

# Riccardo Vicedomini

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9623115/publications.pdf>

Version: 2024-02-01

8  
papers

2,180  
citations

1477746

6  
h-index

1588620

8  
g-index

13  
all docs

13  
docs citations

13  
times ranked

4751  
citing authors

#	ARTICLE	IF	CITATIONS
1	Critical Assessment of Metagenome Interpretation: the second round of challenges. <i>Nature Methods</i> , 2022, 19, 429-440.	9.0	133
2	MyCLADE: a multi-source domain annotation server for sequence functional exploration. <i>Nucleic Acids Research</i> , 2021, 49, W452-W458.	6.5	2
3	Strainberry: automated strain separation in low-complexity metagenomes using long reads. <i>Nature Communications</i> , 2021, 12, 4485.	5.8	25
4	Targeted domain assembly for fast functional profiling of metagenomic datasets with S3A. <i>Bioinformatics</i> , 2020, 36, 3975-3981.	1.8	2
5	A multi-source domain annotation pipeline for quantitative metagenomic and metatranscriptomic functional profiling. <i>Microbiome</i> , 2018, 6, 149.	4.9	41
6	GAM-NGS: genomic assemblies merger for next generation sequencing. <i>BMC Bioinformatics</i> , 2013, 14, S6.	1.2	76
7	Assemblathon 2: evaluating de novo methods of genome assembly in three vertebrate species. <i>GigaScience</i> , 2013, 2, 10.	3.3	582
8	The Norway spruce genome sequence and conifer genome evolution. <i>Nature</i> , 2013, 497, 579-584.	13.7	1,303