

E Jane Parmley

List of Publications by Year in descending order

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32
papers

672
citations

623734

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24
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all docs

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docs citations

32
times ranked

1019
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrating Whole-Genome Sequencing Data Into Quantitative Risk Assessment of Foodborne Antimicrobial Resistance: A Review of Opportunities and Challenges. <i>Frontiers in Microbiology</i> , 2019, 10, 1107.	3.5	73
2	Wild Bird Influenza Survey, Canada, 2005. <i>Emerging Infectious Diseases</i> , 2008, 14, 84-87.	4.3	65
3	A Whole-Genome Sequencing Approach To Study Cefoxitin-Resistant <i>Salmonella enterica</i> Serovar Heidelberg Isolates from Various Sources. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	59
4	Ciprofloxacin-Resistant <i>Campylobacter</i> spp. in Retail Chicken, Western Canada. <i>Emerging Infectious Diseases</i> , 2013, 19, 1121-1124.	4.3	47
5	PREVALENCE AND CHARACTERISTICS OF <i>ESCHERICHIA COLI</i> AND <i>SALMONELLA</i> SPP. IN THE FECES OF WILD URBAN NORWAY AND BLACK RATS (<i>RATTUS NORVEGICUS</i> AND <i>RATTUS RATTUS</i>) FROM AN INNER-CITY NEIGHBORHOOD OF VANCOUVER, CANADA. <i>Journal of Wildlife Diseases</i> , 2015, 51, 589-600.	0.8	45
6	Factors potentially linked with the occurrence of antimicrobial resistance in selected bacteria from cattle, chickens and pigs: A scoping review of publications for use in modelling of antimicrobial resistance (IAM.AMR Project). <i>Zoonoses and Public Health</i> , 2018, 65, 957-971.	2.2	37
7	Evidence for action: a One Health learning platform on interventions to tackle antimicrobial resistance. <i>Lancet Infectious Diseases</i> , The, 2020, 20, e307-e311.	9.1	37
8	Evaluating the Integration of One Health in Surveillance Systems for Antimicrobial Use and Resistance: A Conceptual Framework. <i>Frontiers in Veterinary Science</i> , 2021, 8, 611931.	2.2	31
9	Genomic Epidemiology of Major Extraintestinal Pathogenic <i>Escherichia coli</i> Lineages Causing Urinary Tract Infections in Young Women Across Canada. <i>Open Forum Infectious Diseases</i> , 2019, 6, ofz431.	0.9	30
10	AMR-Intervene: a social-ecological framework to capture the diversity of actions to tackle antimicrobial resistance from a One Health perspective. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 1-21.	3.0	29
11	The ecology of avian influenza viruses in wild dabbling ducks (<i>Anas</i> spp.) in Canada. <i>PLoS ONE</i> , 2017, 12, e0176297.	2.5	23
12	Antimicrobial Resistance of <i>Campylobacter</i> in Broiler Chicken Along the Food Chain in Canada. <i>Foodborne Pathogens and Disease</i> , 2020, 17, 512-520.	1.8	22
13	Genomic Investigation of the Emergence of Invasive Multidrug-Resistant <i>Salmonella enterica</i> Serovar Dublin in Humans and Animals in Canada. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	21
14	Changes in antimicrobial resistance levels among and in Ontario broiler chickens between 2003 and 2015. <i>Canadian Journal of Veterinary Research</i> , 2018, 82, 163-177.	0.2	16
15	<i>Salmonella</i> , <i>Campylobacter</i> , <i>Clostridium difficile</i> , and antimicrobial resistant <i>Escherichia coli</i> in the faeces of sympatric meso-mammals in southern Ontario, Canada. <i>Zoonoses and Public Health</i> , 2019, 66, 406-416.	2.2	14
16	Characterizing social-ecological context and success factors of antimicrobial resistance interventions across the One Health spectrum: analysis of 42 interventions targeting <i>E. coli</i> . <i>BMC Infectious Diseases</i> , 2021, 21, 873.	2.9	13
17	Complete Genome and Plasmid Sequences of Three Canadian Isolates of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Heidelberg from Human and Food Sources. <i>Genome Announcements</i> , 2016, 4, .	0.8	11
18	Identifying non-traditional stakeholders with whom to engage, when mitigating antimicrobial resistance in foodborne pathogens (Canada). <i>BMC Research Notes</i> , 2018, 11, 170.	1.4	11

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19	Bait Trapping Linked to Higher Avian Influenza Virus Detection in Wild Ducks. <i>Journal of Wildlife Diseases</i> , 2012, 48, 444-448.	0.8	10
20	Complete Genome Sequences of 17 Canadian Isolates of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Heidelberg from Human, Animal, and Food Sources. <i>Genome Announcements</i> , 2016, 4, .	0.8	10
21	Comparison of annual and regional variation in multidrug resistance using various classification metrics for generic <i>Escherichia coli</i> isolated from chicken abattoir surveillance samples in Canada. <i>Preventive Veterinary Medicine</i> , 2018, 154, 9-17.	1.9	10
22	Antimicrobial Resistance Profiles of <i>Escherichia coli</i> and <i>Salmonella</i> Isolates in Canadian Broiler Chickens and Their Products. <i>Foodborne Pathogens and Disease</i> , 2020, 17, 672-678.	1.8	9
23	Antimicrobial resistance of <i>Salmonella</i> and generic <i>Escherichia coli</i> isolated from surface water samples used for recreation and a source of drinking water in southwestern Ontario, Canada. <i>Zoonoses and Public Health</i> , 2020, 67, 566-575.	2.2	9
24	Measures used to assess the burden of ESBL-producing <i>Escherichia coli</i> infections in humans: a scoping review. <i>JAC-Antimicrobial Resistance</i> , 2021, 3, dlaa104.	2.1	9
25	Avian Pathogenicity Genes and Antibiotic Resistance in <i>Escherichia coli</i> Isolates from Wild Norway Rats (<i>Rattus norvegicus</i>) in British Columbia, Canada. <i>Journal of Wildlife Diseases</i> , 2016, 52, 418-421.	0.8	8
26	A comparison of modelling options to assess annual variation in susceptibility of generic <i>Escherichia coli</i> isolates to ceftiofur, ampicillin and nalidixic acid from retail chicken meat in Canada. <i>Preventive Veterinary Medicine</i> , 2018, 160, 123-135.	1.9	7
27	Targeting discriminatory SNPs in <i>Salmonella enterica</i> serovar Heidelberg genomes using RNase H2-dependent PCR. <i>Journal of Microbiological Methods</i> , 2019, 157, 81-87.	1.6	5
28	The Influence of Sociodemographic Factors on the Engagement of Citizens in the Detection of Dead Corvids During the Emergence of West Nile Virus in Ontario, Canada. <i>Frontiers in Veterinary Science</i> , 2019, 6, 483.	2.2	5
29	Using whole-genome sequence data to examine the epidemiology of <i>Salmonella</i> , <i>Escherichia coli</i> and associated antimicrobial resistance in raccoons (<i>Procyon lotor</i>), swine manure pits, and soil samples on swine farms in southern Ontario, Canada. <i>PLoS ONE</i> , 2021, 16, e0260234.	2.5	3
30	Canadian wildlife health surveillance patterns, challenges and opportunities identified by a scoping review. <i>Facets</i> , 2022, 7, 25-44.	2.4	2
31	Rural Raccoons (<i>Procyon lotor</i>) Not Likely to Be a Major Driver of Antimicrobial Resistant Human <i>Salmonella</i> Cases in Southern Ontario, Canada: A One Health Epidemiological Assessment Using Whole-Genome Sequence Data. <i>Frontiers in Veterinary Science</i> , 2022, 9, 840416.	2.2	1
32	Using whole-genome sequence data to examine the epidemiology of antimicrobial resistance in <i>Escherichia coli</i> from wild meso-mammals and environmental sources on swine farms, conservation areas, and the Grand River watershed in southern Ontario, Canada. <i>PLoS ONE</i> , 2022, 17, e0266829.	2.5	0