## Jane W Marsh

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genomic Diversity of Hospital-Acquired Infections Revealed through Prospective Whole-Genome Sequencing-Based Surveillance. MSystems, 2022, 7, .	3.8	10
2	SARS-CoV-2 N gene mutations impact detection by clinical molecular diagnostics: reports in two cities in the United States. Diagnostic Microbiology and Infectious Disease, 2021, 101, 115468.	1.8	17
3	Outbreak of Vancomycin-resistant Enterococcus faecium in Interventional Radiology: Detection Through Whole-genome Sequencing-based Surveillance. Clinical Infectious Diseases, 2020, 70, 2336-2343.	5.8	43
4	Transmission Dynamics and Microevolution of Neisseria meningitidis During Carriage and Invasive Disease in High School Students in Georgia and Maryland, 2006–2007. Journal of Infectious Diseases, 2020, 223, 2038-2047.	4.0	6
5	Automated data mining of the electronic health record for investigation of healthcare-associated outbreaks. Infection Control and Hospital Epidemiology, 2019, 40, 314-319.	1.8	40
6	<i>Clostridioides difficile</i> : a potential source of NpmA in the clinical environment. Journal of Antimicrobial Chemotherapy, 2019, 74, 521-523.	3.0	13
7	Multi-locus variable number tandem repeat analysis for investigation of the genetic association of Clostridium difficile isolates from food, food animals and humans. Anaerobe, 2011, 17, 156-160.	2.1	25
8	Multilocus Variable-Number Tandem-Repeat Analysis and Multilocus Sequence Typing Reveal Genetic Relationships among Clostridium difficile Isolates Genotyped by Restriction Endonuclease Analysis. Journal of Clinical Microbiology, 2010, 48, 412-418.	3.9	43
9	Deletion of fetA Gene Sequences in Serogroup B and C Neisseria meningitidis Isolates. Journal of Clinical Microbiology, 2007, 45, 1333-1335.	3.9	27
10	Metastatic Lung Disease to the Central Nervous System: in vitro Response to Chemotherapeutic Agents. Journal of Neuro-Oncology, 2004, 66, 81-90.	2.9	2