Jolene R Bowers

List of Publications by Year
in descending order

19 papers	**790** citations	759233 **12** h-index	839539 **18** g-index
19 all docs	**19** docs citations	**19** times ranked	**1334** citing authors

#	Article	IF	Citations
1	Genomic investigation of a household SARS-CoV-2 disease cluster in Arizona involving a cat, dog, and pet owner. One Health, 2021, 13, 100333.	3.4	15
2	Sequencing the pandemic: rapid and high-throughput processing and analysis of COVID-19 clinical samplesÂfor 21stÂcentury public health. F1000Research, 2021, 10, 48.	1.6	6
3	Normalization of SARS-CoV-2 viral load via RT-qPCR provides higher-resolution data for comparison across time and between patients. Virus Research, 2021, 306, 198604.	2.2	7
4	The detection of Coccidioides from ambient air in Phoenix, Arizona: Evidence of uneven distribution and seasonality. Medical Mycology, 2020, 58, 552-559.	0.7	23
5	Suspected Locally Acquired Coccidioidomycosis in Human, Spokane, Washington, USA. Emerging Infectious Diseases, 2020, 26, 606-609.	4.3	10
6	<i>Rhizopus microsporus</i> Infections Associated with Surgical Procedures, Argentina, 2006â€"2014. Emerging Infectious Diseases, 2020, 26, 937-944.	4.3	11
7	Coyotes as Reservoirs for <i>Onchocerca lupi</i>, United States, 2015â€"2018. Emerging Infectious Diseases, 2020, 26, 2989-2993.	4.3	17
8	Genomic Analyses of Acute Flaccid Myelitis Cases among a Cluster in Arizona Provide Further Evidence of Enterovirus D68 Role. MBio, 2019, 10, .	4.1	15
9	Improved Subtyping of Staphylococcus aureus Clonal Complex 8 Strains Based on Whole-Genome Phylogenetic Analysis. MSphere, 2018, 3, .	2.9	32
10	KlebSeq, a Diagnostic Tool for Surveillance, Detection, and Monitoring of Klebsiella pneumoniae. Journal of Clinical Microbiology, 2016, 54, 2582-2596.	3.9	50
11	Genomic Analysis of the Emergence and Rapid Global Dissemination of the Clonal Group 258 Klebsiella pneumoniae Pandemic. PLoS ONE, 2015, 10, e0133727.	2.5	195
12	Dust Devil: The Life and Times of the Fungus That Causes Valley Fever. PLoS Pathogens, 2015, 11, e1004762.	4.7	49
13	Using Whole Genome Analysis to Examine Recombination across Diverse Sequence Types of Staphylococcus aureus. PLoS ONE, 2015, 10, e0130955.	2.5	40
14	Comparison of TaqMan PCR Assays for Detection of the Melioidosis Agent Burkholderia pseudomallei in Clinical Specimens. Journal of Clinical Microbiology, 2012, 50, 2059-2062.	3.9	44
15	Towards a rapid molecular diagnostic for melioidosis: Comparison of DNA extraction methods from clinical specimens. Journal of Microbiological Methods, 2012, 88, 179-181.	1.6	26
16	Dominance of multidrug resistant CC271 clones in macrolide-resistant streptococcus pneumoniae in Arizona. BMC Microbiology, 2012, 12, 12.	3.3	31
17	BurkDiff: A Real-Time PCR Allelic Discrimination Assay for Burkholderia Pseudomallei and B. mallei. PLoS ONE, 2010, 5, e15413.	2.5	38
18	Phylogeography of <i>Francisella tularensis</i> : Global Expansion of a Highly Fit Clone. Journal of Bacteriology, 2009, 191, 2474-2484.	2.2	176

#	Article	IF	Citations
19	Methods for sequencing the pandemic: benefits of rapid or high-throughput processing. F1000Research, 0, 10, 48.	1.6	5