Deirdre A Collins

List of Publications by Year
in descending order

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36	930	430442	454577
papers	citations	18 h-index	30 g-index

37	37	37	1068
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Epidemiology of Clostridium difficile infection in Asia. Antimicrobial Resistance and Infection Control, 2013, 2, 21.	1.5	186
2	Genome Analysis of Clostridium difficile PCR Ribotype 014 Lineage in Australian Pigs and Humans Reveals a Diverse Genetic Repertoire and Signatures of Long-Range Interspecies Transmission. Frontiers in Microbiology, 2016, 7, 2138.	1.5	117
3	Clostridium difficile ribotype 017 â€“ characterization, evolution and epidemiology of the dominant strain in Asia. Emerging Microbes and Infections, 2019, 8, 796-807.	3.0	61
4	Epidemiology of Clostridium difficile infection in two tertiary-care hospitals in Perth, Western Australia: a cross-sectional study. New Microbes and New Infections, 2014, 2, 64-71.	0.8	54
5	Clostridioides difficile infection in the Asia-Pacific region. Emerging Microbes and Infections, 2020, 9, 42-52.	3.0	47
6	Exposure to environmental microbiota explains persistent abdominal pain and irritable bowel syndrome after a major flood. Gut Pathogens, 2017, 9, 75.	1.6	33
7	Prevalence and molecular epidemiology of Clostridium difficile infection in Indonesia. New Microbes and New Infections, 2017, 18, 34-37.	0.8	31
8	Incorrect diagnosis of Clostridium difficile infection in a university hospital in Japan. Journal of Infection and Chemotherapy, 2015, 21, 718-722.	0.8	30
9	Antimicrobial resistance in Clostridium difficile ribotype 017. Expert Review of Anti-Infective Therapy, 2020, 18, 17-25.	2.0	28
10	Antimicrobial Susceptibilities of Clostridium difficile Isolates from 12 Asia-Pacific Countries in 2014 and 2015. Antimicrobial Agents and Chemotherapy, 2020, 64, .	1.4	26
11	Clonal Origins of Vibrio cholerae O1 El Tor Strains, Papua New Guinea, 2009â€“2011. Emerging Infectious Diseases, 2011, 17, 2063-5.	2.0	24
12	Laboratory-based surveillance of Clostridium difficile strains circulating in the Australian healthcare setting in 2012. Pathology, 2017, 49, 309-313.	0.3	24
13	High Prevalence of Toxigenic and Nontoxigenic Clostridium difficile Strains in Malaysia. Journal of Clinical Microbiology, 2018, 56, .	1.8	24
14	The emergence of community-onset Clostridium difficile infection in a tertiary hospital in Singapore: A cause for concern. International Journal of Antimicrobial Agents, 2014, 43, 47-51.	1.1	22
15	Community-associated Clostridium difficile infection in emergency department patients in Western Australia. Anaerobe, 2017, 48, 121-125.	1.0	22
16	Molecular methods for detecting and typing of Clostridium difficile. Pathology, 2015, 47, 211-218.	0.3	21
17	Predictors of pneumococcal carriage and the effect of the 13-valent pneumococcal conjugate vaccination in the Western Australian Aboriginal population. Pneumonia (Nathan Qld), 2017, 9, 14.	2.5	21
18	Laboratory-based surveillance of Clostridium difficile circulating in Australia, September â€“ November 2010. Pathology, 2016, 48, 257-260.	0.3	20

#	Article	IF	Citations
19	Different molecular characteristics and antimicrobial resistance profiles of Clostridium difficile in the Asia-Pacific region. Emerging Microbes and Infections, 2019, 8, 1553-1562.	3.0	17
20	High Nasopharyngeal Carriage of Non-Vaccine Serotypes in Western Australian Aboriginal People Following 10 Years of Pneumococcal Conjugate Vaccination. PLoS ONE, 2013, 8, e82280.	1.1	16
21	Whole-genome sequencing links Clostridium (Clostridioides) difficile in a single hospital to diverse environmental sources in the community. Journal of Applied Microbiology, 2022, 133, 1156-1168.	1.4	13
22	Routine detection of Clostridium difficile in Western Australia. Anaerobe, 2016, 37, 34-37.	1.0	12
23	Clostridium difficile in Asia: Opportunities for One Health Management. Tropical Medicine and Infectious Disease, 2019, 4, 7.	0.9	12
24	Prevalence ofClostridium difficileinfection and colonization in a tertiary hospital and elderly community of North-Eastern Peninsular Malaysia. Epidemiology and Infection, 2017, 145, 3012-3019.	1.0	11
25	Recurrence ofClostridium difficileinfection in the Western Australian population. Epidemiology and Infection, 2019, 147, e153.	1.0	10
26	Clostridium difficile Guidelines. Clinical Infectious Diseases, 2018, 67, 1639.	2.9	8
27	Genetically related Clostridium difficile from water sources and human <scp>CDI</scp> cases revealed by wholeâ€genome sequencing. Environmental Microbiology, 2022, 24, 1221-1230.	1.8	7
28	Ridinilazole: a novel, narrow-spectrum antimicrobial agent targeting Clostridium (Clostridioides) difficile. Letters in Applied Microbiology, 2022, 75, 526-536.	1.0	6
29	Evaluation of the antimicrobial activity of ridinilazole and six comparators against Chinese, Japanese and South Korean strains of Clostridioides difficile. Journal of Antimicrobial Chemotherapy, 2021, 76, 967-972.	1.3	4
30	Clostridioides difficile colonization and infection in a cohort of Australian adults with cystic fibrosis. Journal of Hospital Infection, 2021, 113, 44-51.	1.4	4
31	Wave 2 strains of atypical Vibrio cholerae El Tor caused the 2009â€“2011 cholera outbreak in Papua New Guinea. Microbial Genomics, 2019, 5, .	1.0	4
32	Clostridioides (Clostridium) difficile in children with diarrhoea in Vietnam. Anaerobe, 2022, , 102550.	1.0	4
33	Global evolutionary dynamics and resistome analysis of Clostridioides difficile ribotype 017. Microbial Genomics, 2022, 8, .	1.0	4
34	Evaluation of the Cepheid Â® Xpert Â® C.Âdifficile binary toxin (BT) diagnostic assay. Anaerobe, 2018, 51, 12-16.	1.0	3
35	Linkage study of surveillance and hospital admission data to investigate Clostridium difficile infection in hospital patients in Perth, Western Australia. Anaerobe, 2022, 74, 102528.	1.0	3
36	Microbiological evaluation of the ability of the DEKO-190 Washer/Disinfector to remove Clostridium difficile spores from bedpan surfaces. Infection, Disease and Health, 2019, 24, 208-211.	0.5	1