Dana L Carper

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4673789/publications.pdf

Version: 2024-02-01

13 papers	363 citations	9 h-index	1125743 13 g-index
14	14	14	545
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Evidence for foliar endophytic nitrogen fixation in a widely distributed subalpine conifer. New Phytologist, 2016, 210, 657-668.	7.3	135
2	amPEPpy 1.0: a portable and accurate antimicrobial peptide prediction tool. Bioinformatics, 2020, 37, 2058-2060.	4.1	43
3	Bacterial endophyte communities in Pinus flexilis are structured by host age, tissue type, and environmental factors. Plant and Soil, 2018, 428, 335-352.	3.7	32
4	Plant–Microbe Interactions: From Genes to Ecosystems Using <i>Populus</i> as a Model System. Phytobiomes Journal, 2021, 5, 29-38.	2.7	31
5	FAST: FAST Analysis of Sequences Toolbox. Frontiers in Genetics, 2015, 6, 172.	2.3	29
6	Subalpine conifers in different geographical locations host highly similar foliar bacterial endophyte communities. FEMS Microbiology Ecology, 2016, 92, fiw124.	2.7	23
7	Habitatâ€ndapted microbial communities mediate <i>Sphagnum</i> peatmoss resilience to warming. New Phytologist, 2022, 234, 2111-2125.	7.3	18
8	Cultivating the Bacterial Microbiota of <i>Populus</i> Roots. MSystems, 2021, 6, e0130620.	3.8	17
9	Formation, characterization and modeling of emergent synthetic microbial communities. Computational and Structural Biotechnology Journal, 2021, 19, 1917-1927.	4.1	12
10	Metaproteomics reveals insights into microbial structure, interactions, and dynamic regulation in defined communities as they respond to environmental disturbance. BMC Microbiology, 2021, 21, 308.	3.3	11
11	DISCo-microbe: design of an identifiable synthetic community of microbes. PeerJ, 2020, 8, e8534.	2.0	7
12	Development of an Experimental Approach to Achieve Spatially Resolved Plant Root-Associated Metaproteomics Using an Agar-Plate System. Molecular Plant-Microbe Interactions, 2022, 35, 639-649.	2.6	3
13	Draft Genome Sequence of <i>Tumebacillus</i> sp. Strain BK434, Isolated from the Roots of Eastern Cottonwood. Microbiology Resource Announcements, 2020, 9, .	0.6	1