <i>Journal of Virology</i> , 2013, 87, 8419-8428.	1.5	84

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19	Bipartite Network Analysis of the Archaeal Virosphere: Evolutionary Connections between Viruses and Capsidless Mobile Elements. <i>Journal of Virology</i> , 2016, 90, 11043-11055.	1.5	84
20	Taxonomy of prokaryotic viruses: update from the ICTV bacterial and archaeal viruses subcommittee. <i>Archives of Virology</i> , 2016, 161, 1095-1099.	0.9	83
21	Diversity of virus-host systems in hypersaline Lake Retba, Senegal. <i>Environmental Microbiology</i> , 2011, 13, 1956-1972.	1.8	82
22	Archaeal virus with exceptional virion architecture and the largest single-stranded DNA genome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 13386-13391.	3.3	78
23	pING Family of Conjugative Plasmids from the Extremely Thermophilic Archaeon <i>Sulfolobus islandicus</i> : Insights into Recombination and Conjugation in Crenarchaeota. <i>Journal of Bacteriology</i> , 2000, 182, 7014-7020.	1.0	74
24	Viruses of hyperthermophilic Crenarchaea. <i>Trends in Microbiology</i> , 2005, 13, 535-542.	3.5	74
25	Unification of the Globally Distributed Spindle-Shaped Viruses of the Archaea. <i>Journal of Virology</i> , 2014, 88, 2354-2358.	1.5	72
26	Familial Relationships in Hyperthermo- and Acidophilic Archaeal Viruses. <i>Journal of Virology</i> , 2010, 84, 4747-4754.	1.5	66
27	First Insights into the Entry Process of Hyperthermophilic Archaeal Viruses. <i>Journal of Virology</i> , 2013, 87, 13379-13385.	1.5	66
28	Eukaryotic-Like Virus Budding in Archaea. <i>MBio</i> , 2016, 7, .	1.8	65
29	Conjugation in Archaea: Frequent Occurrence of Conjugative Plasmids in <i>Sulfolobus</i> . <i>Plasmid</i> , 1998, 40, 190-202.	0.4	61
30	Analysis of metagenomic data reveals common features of halophilic viral communities across continents. <i>Environmental Microbiology</i> , 2016, 18, 889-903.	1.8	59
31	Taxonomy of prokaryotic viruses: 2016 update from the ICTV bacterial and archaeal viruses subcommittee. <i>Archives of Virology</i> , 2017, 162, 1153-1157.	0.9	57
32	Provirus Induction in Hyperthermophilic Archaea: Characterization of <i>Aeropyrum pernix</i> Spindle-Shaped Virus 1 and <i>Aeropyrum pernix</i> Ovoid Virus 1. <i>Journal of Bacteriology</i> , 2011, 193, 5412-5419.	1.0	56
33	Genomics and genetics of <i>Sulfolobus islandicus</i> LAL14/1, a model hyperthermophilic archaeon. <i>Open Biology</i> , 2013, 3, 130010.	1.5	55
34	<i>Sulfolobus</i> Spindle-Shaped Virus 1 Contains Glycosylated Capsid Proteins, a Cellular Chromatin Protein, and Host-Derived Lipids. <i>Journal of Virology</i> , 2015, 89, 11681-11691.	1.5	54
35	Structure and Genome Organization of AFV2, a Novel Archaeal Lipothrixvirus with Unusual Terminal and Core Structures. <i>Journal of Bacteriology</i> , 2005, 187, 3855-3858.	1.0	51
36	Virus-borne mini-CRISPR arrays are involved in interviral conflicts. <i>Nature Communications</i> , 2019, 10, 5204.	5.8	50

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37	Dark matter in archaeal genomes: a rich source of novel mobile elements, defense systems and secretory complexes. <i>Extremophiles</i> , 2014, 18, 877-893.	0.9	48
38	An extensively glycosylated archaeal pilus survives extreme conditions. <i>Nature Microbiology</i> , 2019, 4, 1401-1410.	5.9	46
39	A new proposed taxon for double-stranded DNA viruses, the order "Ligamenvirales". <i>Archives of Virology</i> , 2012, 157, 791-795.	0.9	45
40	Self-assembly of the general membrane-remodeling protein PVAP into sevenfold virus-associated pyramids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 3829-3834.	3.3	45
41	Diversity, taxonomy, and evolution of archaeal viruses of the class Caudoviricetes. <i>PLoS Biology</i> , 2021, 19, e3001442.	2.6	44
42	Viruses in acidic geothermal environments of the Kamchatka Peninsula. <i>Research in Microbiology</i> , 2008, 159, 358-366.	1.0	41
43	Metagenomic analyses of novel viruses and plasmids from a cultured environmental sample of hyperthermophilic neutrophiles. <i>Environmental Microbiology</i> , 2010, 12, 2918-2930.	1.8	39
44	Integrated mobile genetic elements in Thaumarchaeota. <i>Environmental Microbiology</i> , 2019, 21, 2056-2078.	1.8	38
45	Genomics and biology of Rudiviruses, a model for the study of virus-host interactions in Archaea. <i>Biochemical Society Transactions</i> , 2013, 41, 443-450.	1.6	37
46	Model for a novel membrane envelope in a filamentous hyperthermophilic virus. <i>ELife</i> , 2017, 6, .	2.8	37
47	Abundant Lysine Methylation and N-Terminal Acetylation in <i>Sulfolobus islandicus</i> Revealed by Bottom-Up and Top-Down Proteomics. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 3388-3404.	2.5	36
48	Structure and assembly of archaeal viruses. <i>Advances in Virus Research</i> , 2020, 108, 127-164.	0.9	35
49	A virus of hyperthermophilic archaea with a unique architecture among DNA viruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 2478-2483.	3.3	32
50	Unique architecture of thermophilic archaeal virus APBV1 and its genome packaging. <i>Nature Communications</i> , 2017, 8, 1436.	5.8	31
51	Structures of filamentous viruses infecting hyperthermophilic archaea explain DNA stabilization in extreme environments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 19643-19652.	3.3	29
52	New virus isolates from Italian hydrothermal environments underscore the biogeographic pattern in archaeal virus communities. <i>ISME Journal</i> , 2020, 14, 1821-1833.	4.4	29
53	Novel haloarchaeal viruses from Lake Retba infecting <i>Haloferax</i> and <i>Halorubrum</i> species. <i>Environmental Microbiology</i> , 2019, 21, 2129-2147.	1.8	28
54	Exceptional virion release mechanism: one more surprise from archaeal viruses. <i>Current Opinion in Microbiology</i> , 2011, 14, 315-320.	2.3	26

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55	Structural conservation in a membrane-enveloped filamentous virus infecting a hyperthermophilic acidophile. <i>Nature Communications</i> , 2018, 9, 3360.	5.8	24
56	The structures of two archaeal type IV pili illuminate evolutionary relationships. <i>Nature Communications</i> , 2020, 11, 3424.	5.8	24
57	A packing for A-form DNA in an icosahedral virus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 22591-22597.	3.3	23
58	<i>Adnaviria</i> : a New Realm for Archaeal Filamentous Viruses with Linear A-Form Double-Stranded DNA Genomes. <i>Journal of Virology</i> , 2021, 95, e0067321.	1.5	22
59	A Novel Type of Polyhedral Viruses Infecting Hyperthermophilic Archaea. <i>Journal of Virology</i> , 2017, 91, .	1.5	21
60	Archaeal viruses: living fossils of the ancient virosphere?. <i>Annals of the New York Academy of Sciences</i> , 2015, 1341, 35-40.	1.8	20
61	New archaeal viruses discovered by metagenomic analysis of viral communities in enrichment cultures. <i>Environmental Microbiology</i> , 2019, 21, 2002-2014.	1.8	18
62	Protein-Protein Interactions Leading to Recruitment of the Host DNA Sliding Clamp by the Hyperthermophilic <i>Sulfolobus islandicus</i> Rod-Shaped Virus 2. <i>Journal of Virology</i> , 2014, 88, 7105-7108.	1.5	16
63	Evolution of an archaeal virus nucleocapsid protein from the CRISPR-associated Cas4 nuclease. <i>Biology Direct</i> , 2015, 10, 65.	1.9	16
64	Going to extremes – a metagenomic journey into the dark matter of life. <i>FEMS Microbiology Letters</i> , 2021, 368, .	0.7	16
65	Structure of a filamentous virus uncovers familial ties within the archaeal virosphere. <i>Virus Evolution</i> , 2020, 6, veaa023.	2.2	13
66	Mysterious hexagonal pyramids on the surface of <i>Pyrobaculum</i> cells. <i>Biochimie</i> , 2015, 118, 365-367.	1.3	10
67	DNA-Interacting Characteristics of the Archaeal Rudiviral Protein SIRV2_Gp1. <i>Viruses</i> , 2017, 9, 190.	1.5	10
68	Centennial celebration of the bacteriophage research. <i>Research in Microbiology</i> , 2018, 169, 479-480.	1.0	9
69	ICTV Virus Taxonomy Profile: Bicaudaviridae. <i>Journal of General Virology</i> , 2018, 99, 864-865.	1.3	9
70	Rolling-circle replication initiation protein of haloarchaeal sphaerolipovirus SNJ1 is homologous to bacterial transposases of the IS91 family insertion sequences. <i>Journal of General Virology</i> , 2018, 99, 416-421.	1.3	8
71	A filamentous archaeal virus is enveloped inside the cell and released through pyramidal portals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	7
72	Egress of archaeal viruses. <i>Cellular Microbiology</i> , 2021, 23, e13394.	1.1	7

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73	ICTV Virus Taxonomy Profile: Guttaviridae. Journal of General Virology, 2018, 99, 290-291.	1.3	7
74	ICTV Virus Taxonomy Profile: Tristromaviridae. Journal of General Virology, 2019, 100, 135-136.	1.3	7
75	New insights into the diversity and evolution of the archaeal mobilome from three complete genomes of <i>Saccharolobus shibatae</i> . Environmental Microbiology, 2021, 23, 4612-4630.	1.8	5
76	ICTV Virus Taxonomy Profile: Globuloviridae. Journal of General Virology, 2018, 99, 1357-1358.	1.3	5
77	ICTV Virus Taxonomy Profile: Ampullaviridae. Journal of General Virology, 2018, 99, 288-289.	1.3	4
78	ICTV Virus Taxonomy Profile: Spiraviridae. Journal of General Virology, 2020, 101, 240-241.	1.3	3
79	ICTV Virus Taxonomy Profile: Clavaviridae. Journal of General Virology, 2019, 100, 1267-1268.	1.3	2
80	ICTV Virus Taxonomy Profile: Portogloboviridae. Journal of General Virology, 2021, 102, .	1.3	1