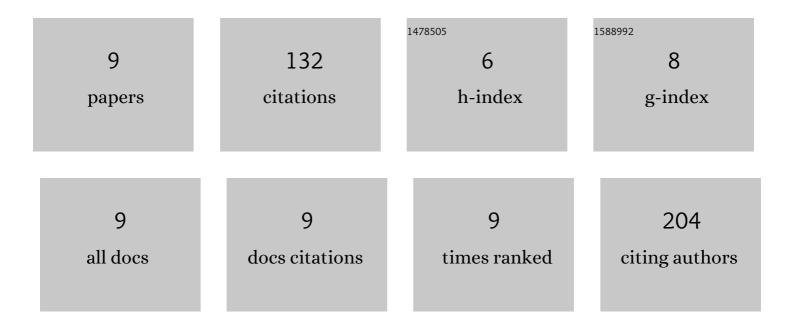
Xiangyu Yao

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Functional Characterization of Seven γ-Glutamylpolyamine Synthetase Genes and the <i>bauRABCD</i> Locus for Polyamine and β-Alanine Utilization in Pseudomonas aeruginosa PAO1. Journal of Bacteriology, 2011, 193, 3923-3930.	2.2	44
2	Regulation of the dauBAR operon and characterization of d-amino acid dehydrogenase DauA in arginine and lysine catabolism of Pseudomonas aeruginosa PAO1. Microbiology (United Kingdom), 2010, 156, 60-71.	1.8	27
3	Functional Characterization of the potRABCD Operon for Spermine and Spermidine Uptake and Regulation in Staphylococcus aureus. Current Microbiology, 2014, 69, 75-81.	2.2	20
4	A PBP 2 Mutant Devoid of the Transpeptidase Domain Abolishes Spermine–β-Lactam Synergy in Staphylococcus aureus Mu50. Antimicrobial Agents and Chemotherapy, 2012, 56, 83-91.	3.2	14
5	Characterization of Staphylococcus aureus Responses to Spermine Stress. Current Microbiology, 2014, 69, 394-403.	2.2	12
6	A Novel Pseudomonas aeruginosa Strain with an <i>oprD</i> Mutation in Relation to a Nosocomial Respiratory Infection Outbreak in an Intensive Care Unit. Journal of Clinical Microbiology, 2014, 52, 4388-4390.	3.9	7
7	Î ³ -Glutamyl Spermine Synthetase PauA2 as a Potential Target of Antibiotic Development against Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2012, 56, 5309-5314.	3.2	6
8	Transcriptomic data for analyzing global gene expression patterns in Methicillin-resistance Staphylococcus aureus in response to spermine and oxacillin stress. Data in Brief, 2018, 21, 2230-2236.	1.0	2
9	Spermine and oxacillin stress response on the cell wall synthesis and the global gene expression analysis in Methicillin-resistance Staphylococcus aureus. Genes and Genomics, 2019, 41, 43-59.	1.4	Ο