

Nguyet Kong

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

419
citations

932766

10
h-index

887659

17
g-index

19
all docs

19
docs citations

19
times ranked

668
citing authors

#	ARTICLE	IF	CITATIONS
1	Salmonella Degrades the Host Glycocalyx Leading to Altered Infection and Glycan Remodeling. Scientific Reports, 2016, 6, 29525.	1.6	66
2	Fallacy of the Unique Genome: Sequence Diversity within Single <i>Helicobacter pylori</i> Strains. MBio, 2017, 8, .	1.8	64
3	A Syst-OMICS Approach to Ensuring Food Safety and Reducing the Economic Burden of Salmonellosis. Frontiers in Microbiology, 2017, 8, 996.	1.5	42
4	Complete Genome Sequences of a Clinical Isolate and an Environmental Isolate of <i>Vibrio parahaemolyticus</i> . Genome Announcements, 2015, 3, .	0.8	32
5	Comparative Genomics Reveals the Diversity of Restriction-Modification Systems and DNA Methylation Sites in <i>Listeria monocytogenes</i> . Applied and Environmental Microbiology, 2017, 83, .	1.4	31
6	Food authentication from shotgun sequencing reads with an application on high protein powders. Npj Science of Food, 2019, 3, 24.	2.5	31
7	Automation of PacBio SMRTbell NGS library preparation for bacterial genome sequencing. Standards in Genomic Sciences, 2017, 12, 27.	1.5	24
8	Prebiotic Oligosaccharides Potentiate Host Protective Responses against <i>L. Monocytogenes</i> Infection. Pathogens, 2017, 6, 68.	1.2	22
9	Monitoring the microbiome for food safety and quality using deep shotgun sequencing. Npj Science of Food, 2021, 5, 3.	2.5	22
10	Large-Scale Release of <i>Campylobacter</i> Draft Genomes: Resources for Food Safety and Public Health from the 100K Pathogen Genome Project. Genome Announcements, 2017, 5, .	0.8	15
11	Genomic characterization of a diazotrophic microbiota associated with maize aerial root mucilage. PLoS ONE, 2020, 15, e0239677.	1.1	13
12	Draft Genome Sequences of 1,183 <i>Salmonella</i> Strains from the 100K Pathogen Genome Project. Genome Announcements, 2017, 5, .	0.8	12
13	Draft Genome Sequences of <i>Campylobacter jejuni</i> Strains That Cause Abortion in Livestock. Genome Announcements, 2016, 4, .	0.8	11
14	Draft Genome Sequence of Multidrug-Resistant Abortive <i>Campylobacter jejuni</i> from Northern California. Genome Announcements, 2017, 5, .	0.8	9
15	100K Pathogen Genome Project: 306 <i>Listeria</i> Draft Genome Sequences for Food Safety and Public Health. Genome Announcements, 2017, 5, .	0.8	7
16	<i>Shigella</i> Draft Genome Sequences: Resources for Food Safety and Public Health. Genome Announcements, 2017, 5, .	0.8	5
17	Draft Genome Sequence of <i>Bacillus velezensis</i> CE2, Which Genetically Encodes a Novel Multicomponent Lantibiotic. Microbiology Resource Announcements, 2019, 8, .	0.3	4