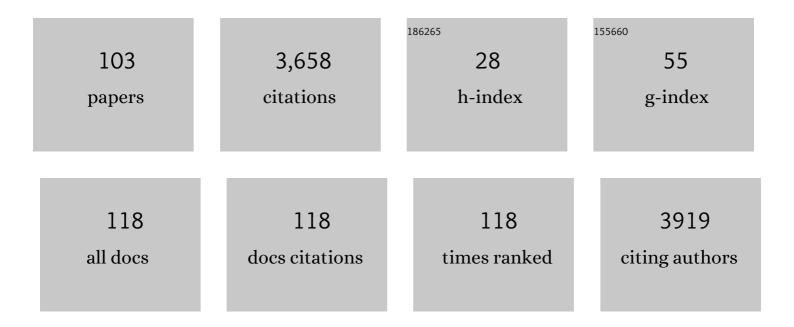
Maria Soledad Ramirez

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Characterization of the genetic structure of mcr-1 gene among Escherichia coli isolates recovered from surface waters and sediments from Ecuador. Science of the Total Environment, 2022, 806, 150566.	8.0	7
2	Amikacin potentiator activity of zinc complexed to a pyrithione derivative with enhanced solubility. Scientific Reports, 2022, 12, 285.	3.3	2
3	Human Serum Proteins and Susceptibility of Acinetobacter baumannii to Cefiderocol: Role of Iron Transport. Biomedicines, 2022, 10, 600.	3.2	8
4	Staphylococcus aureus Î \pm -Toxin Effect on Acinetobacter baumannii Behavior. Biology, 2022, 11, 570.	2.8	4
5	Acinetobacter baumannii response to cefiderocol challenge in human urine. Scientific Reports, 2022, 12, .	3.3	9
6	Light Modulates Important Pathogenic Determinants and Virulence in ESKAPE Pathogens Acinetobacter baumannii, Pseudomonas aeruginosa, and Staphylococcus aureus. Journal of Bacteriology, 2021, 203, .	2.2	16
7	Cerebrospinal fluid (CSF) augments metabolism and virulence expression factors in Acinetobacter baumannii. Scientific Reports, 2021, 11, 4737.	3.3	16
8	Zinc: Multidimensional Effects on Living Organisms. Biomedicines, 2021, 9, 208.	3.2	33
9	Staphylococcus aureus Potentiates the Hemolytic Activity of Burkholderia cepacia Complex (Bcc) Bacteria. Current Microbiology, 2021, 78, 1864-1870.	2.2	3
10	Full characterization of plasmids from Achromobacter ruhlandii isolates recovered from a single patient with cystic fibrosis (CF). Revista Argentina De Microbiologia, 2021, , .	0.7	0
11	Silencing Antibiotic Resistance with Antisense Oligonucleotides. Biomedicines, 2021, 9, 416.	3.2	13
12	Human Pleural Fluid and Human Serum Albumin Modulate the Behavior of a Hypervirulent and Multidrug-Resistant (MDR) Acinetobacter baumannii Representative Strain. Pathogens, 2021, 10, 471.	2.8	17
13	A New Twist: The Combination of Sulbactam/Avibactam Enhances Sulbactam Activity against Carbapenem-Resistant Acinetobacter baumannii (CRAB) Isolates. Antibiotics, 2021, 10, 577.	3.7	6
14	Interaction of Acinetobacter baumannii with Human Serum Albumin: Does the Host Determine the Outcome?. Antibiotics, 2021, 10, 833.	3.7	5
15	Involvement of the Histone-Like Nucleoid Structuring Protein (H-NS) in Acinetobacter baumannii's Natural Transformation. Pathogens, 2021, 10, 1083.	2.8	4
16	Inhibition of Aminoglycoside 6′-N-acetyltransferase Type Ib (AAC(6′)-Ib): Structure–Activity Relationship of Substituted Pyrrolidine Pentamine Derivatives as Inhibitors. Biomedicines, 2021, 9, 1218.	3.2	1
17	Effect of Serum Albumin, a Component of Human Pleural Fluid, on Transcriptional and Phenotypic Changes on Acinetobacter baumannii A118. Current Microbiology, 2021, 78, 3829-3834.	2.2	2
18	Histone-like nucleoid-structuring protein (H-NS) regulatory role in antibiotic resistance in Acinetobacter baumannii. Scientific Reports, 2021, 11, 18414.	3.3	8

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19	Amikacin in combination with zinc pyrithione prevents growth of a multidrug-resistant carbapenem-resistant Klebsiella pneumoniae isolate. International Journal of Antimicrobial Agents, 2021, 58, 106442.	2.5	4
20	Interplay between Meropenem and Human Serum Albumin on Expression of Carbapenem Resistance Genes and Natural Competence in Acinetobacter baumannii. Antimicrobial Agents and Chemotherapy, 2021, 65, e0101921.	3.2	10
21	An Acinetobacter non-baumannii Population Study: Antimicrobial Resistance Genes (ARGs). Antibiotics, 2021, 10, 16.	3.7	14
22	Aminoglycoside 6′-N-acetyltransferase Type Ib [AAC(6′)-Ib]-Mediated Aminoglycoside Resistance: Phenotypic Conversion to Susceptibility by Silver Ions. Antibiotics, 2021, 10, 29.	3.7	9
23	Diversity of Achromobacter species recovered from patients with cystic fibrosis, in Argentina. Revista Argentina De Microbiologia, 2020, 52, 13-18.	0.7	24
24	Genomic Analysis of two NDM-1 Providencia stuartii Strains Recovered from a Single Patient. Current Microbiology, 2020, 77, 4029-4036.	2.2	5
25	Characterisation of ST25 NDM-1-producing Acinetobacter spp. strains leading the increase in NDM-1 emergence in Argentina. Journal of Global Antimicrobial Resistance, 2020, 23, 108-110.	2.2	9
26	Functional Analysis of the Acinetobacter baumannii XerC and XerD Site-Specific Recombinases: Potential Role in Dissemination of Resistance Genes. Antibiotics, 2020, 9, 405.	3.7	19
27	Genomic analysis of two Acinetobacter baumannii strains belonging to two different sequence types (ST172 and ST25). Journal of Global Antimicrobial Resistance, 2020, 23, 154-161.	2.2	6
28	The H-NS Regulator Plays a Role in the Stress Induced by Carbapenemase Expression in Acinetobacter baumannii. MSphere, 2020, 5, .	2.9	10
29	Carbapenemases: Transforming Acinetobacter baumannii into a Yet More Dangerous Menace. Biomolecules, 2020, 10, 720.	4.0	124
30	Crucial Role of the Accessory Genome in the Evolutionary Trajectory of Acinetobacter baumannii Global Clone 1. Frontiers in Microbiology, 2020, 11, 342.	3.5	16
31	Distinct Mechanisms of Dissemination of NDM-1 Metallo-β-Lactamase in <i>Acinetobacter</i> Species in Argentina. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	25
32	Human Pleural Fluid Elicits Pyruvate and Phenylalanine Metabolism in Acinetobacter baumannii to Enhance Cytotoxicity and Immune Evasion. Frontiers in Microbiology, 2019, 10, 1581.	3.5	30
33	Identification of Potential Virulence Factors in the Model Strain Acinetobacter baumannii A118. Frontiers in Microbiology, 2019, 10, 1599.	3.5	28
34	Disruption of hmgA by DNA Duplication is Responsible for Hyperpigmentation in a Vibrio anguillarum Strain. Scientific Reports, 2019, 9, 14589.	3.3	2
35	Small Klebsiella pneumoniae Plasmids: Neglected Contributors to Antibiotic Resistance. Frontiers in Microbiology, 2019, 10, 2182.	3.5	23
36	Genetic and Phenotypic Features of a Novel Acinetobacter Species, Strain A47, Isolated From the Clinical Setting. Frontiers in Microbiology, 2019, 10, 1375.	3.5	4

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37	Restoration of susceptibility to amikacin by 8-hydroxyquinoline analogs complexed to zinc. PLoS ONE, 2019, 14, e0217602.	2.5	18
38	ICE SXT vs. ICESh95: Co-existence of Integrative and Conjugative Elements and Competition for a New Host. Scientific Reports, 2019, 9, 8045.	3.3	3
39	A Study on Acinetobacter baumannii and Staphylococcus aureus Strains Recovered from the Same Infection Site of a Diabetic Patient. Current Microbiology, 2019, 76, 842-847.	2.2	27
40	Whole Genome Sequence Analysis of Burkholderia contaminans FFH2055 Strain Reveals the Presence of Putative β-Lactamases. Current Microbiology, 2019, 76, 485-494.	2.2	2
41	Human pleural fluid triggers global changes in the transcriptional landscape of Acinetobacter baumannii as an adaptive response to stress. Scientific Reports, 2019, 9, 17251.	3.3	27
42	Interspecies DNA acquisition by a naturally competent Acinetobacter baumannii strain. International Journal of Antimicrobial Agents, 2019, 53, 483-490.	2.5	14
43	Human fluids alter DNA-acquisition in Acinetobacter baumannii. Diagnostic Microbiology and Infectious Disease, 2019, 93, 183-187.	1.8	20
44	Effect of Host Human Products on Natural Transformation in Acinetobacter baumannii. Current Microbiology, 2019, 76, 950-953.	2.2	21
45	Characterisation of OXA-258 enzymes and AxyABM efflux pump in Achromobacter ruhlandii. Journal of Global Antimicrobial Resistance, 2018, 14, 233-237.	2.2	7
46	Whole-Genome Analysis of an Extensively Drug-Resistance Empedobacter falsenii Strain Reveals Distinct Features and the Presence of a Novel Metallo-ß-Lactamase (EBR-2). Current Microbiology, 2018, 75, 1084-1089.	2.2	6
47	The effect of sub-inhibitory concentrations of antibiotics on natural transformation in Acinetobacter baumannii. International Journal of Antimicrobial Agents, 2018, 51, 809-810.	2.5	16
48	Genome sequence analysis of an extensively drug-resistant Acinetobacter baumannii indigo-pigmented strain depicts evidence of increase genome plasticity. Scientific Reports, 2018, 8, 16961.	3.3	28
49	Human serum albumin alters specific genes that can play a role in survival and persistence in Acinetobacter baumannii. Scientific Reports, 2018, 8, 14741.	3.3	47
50	Whole-genome analysis and description of an outbreak due to carbapenem-resistant Ochrobactrum anthropi causing pseudo-bacteraemias. New Microbes and New Infections, 2018, 26, 100-106.	1.6	3
51	Clinical cases of VIM-producing Pseudomonas mendocina from two burned patients. Journal of Global Antimicrobial Resistance, 2018, 14, 273-274.	2.2	14
52	Genomics helps to decipher the resistance mechanisms present in a Pseudomonas chlororaphis strain recovered in an HIV patient. New Microbes and New Infections, 2018, 25, 45-47.	1.6	7
53	Comparison between disk diffusion and agar dilution methods to determine in vitro susceptibility of Corynebacterium spp. clinical isolates and update of their susceptibility. Journal of Global Antimicrobial Resistance, 2018, 14, 246-252.	2.2	20
54	Antimicrobial susceptibility of clinical isolates of Actinomyces and related genera reveals an unusual clindamycin resistance among Actinomyces urogenitalis strains. Journal of Global Antimicrobial Resistance, 2017, 8, 115-120.	2.2	31

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55	Light Modulates Metabolic Pathways and Other Novel Physiological Traits in the Human Pathogen Acinetobacter baumannii. Journal of Bacteriology, 2017, 199, .	2.2	49
56	Genetic analysis of a PER-2-producing Shewanella sp. strain harbouring a variety of mobile genetic elements and antibiotic resistance determinants. Journal of Global Antimicrobial Resistance, 2017, 11, 81-86.	2.2	10
57	Amikacin: Uses, Resistance, and Prospects for Inhibition. Molecules, 2017, 22, 2267.	3.8	156
58	ISCR2 and IS26: Two Insertion Sequences Highly Dispersed among Acinetobacter spp. Clinical Strainss. Journal of Bacteriology & Mycology Open Access, 2017, 4, .	0.2	2
59	Matrix-assisted Laser Desorption Ionization-Time-of-Flight Mass Spectrometry (MALDI-TOF MS) as a Reliable Tool to Identify Species of Catalase-negative Gram-positive Cocci not Belonging to the Streptococcus Genus. Open Microbiology Journal, 2016, 10, 202-208.	0.7	9
60	The Genetic Analysis of an Acinetobacter johnsonii Clinical Strain Evidenced the Presence of Horizontal Genetic Transfer. PLoS ONE, 2016, 11, e0161528.	2.5	35
61	Whole-Genome Comparative Analysis of Two Carbapenem-Resistant ST-258Klebsiella pneumoniaeStrains Isolated during a North-Eastern Ohio Outbreak: Differences within the High Heterogeneity Zones. Genome Biology and Evolution, 2016, 8, 2036-2043.	2.5	28
62	Serum Albumin and Ca ²⁺ Are Natural Competence Inducers in the Human Pathogen Acinetobacter baumannii. Antimicrobial Agents and Chemotherapy, 2016, 60, 4920-4929.	3.2	58
63	Analysis of class 2 integrons as a marker for multidrug resistance among Gram negative bacilli. AIMS Genetics, 2016, 03, 196-204.	1.9	0
64	More Than Just Light: Clinical Relevance of Light Perception in the Nosocomial Pathogen <i>Acinetobacter baumannii</i> and Other Members of the Genus <i>Acinetobacter</i> . Photochemistry and Photobiology, 2015, 91, 1291-1301.	2.5	38
65	Draft Genome of the Multidrug-Resistant Acinetobacter baumannii Strain A155 Clinical Isolate. Genome Announcements, 2015, 3, .	0.8	21
66	Inhibition of Aminoglycoside 6′-N-Acetyltransferase Type Ib-Mediated Amikacin Resistance in Klebsiella pneumoniae by Zinc and Copper Pyrithione. Antimicrobial Agents and Chemotherapy, 2015, 59, 5851-5853.	3.2	34
67	White and blue light induce reduction in susceptibility to minocycline and tigecycline in Acinetobacter spp. and other bacteria of clinical importance. Journal of Medical Microbiology, 2015, 64, 525-537.	1.8	46
68	Evaluation of matrix-assisted laser desorption ionization-time-of-flight mass spectrometry for species identification of Nonfermenting Gram-Negative Bacilli. Journal of Microbiological Methods, 2015, 112, 24-27.	1.6	37
69	Draft Genome Sequence of a Taxonomically Unique Acinetobacter Clinical Strain with Proteolytic and Hemolytic Activities. Genome Announcements, 2015, 3, .	0.8	10
70	Draft Genome Sequence of Empedobacter (Formerly Wautersiella) falsenii comb. nov. Wf282, a Strain Isolated from a Cervical Neck Abscess. Genome Announcements, 2015, 3, .	0.8	8
71	A Taxonomically Unique Acinetobacter Strain with Proteolytic and Hemolytic Activities Recovered from a Patient with a Soft Tissue Injury. Journal of Clinical Microbiology, 2015, 53, 349-351.	3.9	13
72	Comparison of the Bruker MALDI-TOF Mass Spectrometry System and Conventional Phenotypic Methods for Identification of Gram-Positive Rods. PLoS ONE, 2014, 9, e106303.	2.5	77

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73	Bacteremia caused by an Acinetobacter junii strain harboring class 1 integron and diverse DNA mobile elements. Journal of Infection in Developing Countries, 2014, 8, 666-669.	1.2	6
74	Draft Genome Sequence of an International Clonal Lineage 1 Acinetobacter baumannii Strain from Argentina. Genome Announcements, 2014, 2, .	0.8	7
75	Whole-Genome Sequence Analysis of the Naturally Competent Acinetobacter baumannii Clinical Isolate A118. Genome Biology and Evolution, 2014, 6, 2235-2239.	2.5	85
76	Inhibition of Aminoglycoside 6′- <i>N</i> -Acetyltransferase Type Ib by Zinc: Reversal of Amikacin Resistance in Acinetobacter baumannii and Escherichia coli by a Zinc Ionophore. Antimicrobial Agents and Chemotherapy, 2014, 58, 4238-4241.	3.2	43
77	Presence of OXA-Type Enzymes in Achromobacter insuavis and A. dolens. Current Microbiology, 2014, 69, 501-506.	2.2	11
78	Plasmid-Mediated Antibiotic Resistance and Virulence in Gram-Negatives: the <i>Klebsiella pneumoniae</i> Paradigm. Microbiology Spectrum, 2014, 2, 1-15.	3.0	93
79	Draft Genome Sequence of an Extensively Drug-Resistant Acinetobacter baumannii Indigo-Pigmented Strain. Genome Announcements, 2014, 2, .	0.8	6
80	Distribution of Allelic Variants of the Chromosomal Gene bla OXA-114-like in Achromobacter xylosoxidans Clinical Isolates. Current Microbiology, 2013, 67, 596-600.	2.2	7
81	Outbreak of Extensively Drug-Resistant Acinetobacter baumannii Indigo-Pigmented Strains. Journal of Clinical Microbiology, 2013, 51, 3726-3730.	3.9	21
82	Acinetobacter baumannii extensively drug resistant lineages in Buenos Aires hospitals differ from the international clones l–III. Infection, Genetics and Evolution, 2013, 14, 294-301.	2.3	74
83	Spreading of AbaR-type Genomic Islands in Multidrug Resistance Acinetobacter baumannii Strains Belonging to Different Clonal Complexes. Current Microbiology, 2013, 67, 9-14.	2.2	53
84	Genome Sequences of Two Klebsiella pneumoniae Isolates from Different Geographical Regions, Argentina (Strain JHCK1) and the United States (Strain VA360). Genome Announcements, 2013, 1, .	0.8	13
85	Emergence and Spread of Plasmid-Borne <i>tet</i> (B)::IS <i>CR2</i> in Minocycline-Resistant Acinetobacter baumannii Isolates. Antimicrobial Agents and Chemotherapy, 2013, 57, 651-654.	3.2	44
86	OXA-258 from Achromobacter ruhlandii: a Species-Specific Marker. Journal of Clinical Microbiology, 2013, 51, 1602-1605.	3.9	14
87	Rise and dissemination of aminoglycoside resistance: the aac(6′)-Ib paradigm. Frontiers in Microbiology, 2013, 4, 121.	3.5	133
88	Achromobacter xylosoxidans: An Emerging Pathogen Carrying Different Elements Involved in Horizontal Genetic Transfer. Current Microbiology, 2012, 65, 673-678.	2.2	41
89	Transposons and integrons in colistin-resistant clones of Klebsiella pneumoniae and Acinetobacter baumannii with epidemic or sporadic behaviour. Journal of Medical Microbiology, 2012, 61, 1417-1420.	1.8	52
90	Acinetobacter baumannii is Able to Gain and Maintain a Plasmid Harbouring In35 Found in Enterobacteriaceae Isolates From Argentina. Current Microbiology, 2012, 64, 211-213.	2.2	18

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91	Class 2 Integrons Dissemination Among Multidrug Resistance (MDR) Clones of Acinetobacter baumannii. Current Microbiology, 2012, 64, 290-293.	2.2	11
92	Increasing frequency of class 1 and 2 integrons in multidrug-resistant clones of Acinetobacter baumannii reveals the need for continuous molecular surveillance. International Journal of Antimicrobial Agents, 2011, 37, 175-177.	2.5	11
93	Genomic Analysis ofAcinetobacter baumanniiA118 by Comparison of Optical Maps: Identification of Structures Related to Its Susceptibility Phenotype. Antimicrobial Agents and Chemotherapy, 2011, 55, 1520-1526.	3.2	48
94	Novel Insights about Class 2 Integrons from Experimental and Genomic Epidemiology. Antimicrobial Agents and Chemotherapy, 2010, 54, 699-706.	3.2	93
95	Naturally Competent <i>Acinetobacter baumannii</i> Clinical Isolate as a Convenient Model for Genetic Studies. Journal of Clinical Microbiology, 2010, 48, 1488-1490.	3.9	95
96	Reservoir of Antimicrobial Resistance Determinants Associated with Horizontal Gene Transfer in Clinical Isolates of the Genus <i>Shewanella</i> . Antimicrobial Agents and Chemotherapy, 2010, 54, 4516-4517.	3.2	16
97	Aminoglycoside modifying enzymes. Drug Resistance Updates, 2010, 13, 151-171.	14.4	1,007
98	Functional characterization of Tn1331 gene cassettes. Journal of Antimicrobial Chemotherapy, 2008,		96
	62, 669-673.	3.0	26
99	62, 669-673. Polyclonal spread of blaOXA-23 and blaOXA-58 in Acinetobacter baumannii isolates from Argentina. Journal of Infection in Developing Countries, 2008, 2, 235-40.	3.0 1.2	45
99 100	Polyclonal spread of blaOXA-23 and blaOXA-58 in Acinetobacter baumannii isolates from Argentina.		
	Polyclonal spread of blaOXA-23 and blaOXA-58 in Acinetobacter baumannii isolates from Argentina. Journal of Infection in Developing Countries, 2008, 2, 235-40. Novel Rearrangement of a Class 2 Integron in Two Non-Epidemiologically Related Isolates of	1.2	45
100	Polyclonal spread of blaOXA-23 and blaOXA-58 in Acinetobacter baumannii isolates from Argentina. Journal of Infection in Developing Countries, 2008, 2, 235-40. Novel Rearrangement of a Class 2 Integron in Two Non-Epidemiologically Related Isolates of Acinetobacter baumannii. Antimicrobial Agents and Chemotherapy, 2005, 49, 5179-5181. Class 2 Integron with a Novel Cassette Array in a Burkholderia cenocepacia Isolate. Antimicrobial	1.2 3.2	45 63