

Norihiko Takemoto

List of Publications by Year in descending order

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15
papers

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933447

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365
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#	ARTICLE	IF	CITATIONS
1	Predominance of ST8 and CC1/spa-t1784 methicillin-resistant <i>Staphylococcus aureus</i> isolates in Japan and their genomic characteristics. <i>Journal of Global Antimicrobial Resistance</i> , 2022, 28, 195-202.	2.2	9
2	Comparative genome analysis of three Group A <i>Streptococcus dysgalactiae</i> subsp. <i>equisimilis</i> strains isolated in Japan. <i>Journal of Medical Microbiology</i> , 2021, 70, .	1.8	1
3	Novel Hyaluronate Lyase Involved in Pathogenicity of <i>Streptococcus dysgalactiae</i> subsp. <i>equisimilis</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 552418.	3.5	1
4	Lipopolysaccharide-Deficient <i>Acinetobacter baumannii</i> Due to Colistin Resistance Is Killed by Neutrophil-Produced Lysozyme. <i>Frontiers in Microbiology</i> , 2020, 11, 573.	3.5	26
5	Epidemiology of <i>Enterobacter cloacae</i> strains producing a carbapenemase or metallo-beta-lactamase in Vietnamese clinical settings in 2014–2017. <i>Journal of Medical Microbiology</i> , 2020, 69, 530-536.	1.8	4
6	Complete Genome Sequences of <i>Staphylococcus argenteus</i> TWCC 58113, Which Bears Two Plasmids. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	0
7	Bacterial EndoMS/NucS acts as a clamp-mediated mismatch endonuclease to prevent asymmetric accumulation of replication errors. <i>Nucleic Acids Research</i> , 2018, 46, 6152-6165.	14.5	33
8	Severe invasive streptococcal infection by <i>Streptococcus pyogenes</i> and <i>Streptococcus dysgalactiae</i> subsp. <i>equisimilis</i> . <i>Microbiology and Immunology</i> , 2016, 60, 1-9.	1.4	31
9	RNase III mediated cleavage of the coding region of <i>mraZ</i> mRNA is required for efficient cell division in <i>Corynebacterium glutamicum</i> . <i>Molecular Microbiology</i> , 2016, 99, 1149-1166.	2.5	21
10	Identification and expression analysis of a gene encoding a shikimate transporter of <i>Corynebacterium glutamicum</i> . <i>Microbiology (United Kingdom)</i> , 2015, 161, 254-263.	1.8	19
11	Rho and RNase play a central role in FMN riboswitch regulation in <i>Corynebacterium glutamicum</i> . <i>Nucleic Acids Research</i> , 2015, 43, 520-529.	14.5	37
12	The physiological role of riboflavin transporter and involvement of FMN-riboswitch in its gene expression in <i>Corynebacterium glutamicum</i> . <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 4159-4168.	3.6	12
13	Chorismate-dependent transcriptional regulation of quinate/shikimate utilization genes by <i>LysR</i> -type transcriptional regulator <i>QsuR</i> in <i>Corynebacterium glutamicum</i> : carbon flow control at metabolic branch point. <i>Molecular Microbiology</i> , 2014, 92, 356-368.	2.5	28
14	Genome-Wide Analysis of the Role of Global Transcriptional Regulator GntR1 in <i>Corynebacterium glutamicum</i> . <i>Journal of Bacteriology</i> , 2014, 196, 3249-3258.	2.2	10
15	High cGMP synthetic activity in carp cones. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 11788-11793.	7.1	55