

Laura Steindler

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2484821/publications.pdf>

Version: 2024-02-01

32
papers

2,228
citations

430754

18
h-index

395590

33
g-index

38
all docs

38
docs citations

38
times ranked

2645
citing authors

#	ARTICLE	IF	CITATIONS
1	Petrosia ficiformis (Poiret, 1789): an excellent model for holobiont and biotechnological studies. <i>Current Opinion in Biotechnology</i> , 2022, 74, 61-65.	3.3	6
2	Spatiotemporal Variation of Microbial Communities in the Ultra-Oligotrophic Eastern Mediterranean Sea. <i>Frontiers in Microbiology</i> , 2022, 13, 867694.	1.5	7
3	Lineage-specific energy and carbon metabolism of sponge symbionts and contributions to the host carbon pool. <i>ISME Journal</i> , 2022, 16, 1163-1175.	4.4	13
4	Draft Genome Sequence of Terrestrial <i>Streptomyces</i> sp. Strain VITNK9, Isolated from Vellore, Tamil Nadu, India, Exhibiting Antagonistic Activity against Fish Pathogens. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.3	2
5	Microbial rhodopsins are increasingly favoured over chlorophyll in High Nutrient Low Chlorophyll waters. <i>Environmental Microbiology Reports</i> , 2021, 13, 401-406.	1.0	11
6	Particle-associated and free-living bacterial communities in an oligotrophic sea are affected by different environmental factors. <i>Environmental Microbiology</i> , 2021, 23, 4295-4308.	1.8	35
7	Contribution of Maternal and Paternal Transmission to Bacterial Colonization in <i>Nematostella vectensis</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 726795.	1.5	11
8	Sponge microbiome stability during environmental acquisition of highly specific photosymbionts. <i>Environmental Microbiology</i> , 2020, 22, 3593-3607.	1.8	20
9	Characterization of sponge-associated <i>Verrucomicrobia</i> : microcompartment-based sugar utilization and enhanced toxin-antitoxin modules as features of host-associated <i>Opitutales</i> . <i>Environmental Microbiology</i> , 2020, 22, 4669-4688.	1.8	26
10	Identification of Quorum Sensing Activators and Inhibitors in The Marine Sponge <i>Sarcotragus spinosulus</i> . <i>Marine Drugs</i> , 2020, 18, 127.	2.2	17
11	Isolation, Genomic and Metabolomic Characterization of <i>Streptomyces tendae</i> VITAKN with Quorum Sensing Inhibitory Activity from Southern India. <i>Microorganisms</i> , 2020, 8, 121.	1.6	17
12	Genomic Insights Into the Lifestyles of Thaumarchaeota Inside Sponges. <i>Frontiers in Microbiology</i> , 2020, 11, 622824.	1.5	16
13	Life at Home and on the Roam: Genomic Adaptions Reflect the Dual Lifestyle of an Intracellular, Facultative Symbiont. <i>MSystems</i> , 2019, 4, .	1.7	30
14	Identification and chemical characterization of N-acyl-homoserine lactone quorum sensing signals across sponge species and time. <i>FEMS Microbiology Ecology</i> , 2018, 94, .	1.3	13
15	Pesticide-mediated trophic cascade and an ecological trap for mosquitoes. <i>Ecosphere</i> , 2018, 9, e02179.	1.0	17
16	Surface properties of SAR11 bacteria facilitate grazing avoidance. <i>Nature Microbiology</i> , 2017, 2, 1608-1615.	5.9	44
17	The sponge microbiome project. <i>GigaScience</i> , 2017, 6, 1-7.	3.3	193
18	Metagenomic analysis reveals unusually high incidence of proteorhodopsin genes in the ultraoligotrophic eastern Mediterranean Sea. <i>Environmental Microbiology</i> , 2017, 19, 1077-1090.	1.8	31

#	ARTICLE	IF	CITATIONS
19	Quorum Sensing Inhibitors from the Sea Discovered Using Bacterial N-acyl-homoserine Lactone-Based Biosensors. <i>Marine Drugs</i> , 2017, 15, 53.	2.2	68
20	Plakofuranolactone as a Quorum Quenching Agent from the Indonesian Sponge <i>Plakortis cf. lita</i> . <i>Marine Drugs</i> , 2017, 15, 59.	2.2	28
21	In Search of Alternative Antibiotic Drugs: Quorum-Quenching Activity in Sponges and their Bacterial Isolates. <i>Frontiers in Microbiology</i> , 2016, 7, 416.	1.5	66
22	Isolation of Marine <i>Paracoccus</i> sp. Ss63 from the Sponge <i>Sarcotragus</i> sp. and Characterization of its Quorum-Sensing Chemical-Signaling Molecules by LC-MS/MS Analysis. <i>Israel Journal of Chemistry</i> , 2016, 56, 330-340.	1.0	16
23	Diversity, structure and convergent evolution of the global sponge microbiome. <i>Nature Communications</i> , 2016, 7, 11870.	5.8	594
24	A New N -Acyl Homoserine Lactone Synthase in an Uncultured Symbiont of the Red Sea Sponge <i>Theonella swinhoei</i> . <i>Applied and Environmental Microbiology</i> , 2016, 82, 1274-1285.	1.4	30
25	Pyrosequencing analysis of aerobic anoxygenic phototrophic bacterial community structure in the oligotrophic western Pacific Ocean. <i>FEMS Microbiology Letters</i> , 2015, 362, fnv034.	0.7	14
26	Lifestyle Evolution in Cyanobacterial Symbionts of Sponges. <i>MBio</i> , 2015, 6, e00391-15.	1.8	103
27	Biogeography rather than association with cyanobacteria structures symbiotic microbial communities in the marine sponge <i>Petrosia ficiformis</i> . <i>Frontiers in Microbiology</i> , 2014, 5, 529.	1.5	68
28	Energy Starved Candidatus <i>Pelagibacter Ubique</i> Substitutes Light-Mediated ATP Production for Endogenous Carbon Respiration. <i>PLoS ONE</i> , 2011, 6, e19725.	1.1	190
29	Detection of quorum-sensing N-acyl homoserine lactone signal molecules by bacterial biosensors. <i>FEMS Microbiology Letters</i> , 2007, 266, 1-9.	0.7	349
30	Differential Gene Expression in a Marine Sponge in Relation to Its Symbiotic State. <i>Marine Biotechnology</i> , 2007, 9, 543-549.	1.1	33
31	Transmission, plasticity and the molecular identification of cyanobacterial symbionts in the Red Sea sponge <i>Diacarnus erythraenus</i> . <i>Marine Biology</i> , 2005, 148, 35-41.	0.7	50
32	16S rRNA Phylogeny of Sponge-Associated Cyanobacteria. <i>Applied and Environmental Microbiology</i> , 2005, 71, 4127-4131.	1.4	102