

Jian Zhuge

List of Publications by Year in descending order

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Version: 2024-02-01

33
papers

1,331
citations

411340

20
h-index

445137

33
g-index

33
all docs

33
docs citations

33
times ranked

2524
citing authors

#	ARTICLE	IF	CITATIONS
1	Biennial Upsurge and Molecular Epidemiology of Enterovirus D68 Infection in New York, USA, 2014 to 2018. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	1.8	15
2	Complete Genome Sequences of Four Toxigenic <i>Clostridium difficile</i> Clinical Isolates from Patients of the Lower Hudson Valley, New York, USA. <i>Genome Announcements</i> , 2018, 6, .	0.8	5
3	Estrogen-dependent epigenetic regulation of soluble epoxide hydrolase via DNA methylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 613-618.	3.3	34
4	Evolution and mutations predisposing to daptomycin resistance in vancomycin-resistant <i>Enterococcus faecium</i> ST736 strains. <i>PLoS ONE</i> , 2018, 13, e0209785.	1.1	27
5	Enterovirus D68 Subclade B3 Strain Circulating and Causing an Outbreak in the United States in 2016. <i>Scientific Reports</i> , 2017, 7, 1242.	1.6	67
6	Emergence and Evolution of Multidrug-Resistant <i>Klebsiella pneumoniae</i> with both <i>bla</i> _{KPC} and <i>bla</i> _{CTX-M} Integrated in the Chromosome. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	1.4	66
7	Complete Genome Sequence of a Colistin-Resistant <i>Escherichia coli</i> Strain Harboring <i>mcr-1</i> on an IncHI2 Plasmid in the United States. <i>Genome Announcements</i> , 2017, 5, .	0.8	23
8	Complete Genome Sequences of Nine Enterovirus D68 Strains from Patients of the Lower Hudson Valley, New York, 2016. <i>Genome Announcements</i> , 2016, 4, .	0.8	8
9	Insights into <i>Borrelia miyamotoi</i> infection from an untreated case demonstrating relapsing fever, monocytosis and a positive C6 Lyme serology. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 86, 93-96.	0.8	35
10	Recurrent TERT promoter mutations in urothelial carcinoma and potential clinical applications. <i>Annals of Diagnostic Pathology</i> , 2016, 21, 7-11.	0.6	38
11	Assessing next-generation sequencing and 4 bioinformatics tools for detection of Enterovirus D68 and other respiratory viruses in clinical samples. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 85, 26-29.	0.8	8
12	Whole-Genome Sequence Analysis Reveals the Enterovirus D68 Isolates during the United States 2014 Outbreak Mainly Belong to a Novel Clade. <i>Scientific Reports</i> , 2015, 5, 15223.	1.6	50
13	Utilization of a real-time PCR assay for diagnosis of <i>Babesia microti</i> infection in clinical practice. <i>Ticks and Tick-borne Diseases</i> , 2015, 6, 376-382.	1.1	43
14	Comparison of a quantitative PCR assay with peripheral blood smear examination for detection and quantitation of <i>Babesia microti</i> infection in humans. <i>Diagnostic Microbiology and Infectious Disease</i> , 2015, 82, 109-113.	0.8	34
15	Evaluation of a Real-Time Reverse Transcription-PCR Assay for Detection of Enterovirus D68 in Clinical Samples from an Outbreak in New York State in 2014. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1915-1920.	1.8	24
16	Heterogeneity of TERT promoter mutations status in squamous cell carcinomas of different anatomical sites. <i>Annals of Diagnostic Pathology</i> , 2015, 19, 146-148.	0.6	34
17	Distinguishing Nested Variants of Urothelial Carcinoma From Benign Mimickers by TERT Promoter Mutation. <i>American Journal of Surgical Pathology</i> , 2015, 39, 127-131.	2.1	78
18	High frequency of TERT promoter mutation in small cell carcinoma of bladder, but not in small cell carcinoma of other origins. <i>Journal of Hematology and Oncology</i> , 2014, 7, 47.	6.9	66

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19	Identification of a Novel Clone, ST736, among <i>Enterococcus faecium</i> Clinical Isolates and Its Association with Daptomycin Nonsusceptibility. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 4848-4854.	1.4	15
20	B-Raf and the inhibitors: from bench to bedside. <i>Journal of Hematology and Oncology</i> , 2013, 6, 30.	6.9	80
21	Sensitive detection of BRAF V600E mutation by Amplification Refractory Mutation System (ARMS)-PCR. <i>Biomarker Research</i> , 2013, 1, 3.	2.8	64
22	CTX-M β -Lactamase-producing <i>Klebsiella pneumoniae</i> in Suburban New York City, New York, USA. <i>Emerging Infectious Diseases</i> , 2013, 19, 1803-1810.	2.0	46
23	Ethanol Induction of CYP2A5: Permissive Role for CYP2E1. <i>Drug Metabolism and Disposition</i> , 2011, 39, 330-336.	1.7	40
24	Inhibition of the mitochondrial permeability transition by cyclosporin A prevents pyrazole plus lipopolysaccharide-induced liver injury in mice. <i>Free Radical Biology and Medicine</i> , 2009, 46, 406-413.	1.3	11
25	Overexpression of CYP2E1 induces HepG2 cells death by the AMP kinase activator 5- α -aminoimidazole-4-carboxamide-1- β -D-ribofuranoside (AICAR). <i>Cell Biology and Toxicology</i> , 2009, 25, 253-263.	2.4	6
26	A decrease in S-adenosyl-L-methionine potentiates arachidonic acid cytotoxicity in primary rat hepatocytes enriched in CYP2E1. <i>Molecular and Cellular Biochemistry</i> , 2008, 314, 105-112.	1.4	3
27	Cytochrome P450 2E1 contributes to ethanol-induced fatty liver in mice. <i>Hepatology</i> , 2008, 47, 1483-1494.	3.6	262
28	Depletion of S-adenosyl-L-methionine with cycloleucine potentiates cytochrome P450 2E1 toxicity in primary rat hepatocytes. <i>Archives of Biochemistry and Biophysics</i> , 2007, 466, 177-185.	1.4	23
29	Serum deprivation-induced HepG2 cell death is potentiated by CYP2E1. <i>Free Radical Biology and Medicine</i> , 2006, 40, 63-74.	1.3	61
30	Increased toxicity by transforming growth factor-beta 1 in liver cells overexpressing CYP2E1. <i>Free Radical Biology and Medicine</i> , 2006, 41, 1100-1112.	1.3	28
31	Stable expression of human cytochrome P450 2D6*10 in HepG2 cells. <i>World Journal of Gastroenterology</i> , 2004, 10, 234.	1.4	12
32	Three new alternative splicing variants of human cytochrome P450 2D6 mRNA in human extratumoral liver tissue. <i>World Journal of Gastroenterology</i> , 2004, 10, 3356.	1.4	9
33	Heterologous expression of human cytochrome P450 2E1 in HepG2 cell line. <i>World Journal of Gastroenterology</i> , 2003, 9, 2732.	1.4	16