

Qi Yang

List of Publications by Year in descending order

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13
papers

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citations

1163117

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13
docs citations

13
times ranked

623
citing authors

#	ARTICLE	IF	CITATIONS
1	Factors affecting the isolation and diversity of marine sponge-associated bacteria. Applied Microbiology and Biotechnology, 2022, 106, 1729-1744.	3.6	4
2	Marine Sponge Endosymbionts: Structural and Functional Specificity of the Microbiome within <i>Euryspongia arenaria</i> Cells. Microbiology Spectrum, 2022, 10, e0229621.	3.0	5
3	Antimicrobial Activities of Marine Sponge-Associated Bacteria. Microorganisms, 2021, 9, 171.	3.6	17
4	Cultivation of fractionated cells from a bioactive-alkaloid-bearing marine sponge <i>Axinella</i> sp.. In Vitro Cellular and Developmental Biology - Animal, 2021, 57, 539-549.	1.5	2
5	Commercial cultivation, industrial application, and potential halocarbon biosynthesis pathway of <i>Asparagopsis</i> sp.. Algal Research, 2021, 56, 102319.	4.6	16
6	Spiroetherones A and B, sesquiterpene naphthoquinones, as angiogenesis inhibitors from the marine sponge <i>Dysidea etheria</i> . Organic Chemistry Frontiers, 2020, 7, 368-373.	4.5	12
7	Role of Dietary Nutrients in the Modulation of Gut Microbiota: A Narrative Review. Nutrients, 2020, 12, 381.	4.1	265
8	Untapped sponge microbiomes: structure specificity at host order and family levels. FEMS Microbiology Ecology, 2019, 95, .	2.7	14
9	Response of Sponge Microbiomes to Environmental Variations. , 2019, , 181-247.		4
10	Uncovering the hidden marine sponge microbiome by applying a multi-primer approach. Scientific Reports, 2019, 9, 6214.	3.3	12
11	A new deep-water <i>Tethya</i> (Porifera, Tethyida, Tethyidae) from the Great Australian Bight and an updated <i>Tethyida</i> phylogeny. European Journal of Taxonomy, 2019, , .	0.6	0
12	Development of a multilocus-based approach for sponge (phylum Porifera) identification: refinement and limitations. Scientific Reports, 2017, 7, 41422.	3.3	18
13	Sponge-associated actinobacterial diversity: validation of the methods of actinobacterial DNA extraction and optimization of 16S rRNA gene amplification. Applied Microbiology and Biotechnology, 2015, 99, 8731-8740.	3.6	12