

# Micheli Filippi

## List of Publications by Year in descending order

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Version: 2024-02-01

11  
papers

218  
citations

1478505

6  
h-index

1372567

10  
g-index

12  
all docs

12  
docs citations

12  
times ranked

505  
citing authors

#	ARTICLE	IF	CITATIONS
1	Pervasive transmission of E484K and emergence of VUI-NP13L with evidence of SARS-CoV-2 co-infection events by two different lineages in Rio Grande do Sul, Brazil. <i>Virus Research</i> , 2021, 296, 198345.	2.2	105
2	Early detection of SARS-CoV-2 P.1 variant in Southern Brazil and reinfection of the same patient by P.2. <i>Revista Do Instituto De Medicina Tropical De Sao Paulo</i> , 2021, 63, e58.	1.1	31
3	Low circulation of Influenza A and coinfection with SARS-CoV-2 among other respiratory viruses during the COVID-19 pandemic in a region of southern Brazil. <i>Journal of Medical Virology</i> , 2021, 93, 4392-4398.	5.0	22
4	Human mastadenovirus in water, sediment, sea surface microlayer, and bivalve mollusk from southern Brazilian beaches. <i>Marine Pollution Bulletin</i> , 2019, 142, 335-349.	5.0	18
5	Early introduction, dispersal and evolution of Delta SARS-CoV-2 in Southern Brazil, late predominance of AY.99.2 and AY.101 related lineages. <i>Virus Research</i> , 2022, 311, 198702.	2.2	15
6	Predominance of SARS-CoV-2 P.1 (Gamma) lineage inducing the recent COVID-19 wave in southern Brazil and the finding of an additional S: D614A mutation. <i>Infection, Genetics and Evolution</i> , 2021, 96, 105134.	2.3	11
7	Soil contamination of a public park by human and canine mastadenovirus, as well as hookworms and <i>Toxocara</i> spp eggs. <i>Revista Do Instituto De Medicina Tropical De Sao Paulo</i> , 2019, 61, e60.	1.1	7
8	Brief dispersion of a putative B.1.1.28-derived SARS-CoV-2 lineage harboring additional N234P and E471Q spike protein mutations in individuals crossing the Argentina-Brazil border. <i>Travel Medicine and Infectious Disease</i> , 2022, 49, 102390.	3.0	3
9	Reinfection cases by closely related SARS-CoV-2 lineages in Southern Brazil. <i>Brazilian Journal of Microbiology</i> , 2021, 52, 1881-1885.	2.0	2
10	Viral isolation allows characterization of early samples of SARS-CoV-2 lineage B1.1.33 with unique mutations (S: H655Y and T63N) circulating in Southern Brazil in 2020. <i>Brazilian Journal of Microbiology</i> , 2022, 53, 1313-1319.	2.0	2
11	Emerging Porcine adenovirus PAdV-SVN1 and other enteric viruses in samples of industrialized meat by-products. <i>Ciencia Rural</i> , 2020, 50, .	0.5	0