

Abundant SAR11 viruses in the ocean

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Citation Report

#	ARTICLE	IF	CITATIONS
1	SAR11 viruses and defensive host strains. <i>Nature</i> , 2013, 499, E3-E4.	13.7	39
2	Giovannoni et al. reply. <i>Nature</i> , 2013, 499, E4-E5.	13.7	31
3	The global virome: not as big as we thought?. <i>Current Opinion in Virology</i> , 2013, 3, 566-571.	2.6	64
4	Patterns of ecological specialization among microbial populations in the <scp>R</scp>ed <scp>S</scp>ea and diverse oligotrophic marine environments. <i>Ecology and Evolution</i> , 2013, 3, 1780-1797.	0.8	45
5	Killers of the winners. <i>Nature</i> , 2013, 494, 320-321.	13.7	3
6	Single-cell genomics reveal low recombination frequencies in freshwater bacteria of the SAR11 clade. <i>Genome Biology</i> , 2013, 14, R130.	13.9	74
7	Preparation of Metagenomic Libraries from Naturally Occurring Marine Viruses. <i>Methods in Enzymology</i> , 2013, 531, 143-165.	0.4	33
8	Spatiotemporal distributions of rare bacterioplankton populations indicate adaptive strategies in the oligotrophic ocean. <i>Aquatic Microbial Ecology</i> , 2013, 71, 1-13.	0.9	90
9	Cheats never prosper. <i>Nature</i> , 2013, 494, 321-322.	13.7	3
10	Coevolutionary diversification creates nested-modular structure in phageâ€“bacteria interaction networks. <i>Interface Focus</i> , 2013, 3, 20130033.	1.5	73
11	Lineage specific gene family enrichment at the microscale in marine systems. <i>Current Opinion in Microbiology</i> , 2013, 16, 605-617.	2.3	16
12	Insight into the unknown marine virus majority. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 12166-12167.	3.3	12
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16	Metabolic reprogramming by viruses in the sunlit and dark ocean. <i>Genome Biology</i> , 2013, 14, R123.	13.9	208
17	Regulation of bacterial metabolic activity by dissolved organic carbon and viruses. <i>Journal of Geophysical Research G: Biogeosciences</i> , 2013, 118, 1573-1583.	1.3	23
18	Prevalence of psbA-containing cyanobacterial podoviruses in the ocean. <i>Scientific Reports</i> , 2013, 3, 3207.	1.6	12

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20	The Pacific Ocean Virome (POV): A Marine Viral Metagenomic Dataset and Associated Protein Clusters for Quantitative Viral Ecology. PLoS ONE, 2013, 8, e57355.	1.1	336
21	Optimal Defense Strategies in an Idealized Microbial Food Web under Trade-Off between Competition and Defense. PLoS ONE, 2014, 9, e101415.	1.1	29
22	Evidence for metaviromic islands in marine phages. Frontiers in Microbiology, 2014, 5, 27.	1.5	39
23	The distribution and impact of viral lineages in domains of life. Frontiers in Microbiology, 2014, 5, 194.	1.5	42
24	Diversity and abundance of Pelagibacterales (SAR11) in the Baltic Sea salinity gradient. Systematic and Applied Microbiology, 2014, 37, 601-604.	1.2	58
25	Tales from a thousand and one phages. Bacteriophage, 2014, 4, e28265.	1.9	17
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28	Dispersal network structure and infection mechanism shape diversity in a coevolutionary bacteria-phage system. ISME Journal, 2014, 8, 504-514.	4.4	9
29	Temperature dependent bacteriophages of a tropical bacterial pathogen. Frontiers in Microbiology, 2014, 5, 599.	1.5	63
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31	Unprecedented evidence for high viral abundance and lytic activity in coral reef waters of the South Pacific Ocean. Frontiers in Microbiology, 2014, 5, 493.	1.5	32
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44	Classification and quantification of bacteriophage taxa in human gut metagenomes. <i>ISME Journal</i> , 2014, 8, 1391-1402.	4.4	127
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