Andrs Mauricio Caraballo-Rodriguez

List of Publications by Citations

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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.

 38
 7,122
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 ext. citations
 avg, IF
 L-index

#	Paper	IF	Citations
38	Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. <i>Nature Biotechnology</i> , 2019 , 37, 852-857	44.5	4050
37	Sharing and community curation of mass spectrometry data with Global Natural Products Social Molecular Networking. <i>Nature Biotechnology</i> , 2016 , 34, 828-837	44.5	1566
36	Feature-based molecular networking in the GNPS analysis environment. <i>Nature Methods</i> , 2020 , 17, 905	- 9:0:8 6	207
35	Propagating annotations of molecular networks using in silico fragmentation. <i>PLoS Computational Biology</i> , 2018 , 14, e1006089	5	139
34	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science		138
33	Reproducible molecular networking of untargeted mass spectrometry data using GNPS. <i>Nature Protocols</i> , 2020 , 15, 1954-1991	18.8	125
32	MolNetEnhancer: Enhanced Molecular Networks by Integrating Metabolome Mining and Annotation Tools. <i>Metabolites</i> , 2019 , 9,	5.6	101
31	Chemical signaling involved in plant-microbe interactions. <i>Chemical Society Reviews</i> , 2018 , 47, 1652-170) 4 58.5	90
30	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science 2018 ,		78
29	Mass spectrometry searches using MASST. <i>Nature Biotechnology</i> , 2020 , 38, 23-26	44.5	74
28	The extracellular matrix protects Bacillus subtilis colonies from Pseudomonas invasion and modulates plant co-colonization. <i>Nature Communications</i> , 2019 , 10, 1919	17.4	59
27	A Convolutional Neural Network-Based Approach for the Rapid Annotation of Molecularly Diverse Natural Products. <i>Journal of the American Chemical Society</i> , 2020 , 142, 4114-4120	16.4	57
26	Natural products as mediators of disease. <i>Natural Product Reports</i> , 2017 , 34, 194-219	15.1	47
25	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science		36
24	Untargeted mass spectrometry-based metabolomics approach unveils molecular changes in raw and processed foods and beverages. <i>Food Chemistry</i> , 2020 , 302, 125290	8.5	34
23	Feature-based Molecular Networking in the GNPS Analysis Environment		29
22	Ion identity molecular networking for mass spectrometry-based metabolomics in the GNPS environment. <i>Nature Communications</i> , 2021 , 12, 3832	17.4	22

(2021-2016)

21	Endophytic Actinobacteria from the Brazilian Medicinal Plant Lychnophora ericoides Mart. and the Biological Potential of Their Secondary Metabolites. <i>Chemistry and Biodiversity</i> , 2016 , 13, 727-36	2.5	21
20	Expanding the Chemical Repertoire of the Endophyte Streptomyces albospinus RLe7 Reveals Amphotericin B as an Inducer of a Fungal Phenotype. <i>Journal of Natural Products</i> , 2017 , 80, 1302-1309	4.9	15
19	Molecular inter-kingdom interactions of endophytes isolated from Lychnophora ericoides. <i>Scientific Reports</i> , 2017 , 7, 5373	4.9	14
18	Ion Identity Molecular Networking in the GNPS Environment		11
17	MolNetEnhancer: enhanced molecular networks by integrating metabolome mining and annotation to	ols	10
16	Integrating genomics and metabolomics for scalable non-ribosomal peptide discovery. <i>Nature Communications</i> , 2021 , 12, 3225	17.4	8
15	Wildlife-microbiome interactions and disease: exploring opportunities for disease mitigation across ecological scales. <i>Drug Discovery Today: Disease Models</i> , 2018 , 28, 105-115	1.3	8
14	Reproducible Molecular Networking Of Untargeted Mass Spectrometry Data Using GNPS.		7
13	Virulence as a Side Effect of Interspecies Interaction in Coral Pathogens. MBio, 2020, 11,	7.8	7
12	Cryptic Species Account for the Seemingly Idiosyncratic Secondary Metabolism of Specimens Collected in Palau. <i>Journal of Natural Products</i> , 2020 , 83, 693-705	4.9	6
11	Amphotericin B as an inducer of griseofulvin-containing guttate in the endophytic fungus Xylaria cubensis FLe9. <i>Chemoecology</i> , 2017 , 27, 177-185	2	6
10	Protocol for community-created public MS/MS reference spectra within the Global Natural Products Social Molecular Networking infrastructure. <i>Rapid Communications in Mass Spectrometry</i> , 2020 , 34, e8725	2.2	5
9	Extracellular matrix components are required to protectBacillus subtilisfrom T6SS-dependentPseudomonasinvasion and modulate co-colonization of plants		3
8	Protocol for Community-created Public MS/MS Reference Library Within the GNPS Infrastructure		3
7	Nerpa: A Tool for Discovering Biosynthetic Gene Clusters of Bacterial Nonribosomal Peptides. <i>Metabolites</i> , 2021 , 11,	5.6	2
6	Metabolites from Microbes Isolated from the Skin of the Panamanian Rocket Frog (Anura: Dendrobatidae). <i>Metabolites</i> , 2020 , 10,	5.6	2
5	Chemical interplay and complementary adaptative strategies toggle bacterial antagonism and co-existence. <i>Cell Reports</i> , 2021 , 36, 109449	10.6	2
4	Chemical Proportionality within Molecular Networks. <i>Analytical Chemistry</i> , 2021 , 93, 12833-12839	7.8	2

Untargeted Metabolomics Sheds Light on the Diversity of Major Classes of Secondary Metabolites in the Malpighiaceae Botanical Family.. Frontiers in Plant Science, **2022**, 13, 854842

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2 Chemical interplay and complementary adaptative strategies toggle bacterial antagonism and co-existence

Chemical Gradients of Plant Substrates in an Fungus Garden. MSystems, **2021**, 6, e0060121

7.6