Bruno Gomez-Gil

List of Publications by Year
in descending order

130	4,658	109321	114465
papers	citations	35 h-index	63 g-index

134	134	134	3723
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The use and selection of probiotic bacteria for use in the culture of larval aquatic organisms. Aquaculture, 2000, 191, 259-270.	3.5	354
2	Field and Experimental Evidence of Vibrio parahaemolyticus as the Causative Agent of Acute Hepatopancreatic Necrosis Disease of Cultured Shrimp (Litopenaeus vannamei) in Northwestern Mexico. Applied and Environmental Microbiology, 2015, 81, 1689-1699.	3.1	274
3	Updating the Vibrio clades defined by multilocus sequence phylogeny: proposal of eight new clades, and the description of Vibrio tritonius sp. nov.. Frontiers in Microbiology, 2013, 4, 414.	3.5	264
4	Molecular identification of Vibrio harveyi-related isolates associated with diseased aquatic organisms. Microbiology (United Kingdom), 2004, 150, 1769-1777.	1.8	180
5	Comparative genomic analyses identify the Vibrio harveyi genome sequenced strains BAAâ€1116 and HY01 as Vibrio campbellii. Environmental Microbiology Reports, 2010, 2, 81-89.	2.4	153
6	TheVibriocore group induces yellow band disease in Caribbean and Indo-Pacific reef-building corals. Journal of Applied Microbiology, 2008, 105, 1658-1671.	3.1	150
7	Vibrios Associated with Litopenaeus vannamei Larvae, Postlarvae, Broodstock, and Hatchery Probionts. Applied and Environmental Microbiology, 1999, 65, 2592-2597.	3.1	141
8	Species of Vibrio isolated from hepatopancreas, haemolymph and digestive tract of a population of healthy juvenile Penaeus vannamei. Aquaculture, 1998, 163, 1-9.	3.5	134
9	Cadmium and zinc removal from aqueous solutions by Bacillus jeotgali: pH, salinity and temperature effects. Bioresource Technology, 2008, 99, 3864-3870.	9.6	119
10	Multilocus Sequence Analysis Reveals that Vibrio harveyi and V. campbellii Are Distinct Species. Applied and Environmental Microbiology, 2007, 73, 4279-4285.	3.1	116
11	Phenotypic diversity amongst Vibrio isolates from marine aquaculture systems. Aquaculture, 2003, 219, 9-20.	3.5	112
12	Virulence of luminous vibrios to Artemia franciscana nauplii. Diseases of Aquatic Organisms, 2003, 53, 231-240.	1.0	100
13	Beneficial effects of four Bacillus strains on the larval cultivation of Litopenaeus vannamei. Aquaculture, 2011, 321, 136-144.	3.5	95
14	In vitro susceptibility to 15 antibiotics of vibrios isolated from penaeid shrimps in Northwestern Mexico. International Journal of Antimicrobial Agents, 2001, 17, 383-387.	2.5	89
15	Vibrio kanaloae sp. nov., Vibrio pomeroyi sp. nov. and Vibrio chagasii sp. nov., from sea water and marine animals. International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 753-759.	1.7	86
16	Plasmid profiling and antibiotic resistance ofVibriostrains isolated from cultured penaeid shrimp. FEMS Microbiology Letters, 2002, 213, 7-12.	1.8	84
17	Vibrio rotiferianus sp. nov., isolated from cultures of the rotifer Brachionus plicatilis. International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 239-243.	1.7	83
18	Draft Genome Sequence of Vibrio parahaemolyticus Strain M0605, Which Causes Severe Mortalities of Shrimps in Mexico. Genome Announcements, 2014, 2, .	0.8	81

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19	Vibrio neptunius sp. nov., Vibrio brasiliensis sp. nov. and Vibrio xuii sp. nov., isolated from the marine aquaculture environment (bivalves, fish, rotifers and shrimps). International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 245-252.	1.7	75
20	Virulence of Vibrio harveyi responsible for the â€œBright-redâ€•Syndrome in the Pacific white shrimp Litopenaeus vannamei. Journal of Invertebrate Pathology, 2012, 109, 307-317.	3.2	70
21	Bioencapsulation of Two DifferentVibrioSpecies in Nauplii of the Brine Shrimp (Artemia) Tj ETQq1 1 0.784314 rgBT /Overloc	3.1	65
22	Probiotics in the intestinal tract of juvenile whiteleg shrimp Litopenaeus vannamei: modulation of the bacterial community. World Journal of Microbiology and Biotechnology, 2013, 29, 257-265.	3.6	64
23	Outbreak of gastroenteritis caused by the pandemic Vibrio parahaemolyticus O3Ã¢Â€Âƒ:Ã¢Â€ÂƒK6 in Mexico. FEMS Microbiology Letters, 2006, 265, 76-80.	1.8	61
24	Effect of temperature and dietary lipid proportion on gut microbiota in yellowtail kingfish Seriola lalandi juveniles. Aquaculture, 2018, 497, 269-277.	3.5	59
25	Culture of Vibrio alginolyticus C7b, a potential probiotic bacterium, with the microalga Chaetoceros muelleri. Aquaculture, 2002, 211, 43-48.	3.5	57
26	â€˜Bright-redâ€™ syndrome in Pacific white shrimp Litopenaeus vannamei is caused by Vibrio harveyi. Diseases of Aquatic Organisms, 2010, 92, 11-19.	1.0	56
27	Identification and virulence of Aeromonas dhakensis , Pseudomonas mosselii and Microbacterium paraoxydans isolated from Nile tilapia, Oreochromis niloticus , cultivated in Mexico. Journal of Applied Microbiology, 2013, 115, 654-662.	3.1	55
28	Bactericidal effect of lactoferrin and lactoferrin chimera against halophilic Vibrio parahaemolyticus. Biochimie, 2009, 91, 133-140.	2.6	52
29	Genomic diversity of vibrios associated with the Brazilian coralMussismilia hispidaand its sympatric zoanthids (Palythoa caribaeorum,Palythoa variabilisandZoanthus) Tj ETQq1 1 0.784314 rgBT /Overlock 10	3.3	51
30	Cu and Pb biosorption on Bacillus thioparans strain U3 in aqueous solution: Kinetic and equilibrium studies. Chemical Engineering Journal, 2012, 181-182, 352-359.	12.7	50
31	Advanced Microbial Taxonomy Combined with Genome-Based-Approaches Reveals that Vibrio astriarenae sp. nov., an Agarolytic Marine Bacterium, Forms a New Clade in Vibrionaceae. PLoS ONE, 2015, 10, e0136279.	2.5	47
32	Photobacterium swingsii sp. nov., isolated from marine organisms. International Journal of Systematic and Evolutionary Microbiology, 2011, 61, 315-319.	1.7	43
33	Enterovibrio norvegicus gen. nov., sp. nov., isolated from the gut of turbot (Scophthalmus maximus) larvae: a new member of the family Vibrionaceae. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 2015-2022.	1.7	41
34	Detection and Identification of tdh - and trh -Positive Vibrio parahaemolyticus Strains from Four Species of Cultured Bivalve Molluscs on the Spanish Mediterranean Coast. Applied and Environmental Microbiology, 2009, 75, 7574-7577.	3.1	40
35	Doing More with Less: A Comparison of 16S Hypervariable Regions in Search of Defining the Shrimp Microbiota. Microorganisms, 2020, 8, 134.	3.6	37
36	Enterovibrio norvegicus gen. nov., sp. nov., isolated from the gut of turbot (Scophthalmus maximus) larvae: a new member of the family Vibrionaceae.. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 2015-2022.	1.7	37

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37	Vibrio hispanicus sp. nov., isolated from Artemia sp. and sea water in Spain. International Journal of Systematic and Evolutionary Microbiology, 2004, 54, 261-265.	1.7	36
38	Vibrio sinaloensis sp. nov., isolated from the spotted rose snapper, Lutjanus guttatus Steindachner, 1869. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 1621-1624.	1.7	32
39	Overfeeding a High-Fat Diet Promotes Sex-Specific Alterations on the Gut Microbiota of the Zebrafish (Danio rerio). Zebrafish, 2019, 16, 268-279.	1.1	32
40	Effects of enrofloxacin and florfenicol on survival and bacterial population in an experimental infection with luminescent Vibrio campbellii in shrimp larvae of Litopenaeus vannamei. Aquaculture, 2006, 255, 48-54.	3.5	31
41	Vibrio mexicanus sp. nov., isolated from a cultured oyster Crassostrea corteziensis. Antonie Van Leeuwenhoek, 2015, 108, 355-364.	1.7	30
42	Vibrios of the spotted rose snapper Lutjanus guttatus Steindachner, 1869 from northwestern Mexico. Journal of Applied Microbiology, 2007, 102, 1518-1526.	3.1	29
43	Thaumasiovibrio occultus gen. nov. sp. nov. and Thaumasiovibrio subtropicus sp. nov. within the family Vibrionaceae, isolated from coral reef seawater off Ishigaki Island, Japan. Systematic and Applied Microbiology, 2017, 40, 290-296.	2.8	28
44	Isolation of Vibrionaceae from wild blue mussel (Mytilus edulis) adults and their impact on blue mussel larviculture. FEMS Microbiology Ecology, 2017, 93, .	2.7	26
45	Virulence of the fish pathogen Aeromonas dhakensis: genes involved, characterization and histopathology of experimentally infected hybrid tilapia. Diseases of Aquatic Organisms, 2018, 129, 107-116.	1.0	26
46	Vibrio Clade 3.0: New Vibrionaceae Evolutionary Units Using Genome-Based Approach. Current Microbiology, 2022, 79, 10.	2.2	26
47	Vibrio pacinii sp. nov., from cultured aquatic organisms. International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 1569-1573.	1.7	25
48	Vibrio alfacsensis sp. nov., isolated from marine organisms. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 2955-2961.	1.7	24
49	Photobacterium sanguinicancri sp. nov. isolated from marine animals. Antonie Van Leeuwenhoek, 2016, 109, 817-825.	1.7	24
50	OTUs and ASVs Produce Comparable Taxonomic and Diversity from Shrimp Microbiota 16S Profiles Using Tailored Abundance Filters. Genes, 2021, 12, 564.	2.4	23
51	Diversity of vibrios in the haemolymph of the spider crab Maja brachydactyla. Journal of Applied Microbiology, 2010, 109, 918-926.	3.1	22
52	Relationship of aquatic environmental factors with the abundance of Vibrio cholerae, Vibrio parahaemolyticus, Vibrio mimicus and Vibrio vulnificus in the coastal area of Guaymas, Sonora, Mexico. Journal of Water and Health, 2013, 11, 700-712.	2.6	22
53	Vibrio sonorensis sp. nov. isolated from a cultured oyster Crassostrea gigas. Antonie Van Leeuwenhoek, 2016, 109, 1447-1455.	1.7	22
54	Assessment of microbial dynamics and antioxidant enzyme gene expression following probiotic administration in farmed Pacific white shrimp (Litopenaeus vannamei). Aquaculture, 2020, 519, 734907.	3.5	22

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55	Gut microbiota shifts in the giant tiger shrimp, Penaeus monodon, during the postlarvae, juvenile, and adult stages. Aquaculture International, 2020, 28, 1421-1433.	2.2	22
56	The Alpha Variant (B.1.1.7) of SARS-CoV-2 Failed to Become Dominant in Mexico. Microbiology Spectrum, 2022, 10, e0224021.	3.0	21
57	Vibrio ishigakensis sp. nov., in Halioticoli clade isolated from seawater in Okinawa coral reef area, Japan. Systematic and Applied Microbiology, 2016, 39, 330-335.	2.8	20
58	Standardization of the bioencapsulation of enrofloxacin and oxytetracycline in Artemia franciscana Kellogg, 1906. Aquaculture, 2001, 196, 1-12.	3.5	19
59	Assessment of fluorescent-labeled bacteria for evaluation of in vivo uptake of bacteria (Vibrio spp.) by crustacean larvae. Journal of Microbiological Methods, 2003, 52, 101-114.	1.6	18
60	Evaluation of the susceptibility of the cultured shrimp Litopenaeus vannamei to vibriosis when orally exposed to the insecticide methyl parathion. Chemosphere, 2005, 60, 126-134.	8.2	18
61	Vibrio taketomensis sp. nov. by genome taxonomy. Systematic and Applied Microbiology, 2020, 43, 126048.	2.8	17
62	Veronia nyctiphanis gen. nov., sp. nov., Isolated from the Stomach of the Euphausiid Nyctiphanes simplex (Hansen, 1911) in the Gulf of California, and Reclassification of Enterovibrio pacificus as Veronia pacifica comb. nov.. Current Microbiology, 2021, 78, 3782-3790.	2.2	16
63	Vibrio plantisponsor sp. nov., a diazotrophic bacterium isolated from a mangrove associated wild rice (Porteresia coarctata Tateoka). Systematic and Applied Microbiology, 2011, 34, 487-493.	2.8	15
64	The Famlily Vibrionaceae. , 2014, , 659-747.		15
65	Inhibition of Batrachochytrium dendrobatidis Infection by Skin Bacterial Communities in Wild Amphibian Populations. Microbial Ecology, 2021, 82, 666-676.	2.8	14
66	Molecular and Genomic Characterization of Vibrio mimicus Isolated from a Frozen Shrimp Processing Facility in Mexico. PLoS ONE, 2016, 11, e0144885.	2.5	13
67	Occurrence and Abundance of Pathogenic Vibrio Species in Raw Oysters at Retail Seafood Markets in Northwestern Mexico. Journal of Food Protection, 2019, 82, 2094-2099.	1.7	13
68	Development of a bath challenge for the marine shrimp Penaeus vannamei Boone, 1931. Aquaculture, 1998, 169, 283-290.	3.5	12
69	Vibrio barjaei sp. nov., a new species of the Mediterranei clade isolated in a shellfish hatchery. Systematic and Applied Microbiology, 2016, 39, 553-556.	2.8	12
70	Phylogenomic Analysis Supports Two Possible Origins for Latin American Strains of Vibrio parahaemolyticus Associated with Acute Hepatopancreatic Necrosis Disease (AHPND). Current Microbiology, 2020, 77, 3851-3860.	2.2	12
71	Use of bacteriophage vB_Pd_PDCCâ€1 as biological control agent of Photobacterium damselae subsp. damselae during hatching of longfin yellowtail (Seriola rivoliana) eggs. Journal of Applied Microbiology, 2020, 129, 1497-1510.	3.1	12
72	Unique and conserved genome regions in Vibrio harveyi and related species in comparison with the shrimp pathogen Vibrio harveyi CAIM 1792. Microbiology (United Kingdom), 2015, 161, 1762-1779.	1.8	12

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73	Dominance of Three Sublineages of the SARS-CoV-2 Delta Variant in Mexico. Viruses, 2022, 14, 1165.	3.3	12
74	Vibrio crosai sp. nov., isolated from a cultured oyster Crassostrea gigas. Antonie Van Leeuwenhoek, 2014, 106, 457-463.	1.7	11
75	Molecular variability and genetic structure of white spot syndrome virus strains from northwest Mexico based on the analysis of genomes. FEMS Microbiology Letters, 2018, 365, .	1.8	11
76	Photobacterium lucens sp. nov., Isolated from a Cultured Shrimp Penaeus vannamei.. Current Microbiology, 2020, 77, 1111-1116.	2.2	11
77	Quantification of Vibrio species in oysters from the Gulf of Mexico with two procedures based on MPN and PCR. Environmental Monitoring and Assessment, 2016, 188, 602.	2.7	10
78	Artisanal Fresco cheese from Sonora: Physicochemical composition, microbial quality, and bacterial characterization by highâ€throughput sequencing. International Journal of Dairy Technology, 2021, 74, 359-370.	2.8	10
79	Inhibition ofSalmonellaspp. isolated from mango using bacteriocin-like produced by lactobacilli InhibiciÃ³n deSalmonellaspp. aislada de mango usando sustancias tipo bacteriocinas producidas por lactobacilos. CYTA - Journal of Food, 2009, 7, 181-187.	1.9	9
80	Silver nanoparticles are lethal to the ciliate model Tetrahymena and safe to the pike silverside Chirostoma estor. Experimental Parasitology, 2020, 209, 107825.	1.2	9
81	Plasmid profiling and antibiotic resistance of Vibrio strains isolated from cultured penaeid shrimp. FEMS Microbiology Letters, 2002, 213, 7-12.	1.8	9
82	International Committee on Systematics of Prokaryotes Subcommittee on the taxonomy of Aeromonadaceae, Vibrionaceae and related organisms Minutes of the meeting, 13 November 2017, Chicago, USA. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2111-2112.	1.7	9
83	Probiotic modulation of the gut bacterial community of juvenile Litopenaeus vannamei challenged with Vibrio parahaemolyticus CAIM 170. Latin American Journal of Aquatic Research, 2017, 43, 766-775.	0.6	9
84	Delivery of Bioencapsulated Oxytetracycline to the Marine Shrimp Penaeus monodon. Journal of the World Aquaculture Society, 1998, 29, 249-251.	2.4	8
85	Draft Genome Sequence of the Shrimp Pathogen Vibrio harveyi CAIM 1792. Journal of Bacteriology, 2012, 194, 2104-2104.	2.2	8
86	PathogenicVibrio parahaemolyticusisolated from biofouling on commercial vessels and harbor structures. Biofouling, 2015, 31, 275-282.	2.2	8
87	Draft genome sequence of Pseudoalteromonas piscicida strain 36Y_RITHPW, a hypersaline seawater isolate from the south coast of Sonora, Mexico. Journal of Global Antimicrobial Resistance, 2019, 16, 83-86.	2.2	8
88	Streptococcus penaeicida sp. nov., isolated from a diseased farmed Pacific white shrimp (Penaeus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.7	8
89	Draft Genome Sequence of Salmonella enterica subsp. enterica Serotype Oranienburg Strain S-76, Isolated from an Aquatic Environment. Genome Announcements, 2013, 1, .	0.8	7
90	Effect of methyl parathion on the susceptibility of shrimp Litopenaeus vannamei to experimental vibriosis. Diseases of Aquatic Organisms, 2003, 57, 265-270.	1.0	7

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91	Oral Challenge Of Postlarvae Of Litopenaeus Vannamei Through Bioencapsulation Of Vibrio Parahaemolyticus In Artemia Franciscana. Ciencias Marinas, 2000, 26, 65-77.	0.4	7
92	Genomic and biological characterization of the novel phages vB_VpaP_AL-1 and vB_VpaS_AL-2 infecting Vibrio parahaemolyticus associated with acute hepatopancreatic necrosis disease (AHPND). Virus Research, 2022, 312, 198719.	2.2	7
93	Therapeutic effects of enrofloxacin in an experimental infection with a luminescent Vibrio harveyi in Artemia franciscana Kellog 1906. Aquaculture, 2003, 220, 37-42.	3.5	6
94	Draft Genome Sequence of Salmonella enterica subsp. enterica Serotype Saintpaul Strain S-70, Isolated from an Aquatic Environment. Genome Announcements, 2013, 1, .	0.8	6
95	Soy protein concentrate effects on gut microbiota structure and digestive physiology of Totoaba macdonaldi. Journal of Applied Microbiology, 2021, , .	3.1	6
96	Core and Accessory Genome Analysis of Vibrio mimicus. Microorganisms, 2021, 9, 191.	3.6	6
97	Water microbiome dynamics of Pacific white shrimp Penaeus vannamei infected with Vibrio parahaemolyticus strains responsible for acute hepatopancreatic necrosis disease. Aquaculture, 2022, 551, 737871.	3.5	6
98	Effect of functional diets on intestinal microbiota and resistance to Vibrio parahaemolyticus causing acute hepatopancreatic necrosis disease (AHPND) of Pacific white shrimp (Penaeus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 45	3.1	6
99	Genomic Profiling of Antibiotic-Resistant Escherichia coli Isolates from Surface Water of Agricultural Drainage in North-Western Mexico: Detection of the International High-Risk Lineages ST410 and ST617. Microorganisms, 2022, 10, 662.	3.6	6
100	Exploring the Genome of Cheese Starter Lactic Acid Bacterium Lactococcus lactis subsp. lactis CECT 4433. Genome Announcements, 2014, 2, .	0.8	5
101	Effect of pH on the bacterial community present in larvae and spat of Crassostrea gigas. Latin American Journal of Aquatic Research, 2019, 47, 513-523.	0.6	5
102	Tratamientos profilÃ¡cticos para desinfectar la superficie de huevos del pargo flamenco Lutjanus guttatus. Revista De Biologia Marina Y Oceanografia, 2012, 47, 155-160.	0.2	5
103	A review on the use of microorganisms as probiotics. Revista Latinoamericana De MicrobiologÃa, 1998, 40, 166-72.	0.1	5
104	A comparison between total viable count by spread plating and AquaPlakÂ® for enumeration of bacteria in water from a shrimp farm. Journal of Microbiological Methods, 1997, 30, 217-220.	1.6	4
105	Draft Genome Sequence of Vibrio mimicus Strain CAIM 602 T. Genome Announcements, 2013, 1, e0008413.	0.8	4
106	Toxigenic V. cholerae, V. parahaemolyticus, and V. vulnificus in oysters from the Gulf of Mexico and sold in Mexico City. International Journal of Environmental Health Research, 2019, 29, 430-440.	2.7	4
107	Genomic characterization of closely related species in the Rumoiensis clade infers ecogenomic signatures to nonâ€marine environments. Environmental Microbiology, 2020, 22, 3205-3217.	3.8	4
108	Genomic taxonomy of the Mediterranei clade of the genus Vibrio (Gammaproteobacteria). Antonie Van Leeuwenhoek, 2020, 113, 851-859.	1.7	4

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109	A metagenomic assessment of microbial communities in anaerobic bioreactors and sediments: Taxonomic and functional relationships. Anaerobe, 2021, 68, 102296.	2.1	4
110	Use of Corn Husk Meal in the Development of a Functional Diet for Nile tilapia (Oreochromis) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 707 Waste and Biomass Valorization, 2021, 12, 4355.	3.4	4
111	Prevalence and Genomic Diversity of Salmonella enterica Recovered from River Water in a Major Agricultural Region in Northwestern Mexico. Microorganisms, 2022, 10, 1214.	3.6	4
112	Vibrio tetraodonis sp. nov.: genomic insights on the secondary metabolites repertoire. Archives of Microbiology, 2021, 203, 399-404.	2.2	3
113	Reciprocal effect of temperature and dietary lipids on metabolic performance and gut microbiota of Yellowtail kingfish (Seriola lalandi) juveniles. Aquaculture Research, 2021, 52, 6189-6204.	1.8	3
114	Genomic stability among O3:K6â€‰V. parahaemolyticus pandemic strains isolated between 1996 to 2012 in American countries. BMC Genomic Data, 2021, 22, 38.	1.7	3
115	Isolation, Enumeration, and Preservation of the Vibrionaceae. , 0, , 13-26.		3
116	Spatiotemporal distribution of Vibrio parahaemolyticus in relation to environmental parameters in a coastal lagoon on the Pacific coast of northwestern Mexico. Ciencias Marinas, 2018, 44, 141-153.	0.4	3
117	BACTERIAL COMMUNITIES OF THE OYSTERS Crassostrea corteziensis AND C. sikamea OF COSPITA BAY, SINALOA, MEXICO. Revista Internacional De Contaminacion Ambiental, 2018, 34, 203-213.	0.4	3
118	Global m6A RNA Methylation in SARS-CoV-2 Positive Nasopharyngeal Samples in a Mexican Population: A First Approximation Study. Epigenomes, 2022, 6, 16.	1.8	3
119	Bacterial Fish Diseases and Molecular Tools for Bacterial Fish Pathogens Detection. , 2009, , 73-99.		2
120	Draft Genome Sequences of Two Vibrionaceae Species, Vibrio ponticus C121 and Photobacterium aphoticum C119, Isolated as Coral Reef Microbiota. Genome Announcements, 2014, 2, .	0.8	2
121	A preliminary study of the effect of total fishmeal replacement with different dietary sources on the gut microbiota of spotted rose snapper juvenile (Lutjanus guttatus Steindachner, 1869). Aquaculture Research, 2020, 51, 4771-4784.	1.8	2
122	Experimental infection of the white snook Centropomus viridis Lockington (1877) with Vibrio ponticus: Histopathological manifestations and screening for putative virulence genes. Aquaculture, 2020, 528, 735599.	3.5	2
123	Genomic and molecular evolutionary dynamics of transcriptional response regulator genes in bacterial species of the Harveyi clade of Vibrio. Gene, 2021, 783, 145577.	2.2	2
124	BIOSORPTION OF CADMIUM AND LEAD USING SUSPENDED AND IMMOBILIZED Enterobacter cloacae AT DIFFERENT PH. Revista Internacional De Contaminacion Ambiental, 2019, 35, 259-264.	0.4	2
125	Vibrio hispanicus sp. nov., isolated from Artemia sp. and sea water in Spain. International Journal of Systematic and Evolutionary Microbiology, 2004, 54, 629-629.	1.7	1
126	Probiotics in the Larval Culture of Aquatic Organisms. , 2011, , 31-46.		1

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127	Draft Genome Sequence of a Mexican Community-Associated Methicillin-Resistant Staphylococcus epidermidis Strain. Genome Announcements, 2017, 5, .	0.8	0
128	Draft Genome Sequence of Escherichia coli Strain M15-4, a Typical Enteropathogenic E. coli Strain Isolated in Mexico. Genome Announcements, 2018, 6, .	0.8	0
129	Whole-genome sequencing of Staphylococcus aureus L401, a mecA-negative community-associated methicillin-resistant strain isolated from a healthy carrier. Journal of Global Antimicrobial Resistance, 2019, 17, 260-262.	2.2	0
130	Shrimp Diseases and Molecular Diagnostic Methods. , 2009, , 101-131.		0