

Irene Sanchez-Andrea

List of Publications by Year in descending order

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Version: 2024-02-01

45
papers

2,140
citations

279487

23
h-index

243296

44
g-index

47
all docs

47
docs citations

47
times ranked

2459
citing authors

#	ARTICLE	IF	CITATIONS
1	Sulfate reduction at low pH to remediate acid mine drainage. <i>Journal of Hazardous Materials</i> , 2014, 269, 98-109.	6.5	288
2	Microbial Diversity in Anaerobic Sediments at Río Tinto, a Naturally Acidic Environment with a High Heavy Metal Content. <i>Applied and Environmental Microbiology</i> , 2011, 77, 6085-6093.	1.4	205
3	The reductive glycine pathway allows autotrophic growth of <i>Desulfovibrio desulfuricans</i> . <i>Nature Communications</i> , 2020, 11, 5090.	5.8	152
4	The bacterial sulfur cycle in expanding dysoxic and euxinic marine waters. <i>Environmental Microbiology</i> , 2021, 23, 2834-2857.	1.8	145
5	<i>Desulfosporosinus acididurans</i> sp. nov.: an acidophilic sulfate-reducing bacterium isolated from acidic sediments. <i>Extremophiles</i> , 2015, 19, 39-47.	0.9	128
6	Prospects for harnessing biocide resistance for bioremediation and detoxification. <i>Science</i> , 2018, 360, 743-746.	6.0	114
7	Towards sustainable feedstocks: A guide to electron donors for microbial carbon fixation. <i>Current Opinion in Biotechnology</i> , 2018, 50, 195-205.	3.3	80
8	Enrichment and isolation of acidophilic sulfate-reducing bacteria from Tinto River sediments. <i>Environmental Microbiology Reports</i> , 2013, 5, 672-678.	1.0	75
9	Quantification of Tinto River Sediment Microbial Communities: Importance of Sulfate-Reducing Bacteria and Their Role in Attenuating Acid Mine Drainage. <i>Applied and Environmental Microbiology</i> , 2012, 78, 4638-4645.	1.4	74
10	Isolation and genetic identification of PAH degrading bacteria from a microbial consortium. <i>Biodegradation</i> , 2009, 20, 789-800.	1.5	72
11	Sulfur Reduction in Acid Rock Drainage Environments. <i>Environmental Science & Technology</i> , 2015, 49, 11746-11755.	4.6	59
12	Insight into the sulfur metabolism of <i>Desulfurella amilsii</i> by differential proteomics. <i>Environmental Microbiology</i> , 2019, 21, 209-225.	1.8	57
13	Anaerobic Degradation of Sulfated Polysaccharides by Two Novel Kiritimatiellales Strains Isolated From Black Sea Sediment. <i>Frontiers in Microbiology</i> , 2019, 10, 253.	1.5	56
14	Bioremediation of acid mine drainage coupled with domestic wastewater treatment. <i>Water Science and Technology</i> , 2012, 66, 2425-2431.	1.2	48
15	<i>Desulfurella amilsii</i> sp. nov., a novel acidotolerant sulfur-respiring bacterium isolated from acidic river sediments. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 1249-1253.	0.8	47
16	Screening of anaerobic activities in sediments of an acidic environment: Tinto River. <i>Extremophiles</i> , 2012, 16, 829-839.	0.9	38
17	<i>Microbacter margulisiae</i> gen. nov., sp. nov., a propionigenic bacterium isolated from sediments of an acid rock drainage pond. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 3936-3942.	0.8	37
18	<i>Ercella succinigenes</i> gen. nov., sp. nov., an anaerobic succinate-producing bacterium. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 2449-2454.	0.8	36

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19	Genome Sequence of <i>Desulfurella amilsii</i> Strain TR1 and Comparative Genomics of Desulfurellaceae Family. <i>Frontiers in Microbiology</i> , 2017, 8, 222.	1.5	35
20	Redox Sensing within the Genus <i>Shewanella</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 2568.	1.5	32
21	<i>Pontiella desulfatans</i> gen. nov., sp. nov., and <i>Pontiella sulfatireligans</i> sp. nov., Two Marine Anaerobes of the Pontiaceae fam. nov. Producing Sulfated Glycosaminoglycan-like Exopolymers. <i>Microorganisms</i> , 2020, 8, 920.	1.6	31
22	From <i>R</i> hizosphere Tinto to Mars. <i>Advances in Applied Microbiology</i> , 2011, 77, 41-70.	1.3	28
23	Description of <i>Trichococcus ilyis</i> sp. nov. by combined physiological and in silico genome hybridization analyses. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 3957-3963.	0.8	27
24	Enrichment of sulfidogenic bacteria from the human intestinal tract. <i>FEMS Microbiology Letters</i> , 2017, 364, .	0.7	25
25	A case in support of implementing innovative bio-processes in the metal mining industry. <i>FEMS Microbiology Letters</i> , 2016, 363, fnw106.	0.7	23
26	Ecophysiology and Application of Acidophilic Sulfur-Reducing Microorganisms. <i>Grand Challenges in Biology and Biotechnology</i> , 2016, , 141-175.	2.4	22
27	Co-culture of a Novel Fermentative Bacterium, <i>Lucifera butyrlica</i> gen. nov. sp. nov., With the Sulfur Reducer <i>Desulfurella amilsii</i> for Enhanced Sulfidogenesis. <i>Frontiers in Microbiology</i> , 2018, 9, 3108.	1.5	22
28	Lysine and novel hydroxylysine lipids in soil bacteria: amino acid membrane lipid response to temperature and pH in <i>Pseudopedobacter saltans</i> . <i>Frontiers in Microbiology</i> , 2015, 6, 637.	1.5	21
29	Dissimilatory reduction of sulfate and zero-valent sulfur at low pH and its significance for bioremediation and metal recovery. <i>Advances in Microbial Physiology</i> , 2019, 75, 205-231.	1.0	20
30	Biosulfidogenesis Mediates Natural Attenuation in Acidic Mine Pit Lakes. <i>Microorganisms</i> , 2020, 8, 1275.	1.6	19
31	Microbial Geochemistry of the Acidic Saline Pit Lake of Brunita Mine (La Unión, <i>Á</i> SE Spain). <i>Mine Water and the Environment</i> , 2020, 39, 535-555.	0.9	18
32	Anaerobic microbial methanol conversion in marine sediments. <i>Environmental Microbiology</i> , 2021, 23, 1348-1362.	1.8	15
33	Organic Matter Type Defines the Composition of Active Microbial Communities Originating From Anoxic Baltic Sea Sediments. <i>Frontiers in Microbiology</i> , 2021, 12, 628301.	1.5	13
34	Bacterial glycerol oxidation coupled to sulfate reduction at neutral and acidic pH. <i>Journal of General and Applied Microbiology</i> , 2018, 64, 1-8.	0.4	11
35	An integrated green methodology for the continuous biological removal and fixation of arsenic from acid wastewater through the GAC-catalyzed As(III) oxidation. <i>Chemical Engineering Journal</i> , 2021, 421, 127758.	6.6	11
36	<i>Eubacterium maltosivorans</i> sp. nov., a novel human intestinal acetogenic and butyrogenic bacterium with a versatile metabolism. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3546-3550.	0.8	11

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37	Novel haloalkaliphilic methanotrophic bacteria: An attempt for enhancing methane bio-refinery. <i>Journal of Environmental Management</i> , 2019, 231, 1091-1099.	3.8	9
38	Comparative genomics and proteomics of <i>Eubacterium maltosivorans</i> : functional identification of trimethylamine methyltransferases and bacterial microcompartments in a human intestinal bacterium with a versatile lifestyle. <i>Environmental Microbiology</i> , 2022, 24, 517-534.	1.8	8
39	Effects of metals on activity and community of sulfate-reducing bacterial enrichments and the discovery of a new heavy metal-resistant SRB from Santos Port sediment (São Paulo, Brazil). <i>Environmental Science and Pollution Research</i> , 2022, 29, 922-935.	2.7	7
40	Sulfur Reduction at Hyperthermoacidophilic Conditions with Mesophilic Anaerobic Sludge as the Inoculum. <i>Environmental Science & Technology</i> , 2020, 54, 14656-14663.	4.6	6
41	Acetate Degradation at Low pH by the Moderately Acidophilic Sulfate Reducer <i>Acididesulfobacillus acetoxydans</i> gen. nov. sp. nov.. <i>Frontiers in Microbiology</i> , 2022, 13, 816605.	1.5	6
42	Acetotrophic sulfate-reducing consortia develop active biofilms on zeolite and glass beads in batch cultures at initial pH 3. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 5213-5227.	1.7	3
43	Microbial Communities in Peruvian Acid Mine Drainages: Low-Abundance Sulfate-Reducing Bacteria With High Metabolic Activity. <i>Geomicrobiology Journal</i> , 2022, 39, 867-883.	1.0	3
44	In search of sulfate-reducing consortia able to degrade acetate under acidic conditions. <i>Journal of Chemical Technology and Biotechnology</i> , 2021, 96, 1228-1236.	1.6	2
45	Editorial overview: Microbial environmental biotechnology. <i>Current Opinion in Biotechnology</i> , 2018, 50, vii-ix.	3.3	0