Yu-Tzu Lin

List of Publications by Year in descending order

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ΥΠ-ΤΖΗ ΓΙΝ

#	Article	IF	CITATIONS
1	AmpD _I Is Involved in Expression of the Chromosomal L1 and L2 β-Lactamases of <i>Stenotrophomonas maltophilia</i> . Antimicrobial Agents and Chemotherapy, 2009, 53, 2902-2907.	3.2	60
2	Emergence of a small colony variant of vancomycin-intermediate <i>Staphylococcus aureus</i> in a patient with septic arthritis during long-term treatment with daptomycin. Journal of Antimicrobial Chemotherapy, 2016, 71, 1807-1814.	3.0	34
3	AmpN-AmpG Operon Is Essential for Expression of L1 and L2 β-Lactamases in <i>Stenotrophomonas maltophilia</i> . Antimicrobial Agents and Chemotherapy, 2010, 54, 2583-2589.	3.2	32
4	Rapid antibiotic susceptibility testing of bacteria from patients' blood via assaying bacterial metabolic response with surface-enhanced Raman spectroscopy. Scientific Reports, 2020, 10, 12538.	3.3	30
5	Distribution of antibiotic resistance genes among Staphylococcus species isolated from ready-to-eat foods. Journal of Food and Drug Analysis, 2019, 27, 841-848.	1.9	28
6	Skin Commensal Staphylococci May Act as Reservoir for Fusidic Acid Resistance Genes. PLoS ONE, 2015, 10, e0143106.	2.5	28
7	A novel fusidic acid resistance determinant, fusF, in Staphylococcus cohnii. Journal of Antimicrobial Chemotherapy, 2015, 70, 416-419.	3.0	26
8	A Novel Staphylococcal Cassette Chromosomal Element, SCC <i>fusC</i> , Carrying <i>fusC</i> and <i>speG</i> in Fusidic Acid-Resistant Methicillin-Resistant Staphylococcus aureus. Antimicrobial Agents and Chemotherapy, 2014, 58, 1224-1227.	3.2	24
9	Effects of toluidine blue O (TBO)-photodynamic inactivation on community-associated methicillin-resistant Staphylococcus aureus isolates. Journal of Microbiology, Immunology and Infection, 2017, 50, 46-54.	3.1	23
10	Clinical characteristics of patients with bacteraemia due to the emergence of mcr-1-harbouring Enterobacteriaceae in humans and pigs in Taiwan. International Journal of Antimicrobial Agents, 2018, 52, 651-657.	2.5	19
11	Molecular Evolutionary Pathways toward Two Successful Community-Associated but Multidrug-Resistant ST59 Methicillin-Resistant Staphylococcus aureus Lineages in Taiwan: Dynamic Modes of Mobile Genetic Element Salvages. PLoS ONE, 2016, 11, e0162526.	2.5	19
12	New Structure of Phage-Related Islands Carrying <i>fusB</i> and a Virulence Gene in Fusidic Acid-Resistant Staphylococcus epidermidis. Antimicrobial Agents and Chemotherapy, 2013, 57, 5737-5739.	3.2	18
13	Genotypes and phenotypes of Staphylococcus lugdunensis isolates recovered from bacteremia. Journal of Microbiology, Immunology and Infection, 2015, 48, 397-405.	3.1	17
14	Novel Structure of Enterococcus faecium-Originated <i>ermB</i> -Positive Tn <i>1546</i> -Like Element in Staphylococcus aureus. Antimicrobial Agents and Chemotherapy, 2016, 60, 6108-6114.	3.2	15
15	Long-Term Surveillance of Antibiotic Prescriptions and the Prevalence of Antimicrobial Resistance in Non-Fermenting Gram-Negative Bacilli. Microorganisms, 2020, 8, 397.	3.6	11
16	Haplotype distribution of SARS-CoV-2 variants in low and high vaccination rate countries during ongoing global COVID-19 pandemic in early 2021. Infection, Genetics and Evolution, 2022, 97, 105164.	2.3	9
17	Staphylococcus taiwanensis sp. nov., isolated from human blood. International Journal of Systematic and Evolutionary Microbiology, 2022, 72, .	1.7	7
18	A Possible Role of Insertion Sequence IS1216V in Dissemination of Multidrug-Resistant Elements MESPM1 and MES6272-2 between Enterococcus and ST59 Staphylococcus aureus. Microorganisms, 2020, 8, 1905.	3.6	6

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19	Distribution of Staphylococcal Cassette Chromosome (SCC)mecElement Types in Fusidic Acid-Resistant Staphylococcus epidermidis and Identification of a Novel SCC7684Element. Antimicrobial Agents and Chemotherapy, 2016, 60, 5006-5009.	3.2	5
20	Rapid identification of Streptococcus intermedius by multiplex polymerase chain reaction 1 week before culture positivity in a patient with antibiotic-treated thalamic brain abscess. Journal of Microbiology, Immunology and Infection, 2017, 50, 549-551.	3.1	5
21	Genomic comparison between Staphylococcus aureus GN strains clinically isolated from a familial infection case: IS1272 transposition through a novel inverted repeat-replacing mechanism. PLoS ONE, 2017, 12, e0187288.	2.5	5
22	Heterogeneity of Molecular Characteristics among Staphylococcus argenteus Clinical Isolates (ST2250, ST2793, ST1223, and ST2198) in Northern Taiwan. Microorganisms, 2020, 8, 1157.	3.6	5
23	In Vitro and In Vivo Evaluations of β-Lactam/β-Lactamase Mono- and Combined Therapies against Carbapenem-Nonsusceptible Enterobacteriaceae in Taiwan. Microorganisms, 2020, 8, 1981.	3.6	5
24	Potentially conjugative plasmids harboring Tn6636, a multidrug-resistant and composite mobile element, in Staphylococcus aureus. Journal of Microbiology, Immunology and Infection, 2022, 55, 225-233.	3.1	5
25	The extent of molecular variation in novel SARS-CoV-2 after the six-month global spread. Infection, Genetics and Evolution, 2021, 91, 104800.	2.3	5
26	Wide dissemination of SCC fusC in fusidic acid-resistant coagulase-negative staphylococci and implication for its spread to methicillin-resistant staphylococcus aureus in Taiwan. International Journal of Antimicrobial Agents, 2018, 51, 875-880.	2.5	2

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