

Mya Breitbart

List of Publications by Citations

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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.

140 papers	14,617 citations	61 h-index	120 g-index
152 ext. papers	17,520 ext. citations	7 avg, IF	6.53 L-index

#	Paper	IF	Citations
140	Functional metagenomic profiling of nine biomes. <i>Nature</i> , 2008 , 452, 629-32	50.4	726
139	The marine viromes of four oceanic regions. <i>PLoS Biology</i> , 2006 , 4, e368	9.7	726
138	Genomic analysis of uncultured marine viral communities. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 14250-5	11.5	710
137	Metagenomic analyses of an uncultured viral community from human feces. <i>Journal of Bacteriology</i> , 2003 , 185, 6220-3	3.5	595
136	Here a virus, there a virus, everywhere the same virus?. <i>Trends in Microbiology</i> , 2005 , 13, 278-84	12.4	555
135	RNA viral community in human feces: prevalence of plant pathogenic viruses. <i>PLoS Biology</i> , 2006 , 4, e3	9.7	453
134	Laboratory procedures to generate viral metagenomes. <i>Nature Protocols</i> , 2009 , 4, 470-83	18.8	411
133	Metagenomic and small-subunit rRNA analyses reveal the genetic diversity of bacteria, archaea, fungi, and viruses in soil. <i>Applied and Environmental Microbiology</i> , 2007 , 73, 7059-66	4.8	406
132	Consensus statement: Virus taxonomy in the age of metagenomics. <i>Nature Reviews Microbiology</i> , 2017 , 15, 161-168	22.2	375
131	Using pyrosequencing to shed light on deep mine microbial ecology. <i>BMC Genomics</i> , 2006 , 7, 57	4.5	352
130	Marine viruses: truth or dare. <i>Annual Review of Marine Science</i> , 2012 , 4, 425-48	15.4	344
129	Diversity of bacteria associated with the Caribbean coral <i>Montastraea franksi</i> . <i>Coral Reefs</i> , 2001 , 20, 85-94	4.2	310
128	Viral and microbial community dynamics in four aquatic environments. <i>ISME Journal</i> , 2010 , 4, 739-51	11.9	305
127	Viral diversity and dynamics in an infant gut. <i>Research in Microbiology</i> , 2008 , 159, 367-73	4	234
126	Diversity and population structure of a near-shore marine-sediment viral community. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2004 , 271, 565-74	4.4	227
125	Metagenomic analysis of viruses in reclaimed water. <i>Environmental Microbiology</i> , 2009 , 11, 2806-20	5.2	218
124	Phage puppet masters of the marine microbial realm. <i>Nature Microbiology</i> , 2018 , 3, 754-766	26.6	216

123	Biodiversity and biogeography of phages in modern stromatolites and thrombolites. <i>Nature</i> , 2008 , 452, 340-3	50.4	212
122	A field guide to eukaryotic circular single-stranded DNA viruses: insights gained from metagenomics. <i>Archives of Virology</i> , 2012 , 157, 1851-71	2.6	198
121	Transposases are the most abundant, most ubiquitous genes in nature. <i>Nucleic Acids Research</i> , 2010 , 38, 4207-17	20.1	196
120	Exploring the viral world through metagenomics. <i>Current Opinion in Virology</i> , 2011 , 1, 289-97	7.5	189
119	Densovirus associated with sea-star wasting disease and mass mortality. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 17278-83	11.5	187
118	Role of elevated organic carbon levels and microbial activity in coral mortality. <i>Marine Ecology - Progress Series</i> , 2006 , 314, 119-125	2.6	184
117	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
116	Minimum Information about an Uncultivated Virus Genome (MIUViG). <i>Nature Biotechnology</i> , 2019 , 37, 29-37	44.5	180
115	Exploring the Vast Diversity of Marine Viruses. <i>Oceanography</i> , 2007 , 20, 135-139	2.3	173
114	FastGroupII: a web-based bioinformatics platform for analyses of large 16S rDNA libraries. <i>BMC Bioinformatics</i> , 2006 , 7, 57	3.6	173
113	Broad surveys of DNA viral diversity obtained through viral metagenomics of mosquitoes. <i>PLoS ONE</i> , 2011 , 6, e20579	3.7	168
112	Pepper mild mottle virus as an indicator of fecal pollution. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 7261-7	4.8	165
111	Global distribution of nearly identical phage-encoded DNA sequences. <i>FEMS Microbiology Letters</i> , 2004 , 236, 249-256	2.9	165
110	Metagenomic and stable isotopic analyses of modern freshwater microbialites in Cuatro Ciénegas, Mexico. <i>Environmental Microbiology</i> , 2009 , 11, 16-34	5.2	158
109	PHACCS, an online tool for estimating the structure and diversity of uncultured viral communities using metagenomic information. <i>BMC Bioinformatics</i> , 2005 , 6, 41	3.6	142
108	Diverse circular ssDNA viruses discovered in dragonflies (Odonata: Epiprocta). <i>Journal of General Virology</i> , 2012 , 93, 2668-2681	4.9	140
107	The complete genomic sequence of the marine phage Roseophage SIO1 shares homology with nonmarine phages. <i>Limnology and Oceanography</i> , 2000 , 45, 408-418	4.8	135
106	Diverse circovirus-like genome architectures revealed by environmental metagenomics. <i>Journal of General Virology</i> , 2009 , 90, 2418-2424	4.9	134

105	Method for discovering novel DNA viruses in blood using viral particle selection and shotgun sequencing. <i>BioTechniques</i> , 2005 , 39, 729-36	2.5	133
104	Comparative metagenomics of microbial traits within oceanic viral communities. <i>ISME Journal</i> , 2011 , 5, 1178-90	11.9	119
103	A bacterial metapopulation adapts locally to phage predation despite global dispersal. <i>Genome Research</i> , 2008 , 18, 293-7	9.7	119
102	Frequent detection of highly diverse variants of cardiovirus, cosavirus, bocavirus, and circovirus in sewage samples collected in the United States. <i>Journal of Clinical Microbiology</i> , 2009 , 47, 3507-13	9.7	118
101	Phage community dynamics in hot springs. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 1633-40	4.8	118
100	Discovery of a novel single-stranded DNA virus from a sea turtle fibropapilloma by using viral metagenomics. <i>Journal of Virology</i> , 2009 , 83, 2500-9	6.6	109
99	Towards quantitative viromics for both double-stranded and single-stranded DNA viruses. <i>PeerJ</i> , 2016 , 4, e2777	3.1	108
98	Ocean time-series reveals recurring seasonal patterns of viroplankton dynamics in the northwestern Sargasso Sea. <i>ISME Journal</i> , 2012 , 6, 273-84	11.9	106
97	Coral-associated Archaea. <i>Marine Ecology - Progress Series</i> , 2004 , 273, 89-96	2.6	106
96	Eukaryotic viruses in wastewater samples from the United States. <i>Applied and Environmental Microbiology</i> , 2009 , 75, 1402-9	4.8	99
95	A case study of enteric virus removal and insights into the associated risk of water reuse for two wastewater treatment pond systems in Bolivia. <i>Water Research</i> , 2014 , 65, 257-70	12.5	88
94	Dragonfly cyclovirus, a novel single-stranded DNA virus discovered in dragonflies (Odonata: Anisoptera). <i>Journal of General Virology</i> , 2011 , 92, 1302-1308	4.9	88
93	ICTV Virus Taxonomy Profile: Circoviridae. <i>Journal of General Virology</i> , 2017 , 98, 1997-1998	4.9	84
92	Exploring the diversity of plant DNA viruses and their satellites using vector-enabled metagenomics on whiteflies. <i>PLoS ONE</i> , 2011 , 6, e19050	3.7	83
91	Global distribution of nearly identical phage-encoded DNA sequences. <i>FEMS Microbiology Letters</i> , 2004 , 236, 249-56	2.9	80
90	Eukaryotic Circular Rep-Encoding Single-Stranded DNA (CRESS DNA) Viruses: Ubiquitous Viruses With Small Genomes and a Diverse Host Range. <i>Advances in Virus Research</i> , 2019 , 103, 71-133	10.7	80
89	A bioinformatic analysis of ribonucleotide reductase genes in phage genomes and metagenomes. <i>BMC Evolutionary Biology</i> , 2013 , 13, 33	3	79
88	Diversity and distribution of single-stranded DNA phages in the North Atlantic Ocean. <i>ISME Journal</i> , 2011 , 5, 822-30	11.9	78

87	Development of phoH as a novel signature gene for assessing marine phage diversity. <i>Applied and Environmental Microbiology</i> , 2011 , 77, 7730-9	4.8	76
86	Novel anellovirus discovered from a mortality event of captive California sea lions. <i>Journal of General Virology</i> , 2009 , 90, 1256-1261	4.9	74
85	Diversity of virus-host systems in hypersaline Lake Retba, Senegal. <i>Environmental Microbiology</i> , 2011 , 13, 1956-72	5.2	73
84	Evaluation of Filtration and DNA Extraction Methods for Environmental DNA Biodiversity Assessments across Multiple Trophic Levels. <i>Frontiers in Marine Science</i> , 2017 , 4,	4.5	70
83	Pepper mild mottle virus: A plant pathogen with a greater purpose in (waste)water treatment development and public health management. <i>Water Research</i> , 2018 , 144, 1-12	12.5	68
82	Environmental DNA reveals seasonal shifts and potential interactions in a marine community. <i>Nature Communications</i> , 2020 , 11, 254	17.4	66
81	Molecular and microscopic evidence of viruses in marine copepods. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 1375-80	11.5	62
80	Use of fluorescently labeled phage in the detection and identification of bacterial species. <i>Applied Spectroscopy</i> , 2003 , 57, 1138-44	3.1	60
79	Widespread association of a Rickettsiales-like bacterium with reef-building corals. <i>Environmental Microbiology</i> , 2004 , 6, 1137-48	5.2	59
78	Evaluation of marine zooplankton community structure through environmental DNA metabarcoding. <i>Limnology and Oceanography: Methods</i> , 2018 , 16, 209-221	2.6	58
77	Membrane vesicles in sea water: heterogeneous DNA content and implications for viral abundance estimates. <i>ISME Journal</i> , 2017 , 11, 394-404	11.9	58
76	Managing Microbial Risks from Indirect Wastewater Reuse for Irrigation in Urbanizing Watersheds. <i>Environmental Science & Technology</i> , 2016 , 50, 6803-13	10.3	57
75	Genomic analysis of multiple Roseophage SIO1 strains. <i>Environmental Microbiology</i> , 2009 , 11, 2863-73	5.2	54
74	Single-stranded DNA phages: from early molecular biology tools to recent revolutions in environmental microbiology. <i>FEMS Microbiology Letters</i> , 2016 , 363,	2.9	52
73	Novel circular single-stranded DNA viruses identified in marine invertebrates reveal high sequence diversity and consistent predicted intrinsic disorder patterns within putative structural proteins. <i>Frontiers in Microbiology</i> , 2015 , 6, 696	5.7	52
72	Metagenomic analysis of lysogeny in Tampa Bay: implications for prophage gene expression. <i>PLoS ONE</i> , 2008 , 3, e3263	3.7	52
71	Metagenomic identification of a novel anellovirus in Pacific harbor seal (<i>Phoca vitulina richardsii</i>) lung samples and its detection in samples from multiple years. <i>Journal of General Virology</i> , 2011 , 92, 1318-1323	4.9	51
70	Diversity of environmental single-stranded DNA phages revealed by PCR amplification of the partial major capsid protein. <i>ISME Journal</i> , 2014 , 8, 2093-103	11.9	50

69	High global diversity of cycloviruses amongst dragonflies. <i>Journal of General Virology</i> , 2013 , 94, 1827-1840	4.9	48
68	Metagenomic sequencing for virus identification in a public-health setting. <i>Journal of General Virology</i> , 2010 , 91, 2846-56	4.9	48
67	: a Virus Phylum Unifying Seven Families of Rep-Encoding Viruses with Single-Stranded, Circular DNA Genomes. <i>Journal of Virology</i> , 2020 , 94,	6.6	47
66	Widespread occurrence of phage-encoded exotoxin genes in terrestrial and aquatic environments in Southern California. <i>FEMS Microbiology Letters</i> , 2006 , 261, 141-9	2.9	46
65	Comparative metagenomics: natural populations of induced prophages demonstrate highly unique, lower diversity viral sequences. <i>Environmental Microbiology</i> , 2014 , 16, 570-85	5.2	45
64	Discovery of a novel mastrevirus and alphasatellite-like circular DNA in dragonflies (Epirocta) from Puerto Rico. <i>Virus Research</i> , 2013 , 171, 231-7	6.4	43
63	Power law rank-abundance models for marine phage communities. <i>FEMS Microbiology Letters</i> , 2007 , 273, 224-8	2.9	43
62	Phage encoded H-NS: a potential achilles heel in the bacterial defence system. <i>PLoS ONE</i> , 2011 , 6, e20095	3.7	41
61	Variability and host density independence in inductions-based estimates of environmental lysogeny. <i>Nature Microbiology</i> , 2017 , 2, 17064	26.6	40
60	Spatial heterogeneity of bacterial communities in the mucus of <i>Montastraea annularis</i> . <i>Marine Ecology - Progress Series</i> , 2011 , 426, 29-40	2.6	40
59	Diverse and highly recombinant anelloviruses associated with Weddell seals in Antarctica. <i>Virus Evolution</i> , 2017 , 3, vex017	3.7	39
58	Microbial communities associated with skeletal tumors on <i>Porites compressa</i> . <i>FEMS Microbiology Letters</i> , 2005 , 243, 431-6	2.9	38
57	Faecal pollution along the southeastern coast of Florida and insight into the use of pepper mild mottle virus as an indicator. <i>Journal of Applied Microbiology</i> , 2016 , 121, 1469-1481	4.7	36
56	RNA viral metagenome of whiteflies leads to the discovery and characterization of a whitefly-transmitted carlavirus in North America. <i>PLoS ONE</i> , 2014 , 9, e86748	3.7	35
55	Microbial source tracking in shellfish harvesting waters in the Gulf of Nicoya, Costa Rica. <i>Water Research</i> , 2017 , 111, 177-184	12.5	32
54	A Framework for a Marine Biodiversity Observing Network Within Changing Continental Shelf Seascapes. <i>Oceanography</i> , 2014 , 27, 18-23	2.3	32
53	The Ferrojan Horse Hypothesis: Iron-Virus Interactions in the Ocean. <i>Frontiers in Marine Science</i> , 2016 , 3,	4.5	32
52	Unprecedented Diversity of ssDNA Phages from the Family Detected within the Gut of a Protochordate Model Organism (). <i>Viruses</i> , 2018 , 10,	6.2	30

51	Bacterial communities associated with the ctenophores <i>Mnemiopsis leidyi</i> and <i>Beroe ovata</i> . <i>FEMS Microbiology Ecology</i> , 2012 , 82, 90-101	4.3	30
50	Novel cyclovirus discovered in the Florida woods cockroach <i>Eurycotis floridana</i> (Walker). <i>Archives of Virology</i> , 2013 , 158, 1389-92	2.6	30
49	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
48	Begomovirus-Associated Satellite DNA Diversity Captured Through Vector-Enabled Metagenomic (VEM) Surveys Using Whiteflies (Aleyrodidae). <i>Viruses</i> , 2016 , 8,	6.2	30
47	Assessing eukaryotic biodiversity in the Florida Keys National Marine Sanctuary through environmental DNA metabarcoding. <i>Ecology and Evolution</i> , 2019 , 9, 1029-1040	2.8	28
46	Pepper mild mottle virus: Agricultural menace turned effective tool for microbial water quality monitoring and assessing (waste)water treatment technologies. <i>PLoS Pathogens</i> , 2019 , 15, e1007639	7.6	28
45	Metagenomic identification, seasonal dynamics, and potential transmission mechanisms of a <i>Daphnia</i> -associated single-stranded DNA virus in two temperate lakes. <i>Limnology and Oceanography</i> , 2013 , 58, 1605-1620	4.8	28
44	Spatially resolved genomic, stable isotopic, and lipid analyses of a modern freshwater microbialite from Cuatro Ciénegas, Mexico. <i>Astrobiology</i> , 2012 , 12, 685-98	3.7	27
43	Deep sequencing of the viral <i>phoH</i> gene reveals temporal variation, depth-specific composition, and persistent dominance of the same viral <i>phoH</i> genes in the Sargasso Sea. <i>PeerJ</i> , 2015 , 3, e997	3.1	27
42	Vector-Enabled Metagenomic (VEM) Surveys Using Whiteflies (Aleyrodidae) Reveal Novel Begomovirus Species in the New and Old Worlds. <i>Viruses</i> , 2015 , 7, 5553-70	6.2	25
41	Diversity of DNA and RNA Viruses in Indoor Air As Assessed via Metagenomic Sequencing. <i>Environmental Science & Technology</i> , 2018 , 52, 1014-1027	10.3	24
40	Metagenomic identification of a nodavirus and a circular ssDNA virus in semi-purified viral nucleic acids from the hepatopancreas of healthy <i>Farfantepenaeus duorarum</i> shrimp. <i>Diseases of Aquatic Organisms</i> , 2013 , 105, 237-42	1.7	24
39	Discovery of a novel circular DNA virus in the Forbes sea star, <i>Asterias forbesi</i> . <i>Archives of Virology</i> , 2015 , 160, 2349-51	2.6	23
38	Reduction of nutrients, microbes, and personal care products in domestic wastewater by a benchtop electrocoagulation unit. <i>Scientific Reports</i> , 2015 , 5, 9380	4.9	22
37	Multidimensional metrics for estimating phage abundance, distribution, gene density, and sequence coverage in metagenomes. <i>Frontiers in Microbiology</i> , 2015 , 6, 381	5.7	21
36	Affordable Enteric Virus Detection Techniques Are Needed to Support Changing Paradigms in Water Quality Management. <i>Clean - Soil, Air, Water</i> , 2015 , 43, 8-12	1.6	20
35	PhiSiGns: an online tool to identify signature genes in phages and design PCR primers for examining phage diversity. <i>BMC Bioinformatics</i> , 2012 , 13, 37	3.6	19
34	Distinct lineage of vesiculovirus from big brown bats, United States. <i>Emerging Infectious Diseases</i> , 2013 , 19, 1978-80	10.2	19

33	Genomic evolution, recombination, and inter-strain diversity of chelonid alphaherpesvirus 5 from Florida and Hawaii green sea turtles with fibropapillomatosis. <i>PeerJ</i> , 2018 , 6, e4386	3.1	18
32	Disparity between planktonic fish egg and larval communities as indicated by DNA barcoding. <i>Marine Ecology - Progress Series</i> , 2014 , 503, 195-204	2.6	17
31	Removal of Six Estrogenic Endocrine-Disrupting Compounds (EDCs) from Municipal Wastewater Using Aluminum Electrocoagulation. <i>Water (Switzerland)</i> , 2016 , 8, 128	3	17
30	Marine Viruses: Community Dynamics, Diversity and Impact on Microbial Processes443-479		16
29	Water column stratification structures viral community composition in the Sargasso Sea. <i>Aquatic Microbial Ecology</i> , 2015 , 76, 85-94	1.1	13
28	Relationships among microbial indicators of fecal pollution, microbial source tracking markers, and pathogens in Costa Rican coastal waters. <i>Water Research</i> , 2021 , 188, 116507	12.5	13
27	The gut virome of the protochordate model organism, <i>Ciona intestinalis</i> subtype A. <i>Virus Research</i> , 2018 , 244, 137-146	6.4	13
26	Discovery, Prevalence, and Persistence of Novel Circular Single-Stranded DNA Viruses in the Ctenophores <i>Mnemiopsis leidyi</i> and <i>Beroe ovata</i> . <i>Frontiers in Microbiology</i> , 2015 , 6, 1427	5.7	12
25	Isolation and Characterization of a Shewanella Phage-Host System from the Gut of the Tunicate, <i>Ciona intestinalis</i> . <i>Viruses</i> , 2017 , 9,	6.2	11
24	Prokaryotic and Viral Community Composition of Freshwater Springs in Florida, USA. <i>MBio</i> , 2020 , 11,	7.8	8
23	Discovery of Four Novel Circular Single-Stranded DNA Viruses in Fungus-Farming Termites. <i>Genome Announcements</i> , 2018 , 6,		8
22	Regeneration of macronutrients and trace metals during phytoplankton decay: An experimental study. <i>Limnology and Oceanography</i> , 2020 , 65, 1936-1960	4.8	6
21	Development of a Serological Assay for the Sea Lion (<i>Zalophus californianus</i>) Anellovirus, ZcAV. <i>Scientific Reports</i> , 2015 , 5, 9637	4.9	6
20	DNA barcoding reveals clear delineation between spawning sites for neritic versus oceanic fishes in the Gulf of Mexico. <i>Fisheries Oceanography</i> , 2019 , 28, 228-239	2.4	6
19	Genome Sequence of PM2-Like Phage Cr39582, Induced from a <i>Pseudoalteromonas</i> sp. Isolated from the Gut of <i>Ciona robusta</i> . <i>Genome Announcements</i> , 2018 , 6,		6
18	Population genomics of three deep-sea cephalopod species reveals connectivity between the Gulf of Mexico and northwestern Atlantic Ocean. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2020 , 158, 103222	2.5	4
17	Zoonotic Infection With Pigeon Paramyxovirus Type 1 Linked to Fatal Pneumonia. <i>Journal of Infectious Diseases</i> , 2018 , 218, 1037-1044	7	4
16	A Brilliant Impostor?. <i>Oceanography</i> , 2014 , 27, 14-14	2.3	4

15	Phage Ecology and Bacterial Pathogenesis66-91		4
14	Genome Sequence of from Passionfruit and in Florida. <i>Genome Announcements</i> , 2017 , 5,		3
13	Phage Eco-Locator: a web tool for visualization and analysis of phage genomes in metagenomic data sets. <i>BMC Bioinformatics</i> , 2011 , 12,	3.6	3
12	Discovery of a novel potexvirus in the seagrass <i>Thalassia testudinum</i> from Tampa Bay, Florida. <i>Limnology and Oceanography Letters</i> , 2019 , 4, 1-8	7.9	2
11	DNA barcoding of fish eggs collected off northwestern Cuba and across the Florida Straits demonstrates egg transport by mesoscale eddies. <i>Fisheries Oceanography</i> , 2020 , 29, 340-348	2.4	2
10	Prevalence of a vertically transmitted single-stranded DNA virus in spinybacked orbweavers (<i>Gasteracantha cancriformis</i>) from Florida, USA. <i>Journal of General Virology</i> , 2019 , 100, 1253-1265	4.9	2
9	Microviridae2, 1-14		2
8	Near-Complete Genome Sequence of a Novel Single-Stranded RNA Virus Discovered in Indoor Air. <i>Genome Announcements</i> , 2018 , 6,		1
7	MOLECULAR SURVEILLANCE OF PLANT VIRUSES: IDENTIFICATION OF NEW AND EMERGING VIRUSES OF TOMATO BEFORE THEY CAUSE EPIDEMICS. <i>Acta Horticulturae</i> , 2015 , 127-131	0.3	1
6	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
5	DNA Detectives: Outreach Activity Teaching Students to Identify Fish Eggs Using DNA Barcoding. <i>Journal of Microbiology and Biology Education</i> , 2021 , 22,	1.3	1
4	First-year graduate courses foster inclusion. <i>Nature Geoscience</i> , 2021 , 14, 539-540	18.3	1
3	Spatial and Temporal Dynamics of Prokaryotic and Viral Community Assemblages in a Lotic System (Manatee Springs, Florida). <i>Applied and Environmental Microbiology</i> , 2021 , 87, e0064621	4.8	1
2	Biodiversity and Biogeography of Phages in Modern Stromatolites and Thrombolites 2011 , 37-44		
1	Dissolved Inorganic Carbon-Accumulating Complexes from Autotrophic Bacteria from Extreme Environments. <i>Journal of Bacteriology</i> , 2021 , 203, e0037721	3.5	