## Akiyo Nakano

List of Publications by Year in descending order

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Version: 2024-02-01

933447 888059 41 393 10 17 citations h-index g-index papers 41 41 41 417 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Predominance of CTX-M-9 Group Among ESBL-Producing <i>Escherichia coli</i> Isolated from Healthy Individuals in Japan. Microbial Drug Resistance, 2022, , .	2.0	3
2	Comamonas thiooxydans Expressing a Plasmid-Encoded IMP-1 Carbapenemase Isolated From Continuous Ambulatory Peritoneal Dialysis of an Inpatient in Japan. Frontiers in Microbiology, 2022, 13, 808993.	3.5	1
3	Inactivation of various variant types of SARS-CoV-2 by indoor-light-sensitive TiO2-based photocatalyst. Scientific Reports, 2022, 12, 5804.	3.3	29
4	Evaluation of NG-Test CARBA 5 for the detection of carbapenemase-producing Gram-negative bacilli. Journal of Medical Microbiology, 2022, 71, .	1.8	2
5	Emergence and Evolution of Unique Plasmids Harboring <i>bla</i> <sub>IMP-70</sub> and <i>bla</i> <sub>CTX-M-253</sub> in Multidrug-Resistant Providencia rettgeri. Microbiology Spectrum, 2022, 10, .	3.0	2
6	Prevalence of Helicobacter pylori among residents and their environments in the Nara prefecture, Japan. Journal of Infection and Public Health, 2021, 14, 271-275.	4.1	2
7	Acinetobacter baumannii LOS Regulate the Expression of Inflammatory Cytokine Genes and Proteins in Human Mast Cells. Pathogens, 2021, 10, 290.	2.8	5
8	Prevalence and Relatedness of mcr-1-Mediated Colistin-Resistant Escherichia coli Isolated From Livestock and Farmers in Japan. Frontiers in Microbiology, 2021, 12, 664931.	3.5	11
9	Preparation of cerium molybdates and their antiviral activity against bacteriophage $\hat{l}$ and SARS-CoV-2. Materials Letters, 2021, 290, 129510.	2.6	21
10	Dysbiosis in the Salivary Microbiome Associated with IgA Nephropathy—â€A―â€Japanese Cohort Study. Microbes and Environments, 2021, 36, n/a.	1.6	7
11	The Role of nmcR, ampR, and ampD in the Regulation of the Class A Carbapenemase NmcA in Enterobacter ludwigii. Frontiers in Microbiology, 2021, 12, 794134.	3.5	0
12	Persimmon-derived tannin has antiviral effects and reduces the severity of infection and transmission of SARS-CoV-2 in a Syrian hamster model. Scientific Reports, 2021, 11, 23695.	3.3	26
13	A Novel Mismatched PCR-Restriction Fragment Length Polymorphism Assay for Rapid Detection of <i>gyrA</i> and <i>parC</i> Mutations Associated With Fluoroquinolone Resistance in <i>Acinetobacter baumannii</i> Annals of Laboratory Medicine, 2020, 40, 27-32.	2.5	6
14	Filifactor alocis brain abscess identified by 16S ribosomal RNA gene sequencing: A case report. Journal of Infection and Chemotherapy, 2020, 26, 305-307.	1.7	8
15	Molecular and Epidemiological Characteristics of Carbapenemase-Producing Klebsiella pneumoniae Clinical Isolates in Japan. MSphere, 2020, 5, .	2.9	23
16	Molecular characteristics of extended-spectrum β-lactamase-producing Klebsiella pneumoniae in Japan: Predominance of CTX-M-15 and emergence of hypervirulent clones. International Journal of Infectious Diseases, 2020, 98, 281-286.	3.3	16
17	Environmental Presence and Genetic Characteristics of Carbapenemase-Producing <i>Enterobacteriaceae</i> from Hospital Sewage and River Water in the Philippines. Applied and Environmental Microbiology, 2020, 86, .	3.1	39
18	Clinical characteristics and molecular epidemiology of invasive Streptococcus agalactiae infections between 2007 and 2016 in Nara, Japan. PLoS ONE, 2020, 15, e0240590.	2.5	8

#	Article	IF	Citations
19	Development of a loop-mediated isothermal amplification assay for rapid Helicobacter pylori detection. Journal of Microbiological Methods, 2019, 163, 105653.	1.6	7
20	Mycoplasma genitalium Infection and Chronic Inflammation in Human Prostate Cancer: Detection Using Prostatectomy and Needle Biopsy Specimens. Cells, 2019, 8, 212.	4.1	46
21	Prevalence and mechanism of fluoroquinolone resistance in clinical isolates of Proteus mirabilis in Japan. Heliyon, 2019, 5, e01291.	3.2	5
22	Emergence of IMP-34- and OXA-58-Producing Carbapenem-Resistant Acinetobacter colistiniresistens. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	11
23	Comparison of the inoculum size effects of antibiotics on IMP-6 $\hat{l}^2$ -lactamase-producing Enterobacteriaceae co-harboring plasmid-mediated quinolone resistance genes. PLoS ONE, 2019, 14, e0225210.	2.5	2
24	Emergence of VIM-2-producing Citrobacter freundii in Japan. Infectious Diseases, 2018, 50, 862-863.	2.8	6
25	Rapid Identification of <i>bla</i> <sub>IMP-1</sub> and <i>bla</i> <sub>IMP-6</sub> by Multiplex Amplification Refractory Mutation System PCR. Annals of Laboratory Medicine, 2018, 38, 378-380.	2.5	9
26	Suitability of Carbapenem Inactivation Method (CIM) for Detection of IMP Metallo-Î <sup>2</sup> -Lactamase-Producing Enterobacteriaceae. Journal of Clinical Microbiology, 2017, 55, 1220-1222.	3.9	15
27	The TNF-α of mast cells induces pro-inflammatory responses during infection with Acinetobacter baumannii. Immunobiology, 2017, 222, 1025-1034.	1.9	10
28	Role of AmpR in the High Expression of the Plasmid-Encoded AmpC $\hat{l}^2$ -Lactamase CFE-1. MSphere, 2017, 2, .	2.9	10
29	Draft Genome Sequence of <i>Leuconostoc mesenteroides</i> 213M0, Isolated from Traditional Fermented Mare Milk Airag in Bulgan Aimag, Mongolia. Genome Announcements, 2016, 4, .	0.8	0
30	Draft Genome Sequence of <i>Leuconostoc mesenteroides</i> 406 Isolated from the Traditional Fermented Mare Milk Airag in Tuv Aimag, Mongolia. Genome Announcements, 2016, 4, .	0.8	2
31	Draft Genome Sequence of Coccoid Lactobacillus equigenerosi NRIC 0697 <sup>T</sup> Isolated from the Gastrointestinal Tracts of Healthy Thoroughbreds. Genome Announcements, 2016, 4, .	0.8	0
32	Draft Genome Sequence of Streptococcus orisasini SH06, Isolated from a Healthy Thoroughbred Gastrointestinal Tract. Genome Announcements, 2016, 4, .	0.8	1
33	Loop-mediated isothermal amplification: Rapid and sensitive detection of the antibiotic resistance gene ISAba1-blaOXA-51-like in Acinetobacter baumannii. Journal of Microbiological Methods, 2016, 121, 36-40.	1.6	13
34	Draft Genome Sequence of Bifidobacterium aesculapii DSM 26737 <code><sup>T</sup></code> , Isolated from Feces of Baby Common Marmoset. Genome Announcements, 2015, 3, .	0.8	6
35	Complete genome sequence of Bifidobacterium angulatum JCM 7096T isolated from human feces. Journal of Biotechnology, 2015, 211, 10-11.	3.8	5
36	Complete genome sequence of Bifidobacterium bifidum JCM 1255T isolated from feces of a breast-fed infant. Journal of Biotechnology, 2015, 210, 66-67.	3.8	4

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37	Complete genome sequence of Bifidobacterium breve JCM 1192T isolated from infant feces. Journal of Biotechnology, 2015, 210, 81-82.	3.8	5
38	Complete genome sequence of Bifidobacterium catenulatum JCM 1194T isolated from human feces. Journal of Biotechnology, 2015, 210, 25-26.	3.8	7
39	Complete genome sequence of Bifidobacterium pseudocatenulatum JCM 1200T isolated from infant feces. Journal of Biotechnology, 2015, 210, 68-69.	3.8	7
40	Complete Genome Sequence of Parascardovia denticolens JCM 12538 T , Isolated from Human Dental Caries. Genome Announcements, 2015, 3, .	0.8	2
41	Rapid assay for detecting gyrA and parC mutations associated with fluoroquinolone resistance in Enterobacteriaceae. Journal of Microbiological Methods, 2013, 94, 213-216.	1.6	11