

# Pablo Yarza

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11322608/publications.pdf>

Version: 2024-02-01

20  
papers

28,966  
citations

430754

18  
h-index

794469

19  
g-index

21  
all docs

21  
docs citations

21  
times ranked

34591  
citing authors

#	ARTICLE	IF	CITATIONS
1	A phylogenetic framework for the kingdom Fungi based on 18S rRNA gene sequences. <i>Marine Genomics</i> , 2017, 36, 33-39.	0.4	47
2	25 years of serving the community with ribosomal RNA gene reference databases and tools. <i>Journal of Biotechnology</i> , 2017, 261, 169-176.	1.9	679
3	Expanding the World of Marine Bacterial and Archaeal Clades. <i>Frontiers in Microbiology</i> , 2015, 6, 1524.	1.5	122
4	The All-Species Living Tree Project. <i>Methods in Microbiology</i> , 2014, 41, 45-59.	0.4	10
5	Harmonized Phylogenetic Trees for The Prokaryotes. , 2014, , 1-3.		10
6	The SILVA and "All-species Living Tree Project (LTP)" taxonomic frameworks. <i>Nucleic Acids Research</i> , 2014, 42, D643-D648.	6.5	2,667
7	New insights into <i>Oculina patagonica</i> coral diseases and their associated <i>Vibrio</i> spp. communities. <i>ISME Journal</i> , 2014, 8, 1794-1807.	4.4	54
8	Uniting the classification of cultured and uncultured bacteria and archaea using 16S rRNA gene sequences. <i>Nature Reviews Microbiology</i> , 2014, 12, 635-645.	13.6	2,000
9	Culture-Independent Approaches for Studying Viruses from Hypersaline Environments. <i>Applied and Environmental Microbiology</i> , 2012, 78, 1635-1643.	1.4	70
10	Virioplankton Community Structure in Tunisian Solar Salterns. <i>Applied and Environmental Microbiology</i> , 2012, 78, 7429-7437.	1.4	51
11	The SILVA ribosomal RNA gene database project: improved data processing and web-based tools. <i>Nucleic Acids Research</i> , 2012, 41, D590-D596.	6.5	21,425
12	Response of sulfate-reducing bacteria to an artificial oil spill in a coastal marine sediment. <i>Environmental Microbiology</i> , 2011, 13, 1488-1499.	1.8	55
13	Release LTPs104 of the All-Species Living Tree. <i>Systematic and Applied Microbiology</i> , 2011, 34, 169-170.	1.2	146
14	Taxonomic and Functional Metagenomic Profiling of the Microbial Community in the Anoxic Sediment of a Sub-saline Shallow Lake (Laguna de Carrizo, Central Spain). <i>Microbial Ecology</i> , 2011, 62, 824-837.	1.4	51
15	Complete genome sequence of <i>Marinobacter adhaerens</i> type strain (HP15), a diatom-interacting marine microorganism. <i>Standards in Genomic Sciences</i> , 2010, 3, 97-107.	1.5	50
16	Update of the All-Species Living Tree Project based on 16S and 23S rRNA sequence analyses. <i>Systematic and Applied Microbiology</i> , 2010, 33, 291-299.	1.2	441
17	Fine-scale evolution: genomic, phenotypic and ecological differentiation in two coexisting <i>Salinibacter ruber</i> strains. <i>ISME Journal</i> , 2010, 4, 882-895.	4.4	81
18	The metavirome of a hypersaline environment. <i>Environmental Microbiology</i> , 2010, 12, 2965-2976.	1.8	78

#	ARTICLE	IF	CITATIONS
19	Extremely halophilic microbial communities in anaerobic sediments from a solar saltern. Environmental Microbiology Reports, 2010, 2, 258-271.	1.0	44
20	The All-Species Living Tree project: A 16S rRNA-based phylogenetic tree of all sequenced type strains. Systematic and Applied Microbiology, 2008, 31, 241-250.	1.2	884