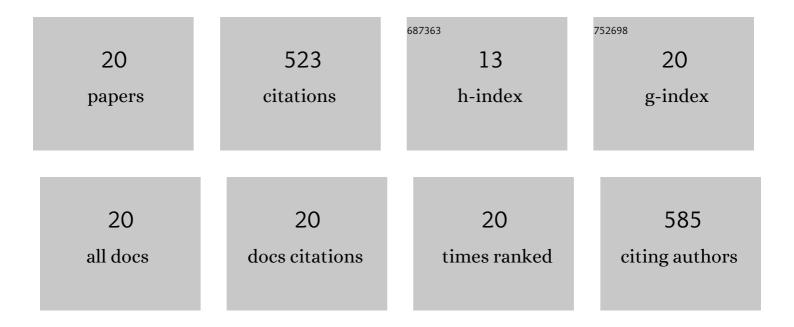
## Alexandre Madi Fialho

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4986982/publications.pdf

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#	Article	lF	CITATIONS
1	Laboratory-based Rotavirus Surveillance During the Introduction of a Vaccination Program, Brazil, 2005–2009. Pediatric Infectious Disease Journal, 2011, 30, S35-S41.	2.0	78
2	Rotavirus Genotype Distribution after Vaccine Introduction, Rio de Janeiro, Brazil. Emerging Infectious Diseases, 2009, 15, 95-97.	4.3	69
3	Assessment of Gastroenteric Viruses Frequency in a Children's Day Care Center in Rio De Janeiro, Brazil: A Fifteen Year Study (1994–2008). PLoS ONE, 2012, 7, e33754.	2.5	59
4	Noroviruses associated with outbreaks of acute gastroenteritis in the State of Rio Grande do Sul, Brazil, 2004–2011. Journal of Clinical Virology, 2014, 61, 345-352.	3.1	38
5	Brazilian P[8],G1, P[8],G5, P[8],G9, and P[4],G2 rotavirus strains: Nucleotide sequence and phylogenetic analysis. Journal of Medical Virology, 2007, 79, 995-1001.	5.0	33
6	G1P[8] species A rotavirus over 27 years – Pre- and post-vaccination eras – in Brazil: Full genomic constellation analysis and no evidence for selection pressure by Rotarix® vaccine. Infection, Genetics and Evolution, 2015, 30, 206-218.	2.3	30
7	Molecular analysis of the NSP4 and VP6 genes of rotavirus strains recovered from hospitalized children in Rio de Janeiro, Brazil. Journal of Medical Microbiology, 2007, 56, 854-859.	1.8	29
8	Detection and molecular characterization of group A rotavirus from hospitalized children in Rio de Janeiro, Brazil, 2004. Memorias Do Instituto Oswaldo Cruz, 2006, 101, 291-294.	1.6	26
9	Prevalence and genomic characterization of G2P[4] group A rotavirus strains during monovalent vaccine introduction in Brazil. Infection, Genetics and Evolution, 2014, 28, 486-494.	2.3	26
10	Rotavirus A in Brazil: Molecular Epidemiology and Surveillance during 2018–2019. Pathogens, 2020, 9, 515.	2.8	20
11	VP7 and VP8* genetic characterization of group A rotavirus genotype G12P[8]: Emergence and spreading in the Eastern Brazilian coast in 2014. Journal of Medical Virology, 2017, 89, 64-70.	5.0	18
12	Human enteric adenovirus F40/41 as a major cause of acute gastroenteritis in children in Brazil, 2018 to 2020. Scientific Reports, 2022, 12, .	3.3	17
13	High genetic diversity of noroviruses in children from a community-based study in Rio de Janeiro, Brazil, 2014-2018. Archives of Virology, 2019, 164, 1427-1432.	2.1	16
14	Factors associated with rotavirus diarrhoea in children living in a socially diverse urban centre in Brazil. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2012, 106, 445-451.	1.8	14
15	Phenotyping of Lewis and secretor HBGA from saliva and detection of new FUT2 gene SNPs from young children from the Amazon presenting acute gastroenteritis and respiratory infection. Infection, Genetics and Evolution, 2019, 70, 61-66.	2.3	12
16	Detection and Molecular Characterization of Human Group C Rotavirus in Brazil. Intervirology, 2011, 54, 261-267.	2.8	10
17	Epidemiology of enteric virus infections in children living in the Amazon region. International Journal of Infectious Diseases, 2021, 108, 494-502.	3.3	9
18	Nosocomial acute gastroenteritis outbreak caused by an equine-like G3P[8] DS-1-like rotavirus and GII.4 Sydney[P16] norovirus at a pediatric hospital in Rio de Janeiro, Brazil, 2019. Human Vaccines and Immunotherapeutics, 2021, 17, 4654-4660.	3.3	7

#	Article	IF	CITATIONS
19	A decade of G3P[8] and G9P[8] rotaviruses in Brazil: Epidemiology and evolutionary analyses. Infection, Genetics and Evolution, 2014, 28, 389-397.	2.3	6
20	Performance of a latex agglutination test in the diagnosis of acute gastroenteritis by rotavirus. Brazilian Journal of Microbiology, 2006, 37, 587-589.	2.0	6