Maria Giufre

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Escherichia coli of human and avian origin: detection of clonal groups associated with fluoroquinolone and multidrug resistance in Italy. Journal of Antimicrobial Chemotherapy, 2012, 67, 860-867.	1.3	94
2	Emergence of NDM-5-producing Escherichia coli sequence type 167 clone in Italy. International Journal of Antimicrobial Agents, 2018, 52, 76-81.	1.1	56
3	Incl1 plasmids associated with the spread of CMY-2, CTX-M-1 and SHV-12 in Escherichia coli of animal and human origin. Clinical Microbiology and Infection, 2013, 19, E238-E240.	2.8	55
4	Bacterial coinfections in COVID-19: an underestimated adversary. Annali Dell'Istituto Superiore Di Sanita, 2020, 56, 359-364.	0.2	55
5	Ten years of Hib vaccination in Italy: Prevalence of non-encapsulated Haemophilus influenzae among invasive isolates and the possible impact on antibiotic resistance. Vaccine, 2011, 29, 3857-3862.	1.7	53
6	Ciprofloxacin-resistant, CTX-M-15-producing Escherichia coli ST131 clone in extraintestinal infections in Italy. Clinical Microbiology and Infection, 2010, 16, 1555-1558.	2.8	49
7	Polymorphism in ftsl gene and Â-lactam susceptibility in Portuguese Haemophilus influenzae strains: clonal dissemination of Â-lactamase-positive isolates with decreased susceptibility to amoxicillin/clavulanic acid. Journal of Antimicrobial Chemotherapy, 2011, 66, 788-796.	1.3	48
8	Haemophilus influenzae in children with cystic fibrosis: Antimicrobial susceptibility, molecular epidemiology, distribution of adhesins and biofilm formation. International Journal of Medical Microbiology, 2012, 302, 45-52.	1.5	47
9	Emergence of Escherichia coli ST131 sub-clone H30 producing VIM-1 and KPC-3 carbapenemases, Italy. Journal of Antimicrobial Chemotherapy, 2014, 69, 2293-2296.	1.3	45
10	Colonization by multidrug-resistant organisms in long-term care facilities in Italy: a point-prevalence study. Clinical Microbiology and Infection, 2017, 23, 961-967.	2.8	45
11	Carriage of Haemophilus influenzae in the oropharynx of young children and molecular epidemiology of the isolates after fifteen years of H. influenzae type b vaccination in Italy. Vaccine, 2015, 33, 6227-6234.	1.7	43
12	First Report of Plasmid-Mediated Quinolone Resistance Determinant <i>qnrS1</i> in an <i>Escherichia coli</i> Strain of Animal Origin in Italy. Antimicrobial Agents and Chemotherapy, 2009, 53, 3112-3114.	1.4	42
13	Increased frequency of the immunoglobulin enhancer HS1,2 allele 2 in coeliac disease. Scandinavian Journal of Gastroenterology, 2004, 39, 1083-1087.	0.6	38
14	Why we need a vaccine for non-typeable <i>Haemophilus influenzae</i> . Human Vaccines and Immunotherapeutics, 2016, 12, 2357-2361.	1.4	38
15	Evolution of human IgH3′EC duplicated structures: both enhancers HS1,2 are polymorphic with variation of transcription factor's consensus sites. Gene, 2005, 346, 105-114.	1.0	36
16	Increasing trend in invasive non-typeable Haemophilus influenzae disease and molecular characterization of the isolates, Italy, 2012–2016. Vaccine, 2018, 36, 6615-6622.	1.7	35
17	Carriage of Haemophilus influenzae is associated with pneumococcal vaccination in Italian children. Vaccine, 2015, 33, 4559-4564.	1.7	34
18	Presence of Multiple Copies of the Capsulation b Locus in InvasiveHaemophilus influenzaeType b (Hib) Strains Isolated from Children with Hib Conjugate Vaccine Failure. Journal of Infectious Diseases, 2005, 192, 819-823.	1.9	32

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19	First Characterization of Heterogeneous Resistance to Imipenem in Invasive Nontypeable Haemophilus influenzae Isolates. Antimicrobial Agents and Chemotherapy, 2007, 51, 3155-3161.	1.4	32
20	Variation in expression of HMW1 and HMW2 adhesins in invasive nontypeable Haemophilus influenzae isolates. BMC Microbiology, 2008, 8, 83.	1.3	25
21	Identification of Haemophilus influenzae Clones Associated with Invasive Disease a Decade after Introduction of H. influenzae Serotype b Vaccination in Italy. Vaccine Journal, 2013, 20, 1223-1229.	3.2	25
22	Extended-spectrum β-lactamase-producing Escherichia coli from extraintestinal infections in humans and from food-producing animals in Italy: a â€~One Health' study. International Journal of Antimicrobial Agents, 2021, 58, 106433.	1.1	24
23	Antimicrobial susceptibility of Haemophilus influenzae strains isolated from invasive disease in Italy. Journal of Antimicrobial Chemotherapy, 2004, 54, 1139-1143.	1.3	22
24	Nontypeable Haemophilus influenzae Meningitis in Children: Phenotypic and Genotypic Characterization of Isolates. Pediatric Infectious Disease Journal, 2007, 26, 577-582.	1.1	22
25	Genital carriage of the genus Haemophilus in pregnancy: species distribution and antibiotic susceptibility. Journal of Medical Microbiology, 2015, 64, 724-730.	0.7	22
26	Haemophilus parainfluenzae meningitis in an adult associated with acute otitis media. New Microbiologica, 2009, 32, 213-5.	0.1	21
27	Emergence of the colistin resistance <i>mcr-1</i> determinant in commensal <i>Escherichia coli</i> from residents of long-term-care facilities in Italy: TableÂ1 Journal of Antimicrobial Chemotherapy, 2016, 71, 2329-2331.	1.3	20
28	Conservation and Diversity of HMW1 and HMW2 Adhesin Binding Domains among Invasive Nontypeable Haemophilus influenzae Isolates. Infection and Immunity, 2006, 74, 1161-1170.	1.0	19
29	Immunoglobulin Enhancer HS1,2 polymorphism: a new powerful anthropogenetic marker. Annals of Human Genetics, 2006, 70, 946-950.	0.3	18
30	Phylogenetic Background and Virulence Genotype of Ciprofloxacinâ€ 5 usceptible and Ciprofloxacinâ€ResistantEscherichia coliStrains of Human and Avian Origin. Journal of Infectious Diseases, 2009, 199, 1209-1217.	1.9	17
31	Contribution of ROB-1 and PBP3 mutations to the resistance phenotype of a β-lactamase-positive amoxicillin/clavulanic acid-resistant Haemophilus influenzae carrying plasmid pB1000 in Italy. Journal of Antimicrobial Chemotherapy, 2011, 66, 96-99.	1.3	17
32	Emergence of Invasive Haemophilus influenzae Type A Disease in Italy. Clinical Infectious Diseases, 2017, 64, 1626-1628.	2.9	17
33	Multidrug-resistant infections in long-term care facilities: extended-spectrum β-lactamase–producing Enterobacteriaceae and hypervirulent antibiotic resistant Clostridium difficile. Diagnostic Microbiology and Infectious Disease, 2018, 91, 275-281.	0.8	17
34	Risk factors for Haemophilus influenzae and pneumococcal respiratory tract colonization in CVID. Journal of Allergy and Clinical Immunology, 2018, 142, 1999-2002.e3.	1.5	17
35	Genetic Diversity of Invasive Strains ofHaemophilus influenzaeType b before and after Introduction of the Conjugate Vaccine in Italy. Clinical Infectious Diseases, 2006, 43, 317-319.	2.9	12
36	Genetic Characterization of the Capsulation Locus of <i>Haemophilus influenzae</i> Serotype e. Journal of Clinical Microbiology, 2010, 48, 1404-1407.	1.8	10

MARIA GIUFRE

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37	Haemophilus influenzaeSerotype e Meningitis in an Infant. Clinical Infectious Diseases, 2004, 38, 1041-1041.	2.9	9
38	Detection of Six Copies of the Capsulation b Locus in a Haemophilus influenzae Type b Strain Isolated from a Splenectomized Patient with Fulminant Septic Shock. Journal of Clinical Microbiology, 2006, 44, 640-642.	1.8	9
39	First report of neonatal bacteremia caused by "Haemophilus quentini―diagnosed by 16S rRNA gene sequencing, Italy. Diagnostic Microbiology and Infectious Disease, 2015, 83, 121-123.	0.8	7
40	Neonatal Invasive <i>Haemophilus influenzae</i> Disease and Genotypic Characterization of the Associated Strains in Italy: Figure 1 Clinical Infectious Diseases, 2015, 61, 1203-1204.	2.9	6
41	Invasive Haemophilus influenzae type b (Hib) disease in children in Italy, after 20Âyears of routine use of conjugate Hib vaccines. Vaccine, 2020, 38, 6533-6538.	1.7	5
42	Dynamics of carbapenemase-producing Enterobacterales intestinal colonisation in the elderly population after hospital discharge, Italy, 2018–2020. International Journal of Antimicrobial Agents, 2022, 59, 106594.	1.1	5
43	Predominance of the fimH30 Subclone Among Multidrug-Resistant Escherichia coli Strains Belonging to Sequence Type 131 in Italy. Journal of Infectious Diseases, 2014, 209, 629-630.	1.9	4
44	Nasopharyngeal carriage of Haemophilus influenzae among adults with co-morbidities. Vaccine, 2022, 40, 826-832.	1.7	4
45	Food Reservoir for <i>Escherichia coli</i> Causing Urinary Tract Infections. Emerging Infectious Diseases, 2010, 16, 1048-1049.	2.0	3
46	Whole-Genome Sequences of Nonencapsulated Haemophilus influenzae Strains Isolated in Italy. Genome Announcements, 2015, 3, .	0.8	3
47	Whole-Genome Sequences of Multidrug-Resistant Escherichia coli Strains Sharing the Same Sequence Type (ST410) and Isolated from Human and Avian Sources in Italy. Genome Announcements, 2015, 3, .	0.8	3
48	Phylogenetic Structure and Comparative Genomics of Multi-National Invasive Haemophilus influenzae Serotype a Isolates. Frontiers in Microbiology, 2022, 13, 856884.	1.5	3
49	First Whole-Genome Sequence of a Haemophilus influenzae Type e Strain Isolated from a Patient with Invasive Disease in Italy. Genome Announcements, 2017, 5, .	0.8	2
50	Variant IS1016 Insertion Elements in Invasive Haemophilus influenzae Type b Isolates Harboring Multiple Copies of the Capsulation b Locus. Clinical Infectious Diseases, 2006, 43, 1225-1226.	2.9	1