

Teresa Nogueira

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10891098/publications.pdf>

Version: 2024-02-01

8
papers

378
citations

1307594

7
h-index

1588992

8
g-index

9
all docs

9
docs citations

9
times ranked

592
citing authors

#	ARTICLE	IF	CITATIONS
1	Horizontal Gene Transfer of the Secretome Drives the Evolution of Bacterial Cooperation and Virulence. <i>Current Biology</i> , 2009, 19, 1683-1691.	3.9	217
2	Antibiotic Resistance Gene Diversity and Virulence Gene Diversity Are Correlated in Human Gut and Environmental Microbiomes. <i>MSphere</i> , 2019, 4, .	2.9	46
3	Antibiotics as both friends and foes of the human gut microbiome: The microbial community approach. <i>Drug Development Research</i> , 2019, 80, 86-97.	2.9	43
4	Rapid Evolution of the Sequences and Gene Repertoires of Secreted Proteins in Bacteria. <i>PLoS ONE</i> , 2012, 7, e49403.	2.5	37
5	Are Virulence and Antibiotic Resistance Genes Linked? A Comprehensive Analysis of Bacterial Chromosomes and Plasmids. <i>Antibiotics</i> , 2022, 11, 706.	3.7	12
6	The Perfect Condition for the Rising of Superbugs: Person-to-Person Contact and Antibiotic Use Are the Key Factors Responsible for the Positive Correlation between Antibiotic Resistance Gene Diversity and Virulence Gene Diversity in Human Metagenomes. <i>Antibiotics</i> , 2021, 10, 605.	3.7	10
7	Metagenomics and Other Omics Approaches to Bacterial Communities and Antimicrobial Resistance Assessment in Aquacultures. <i>Antibiotics</i> , 2021, 10, 787.	3.7	7
8	Using SimulATe to model the effects of antibiotic selective pressure on the dynamics of pathogenic bacterial populations. <i>Biology Methods and Protocols</i> , 2019, 4, bpz004.	2.2	5