Julianne V Kus

List of Publications by Year in descending order

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623734 477307 1,007 47 14 29 citations g-index h-index papers 49 49 49 1301 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Post-traumatic Fungal Keratitis and Endophthalmitis Caused by <i>Coniochaeta Hoffmannii</i> with Late Recurrence following Therapeutic Full-thickness Penetrating Keratoplasty. Ocular Immunology and Inflammation, 2023, 31, 826-829.	1.8	2
2	Impact of coronavirus disease 2019 (COVID-19) pre-test probability on positive predictive value of high cycle threshold severe acute respiratory coronavirus virus 2 (SARS-CoV-2) real-time reverse transcription polymerase chain reaction (RT-PCR) test results. Infection Control and Hospital Epidemiology, 2022, 43, 1179-1183.	1.8	4
3	Linear Regression Equations To Predict \hat{l}^2 -Lactam, Macrolide, Lincosamide, and Fluoroquinolone MICs from Molecular Antimicrobial Resistance Determinants in <i>Streptococcus pneumoniae</i> Antimicrobial Agents and Chemotherapy, 2022, 66, AAC0137021.	3.2	9
4	Mycobacterium genavense Central Nervous System Infection in a Patient with AIDS. Canadian Journal of Neurological Sciences, 2022, , 1-8.	0.5	2
5	Real-Time RT-PCR Allelic Discrimination Assay for Detection of N501Y Mutation in the Spike Protein of SARS-CoV-2 Associated with B.1.1.7 Variant of Concern. Microbiology Spectrum, 2022, 10, e0068121.	3.0	5
6	The 2021 WHO catalogue of Mycobacterium tuberculosis complex mutations associated with drug resistance: a genotypic analysis. Lancet Microbe, The, 2022, 3, e265-e273.	7.3	114
7	Four genomic clades of <i>Candida auris</i> identified in Canada, 2012–2019. Medical Mycology, 2022, 60, .	0.7	7
8	First reported case of Robinsoniella peoriensis pyometra and bloodstream infection: A case report and review of the literature. Anaerobe, 2022, 75, 102581.	2.1	5
9	Development of a Duplex Real-Time PCR Assay for the Differentiation of Blastomyces dermatitidis and <i>Blastomyces gilchristii</i> and a Retrospective Analysis of Culture and Primary Specimens from Blastomycosis Cases from New York (2005 to 2019). Journal of Clinical Microbiology, 2021, 59, .	3.9	5
10	Fatal respiratory diphtheria in a visitor to Canada. Cmaj, 2021, 193, E19-E22.	2.0	2
11	Evaluating the Role of STAT3 in CD4 $<$ sup $>+<$ /sup $>$ T Cells in Susceptibility to Invasive Aspergillosis. Infection and Immunity, 2021, 89, .	2.2	2
12	Phenotypic and Genomic Profiling of Staphylococcus argenteus in Canada and the United States and Recommendations for Clinical Result Reporting. Journal of Clinical Microbiology, 2021, 59, .	3.9	11
13	Practical guidance for clinical laboratories for SARS-CoV-2 serology testing. Canada Communicable Disease Report, 2021, 47, 171-183.	1.3	12
14	Characteristics of SARS-CoV-2 testing for rapid diagnosis of COVID-19 during the initial stages of a global pandemic. PLoS ONE, 2021, 16, e0253941.	2.5	3
15	Antimicrobial susceptibility profiles of invasive isolates of anaerobic bacteria from a large Canadian reference laboratory: 2012–2019. Anaerobe, 2021, 70, 102386.	2.1	17
16	Evaluation of commercial SARS-CoV-2 serological assays in Canadian public health laboratories. Diagnostic Microbiology and Infectious Disease, 2021, 101, 115412.	1.8	10
17	CYP51A polymorphisms of <i> Aspergillus fumigatus < /i > in lung transplant recipients: Prevalence, correlation with phenotype, and impact on outcomes. Medical Mycology, 2021, 59, 728-733.</i>	0.7	3
18	Increased Incidence of Invasive Haemophilus influenzae Disease Driven by Non-Type B Isolates in Ontario, Canada, 2014 to 2018. Microbiology Spectrum, 2021, 9, e0080321.	3.0	7

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19	Population-based incidence of invasive pneumococcal disease in children and adults in Ontario and British Columbia, 2002–2018: A Canadian Immunization Research Network (CIRN) study. Vaccine, 2021, 39, 7545-7553.	3.8	5
20	Whole genome sequencing to study the phylogenetic structure of serotype a <i>Haemophilus influenzae</i> recovered from patients in Canada. Canadian Journal of Microbiology, 2020, 66, 99-110.	1.7	7
21	Antifungal Susceptibility of Clinical Yeast Isolates from a Large Canadian Reference Laboratory and Application of Whole-Genome Sequence Analysis To Elucidate Mechanisms of Acquired Resistance. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	18
22	Identification and characterization of invasive multi-drug-resistant (MDR) Bacteroides genomospecies in Canada. Access Microbiology, 2020, 2, acmi000111.	0.5	5
23	Whole-Genome Sequencing Reveals a Novel Subclade of Pansusceptible Candida auris in Ontario, Canada. Infection Control and Hospital Epidemiology, 2020, 41, s57-s58.	1.8	O
24	Continuing surveillance of invasive <i>Haemophilus influenzae</i> disease in northwestern Ontario emphasizes the importance of serotype a and non-typeable strains as causes of serious disease: a Canadian Immunization Research Network (CIRN) Study. Canadian Journal of Microbiology, 2019, 65, 805-813.	1.7	7
25	Identification and Characterization of $\hat{a} \in \infty$ <i>Haemophilus quentini</i> $\hat{a} \in S$ trains Causing Invasive Disease in Ontario, Canada (2016 to 2018). Journal of Clinical Microbiology, 2019, 57, .	3.9	5
26	Mycobiome Sequencing and Analysis Applied to Fungal Community Profiling of the Lower Respiratory Tract During Fungal Pathogenesis. Frontiers in Microbiology, 2019, 10, 512.	3.5	34
27	Application of whole genome sequencing to query a potential outbreak of Elizabethkingia anophelis in Ontario, Canada. Access Microbiology, 2019, 1, e000017.	0.5	7
28	In Vitro Antibiotic Susceptibility Pattern of Non-diphtheriae Corynebacterium Isolates in Ontario, Canada, from 2011 to 2016. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	36
29	Epidemiology and Geographic Distribution of Blastomycosis, Histoplasmosis, and Coccidioidomycosis, Ontario, Canada, 1990–2015. Emerging Infectious Diseases, 2018, 24, 1257-1266.	4.3	39
30	Incubation time and culture media impact success of identification of Nocardia spp. by MALDI-ToF mass spectrometry. Diagnostic Microbiology and Infectious Disease, 2018, 92, 270-274.	1.8	15
31	Antifungal Susceptibility Testing of Aspergillus spp. by Using a Composite Correlation Index (CCI)-Based Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry Method Appears To Not Offer Benefit over Traditional Broth Microdilution Testing. Journal of Clinical Microbiology, 2017, 55, 2030-2034.	3.9	17
32	Blastomycosis infection in an adolescent patient with Hodgkin lymphoma. Pediatric Blood and Cancer, 2017, 64, e26250.	1.5	0
33	Fungal keratitis caused by Metarhizium anisopliae complex. Medical Mycology Case Reports, 2017, 17, 28-30.	1.3	9
34	<i>Blastomyces gilchristii</i> as Cause of Fatal Acute Respiratory Distress Syndrome. Emerging Infectious Diseases, 2016, 22, 306-308.	4.3	14
35	Draft Genome Sequences of Two " <i>Haemophilus quentini</i> ―Isolates Recovered from Two Different Patients' Blood Cultures. Genome Announcements, 2016, 4, .	0.8	3
36	Bacteremia Due to <i>Pasteurella dagmatis</i> Acquired from a Dog Bite, with a Review of Systemic Infections and Challenges in Laboratory Identification. Canadian Journal of Infectious Diseases and Medical Microbiology, 2015, 26, 273-276.	1.9	4

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37	First draft genome sequence of Aureimonas altamirensis, isolated from patient blood culture. FEMS Microbiology Letters, 2015, 362, .	1.8	10
38	Multilocus Sequence Typing of Mycobacterium xenopi. Journal of Clinical Microbiology, 2014, 52, 3973-3977.	3.9	5
39	Nosocomial Transmission of New Delhi Metallo- \hat{l}^2 -Lactamase-1-Producing Klebsiella pneumoniae in Toronto, Canada. Infection Control and Hospital Epidemiology, 2013, 34, 49-55.	1.8	33
40	Bile Salts Induce Resistance to Polymyxin in Enterohemorrhagic Escherichia coliO157:H7. Journal of Bacteriology, 2011, 193, 4509-4515.	2.2	52
41	Pseudomonas aeruginosa d-Arabinofuranose Biosynthetic Pathway and Its Role in Type IV Pilus Assembly. Journal of Biological Chemistry, 2011, 286, 28128-28137.	3.4	21
42	Novel Proteins That Modulate Type IV Pilus Retraction Dynamics in <i>Pseudomonas aeruginosa</i> Journal of Bacteriology, 2008, 190, 7022-7034.	2.2	46
43	Modification of <i>Pseudomonas aeruginosa</i> Pa5196 Type IV Pilins at Multiple Sites with <scp>d</scp> -Ara <i>f</i> by a Novel GT-C Family Arabinosyltransferase, TfpW. Journal of Bacteriology, 2008, 190, 7464-7478.	2.2	52
44	Glycosylation of <i>Pseudomonas aeruginosa</i> Strain Pa5196 Type IV Pilins with Mycobacterium-Like α-1,5-Linked <scp>d</scp> -Ara <i>f</i> Oligosaccharides. Journal of Bacteriology, 2007, 189, 151-159.	2.2	70
45	Abnormalities in the Pulmonary Innate Immune System in Cystic Fibrosis. American Journal of Respiratory Cell and Molecular Biology, 2006, 34, 364-374.	2.9	36
46	Age-Related Resistance in Arabidopsis Is a Developmentally Regulated Defense Response to Pseudomonas syringae. Plant Cell, 2002, 14, 479-490.	6.6	237
47	Molecular identification of Aspergillus fumigatus complex from lung transplant recipients using multilocus sequencing analysis (MLSA). Jammi, 0, , .	0.5	O