

# Zhengan Wang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9811181/publications.pdf>

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7  
papers

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citations

2258059

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1720034

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#	ARTICLE	IF	CITATIONS
1	Epidemiology, evolution and cryptic susceptibility of methicillin-resistant <i>Staphylococcus aureus</i> in China: a whole-genome-based survey. <i>Clinical Microbiology and Infection</i> , 2022, 28, 85-92.	6.0	31
2	Predominance of ST5-II-t311 clone among healthcare-associated methicillin-resistant <i>Staphylococcus aureus</i> isolates recovered from Zhejiang, China. <i>International Journal of Infectious Diseases</i> , 2018, 71, 107-112.	3.3	17
3	Household Transmission of Community-Associated Methicillin-Resistant <i>Staphylococcus Aureus</i> . <i>Frontiers in Public Health</i> , 2021, 9, 658638.	2.7	8
4	A random forest model based on core genome allelic profiles of MRSA for penicillin plus potassium clavulanate susceptibility prediction. <i>Microbial Genomics</i> , 2021, 7, .	2.0	3
5	The novel fosfomycin resistance gene <i>fosY</i> is present on a genomic island in CC1 methicillin-resistant <i>Staphylococcus aureus</i> . <i>Emerging Microbes and Infections</i> , 2022, 11, 1166-1173.	6.5	3
6	<i>Staphylococcal cassette chromosome mec</i> amplification as a mechanism for ceftobiprole resistance in clinical methicillin-resistant <i>Staphylococcus aureus</i> isolates. <i>Clinical Microbiology and Infection</i> , 2022, 28, 1151.e1-1151.e7.	6.0	3
7	Genetic Characteristics of Multiple Copies of Tn1546-Like Elements in <i>ermB</i> -Positive Methicillin-Resistant <i>Staphylococcus aureus</i> From Mainland China. <i>Frontiers in Microbiology</i> , 2022, 13, 814062.	3.5	1