Christophe Cd Djemiel

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	jvenn: an interactive Venn diagram viewer. BMC Bioinformatics, 2014, 15, 293.	2.6	1,528
2	Inferring microbiota functions from taxonomic genes: a review. GigaScience, 2022, 11, .	6.4	49
3	Nuclear genome sequence of the plastid-lacking cryptomonad Goniomonas avonlea provides insights into the evolution of secondary plastids. BMC Biology, 2018, 16, 137.	3.8	42
4	Characterization of Bacterial and Fungal Community Dynamics by High-Throughput Sequencing (HTS) Metabarcoding during Flax Dew-Retting. Frontiers in Microbiology, 2017, 8, 2052.	3.5	32
5	Multimodal assessment of flax dew retting and its functional impact on fibers and natural fiber composites. Industrial Crops and Products, 2020, 148, 112255.	5.2	25
6	µgreen-db: a reference database for the 23S rRNA gene of eukaryotic plastids and cyanobacteria. Scientific Reports, 2020, 10, 5915.	3.3	17
7	BIOCOM-PIPE: a new user-friendly metabarcoding pipeline for the characterization of microbial diversity from 16S, 18S and 23S rRNA gene amplicons. BMC Bioinformatics, 2020, 21, 492.	2.6	16
8	Potential of Meta-Omics to Provide Modern Microbial Indicators for Monitoring Soil Quality and Securing Food Production. Frontiers in Microbiology, 0, 13, .	3.5	14
9	Targeted Metagenomics of Retting in Flax: The Beginning of the Quest to Harness the Secret Powers of the Microbiota. Frontiers in Genetics, 2020, 11, 581664.	2.3	13
10	ReClustOR: a reâ€clustering tool using an openâ€reference method that improves operational taxonomic unit definition. Methods in Ecology and Evolution, 2020, 11, 168-180.	5.2	8
11	Microbial transfers from permanent grassland ecosystems to milk in dairy farms in the Comté cheese area. Scientific Reports, 2021, 11, 18144.	3.3	8