

# Alfredo Mari

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/4249246/alfredo-mari-publications-by-year.pdf>

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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.

15  
papers

101  
citations

6  
h-index

9  
g-index

17  
ext. papers

173  
ext. citations

6  
avg, IF

1.87  
L-index

#	Paper	IF	Citations
15	Determinants of SARS-CoV-2 transmission to guide vaccination strategy in an urban area.. <i>Virus Evolution</i> , <b>2022</b> , 8, veac002	3.7	0
14	The Leaf Microbiome of Displays Reproducible Dynamics and Patterns throughout the Growing Season.. <i>MBio</i> , <b>2022</b> , e0282521	7.8	2
13	External quality assessment of SARS-CoV-2-sequencing: An ESGMD-SSM pilot trial across 15 European laboratories. <i>Journal of Clinical Microbiology</i> , <b>2021</b> , JCM0169821	9.7	2
12	SARS-CoV-2 outbreak in a tri-national urban area is dominated by a B.1 lineage variant linked to a mass gathering event. <i>PLoS Pathogens</i> , <b>2021</b> , 17, e1009374	7.6	20
11	SARS-CoV-2 N501Y Introductions and Transmissions in Switzerland from Beginning of October 2020 to February 2021-Implementation of Swiss-Wide Diagnostic Screening and Whole Genome Sequencing. <i>Microorganisms</i> , <b>2021</b> , 9,	4.9	16
10	Global Genomic Analysis of SARS-CoV-2 RNA Dependent RNA Polymerase Evolution and Antiviral Drug Resistance. <i>Microorganisms</i> , <b>2021</b> , 9,	4.9	4
9	Obtaining deeper insights into microbiome diversity using a simple method to block host and nontargets in amplicon sequencing. <i>Molecular Ecology Resources</i> , <b>2021</b> , 21, 1952-1965	8.4	1
8	Brief validation of the novel GeneXpert Xpress SARS-CoV-2 PCR assay. <i>Journal of Virological Methods</i> , <b>2020</b> , 284, 113925	2.6	25
7	SARS-CoV-2 phylogeny during the early outbreak in the Basel area, Switzerland: import and spread dominated by a single B.1 lineage variant (C15324T)		6
6	Obtaining deeper insights into microbiome diversity using a simple method to block host and non-targets in amplicon sequencing		1
5	Quantifying SARS-CoV-2 spread in Switzerland based on genomic sequencing data		10
4	Determinants of SARS-CoV-2 transmission to guide vaccination strategy in an urban area		8
3	A novel phyllosphere resident Protomyces species that interacts with the Arabidopsis immune system		1
2	Global surveillance of potential antiviral drug resistance in SARS-CoV-2: proof of concept focussing on the RNA-dependent RNA polymerase		1
1	SARS-CoV-2 N501Y introductions and transmissions in Switzerland from beginning of October 2020 to February 2021 Implementation of Swiss-wide diagnostic screening and whole genome sequencing		4