

# Ivano de Filippis

## List of Publications by Year in descending order

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45  
papers

1,559  
citations

567281

15  
h-index

302126

39  
g-index

45  
all docs

45  
docs citations

45  
times ranked

3126  
citing authors

#	ARTICLE	IF	CITATIONS
1	Flow-Cytometric Method for Viability Analysis of <i>Mycoplasma gallisepticum</i> and Other Cell-Culture-Contaminant Mollicutes. <i>Current Microbiology</i> , 2021, 78, 67-77.	2.2	2
2	Genetic diversity of <i>Listeria monocytogenes</i> serotype 1/2a strains collected in Brazil by Multi- <i>Virulence</i> Locus Sequence Typing. <i>Letters in Applied Microbiology</i> , 2021, 72, 316-324.	2.2	2
3	Molecular surveillance of Brazilian meningococcal isolates serogroup c in the pre and post-men-c-vaccination period: Emergence of ST-3780. <i>Infection, Genetics and Evolution</i> , 2020, 78, 104079.	2.3	2
4	Genetic Relatedness of NDM-Producing <i>Klebsiella pneumoniae</i> Co-Occurring VIM, KPC, and OXA-48 Enzymes from Surveillance Cultures from an Intensive Care Unit. <i>Microbial Drug Resistance</i> , 2020, 26, 1219-1226.	2.0	11
5	Molecular and phenotypical characterization of <i>Cronobacter</i> species isolated with high occurrence from oats and linseeds. <i>FEMS Microbiology Letters</i> , 2019, 366, .	1.8	26
6	Epidemiology and molecular characterization of <i>Neisseria lactamica</i> carried in 11-19 years old students in Salvador, Brazil. <i>International Journal of Medical Microbiology</i> , 2018, 308, 454-458.	3.6	2
7	Characterization of strains of <i>Neisseria meningitidis</i> causing meningococcal meningitis in Mozambique, 2014: Implications for vaccination against meningococcal meningitis. <i>PLoS ONE</i> , 2018, 13, e0197390.	2.5	10
8	The invasive MenC cc103 lineage with penicillin reduced susceptibility persisting in Brazil. <i>International Journal of Medical Microbiology</i> , 2017, 307, 287-290.	3.6	0
9	Isolation, molecular and phenotypic characterization, and antibiotic susceptibility of <i>Cronobacter</i> spp. from Brazilian retail foods. <i>Food Microbiology</i> , 2017, 63, 129-138.	4.2	70
10	Phenotypic characterization of <i>Cronobacter</i> spp. strains isolated from foods and clinical specimens in Brazil. <i>Food Research International</i> , 2017, 102, 61-67.	6.2	28
11	The invasive <i>Neisseria meningitidis</i> MenC CC103 from Brazil is characterized by an accessory gene repertoire. <i>Scientific Reports</i> , 2017, 7, 1617.	3.3	5
12	<i>Cronobacter</i> spp.: infecções, ocorrência e regulação em alimentos – uma revisão no Brasil. <i>Brazilian Journal of Food Technology</i> , 2017, 21, .	0.8	2
13	Detection of antimicrobial resistance genes in beta-lactamase- and carbapenemase-producing <i>Klebsiella pneumoniae</i> by patient surveillance cultures at an intensive care unit in Rio de Janeiro, Brazil. <i>Jornal Brasileiro De Patologia E Medicina Laboratorial</i> , 2016, 52, 284-292.	0.3	9
14	Comparison of PCR-based methods for the simultaneous detection of <i>Neisseria meningitidis</i> , <i>Haemophilus influenzae</i> , and <i>Streptococcus pneumoniae</i> in clinical samples. <i>Brazilian Journal of Infectious Diseases</i> , 2016, 20, 335-341.	0.6	17
15	HIV-1 Genetic Diversity and Transmitted Drug Resistance in Antiretroviral Treatment-Naive Individuals from Amapá State, Northern Brazil. <i>AIDS Research and Human Retroviruses</i> , 2016, 32, 373-376.	1.1	18
16	Detection and sequencing of Zika virus from amniotic fluid of fetuses with microcephaly in Brazil: a case study. <i>Lancet Infectious Diseases</i> , The, 2016, 16, 653-660.	9.1	981
17	Genotypic characteristics of multidrug-resistant <i>Pseudomonas aeruginosa</i> from hospital wastewater treatment plant in Rio de Janeiro, Brazil. <i>Journal of Applied Microbiology</i> , 2015, 118, 1276-1286.	3.1	34
18	Investigação de um surto causado por <i>Cronobacter malonicus</i> em um hospital maternidade em Teresina, Piauí: caracterização e tipificação por eletroforese em gel de campo pulsado. <i>Vigilância Sanitária Em Debate: Sociedade, Ciência &amp; Tecnologia</i> , 2015, .	0.1	3

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19	Methicillin- and vancomycin-resistant <i>Staphylococcus aureus</i> in health care workers and medical devices. <i>Jornal Brasileiro De Patologia E Medicina Laboratorial</i> , 2015, 51, 143-152.	0.3	16
20	Replacement of <i>Neisseria meningitidis</i> C cc11/ET-15 variant by a cc103 hypervirulent clone, Brazil 2005–2011. <i>Diagnostic Microbiology and Infectious Disease</i> , 2013, 76, 524-525.	1.8	4
21	<i>Haemophilus influenzae</i> serotype b and a capsule-deficient type mutant (b <sup>Δ</sup> ) invasive disease in a partially vaccinated child in Brazil. <i>Journal of Medical Microbiology</i> , 2013, 62, 655-657.	1.8	2
22	Extended genetic analysis of Brazilian isolates of <i>Bacillus cereus</i> and <i>Bacillus thuringiensis</i> . <i>Memorias Do Instituto Oswaldo Cruz</i> , 2013, 108, 65-72.	1.6	13
23	The 2010 Meningococcal outbreak in Bahia, Brazil, was caused by 2 different STs belonging to Clonal Complex ST-103. <i>Vigilância Sanitária Em Debate: Sociedade, Ciência &amp; Tecnologia</i> , 2013, 1, .	0.1	1
24	Molecular Epidemiology of <i>Neisseria meningitidis</i> Serogroup B in Brazil. <i>PLoS ONE</i> , 2012, 7, e33016.	2.5	37
25	Molecular characterization and evaluation of antimicrobial susceptibility of enteropathogenic <i>E. coli</i> (EPEC) isolated from minas soft cheese. <i>Food Science and Technology</i> , 2012, 32, 747-753.	1.7	1
26	Fatal meningococcal meningitis in a HIV-infected patient caused by serogroup C <i>Neisseria meningitidis</i> belonging to the non-hypervirulent clonal complex ST-60 (cc60). <i>Brazilian Journal of Infectious Diseases</i> , 2011, 15, 178-180.	0.6	0
27	Fatal meningococcal meningitis in a HIV-infected patient caused by serogroup C <i>Neisseria meningitidis</i> belonging to the non-hypervirulent clonal complex ST-60 (cc60). <i>Brazilian Journal of Infectious Diseases</i> , 2011, 15, 178-180.	0.6	0
28	Isolation of <i>Brevibacillus brevis</i> from tracheal aspirates of a hospitalized patient. <i>Apmis</i> , 2011, 119, 901-902.	2.0	3
29	Presence of qacE <sup>H</sup> 1 Gene and Susceptibility to a Hospital Biocide in Clinical Isolates of <i>Pseudomonas aeruginosa</i> Resistant to Antibiotics. <i>Current Microbiology</i> , 2011, 63, 16-21.	2.2	43
30	PorA VR3 Typing Database: A web-based resource for the determination of PorA VR3 alleles of <i>Neisseria meningitidis</i> . <i>Infection, Genetics and Evolution</i> , 2011, 11, 248-249.	2.3	9
31	Molecular Characterization of Quinolone-Resistant <i>Neisseria gonorrhoeae</i> Isolates from Brazil. <i>Journal of Clinical Microbiology</i> , 2011, 49, 4208-4212.	3.9	25
32	Urinary tract infection caused by nontypable <i>Haemophilus influenzae</i> in the elderly. <i>Journal of Medical Microbiology</i> , 2010, 59, 1132-1133.	1.8	4
33	Changes in <i>Haemophilus influenzae</i> capsule locus: possible emergence of novel variants in Brazil. <i>Diagnostic Microbiology and Infectious Disease</i> , 2010, 68, 97-102.	1.8	1
34	Quest for a broad-range vaccine against <i>Neisseria meningitidis</i> serogroup B: implications of genetic variations of the surface-exposed proteins. <i>Journal of Medical Microbiology</i> , 2009, 58, 1127-1132.	1.8	5
35	Septic arthritis due to <i>Haemophilus influenzae</i> serotype a in the post-vaccination era in Brazil. <i>Journal of Medical Microbiology</i> , 2008, 57, 1311-1312.	1.8	10
36	PorA Variable Antigenic Regions VR1, VR2, and VR3 of <i>Neisseria meningitidis</i> Serogroups B and C Isolated in Brazil from 1999 to 2004. <i>Infection and Immunity</i> , 2007, 75, 3683-3685.	2.2	10

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37	Neisseria meningitidis PorA variable regions: rapid detection of P1 <sup>7</sup> and P1 <sup>19</sup> variants by PCR. Letters in Applied Microbiology, 2007, 45, 426-431.	2.2	1
38	Antimicrobial susceptibility of Haemophilus influenzae isolates collected from 4 centers in Brazil (1990-2003). Diagnostic Microbiology and Infectious Disease, 2006, 54, 57-62.	1.8	3
39	Rapid detection of Neisseria meningitidis in cerebrospinal fluid by one-step polymerase chain reaction of the nspA gene. Diagnostic Microbiology and Infectious Disease, 2005, 51, 85-90.	1.8	29
40	Multilocus sequence typing and repetitive element-based polymerase chain reaction analysis of Neisseria meningitidis isolates in Brazil reveal the emergence of 11 new sequence types genetically related to the ST-32 and ST-41/44 complexes and high prevalence of strains related to hypervirulent lineages. Diagnostic Microbiology and Infectious Disease, 2005, 53, 161-167.	1.8	17
41	Occurrence of Haemophilus influenzae strains in three Brazilian states since the introduction of a conjugate Haemophilus influenzae type b vaccine. Brazilian Journal of Medical and Biological Research, 2005, 38, 777-781.	1.5	20
42	Genetic diversity of Neisseria meningitidis strains isolated in Rio de Janeiro, Brazil, evaluated by multilocus enzyme electrophoresis. Letters in Applied Microbiology, 2004, 39, 232-239.	2.2	1
43	Evaluation of the potential for use in biocatalysis of a lipase from a wild strain of Bacillus megaterium. Journal of Molecular Catalysis B: Enzymatic, 2004, 31, 53-61.	1.8	45
44	PCR Analyses of tRNA Intergenic Spacer, 16S-23S Internal Transcribed Spacer, and Randomly Amplified Polymorphic DNA Reveal Inter- and Intraspecific Relationships of Enterobacter cloacae Strains. Journal of Clinical Microbiology, 2001, 39, 3865-3870.	3.9	34
45	Development of a collection of bacteria causing meningitis in Rio de Janeiro from 1990 to 1991. Memórias Do Instituto Oswaldo Cruz, 1995, 90, 21-24.	1.6	3