

Soo-Jin Yang

List of Publications by Year in descending order

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46
papers

2,595
citations

257101

24
h-index

233125

45
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46
all docs

46
docs citations

46
times ranked

1959
citing authors

#	ARTICLE	IF	CITATIONS
1	Profiles of Non- <i>aureus</i> Staphylococci in Retail Pork and Slaughterhouse Carcasses: Prevalence, Antimicrobial Resistance, and Genetic Determinant of Fusidic Acid Resistance. <i>Food Science of Animal Resources</i> , 2022, 42, 225-239.	1.7	10
2	Co-occurrence of <i>cf</i> -mediated linezolid-resistance in ST398 LA-MRSA and non- <i>aureus</i> staphylococci isolated from a pig farm. <i>Veterinary Microbiology</i> , 2022, 266, 109336.	0.8	10
3	Multilocus sequence type-dependent activity of human and animal cathelicidins against community-, hospital-, and livestock-associated methicillin-resistant <i>Staphylococcus aureus</i> isolates. <i>Journal of Animal Science and Technology</i> , 2022, 64, 515-530.	0.8	2
4	Profiles of coagulase-positive and -negative staphylococci in retail pork: prevalence, antimicrobial resistance, enterotoxigenicity, and virulence factors. <i>Animal Bioscience</i> , 2021, 34, 734-742.	0.8	2
5	Profiles of coagulase-positive and -negative staphylococci in retail pork: prevalence, antimicrobial resistance, enterotoxigenicity, and virulence factors. <i>Animal Bioscience</i> , 2021, 34, 734-742.	0.8	3
6	Genomic Information on Linezolid-Resistant Sequence-Type 398 Livestock-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> Isolated from a Pig. <i>Foodborne Pathogens and Disease</i> , 2021, 18, 378-387.	0.8	4
7	Role of the <i>Staphylococcus aureus</i> Extracellular Loop of GraS in Resistance to Distinct Human Defense Peptides in PMN and Invasive Cardiovascular infections. <i>Infection and Immunity</i> , 2021, 89, e0034721.	1.0	5
8	Type I Interferons Are Involved in the Intracellular Growth Control of <i>Mycobacterium abscessus</i> by Mediating NOD2-Induced Production of Nitric Oxide in Macrophages. <i>Frontiers in Immunology</i> , 2021, 12, 738070.	2.2	9
9	Comparative assessment of genotypic and phenotypic correlates of <i>Staphylococcus pseudintermedius</i> strains isolated from dogs with otitis externa and healthy dogs. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2020, 70, 101376.	0.7	8
10	Genetic Factors Associated with Increased Host Defense Antimicrobial Peptide Resistance in Sequence Type 5 Healthcare-Associated MRSA Clinical Isolates. <i>Biomolecules</i> , 2020, 10, 1415.	1.8	0
11	Species Distribution, Antimicrobial Resistance, and Enterotoxigenicity of Non- <i>aureus</i> Staphylococci in Retail Chicken Meat. <i>Antibiotics</i> , 2020, 9, 809.	1.5	14
12	Livestock-associated methicillin-resistant <i>Staphylococcus aureus</i> in Korea: antimicrobial resistance and molecular characteristics of LA-MRSA strains isolated from pigs, pig farmers, and farm environment. <i>Journal of Veterinary Science</i> , 2020, 21, e2.	0.5	30
13	Occurrence and Characteristics of Methicillin-Resistant and -Susceptible <i>Staphylococcus aureus</i> Isolated from the Beef Production Chain in Korea. <i>Food Science of Animal Resources</i> , 2020, 40, 401-414.	1.7	11
14	Complete genome sequence of a methicillin-resistant <i>Staphylococcus schleiferi</i> strain from canine otitis externa in Korea. <i>Journal of Veterinary Science</i> , 2020, 21, e11.	0.5	3
15	Carriage of <i>Staphylococcus schleiferi</i> from canine otitis externa: antimicrobial resistance profiles and virulence factors associated with skin infection. <i>Journal of Veterinary Science</i> , 2019, 20, e6.	0.5	26
16	Potential role of host defense antimicrobial peptide resistance in increased virulence of health care-associated MRSA strains of sequence type (ST) 5 versus livestock-associated and community-associated MRSA strains of ST72. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2019, 62, 13-18.	0.7	12
17	Prevalence and characteristics of livestock-associated methicillin-susceptible <i>Staphylococcus aureus</i> in the pork production chain in Korea. <i>Journal of Veterinary Science</i> , 2019, 20, e69.	0.5	11
18	Impact of Multiple Single-Nucleotide Polymorphisms Within <i>mprF</i> on Daptomycin Resistance in <i>Staphylococcus aureus</i> . <i>Microbial Drug Resistance</i> , 2018, 24, 1075-1081.	0.9	16

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19	Emergence of Daptomycin-Nonsusceptible Methicillin-Resistant <i>Staphylococcus aureus</i> Clinical Isolates Among Daptomycin-Naive Patients in Korea. <i>Microbial Drug Resistance</i> , 2018, 24, 534-541.	0.9	5
20	Adaptations of Vancomycin-Intermediate Sequence Type 72 Methicillin-Resistant <i>Staphylococcus aureus</i> for Daptomycin Nonsusceptibility. <i>Microbial Drug Resistance</i> , 2018, 24, 1489-1496.	0.9	2
21	Phenotypic and genotypic correlates of daptomycin-resistant methicillin-susceptible <i>Staphylococcus aureus</i> clinical isolates. <i>Journal of Microbiology</i> , 2017, 55, 153-159.	1.3	34
22	Nucleotide-Binding Oligomerization Domain 2 Contributes to Limiting Growth of <i>Mycobacterium abscessus</i> in the Lung of Mice by Regulating Cytokines and Nitric Oxide Production. <i>Frontiers in Immunology</i> , 2017, 8, 1477.	2.2	28
23	Mechanisms of quinolone resistance in <i>Escherichia coli</i> isolated from companion animals, pet-owners, and non-pet-owners. <i>Journal of Veterinary Science</i> , 2017, 18, 449.	0.5	21
24	Isolation and characterization of antimicrobial-resistant <i>Escherichia coli</i> from national horse racetracks and private horse-riding courses in Korea. <i>Journal of Veterinary Science</i> , 2016, 17, 199.	0.5	15
25	Genotypic and Phenotypic Characterization of Methicillin-Resistant <i>Staphylococcus aureus</i> Isolated from Bovine Mastitic Milk in Korea. <i>Journal of Food Protection</i> , 2016, 79, 1725-1732.	0.8	28
26	Dysregulation of <i>mprF</i> and <i>dltABCD</i> expression among daptomycin-non-susceptible MRSA clinical isolates. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 2100-2104.	1.3	44
27	Antimicrobial resistance and virulence profiles of <i>Enterococcus</i> spp. isolated from horses in Korea. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2016, 48, 6-13.	0.7	17
28	The GraS Sensor in <i>Staphylococcus aureus</i> Mediates Resistance to Host Defense Peptides Differing in Mechanisms of Action. <i>Infection and Immunity</i> , 2016, 84, 459-466.	1.0	33
29	Frequency and Distribution of Single-Nucleotide Polymorphisms within <i>mprF</i> in Methicillin-Resistant <i>Staphylococcus aureus</i> Clinical Isolates and Their Role in Cross-Resistance to Daptomycin and Host Defense Antimicrobial Peptides. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 4930-4937.	1.4	102
30	Site-Specific Mutation of the Sensor Kinase GraS in <i>Staphylococcus aureus</i> Alters the Adaptive Response to Distinct Cationic Antimicrobial Peptides. <i>Infection and Immunity</i> , 2014, 82, 5336-5345.	1.0	41
31	Heterogeneity of <i>mprF</i> Sequences in Methicillin-Resistant <i>Staphylococcus aureus</i> Clinical Isolates: Role in Cross-Resistance between Daptomycin and Host Defense Antimicrobial Peptides. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 7462-7467.	1.4	59
32	Phenotypic and Genotypic Characterization of Daptomycin-Resistant Methicillin-Resistant <i>Staphylococcus aureus</i> Strains: Relative Roles of <i>mprF</i> and <i>dlt</i> Operons. <i>PLoS ONE</i> , 2014, 9, e107426.	1.1	105
33	Role of the LytSR Two-Component Regulatory System in Adaptation to Cationic Antimicrobial Peptides in <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 3875-3882.	1.4	46
34	Causal Role of Single Nucleotide Polymorphisms within the <i>mprF</i> Gene of <i>Staphylococcus aureus</i> in Daptomycin Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 5658-5664.	1.4	76
35	Increased Cell Wall Teichoic Acid Production and D-alanylation Are Common Phenotypes among Daptomycin-Resistant Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) Clinical Isolates. <i>PLoS ONE</i> , 2013, 8, e67398.	1.1	86
36	Emergence of Daptomycin Resistance in Daptomycin-Naïve Rabbits with Methicillin-Resistant <i>Staphylococcus aureus</i> Prosthetic Joint Infection Is Associated with Resistance to Host Defense Cationic Peptides and <i>mprF</i> Polymorphisms. <i>PLoS ONE</i> , 2013, 8, e71151.	1.1	76

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37	The <i>Staphylococcus aureus</i> Two-Component Regulatory System, GraRS, Senses and Confers Resistance to Selected Cationic Antimicrobial Peptides. <i>Infection and Immunity</i> , 2012, 80, 74-81.	1.0	159
38	Correlation of Daptomycin Resistance in a Clinical <i>Staphylococcus aureus</i> Strain with Increased Cell Wall Teichoic Acid Production and <i>scpD</i> -Alanylation. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 3922-3928.	1.4	117
39	Cell Wall Thickening Is Not a Universal Accompaniment of the Daptomycin Nonsusceptibility Phenotype in <i>Staphylococcus aureus</i> : Evidence for Multiple Resistance Mechanisms. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 3079-3085.	1.4	128
40	The Bacterial Defensin Resistance Protein MprF Consists of Separable Domains for Lipid Lysinylation and Antimicrobial Peptide Repulsion. <i>PLoS Pathogens</i> , 2009, 5, e1000660.	2.1	283
41	Enhanced Expression of <i>dltABCD</i> Is Associated with the Development of Daptomycin Nonsusceptibility in a Clinical Endocarditis Isolate of <i>Staphylococcus aureus</i> . <i>Journal of Infectious Diseases</i> , 2009, 200, 1916-1920.	1.9	147
42	Analysis of Cell Membrane Characteristics of In Vitro-Selected Daptomycin-Resistant Strains of Methicillin-Resistant <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 2312-2318.	1.4	210
43	Regulation of <i>mprF</i> in Daptomycin-Nonsusceptible <i>Staphylococcus aureus</i> Strains. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 2636-2637.	1.4	117
44	Failures in Clinical Treatment of <i>Staphylococcus aureus</i> Infection with Daptomycin Are Associated with Alterations in Surface Charge, Membrane Phospholipid Asymmetry, and Drug Binding. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 269-278.	1.4	305
45	The role of proton motive force in expression of the <i>Staphylococcus aureus</i> <i>cid</i> and <i>lrg</i> operons. <i>Molecular Microbiology</i> , 2006, 59, 1395-1404.	1.2	80
46	Antimicrobial resistance in <i>Salmonella enterica</i> serovars Enteritidis and Typhimurium isolated from animals in Korea: comparison of phenotypic and genotypic resistance characterization. <i>Veterinary Microbiology</i> , 2002, 86, 295-301.	0.8	55