Xiaoyu Tang

List of Publications by Year in descending order

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XIAOVU TANC

#	Article	IF	CITATIONS
1	Metatranscriptomics reveals different features of methanogenic archaea among global vegetated coastal ecosystems. Science of the Total Environment, 2022, 802, 149848.	8.0	10
2	Mining the Microbial Chemistry behind Tooth Decay. Biochemistry, 2022, 61, 2779-2781.	2.5	0
3	Catabolic protein degradation in marine sediments confined to distinct archaea. ISME Journal, 2022, 16, 1617-1626.	9.8	12
4	Contributions of Human-Associated Archaeal Metabolites to Tumor Microenvironment and Carcinogenesis. Microbiology Spectrum, 2022, 10, e0236721.	3.0	15
5	<i>mucG, mucH,</i> and <i>mucl</i> Modulate Production of Mutanocyclin and Reutericyclins in Streptococcus mutans B04Sm5. Journal of Bacteriology, 2022, 204, e0004222.	2.2	4
6	Identification and Biosynthesis of Pro-Inflammatory Sulfonolipids from an Opportunistic Pathogen <i>Chryseobacterium gleum</i> . ACS Chemical Biology, 2022, 17, 1197-1206.	3.4	12
7	Human Archaea and Associated Metabolites in Health and Disease. Biochemistry, 2022, 61, 2835-2840.	2.5	1
8	Grincamycins P–T: Rearranged Angucyclines from the Marine Sediment-Derived <i>Streptomyces</i> sp. CNZ-748 Inhibit Cell Lines of the Rare Cancer Pseudomyxoma Peritonei. Journal of Natural Products, 2021, 84, 1638-1648.	3.0	9
9	Cariogenic <i>Streptococcus mutans</i> Produces Tetramic Acid Strain-Specific Antibiotics That Impair Commensal Colonization. ACS Infectious Diseases, 2020, 6, 563-571.	3.8	40
10	Pass-back chain extension expands multimodular assembly line biosynthesis. Nature Chemical Biology, 2020, 16, 42-49.	8.0	28
11	Genetic platforms for heterologous expression of microbial natural products. Natural Product Reports, 2019, 36, 1313-1332.	10.3	109
12	Identification of the Bacterial Biosynthetic Gene Clusters of the Oral Microbiome Illuminates the Unexplored Social Language of Bacteria during Health and Disease. MBio, 2019, 10, .	4.1	73
13	<i>Klebsiella</i> and <i>Providencia</i> emerge as lone survivors following long-term starvation of oral microbiota. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 8499-8504.	7.1	30
14	Direct cloning and heterologous expression of natural product biosynthetic gene clusters by transformation-associated recombination. Methods in Enzymology, 2019, 621, 87-110.	1.0	37
15	Engineering Salinispora tropica for heterologous expression of natural product biosynthetic gene clusters. Applied Microbiology and Biotechnology, 2018, 102, 8437-8446.	3.6	24
16	Minimization of the Thiolactomycin Biosynthetic Pathway Reveals that the Cytochrome P450 Enzyme TlmF Is Required for Fiveâ€Membered Thiolactone Ring Formation. ChemBioChem, 2017, 18, 1072-1076.	2.6	18
17	Enzymatic Câ~'H Oxidation–Amidation Cascade in the Production of Natural and Unnatural Thiotetronate Antibiotics with Potentiated Bioactivity. Angewandte Chemie, 2017, 129, 12402-12407.	2.0	5
18	Broad-Host-Range Expression Reveals Native and Host Regulatory Elements That Influence Heterologous Antibiotic Production in Gram-Negative Bacteria. MBio, 2017, 8, .	4.1	39

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19	Enzymatic Câ^'H Oxidation–Amidation Cascade in the Production of Natural and Unnatural Thiotetronate Antibiotics with Potentiated Bioactivity. Angewandte Chemie - International Edition, 2017, 56, 12234-12239.	13.8	15
20	Identification of Thiotetronic Acid Antibiotic Biosynthetic Pathways by Target-directed Genome Mining. ACS Chemical Biology, 2015, 10, 2841-2849.	3.4	238
21	A two-step sulfation in antibiotic biosynthesis requires a type III polyketide synthase. Nature Chemical Biology, 2013, 9, 610-615.	8.0	36
22	Identification of a Napsamycin Biosynthesis Gene Cluster by Genome Mining. ChemBioChem, 2011, 12, 477-487.	2.6	44