

Andrs Mauricio Caraballo-Rodriguez

List of Publications by Year in Descending Order

Source:
<https://exaly.com/author-pdf/238674/andres-mauricio-caraballo-rodriguez-publications-by-year.pdf>
Version: 2024-04-03

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.
The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

38 papers	7,122 citations	19 h-index	43 g-index
43 ext. papers	12,502 ext. citations	13.7 avg, IF	4.63 L-index

#	Paper	IF	Citations
38	Untargeted Metabolomics Sheds Light on the Diversity of Major Classes of Secondary Metabolites in the Malpighiaceae Botanical Family.. <i>Frontiers in Plant Science</i> , 2022 , 13, 854842	6.2	2
37	Nerpa: A Tool for Discovering Biosynthetic Gene Clusters of Bacterial Nonribosomal Peptides. <i>Metabolites</i> , 2021 , 11,	5.6	2
36	Integrating genomics and metabolomics for scalable non-ribosomal peptide discovery. <i>Nature Communications</i> , 2021 , 12, 3225	17.4	8
35	Ion identity molecular networking for mass spectrometry-based metabolomics in the GNPS environment. <i>Nature Communications</i> , 2021 , 12, 3832	17.4	22
34	Chemical interplay and complementary adaptative strategies toggle bacterial antagonism and co-existence. <i>Cell Reports</i> , 2021 , 36, 109449	10.6	2
33	Chemical Gradients of Plant Substrates in an Fungus Garden. <i>MSystems</i> , 2021 , 6, e0060121	7.6	0
32	Chemical Proportionality within Molecular Networks. <i>Analytical Chemistry</i> , 2021 , 93, 12833-12839	7.8	2
31	Reproducible molecular networking of untargeted mass spectrometry data using GNPS. <i>Nature Protocols</i> , 2020 , 15, 1954-1991	18.8	125
30	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
29	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
28	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
27	Mass spectrometry searches using MASST. <i>Nature Biotechnology</i> , 2020 , 38, 23-26	44.5	74
26	Metabolites from Microbes Isolated from the Skin of the Panamanian Rocket Frog (Anura: Dendrobatidae). <i>Metabolites</i> , 2020 , 10,	5.6	2
25	Virulence as a Side Effect of Interspecies Interaction in Coral Pathogens. <i>MBio</i> , 2020 , 11,	7.8	7
24	Feature-based molecular networking in the GNPS analysis environment. <i>Nature Methods</i> , 2020 , 17, 905-908	20.6	207
23	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
22	The extracellular matrix protects <i>Bacillus subtilis</i> colonies from <i>Pseudomonas</i> invasion and modulates plant co-colonization. <i>Nature Communications</i> , 2019 , 10, 1919	17.4	59

21	MolNetEnhancer: Enhanced Molecular Networks by Integrating Metabolome Mining and Annotation Tools. <i>Metabolites</i> , 2019 , 9,	5.6	101
20	Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. <i>Nature Biotechnology</i> , 2019 , 37, 852-857	44.5	4050
19	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science 2018 ,		78
18	Chemical signaling involved in plant-microbe interactions. <i>Chemical Society Reviews</i> , 2018 , 47, 1652-1704	58.5	90
17	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
16	Propagating annotations of molecular networks using in silico fragmentation. <i>PLoS Computational Biology</i> , 2018 , 14, e1006089	5	139
15	Expanding the Chemical Repertoire of the Endophyte <i>Streptomyces albospinus</i> RLe7 Reveals Amphotericin B as an Inducer of a Fungal Phenotype. <i>Journal of Natural Products</i> , 2017 , 80, 1302-1309	4.9	15
14	Amphotericin B as an inducer of griseofulvin-containing guttate in the endophytic fungus <i>Xylaria cubensis</i> FLe9. <i>Chemoecology</i> , 2017 , 27, 177-185	2	6
13	Molecular inter-kingdom interactions of endophytes isolated from <i>Lychnophora ericoides</i> . <i>Scientific Reports</i> , 2017 , 7, 5373	4.9	14
12	Natural products as mediators of disease. <i>Natural Product Reports</i> , 2017 , 34, 194-219	15.1	47
11	Endophytic Actinobacteria from the Brazilian Medicinal Plant <i>Lychnophora ericoides</i> Mart. and the Biological Potential of Their Secondary Metabolites. <i>Chemistry and Biodiversity</i> , 2016 , 13, 727-36	2.5	21
10	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
9	Reproducible Molecular Networking Of Untargeted Mass Spectrometry Data Using GNPS.		7
8	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science		36
7	QIIME 2: Reproducible, interactive, scalable, and extensible microbiome data science		138
6	Extracellular matrix components are required to protect <i>Bacillus subtilis</i> from T6SS-dependent <i>Pseudomonas</i> invasion and modulate co-colonization of plants		3
5	Ion Identity Molecular Networking in the GNPS Environment		11
4	MolNetEnhancer: enhanced molecular networks by integrating metabolome mining and annotation tools		10

3	Protocol for Community-created Public MS/MS Reference Library Within the GNPS Infrastructure	3
2	Feature-based Molecular Networking in the GNPS Analysis Environment	29
1	Chemical interplay and complementary adaptative strategies toggle bacterial antagonism and co-existence	1