

Alfredo Mari

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.

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 | Brief validation of the novel GeneXpert Xpress SARS-CoV-2 PCR assay. <i>Journal of Virological Methods</i> , 2020 , 284, 113925 | 2.6 | 25 |
| 14 | 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> , 2021 , 17, e1009374 | 7.6 | 20 |
| 13 | 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> , 2021 , 9, | 4.9 | 16 |
| 12 | Quantifying SARS-CoV-2 spread in Switzerland based on genomic sequencing data | | 10 |
| 11 | Determinants of SARS-CoV-2 transmission to guide vaccination strategy in an urban area | | 8 |
| 10 | 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 |
| 9 | Global Genomic Analysis of SARS-CoV-2 RNA Dependent RNA Polymerase Evolution and Antiviral Drug Resistance. <i>Microorganisms</i> , 2021 , 9, | 4.9 | 4 |
| 8 | 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 |
| 7 | External quality assessment of SARS-CoV-2-sequencing: An ESGMD-SSM pilot trial across 15 European laboratories. <i>Journal of Clinical Microbiology</i> , 2021 , JCM0169821 | 9.7 | 2 |
| 6 | The Leaf Microbiome of Displays Reproducible Dynamics and Patterns throughout the Growing Season.. <i>MBio</i> , 2022 , e0282521 | 7.8 | 2 |
| 5 | Obtaining deeper insights into microbiome diversity using a simple method to block host and non-targets in amplicon sequencing | | 1 |
| 4 | A novel phyllosphere resident Protomyces species that interacts with the Arabidopsis immune system | | 1 |
| 3 | Obtaining deeper insights into microbiome diversity using a simple method to block host and nontargets in amplicon sequencing. <i>Molecular Ecology Resources</i> , 2021 , 21, 1952-1965 | 8.4 | 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 | Determinants of SARS-CoV-2 transmission to guide vaccination strategy in an urban area.. <i>Virus Evolution</i> , 2022 , 8, veac002 | 3.7 | 0 |