Yin-Ru Chiang

List of Publications by Year in Descending Order

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Version: 2024-04-10

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

35 700 17 26 g-index

38 896 6.2 3.99 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
35	Temporal compositional shifts in an activated sludge microbiome during estrone biodegradation <i>Environmental Science and Pollution Research</i> , 2022 , 1	5.1	O
34	Causal networks of phytoplankton diversity and biomass are modulated by environmental context <i>Nature Communications</i> , 2022 , 13, 1140	17.4	2
33	Omics and mechanistic insights into di-(2-ethylhexyl) phthalate degradation in the O-fluctuating estuarine sediments <i>Chemosphere</i> , 2022 , 299, 134406	8.4	O
32	Bioactive pulvinones from a marine algicolous fungus Aspergillus terreus NTU243 <i>Phytochemistry</i> , 2022 , 200, 113229	4	1
31	Valorization of fish waste and sugarcane bagasse for Alcalase production by Bacillus megaterium via a circular bioeconomy model. <i>Journal of the Taiwan Institute of Chemical Engineers</i> , 2022 , 135, 10435	i8 ^{∙.3}	1
30	Mechanistic and phylogenetic insights into actinobacteria-mediated oestrogen biodegradation in urban estuarine sediments. <i>Microbial Biotechnology</i> , 2021 , 14, 1212-1227	6.3	4
29	Integrated Multi-omics Investigations Reveal the Key Role of Synergistic Microbial Networks in Removing Plasticizer Di-(2-Ethylhexyl) Phthalate from Estuarine Sediments. <i>MSystems</i> , 2021 , 6, e003582	27.6	4
28	Identification of essential Ebxidation genes and corresponding metabolites for oestrogen degradation by actinobacteria. <i>Microbial Biotechnology</i> , 2021 ,	6.3	2
27	Highly Oxygenated Constituents from a Marine Alga-Derived Fungus NTU967. <i>Marine Drugs</i> , 2020 , 18,	6	5
26	Anaerobic Biodegradation of Steroids 2020 , 165-195		
25	Microbial degradation of steroid sex hormones: implications for environmental and ecological studies. <i>Microbial Biotechnology</i> , 2020 , 13, 926-949	6.3	35
24	Retroconversion of estrogens into androgens by bacteria via a cobalamin-mediated methylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 1395-1403	11.5	21
23	Comparative Genome Analysis Reveals Cyanidiococcus gen. nov., A New Extremophilic Red Algal Genus Sister to Cyanidioschyzon (Cyanidioschyzonaceae, Rhodophyta). <i>Journal of Phycology</i> , 2020 , 56, 1428-1442	3	6
22	Long-term warming destabilizes aquatic ecosystems through weakening biodiversity-mediated causal networks. <i>Global Change Biology</i> , 2020 , 26, 6413-6423	11.4	10
21	Biosynthesis of Ascorbic Acid as a Glucose-Induced Photoprotective Process in the Extremophilic Red Alga. <i>Frontiers in Microbiology</i> , 2019 , 10, 3005	5.7	7
20	Metabolites Involved in Aerobic Degradation of the A and B Rings of Estrogen. <i>Applied and Environmental Microbiology</i> , 2019 , 85,	4.8	25
19	Estrogen Degraders and Estrogen Degradation Pathway Identified in an Activated Sludge. <i>Applied and Environmental Microbiology</i> , 2018 , 84,	4.8	40

18	Microbial Functional Responses to Cholesterol Catabolism in Denitrifying Sludge. <i>MSystems</i> , 2018 , 3,	7.6	11	
17	Biochemical Mechanisms and Catabolic Enzymes Involved in Bacterial Estrogen Degradation Pathways. <i>Cell Chemical Biology</i> , 2017 , 24, 712-724.e7	8.2	55	
16	Biochemical Mechanisms and Microorganisms Involved in Anaerobic Testosterone Metabolism in Estuarine Sediments. <i>Frontiers in Microbiology</i> , 2017 , 8, 1520	5.7	11	
15	Anaerobic Biodegradation of Steroids 2017 , 1-32		3	
14	Integrated multi-omics analyses reveal the biochemical mechanisms and phylogenetic relevance of anaerobic androgen biodegradation in the environment. <i>ISME Journal</i> , 2016 , 10, 1967-83	11.9	39	
13	Genomic Insight into the Host-Endosymbiont Relationship of Endozoicomonas montiporae CL-33(T) with its Coral Host. <i>Frontiers in Microbiology</i> , 2016 , 7, 251	5.7	49	
12	Identification of Comamonas testosteroni as an androgen degrader in sewage. <i>Scientific Reports</i> , 2016 , 6, 35386	4.9	26	
11	Substrate uptake and subcellular compartmentation of anoxic cholesterol catabolism in Sterolibacterium denitrificans. <i>Journal of Biological Chemistry</i> , 2015 , 290, 1155-69	5.4	26	
10	Anoxic androgen degradation by the denitrifying bacterium Sterolibacterium denitrificans via the 2,3-seco pathway. <i>Applied and Environmental Microbiology</i> , 2014 , 80, 3442-52	4.8	31	
9	Anaerobic and aerobic cleavage of the steroid core ring structure by Steroidobacter denitrificans. <i>Journal of Lipid Research</i> , 2013 , 54, 1493-504	6.3	43	
8	An oxygenase-independent cholesterol catabolic pathway operates under oxic conditions. <i>PLoS ONE</i> , 2013 , 8, e66675	3.7	21	
7	A novel testosterone catabolic pathway in bacteria. <i>Journal of Bacteriology</i> , 2011 , 193, 4447-55	3.5	27	
6	Initial steps in anoxic testosterone degradation by Steroidobacter denitrificans. <i>Microbiology</i> (<i>United Kingdom</i>), 2010 , 156, 2253-2259	2.9	25	
5	Cholest-4-en-3-one-delta 1-dehydrogenase, a flavoprotein catalyzing the second step in anoxic cholesterol metabolism. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 107-13	4.8	42	
4	Study of anoxic and oxic cholesterol metabolism by Sterolibacterium denitrificans. <i>Journal of Bacteriology</i> , 2008 , 190, 905-14	3.5	69	
3	Initial steps in the anoxic metabolism of cholesterol by the denitrifying Sterolibacterium denitrificans. <i>Journal of Biological Chemistry</i> , 2007 , 282, 13240-9	5.4	56	
2	Genome analysis of the steroid-degrading denitrifying Denitratisoma oestradiolicum DSM 16959 and Denitratisoma sp. strain DHT3		2	
1	Integrated multi-omics investigations reveal the key role of synergistic microbial networks in removing plasticizer di-(2-ethylhexyl) phthalate from estuarine sediments		1	