## Jos Flores-Uribe

## 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.

8 18 13 340 h-index g-index citations papers 12.6 18 486 3.1 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
13	A distinct abundant group of microbial rhodopsins discovered using functional metagenomics. <i>Nature</i> , <b>2018</b> , 558, 595-599	50.4	106
12	Novel Abundant Oceanic Viruses of Uncultured Marine Group II Euryarchaeota. <i>Current Biology</i> , <b>2017</b> , 27, 1362-1368	6.3	54
11	Production of therapeutic proteins in the chloroplast of Chlamydomonas reinhardtii. <i>AMB Express</i> , <b>2014</b> , 4, 57	4.1	47
10	A myovirus encoding both photosystem I and II proteins enhances cyclic electron flow in infected Prochlorococcus cells. <i>Nature Microbiology</i> , <b>2017</b> , 2, 1350-1357	26.6	44
9	MerMAIDs: a family of metagenomically discovered marine anion-conducting and intensely desensitizing channelrhodopsins. <i>Nature Communications</i> , <b>2019</b> , 10, 3315	17.4	33
8	Cyanophage-encoded lipid desaturases: oceanic distribution, diversity and function. <i>ISME Journal</i> , <b>2018</b> , 12, 343-355	11.9	14
7	Heliorhodopsins are absent in diderm (Gram-negative) bacteria: Some thoughts and possible implications for activity. <i>Environmental Microbiology Reports</i> , <b>2019</b> , 11, 419-424	3.7	13
6	Seasonal and diel patterns of abundance and activity of viruses in the Red Sea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2020</b> , 117, 29738-29747	11.5	9
5	A novel uncultured marine cyanophage lineage with lysogenic potential linked to a putative marine Synechococcus 'relic' prophage. <i>Environmental Microbiology Reports</i> , <b>2019</b> , 11, 598-604	3.7	8
4	Closing the gaps on the viral photosystem-I psaDCAB gene organization. <i>Environmental Microbiology</i> , <b>2015</b> , 17, 5100-8	5.2	6
3	An uncultured marine cyanophage encodes an active phycobilisome proteolysis adaptor protein NblA. <i>Environmental Microbiology Reports</i> , <b>2019</b> , 11, 848-854	3.7	3
2	A novel uncultured marine cyanophage lineage with lysogenic potential linked to a putative marineSynechococcus[elic[prophage		1
1	Uncultured marine cyanophages encode for active NblA, phycobilisome proteolysis adaptor protein		1