

Carlos Molina-Santiago

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

1,258
citations

430874

18
h-index

377865

34
g-index

44
all docs

44
docs citations

44
times ranked

2000
citing authors

#	ARTICLE	IF	CITATIONS
1	GNPS Dashboard: collaborative exploration of mass spectrometry data in the web browser. <i>Nature Methods</i> , 2022, 19, 134-136.	19.0	35
2	Mechanisms of resistance to glyphosate: an example of bacterial adaptability to anthropogenic substances. <i>Environmental Microbiology</i> , 2022, 24, 3313-3315.	3.8	1
3	A Noninvasive Method for Time-Lapse Imaging of Microbial Interactions and Colony Dynamics. <i>Microbiology Spectrum</i> , 2022, 10, .	3.0	4
4	Bacterial extracellular matrix as a natural source of biotechnologically multivalent materials. <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 2796-2805.	4.1	10
5	A community resource for paired genomic and metabolomic data mining. <i>Nature Chemical Biology</i> , 2021, 17, 363-368.	8.0	81
6	Chemical interplay and complementary adaptative strategies toggle bacterial antagonism and co-existence. <i>Cell Reports</i> , 2021, 36, 109449.	6.4	28
7	Chemical Proportionality within Molecular Networks. <i>Analytical Chemistry</i> , 2021, 93, 12833-12839.	6.5	22
8	Untargeted mass spectrometry-based metabolomics approach unveils molecular changes in raw and processed foods and beverages. <i>Food Chemistry</i> , 2020, 302, 125290.	8.2	52
9	Chemical fertilization: a short-term solution for plant productivity?. <i>Microbial Biotechnology</i> , 2020, 13, 1311-1313.	4.2	11
10	More than words: the chemistry behind the interactions in the plant holobiont. <i>Environmental Microbiology</i> , 2020, 22, 4532-4544.	3.8	33
11	Dual functionality of the amyloid protein TasA in <i>Bacillus</i> physiology and fitness on the phylloplane. <i>Nature Communications</i> , 2020, 11, 1859.	12.8	59
12	Full Transcriptomic Response of <i>Pseudomonas aeruginosa</i> to an Inulin-Derived Fructooligosaccharide. <i>Frontiers in Microbiology</i> , 2020, 11, 202.	3.5	14
13	Understanding Bacterial Physiology for Improving Full Fitness. <i>Progress in Biological Control</i> , 2020, , 47-60.	0.5	1
14	Microbiomes as the new keystone for life sciences development. <i>Microbial Biotechnology</i> , 2019, 12, 579-581.	4.2	1
15	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.	12.8	102
16	Ruminal metagenomic libraries as a source of relevant hemicellulolytic enzymes for biofuel production. <i>Microbial Biotechnology</i> , 2018, 11, 781-787.	4.2	16
17	The race for antimicrobials in the multidrug resistance era. <i>Microbial Biotechnology</i> , 2018, 11, 976-978.	4.2	3
18	Insights in a novel gram-positive type IV secretion system. <i>Environmental Microbiology</i> , 2018, 20, 2334-2336.	3.8	0

#	ARTICLE	IF	CITATIONS
19	Interspecies cross-talk between co-cultured <i>Pseudomonas putida</i> and <i>Escherichia coli</i> . Environmental Microbiology Reports, 2017, 9, 441-448.	2.4	8
20	Global transcriptional response of solvent-sensitive and solvent-tolerant <i>Pseudomonas putida</i> strains exposed to toluene. Environmental Microbiology, 2017, 19, 645-658.	3.8	36
21	Understanding butanol tolerance and assimilation in <i>Pseudomonas putida</i> ... BIRD: an integrated omics approach. Microbial Biotechnology, 2016, 9, 100-115.	4.2	38
22	A <i>Pseudomonas putida</i> double mutant deficient in butanol assimilation: a promising step for engineering a biological biofuel production platform. FEMS Microbiology Letters, 2016, 363, fnw018.	1.8	16
23	<i>Pseudomonas putida</i> as a platform for the synthesis of aromatic compounds. Microbiology (United Kingdom), 2016, 156, 1-11.	1.8	41
24	Efflux pump-deficient mutants as a platform to search for microbes that produce antibiotics. Microbial Biotechnology, 2015, 8, 716-725.	4.2	9
25	Mechanisms of solvent resistance mediated by interplay of cellular factors in <i>Pseudomonas putida</i> . FEMS Microbiology Reviews, 2015, 39, 555-566.	8.6	143
26	Differential transcriptional response to antibiotics by <i>Pseudomonas putida</i> ... DOT-T1E. Environmental Microbiology, 2015, 17, 3251-3262.	3.8	32
27	Draft whole-genome sequence of the antibiotic-producing soil isolate <i>Pseudomonas</i> sp. strain 250. Environmental Microbiology Reports, 2015, 7, 288-292.	2.4	15
28	Diversity of small RNAs expressed in <i>Pseudomonas</i> species. Environmental Microbiology Reports, 2015, 7, 227-236.	2.4	27
29	Fructooligosaccharides Reduce <i>Pseudomonas aeruginosa</i> PAO1 Pathogenicity through Distinct Mechanisms. PLoS ONE, 2014, 9, e85772.	2.5	25
30	GtrS and GtrR form a two-component system: the central role of 2-ketogluconate in the expression of exotoxin A and glucose catabolic enzymes in <i>Pseudomonas aeruginosa</i> . Nucleic Acids Research, 2014, 42, 7654-7665.	14.5	41
31	Bactericidal and bacteriostatic antibiotics and the Fenton reaction. Microbial Biotechnology, 2014, 7, 194-195.	4.2	5
32	Identification of New Residues Involved in Intramolecular Signal Transmission in a Prokaryotic Transcriptional Repressor. Journal of Bacteriology, 2014, 196, 588-594.	2.2	6
33	Interspecies signalling: <i>Pseudomonas putida</i> efflux pump TtgGH is activated by indole to increase antibiotic resistance. Environmental Microbiology, 2014, 16, 1267-1281.	3.8	77
34	Antibiotic Resistance Determinants in a <i>Pseudomonas putida</i> Strain Isolated from a Hospital. PLoS ONE, 2014, 9, e81604.	2.5	86
35	Antibiotic adjuvants: identification and clinical use. Microbial Biotechnology, 2013, 6, 445-449.	4.2	76
36	Mechanisms of Resistance to Chloramphenicol in <i>Pseudomonas putida</i> KT2440. Antimicrobial Agents and Chemotherapy, 2012, 56, 1001-1009.	3.2	89

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37	Evolution of antibiotic resistance, catabolic pathways and niche colonization. <i>Microbial Biotechnology</i> , 2012, 5, 452-454.	4.2	0
38	Directed evolution, natural products for cancer chemotherapy, and micro-robotics. <i>Microbial Biotechnology</i> , 2011, 4, 314-317.	4.2	0