## Catherine E Ludden

List of Publications by Year in descending order

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CATHEDINE F LUDDEN

#	Article	IF	CITATIONS
1	One Health Genomic Surveillance of Escherichia coli Demonstrates Distinct Lineages and Mobile Genetic Elements in Isolates from Humans versus Livestock. MBio, 2019, 10, .	4.1	130
2	Genomic Surveillance of Enterococcus faecium Reveals Limited Sharing of Strains and Resistance Genes between Livestock and Humans in the United Kingdom. MBio, 2018, 9, .	4.1	63
3	Colonisation with ESBL-producing and carbapenemase-producing Enterobacteriaceae, vancomycin-resistant enterococci, and meticillin-resistant Staphylococcus aureus in a long-term care facility over one year. BMC Infectious Diseases, 2015, 15, 168.	2.9	54
4	Quantifying acquisition and transmission of Enterococcus faecium using genomic surveillance. Nature Microbiology, 2021, 6, 103-111.	13.3	53
5	Production of KPC-2 Carbapenemase by an Escherichia coli Clinical Isolate Belonging to the International ST131 Clone. Antimicrobial Agents and Chemotherapy, 2011, 55, 4935-4936.	3.2	51
6	A One Health Study of the Genetic Relatedness of Klebsiella pneumoniae and Their Mobile Elements in the East of England. Clinical Infectious Diseases, 2020, 70, 219-226.	5.8	46
7	Detection of vancomycin-resistant <i>Enterococcus faecium</i> hospital-adapted lineages in municipal wastewater treatment plants indicates widespread distribution and release into the environment. Genome Research, 2019, 29, 626-634.	5.5	40
8	Defining nosocomial transmission of Escherichia coli and antimicrobial resistance genes: a genomic surveillance study. Lancet Microbe, The, 2021, 2, e472-e480.	7.3	39
9	Sharing of carbapenemase-encoding plasmids between Enterobacteriaceae in UK sewage uncovered by MinION sequencing. Microbial Genomics, 2017, 3, e000114.	2.0	33
10	Genomic surveillance of Escherichia coli ST131 identifies local expansion and serial replacement of subclones. Microbial Genomics, 2020, 6, .	2.0	33
11	Detection of OXA-48 Carbapenemase in the Pandemic Clone Escherichia coli O25b:H4-ST131 in the Course of Investigation of an Outbreak of OXA-48-Producing Klebsiella pneumoniae. Antimicrobial Agents and Chemotherapy, 2012, 56, 4030-4031.	3.2	32
12	Transmission of methicillin-resistant Staphylococcus aureus in long-term care facilities and their related healthcare networks. Genome Medicine, 2016, 8, 102.	8.2	30
13	Genomic surveillance of Escherichia coli in municipal wastewater treatment plants as an indicator of clinically relevant pathogens and their resistance genes. Microbial Genomics, 2019, 5, .	2.0	29
14	Within-host evolution of Enterococcus faecium during longitudinal carriage and transition to bloodstream infection in immunocompromised patients. Genome Medicine, 2017, 9, 119.	8.2	26
15	Dissemination of clonally related multidrug-resistant <i>Klebsiella pneumoniae</i> in Ireland. Epidemiology and Infection, 2016, 144, 443-448.	2.1	16
16	Rapid environmental contamination of a new nursing home with antimicrobial-resistant organisms preceding occupation by residents. Journal of Hospital Infection, 2013, 83, 327-329.	2.9	9
17	A novel therapeutic antibody screening method using bacterial high-content imaging reveals functional antibody binding phenotypes of Escherichia coli ST131. Scientific Reports, 2020, 10, 12414.	3.3	9
18	Characterization of methicillin-resistant <i>Staphylococcus aureus</i> from residents and the environment in a long-term care facility. Epidemiology and Infection, 2015, 143, 2985-2988.	2.1	7

#	Article	IF	CITATIONS
19	Characterising the persistence of RT-PCR positivity and incidence in a community survey of SARS-CoV-2. Wellcome Open Research, 0, 7, 102.	1.8	7