## Jingrang Lu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12037666/publications.pdf

Version: 2024-02-01

471509 713466 1,394 21 17 21 citations h-index g-index papers 21 21 21 1548 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Quantitative PCR for Detection and Enumeration of Genetic Markers of Bovine Fecal Pollution. Applied and Environmental Microbiology, 2008, 74, 745-752.	3.1	183
2	Evaluation of Broiler Litter with Reference to the Microbial Composition as Assessed by Using 16S rRNA and Functional Gene Markers. Applied and Environmental Microbiology, 2003, 69, 901-908.	3.1	176
3	Phylogenetic Diversity and Molecular Detection of Bacteria in Gull Feces. Applied and Environmental Microbiology, 2008, 74, 3969-3976.	3.1	163
4	Genomic Sequence and Evolution of Marine Cyanophage P60: a New Insight on Lytic and Lysogenic Phages. Applied and Environmental Microbiology, 2002, 68, 2589-2594.	3.1	158
5	Distribution, Isolation, Host Specificity, and Diversity of Cyanophages Infecting Marine Synechococcus spp. in River Estuaries. Applied and Environmental Microbiology, 2001, 67, 3285-3290.	3.1	109
6	Identification of chicken-specific fecal microbial sequences using a metagenomic approach. Water Research, 2007, 41, 3561-3574.	11.3	73
7	Microbial Diversity and Host-Specific Sequences of Canada Goose Feces. Applied and Environmental Microbiology, 2009, 75, 5919-5926.	3.1	68
8	Microbial diversities (16S and 18S rRNA gene pyrosequencing) and environmental pathogens within drinking water biofilms grown on the common premise plumbing materials unplasticized polyvinylchloride and copper. FEMS Microbiology Ecology, 2014, 88, 280-295.	2.7	67
9	Identification of Bacterial DNA Markers for the Detection of Human Fecal Pollution in Water. Applied and Environmental Microbiology, 2007, 73, 2416-2422.	3.1	61
10	Turkey fecal microbial community structure and functional gene diversity revealed by 16S rRNA gene and metagenomic sequences. Journal of Microbiology, 2008, 46, 469-477.	2.8	44
11	Eukaryotic diversity in premise drinking water using 18S rDNA sequencing: implications for health risks. Environmental Science and Pollution Research, 2013, 20, 6351-6366.	<b>5.</b> 3	43
12	Distribution and potential significance of a gull fecal marker in urban coastal and riverine areas of southern Ontario, Canada. Water Research, 2011, 45, 3960-3968.	11.3	42
13	Preferential colonization and release of Legionella pneumophila from mature drinking water biofilms grown on copper versus unplasticized polyvinylchloride coupons. International Journal of Hygiene and Environmental Health, 2014, 217, 219-225.	4.3	40
14	Molecular Detection of Campylobacter spp. in California Gull (Larus californicus) Excreta. Applied and Environmental Microbiology, 2011, 77, 5034-5039.	3.1	34
15	Annual variations and effects of temperature on Legionella spp. and other potential opportunistic pathogens in a bathroom. Environmental Science and Pollution Research, 2017, 24, 2326-2336.	<b>5.</b> 3	32
16	Legionella and other opportunistic pathogens in full-scale chloraminated municipal drinking water distribution systems. Water Research, 2021, 205, 117571.	11.3	32
17	Development and Evaluation of a Quantitative PCR Assay Targeting Sandhill Crane (Grus canadensis) Fecal Pollution. Applied and Environmental Microbiology, 2012, 78, 4338-4345.	3.1	27
18	Impacts of Migratory Sandhill Cranes (Grus canadensis) on Microbial Water Quality in the Central Platte River, Nebraska, USA. Water, Air, and Soil Pollution, 2013, 224, 1.	2.4	19

#	Article	IF	CITATIONS
19	Exposure to Synthetic Gray Water Inhibits Amoeba Encystation and Alters Expression of Legionella pneumophila Virulence Genes. Applied and Environmental Microbiology, 2015, 81, 630-639.	3.1	12
20	The Bacterial Community Diversity of Bathroom Hot Tap Water Was Significantly Lower Than That of Cold Tap and Shower Water. Frontiers in Microbiology, 2021, 12, 625324.	3.5	8
21	Comparative Study on the Performance of Anaerobic and Aerobic Biotrickling Filter for Removal of Chloroform. Environmental Engineering Science, 2018, 35, 462-471.	1.6	3