

Tim Lachnit

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3774784/publications.pdf>

Version: 2024-02-01

19
papers

1,446
citations

623188

14
h-index

794141

19
g-index

23
all docs

23
docs citations

23
times ranked

1554
citing authors

#	ARTICLE	IF	CITATIONS
1	Epibacterial community patterns on marine macroalgae are host-specific but temporally variable. <i>Environmental Microbiology</i> , 2011, 13, 655-665.	1.8	328
2	Specific epibacterial communities on macroalgae: phylogeny matters more than habitat. <i>Aquatic Biology</i> , 2009, 5, 181-186.	0.5	203
3	Metaorganisms in extreme environments: do microbes play a role in organismal adaptation?. <i>Zoology</i> , 2018, 127, 1-19.	0.6	194
4	Isolated thallus-associated compounds from the macroalga <i>Fucus vesiculosus</i> mediate bacterial surface colonization in the field similar to that on the natural alga. <i>Biofouling</i> , 2010, 26, 247-255.	0.8	116
5	Compounds associated with algal surfaces mediate epiphytic colonization of the marine macroalga <i>Fucus vesiculosus</i> . <i>FEMS Microbiology Ecology</i> , 2013, 84, 411-420.	1.3	111
6	A Phage Protein Aids Bacterial Symbionts in Eukaryote Immune Evasion. <i>Cell Host and Microbe</i> , 2019, 26, 542-550.e5.	5.1	94
7	Ecology of antifouling resistance in the bladder wrack <i>Fucus vesiculosus</i> : patterns of microfouling and antimicrobial protection. <i>Marine Ecology - Progress Series</i> , 2010, 411, 33-48.	0.9	91
8	Species-Specific Viromes in the Ancestral Holobiont Hydra. <i>PLoS ONE</i> , 2014, 9, e109952.	1.1	53
9	Expanding our Understanding of the Seaweed Holobiont: RNA Viruses of the Red Alga <i>Delisea pulchra</i> . <i>Frontiers in Microbiology</i> , 2015, 6, 1489.	1.5	49
10	rRNA-based profiling of bacteria in the axilla of healthy males suggests right-left asymmetry in bacterial activity. <i>FEMS Microbiology Ecology</i> , 2011, 77, 146-153.	1.3	39
11	Temperate phages as self-replicating weapons in bacterial competition. <i>Journal of the Royal Society Interface</i> , 2017, 14, 20170563.	1.5	39
12	Competing forces maintain the <i>Hydra</i> metaorganism. <i>Immunological Reviews</i> , 2017, 279, 123-136.	2.8	33
13	Microbial ecology in Hydra: Why viruses matter. <i>Journal of Microbiology</i> , 2015, 53, 193-200.	1.3	20
14	Exposure of the Host-Associated Microbiome to Nutrient-Rich Conditions May Lead to Dysbiosis and Disease Development—an Evolutionary Perspective. <i>MBio</i> , 2019, 10, .	1.8	19
15	Lifestyle of sponge symbiont phages by host prediction and correlative microscopy. <i>ISME Journal</i> , 2021, 15, 2001-2011.	4.4	19
16	Novel ssDNA Viruses Detected in the Virome of Bleached, Habitat-Forming Kelp <i>Ecklonia radiata</i> . <i>Frontiers in Marine Science</i> , 2018, 4, .	1.2	14
17	Contrasting distributions of bacteriophages and eukaryotic viruses from contaminated coastal sediments. <i>Environmental Microbiology</i> , 2019, 21, 1929-1941.	1.8	6
18	Can antibiotic-induced changes in the composition of the hair follicle microbiome regulate human hair growth?. <i>Experimental Dermatology</i> , 2021, 30, 1440-1441.	1.4	6

#	ARTICLE	IF	CITATIONS
19	<i>Hydra</i> and the hair follicle – An unconventional comparative biology approach to exploring the human holobiont. <i>BioEssays</i> , 2022, 44, e2100233.	1.2	4