

# Joran Martijn

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1269451/publications.pdf>

Version: 2024-02-01

15  
papers

1,867  
citations

687363

13  
h-index

996975

15  
g-index

19  
all docs

19  
docs citations

19  
times ranked

2555  
citing authors

#	ARTICLE	IF	CITATIONS
1	Complex archaea that bridge the gap between prokaryotes and eukaryotes. <i>Nature</i> , 2015, 521, 173-179.	27.8	995
2	Deep mitochondrial origin outside the sampled alphaproteobacteria. <i>Nature</i> , 2018, 557, 101-105.	27.8	278
3	From archaeon to eukaryote: the evolutionary dark ages of the eukaryotic cell. <i>Biochemical Society Transactions</i> , 2013, 41, 451-457.	3.4	111
4	A progestin (17 $\beta$ ,20 $\alpha$ -dihydroxy-4-pregnen-3-one) stimulates early stages of spermatogenesis in zebrafish. <i>General and Comparative Endocrinology</i> , 2013, 185, 1-9.	1.8	84
5	A <i>Rickettsiales</i> symbiont of amoebae with ancient features. <i>Environmental Microbiology</i> , 2016, 18, 2326-2342.	3.8	73
6	Marine Sediments Illuminate Chlamydiae Diversity and Evolution. <i>Current Biology</i> , 2020, 30, 1032-1048.e7.	3.9	52
7	Single-cell genomics of a rare environmental alphaproteobacterium provides unique insights into Rickettsiaceae evolution. <i>ISME Journal</i> , 2015, 9, 2373-2385.	9.8	51
8	Confident phylogenetic identification of uncultured prokaryotes through long read amplicon sequencing of the 16S-23S rRNA operon. <i>Environmental Microbiology</i> , 2019, 21, 2485-2498.	3.8	46
9	Hikarchaea demonstrate an intermediate stage in the methanogen-to-halophile transition. <i>Nature Communications</i> , 2020, 11, 5490.	12.8	39
10	Comparative and Phylogenomic Evidence That the Alphaproteobacterium HIMB59 Is Not a Member of the Oceanic SAR11 Clade. <i>PLoS ONE</i> , 2013, 8, e78858.	2.5	38
11	The evolutionary origin of host association in the Rickettsiales. <i>Nature Microbiology</i> , 2022, 7, 1189-1199.	13.3	29
12	Close Encounters of the Third Domain: The Emerging Genomic View of Archaeal Diversity and Evolution. <i>Archaea</i> , 2013, 2013, 1-12.	2.3	24
13	New Insights into Ligand-Receptor Pairing and Coevolution of Relaxin Family Peptides and Their Receptors in Teleosts. <i>International Journal of Evolutionary Biology</i> , 2012, 2012, 1-14.	1.0	23
14	Draft Genome Sequence of <i>Candidatus Moanabacter tarae</i> Representing a Novel Marine Verrucomicrobial Lineage. <i>Microbiology Resource Announcements</i> , 2018, 7, .	0.6	9
15	A novel alphaproteobacterium with a small genome identified from the digestive gland of multiple species of abalone. <i>Environmental Microbiology Reports</i> , 2020, 12, 387-395.	2.4	3