

# JesÃ³s Ãngel LÃ³pez Romalde

## List of Publications by Year in descending order

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217  
papers

7,890  
citations

46984

47  
h-index

76872

74  
g-index

228  
all docs

228  
docs citations

228  
times ranked

6227  
citing authors

#	ARTICLE	IF	CITATIONS
1	Prevalence of human bocavirus infections in Europe. A systematic review and meta-analysis. <i>Transboundary and Emerging Diseases</i> , 2022, 69, 2451-2461.	1.3	20
2	World Society for Virology first international conference: Tackling global virus epidemics. <i>Virology</i> , 2022, 566, 114-121.	1.1	2
3	Assessment of animal diseases caused by bacteria resistant to antimicrobials: kept fish species. <i>EFSA Journal</i> , 2022, 20, e07076.	0.9	1
4	Emerging Viruses in Sewage Sludge and Soils. <i>Handbook of Environmental Chemistry</i> , 2022, , 289-305.	0.2	1
5	Health Status of <i>Mytilus chilensis</i> from Intensive Culture Areas in Chile Assessed by Molecular, Microbiological, and Histological Analyses. <i>Pathogens</i> , 2022, 11, 494.	1.2	0
6	Coevolution of Molluscs and Their Microbes. <i>Advances in Environmental Microbiology</i> , 2021, , 513-526.	0.1	0
7	Extended-Spectrum $\hat{2}$ -Lactamase and Carbapenemase Genes are Substantially and Sequentially Reduced during Conveyance and Treatment of Urban Sewage. <i>Environmental Science &amp; Technology</i> , 2021, 55, 5939-5949.	4.6	24
8	A deep-sea bacterium related to coastal marine pathogens. <i>Environmental Microbiology</i> , 2021, 23, 5349-5363.	1.8	4
9	Monitoring Emergence of the SARS-CoV-2 B.1.1.7 Variant through the Spanish National SARS-CoV-2 Wastewater Surveillance System (VATar COVID-19). <i>Environmental Science &amp; Technology</i> , 2021, 55, 11756-11766.	4.6	39
10	Detection of SARS-CoV-2 RNA in bivalve mollusks and marine sediments. <i>Science of the Total Environment</i> , 2021, 786, 147534.	3.9	33
11	Dynamics of integron structures across a wastewater network – Implications to resistance gene transfer. <i>Water Research</i> , 2021, 206, 117720.	5.3	18
12	Metataxonomic analysis of tissue-associated microbiota in grooved carpet-shell (Ruditapes) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 302 T	1.1	2
13	EMBRACE-WATERS statement: Recommendations for reporting of studies on antimicrobial resistance in wastewater and related aquatic environments. <i>One Health</i> , 2021, 13, 100339.	1.5	11
14	Multilocus sequence analysis reveals different lineages of <i>Pseudomonas anguilliseptica</i> associated with disease in farmed lumpfish ( <i>Cyclopterus lumpus</i> L.). <i>PLoS ONE</i> , 2021, 16, e0259725.	1.1	1
15	Comprehensive comparison of chemically enhanced primary treatment and high-rate activated sludge in novel wastewater treatment plant configurations. <i>Water Research</i> , 2020, 169, 115258.	5.3	67
16	<i>Halomonas borealis</i> sp. nov. and <i>Halomonas niordiana</i> sp. nov., two new species isolated from seawater. <i>Systematic and Applied Microbiology</i> , 2020, 43, 126040.	1.2	17
17	Norovirus contamination of sea urchins ( <i>Paracentrotus lividus</i> ): Potential food risk for consumers. <i>Food Control</i> , 2020, 111, 107041.	2.8	11
18	WSV 2019: The First Committee Meeting of the World Society for Virology. <i>Virologica Sinica</i> , 2020, 35, 248-252.	1.2	2

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19	Comparative study of the culturable microbiota present in two different rearing systems, flow-through system (FTS) and recirculation system (RAS), in a great scallop hatchery. <i>Aquaculture Research</i> , 2020, 51, 542-556.	0.9	4
20	Identification of Emerging Hazards in Mussels by the Galician Emerging Food Safety Risks Network (RISEGAL). A First Approach. <i>Foods</i> , 2020, 9, 1641.	1.9	7
21	Making waves: Wastewater-based epidemiology for COVID-19 – approaches and challenges for surveillance and prediction. <i>Water Research</i> , 2020, 186, 116404.	5.3	250
22	Draft Genome Sequence of <i>Aeromonas sobria</i> Strain CHT-30, Isolated from a Diseased Rainbow Trout ( <i>Oncorhynchus mykiss</i> ) in Peru. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	0
23	Comparison of antibiotic-resistant bacteria and antibiotic resistance genes abundance in hospital and community wastewater: A systematic review. <i>Science of the Total Environment</i> , 2020, 743, 140804.	3.9	126
24	Draft Genome Sequence of <i>Yersinia ruckeri</i> Strain FMV-22, Isolated from Diseased Rainbow Trout ( <i>Oncorhynchus mykiss</i> ) in Peru. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	0
25	A Comprehensive Review on Human Aichi Virus. <i>Virologica Sinica</i> , 2020, 35, 501-516.	1.2	40
26	Hepatitis E virus genotype 3 in echinoderms: First report of sea urchin ( <i>Paracentrotus lividus</i> ) contamination. <i>Food Microbiology</i> , 2020, 89, 103415.	2.1	5
27	<i>Aliarcobacter vitoriensis</i> sp. nov., isolated from carrot and urban wastewater. <i>Systematic and Applied Microbiology</i> , 2020, 43, 126091.	1.2	17
28	Detection of Hepatitis E Virus in Shellfish Harvesting Areas from Galicia (Northwestern Spain). <i>Viruses</i> , 2019, 11, 618.	1.5	24
29	Epidemiology of Aichi virus in fecal samples from outpatients with acute gastroenteritis in Northwestern Spain. <i>Journal of Clinical Virology</i> , 2019, 118, 14-19.	1.6	12
30	Spatial ecology of a wastewater network defines the antibiotic resistance genes in downstream receiving waters. <i>Water Research</i> , 2019, 162, 347-357.	5.3	108
31	Clonal relationship among <i>Vibrio parahaemolyticus</i> isolated from Mediterranean mussels ( <i>Mytilus</i> ) Tj ETQq1 1 0.784314 rgBT /Overl... <i>Microbiology</i> , 2019, 84, 103258.	2.1	7
32	Scrutinizing the triad of <i>Vibrio tapetis</i> , the skin barrier and pigmentation as determining factors in the development of skin ulcerations in wild common dab ( <i>Limanda limanda</i> ). <i>Veterinary Research</i> , 2019, 50, 41.	1.1	6
33	Editorial: Microbial Taxonomy, Phylogeny and Biodiversity. <i>Frontiers in Microbiology</i> , 2019, 10, 1324.	1.5	3
34	Human Sapovirus among Outpatients with Acute Gastroenteritis in Spain: A One-Year Study. <i>Viruses</i> , 2019, 11, 144.	1.5	37
35	Development of a novel digital RT-PCR method for detection of human sapovirus in different matrices. <i>Journal of Virological Methods</i> , 2018, 254, 21-24.	1.0	12
36	Hepatitis A Virus Disinfection in Water by Solar Photo-Fenton Systems. <i>Food and Environmental Virology</i> , 2018, 10, 159-166.	1.5	6

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37	Sapovirus in Wastewater Treatment Plants in Tunisia: Prevalence, Removal, and Genetic Characterization. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	1.4	27
38	An overview of 20 years of studies on the prevalence of human enteric viruses in shellfish from Galicia, Spain. <i>Journal of Applied Microbiology</i> , 2018, 124, 943-957.	1.4	24
39	First isolation of <i>Vibrio tapetis</i> and an atypical strain of <i>Aeromonas salmonicida</i> from skin ulcerations in common dab ( <i>Limanda limanda</i> ) in the North Sea. <i>Journal of Fish Diseases</i> , 2018, 41, 329-335.	0.9	13
40	Genetic studies to re-affiliate <i>Edwardsiella tarda</i> fish isolates to <i>Edwardsiella piscicida</i> and <i>Edwardsiella anguillarum</i> species. <i>Systematic and Applied Microbiology</i> , 2018, 41, 30-37.	1.2	58
41	Revisiting the Taxonomy of the Genus <i>Arcobacter</i> : Getting Order From the Chaos. <i>Frontiers in Microbiology</i> , 2018, 9, 2077.	1.5	245
42	Multilocus Variable-Number Tandem-Repeat Analysis of <i>Yersinia ruckeri</i> Confirms the Existence of Host Specificity, Geographic Endemism, and Anthropogenic Dissemination of Virulent Clones. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	1.4	27
43	Population genetic and evolution analysis of controversial genus <i>Edwardsiella</i> by multilocus sequence typing. <i>Molecular Phylogenetics and Evolution</i> , 2018, 127, 513-521.	1.2	11
44	<i>Arcobacter haliotis</i> Tanaka et al. 2017 is a later heterotypic synonym of <i>Arcobacter lekithochrous</i> DiÁguez et al. 2017. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 2851-2854.	0.8	10
45	<i>Neptuniibacter pectenicola</i> sp. nov. and <i>Neptuniibacter marinus</i> sp. nov., two novel species isolated from a Great scallop ( <i>Pecten maximus</i> ) hatchery in Norway and emended description of the genus <i>Neptuniibacter</i> . <i>Systematic and Applied Microbiology</i> , 2017, 40, 80-85.	1.2	17
46	Genome sequence of three <i>Psychrobacter</i> sp. strains with potential applications in bioremediation. <i>Genomics Data</i> , 2017, 12, 7-10.	1.3	21
47	Low prevalence of Aichi virus in molluscan shellfish samples from Galicia (NW Spain). <i>Journal of Applied Microbiology</i> , 2017, 122, 516-521.	1.4	11
48	<i>Kiloniella majae</i> sp. nov., isolated from spider crab ( <i>Maja brachydactyla</i> ) and pullet carpet shell clam ( <i>Venerupis pullastra</i> ). <i>Systematic and Applied Microbiology</i> , 2017, 40, 274-279.	1.2	13
49	Complete Genome Sequence of <i>Arcobacter</i> sp. Strain LFT 1.7 Isolated from Great Scallop ( <i>Pecten maximus</i> ) Tj ETQq1 1 0.784314 rgBT <sub>2</sub> /Overlo 0,8	0,8	0
50	Complete characterization of new isolates of <i>Neptunomonas phycophila</i> leads to emend its description and opens possibilities of biotechnological applications. <i>MicrobiologyOpen</i> , 2017, 6, e00519.	1.2	6
51	Characterization and in vitro evaluation of new bacteriophages for the biocontrol of <i>Escherichia coli</i> . <i>Virus Research</i> , 2017, 227, 171-182.	1.1	36
52	Application of phage therapy during bivalve depuration improves <i>Escherichia coli</i> decontamination. <i>Food Microbiology</i> , 2017, 61, 102-112.	2.1	34
53	From the Gene Sequence to the Phylogeography through the Population Structure: The Cases of <i>Yersinia ruckeri</i> and <i>Vibrio tapetis</i> . , 2017, , .		1
54	Comparative Genomic Analysis of Two <i>Vibrio toranzoniae</i> Strains with Different Virulence Capacity Reveals Clues on Its Pathogenicity for Fish. <i>Frontiers in Microbiology</i> , 2017, 8, 86.	1.5	6

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55	New Insights into Pathogenic Vibrios Affecting Bivalves in Hatcheries: Present and Future Prospects. <i>Frontiers in Microbiology</i> , 2017, 8, 762.	1.5	102
56	Launching a Global Network of Virologists: The World Society for Virology (WSV). <i>Intervirology</i> , 2017, 60, 276-277.	1.2	3
57	Draft Genome Sequences of <i>Neptuniibacter</i> sp. Strains LFT 1.8 and ATR 1.1. <i>Genome Announcements</i> , 2017, 5, .	0.8	0
58	Isolation of <i>Vibrio tapetis</i> from two native fish species ( <i>Genypterus chilensis</i> and <i>Paralichthys</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 627 <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 716-723.	0.8	16
59	<i>Arcobacter lekithochrous</i> sp. nov., isolated from a molluscan hatchery. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1327-1332.	0.8	43
60	<i>Vibrio tapetis</i> from wrasse used for ectoparasite bio-control in salmon farming: phylogenetic analysis and serotyping. <i>Diseases of Aquatic Organisms</i> , 2017, 125, 189-197.	0.5	6
61	Draft Genome Sequence of <i>Vibrio toranzoniae</i> Strain CECT 7225 T. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
62	Draft Genome Sequence of the New Pathogen for Bivalve Larvae <i>Vibrio bivalvicida</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	2
63	Twitter as a Tool for Teaching and Communicating Microbiology: The #microMOOCSEM Initiative. <i>Journal of Microbiology and Biology Education</i> , 2016, 17, 492-494.	0.5	9
64	Hepatitis E virus genotype 3 in mussels ( <i>Mytilus galloprovincialis</i> ), Spain. <i>Food Microbiology</i> , 2016, 58, 13-15.	2.1	55
65	Detection and Molecular Characterization of Hepatitis A Virus from Tunisian Wastewater Treatment Plants with Different Secondary Treatments. <i>Applied and Environmental Microbiology</i> , 2016, 82, 3834-3845.	1.4	22
66	Bacteriophages with potential to inactivate <i>Salmonella</i> Typhimurium: Use of single phage suspensions and phage cocktails. <i>Virus Research</i> , 2016, 220, 179-192.	1.1	90
67	Human Sapovirus in Mussels from R�a do Burgo, A Coru�a (Spain). <i>Food and Environmental Virology</i> , 2016, 8, 187-193.	1.5	10
68	<i>Vibrio barjaei</i> sp. nov., a new species of the Mediterranean clade isolated in a shellfish hatchery. <i>Systematic and Applied Microbiology</i> , 2016, 39, 553-556.	1.2	12
69	<i>Vibrio sonorensis</i> sp. nov. isolated from a cultured oyster <i>Crassostrea gigas</i> . <i>Antonie Van Leeuwenhoek</i> , 2016, 109, 1447-1455.	0.7	22
70	Application of bacteriophages during depuration reduces the load of <i>Salmonella</i> Typhimurium in cockles. <i>Food Research International</i> , 2016, 90, 73-84.	2.9	18
71	<i>Photobacterium sanguinicancri</i> sp. nov. isolated from marine animals. <i>Antonie Van Leeuwenhoek</i> , 2016, 109, 817-825.	0.7	24
72	Prevalence and Genetic Diversity of Human Sapoviruses in Shellfish from Commercial Production Areas in Galicia, Spain. <i>Applied and Environmental Microbiology</i> , 2016, 82, 1167-1172.	1.4	19

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73	<i>Vibrio bivalvicida</i> sp. nov., a novel larval pathogen for bivalve molluscs reared in a hatchery. <i>Systematic and Applied Microbiology</i> , 2016, 39, 8-13.	1.2	26
74	Reclassification of the larval pathogen for marine bivalves <i>Vibrio tubiashii</i> subsp. <i>europaeus</i> as <i>Vibrio europaeus</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 4791-4796.	0.8	20
75	Characterization of the microbiota associated to <i>Pecten maximus</i> gonads using 454-pyrosequencing. <i>International Microbiology</i> , 2016, 19, 93-99.	1.1	18
76	<i>Marinomonas gallaica</i> sp. nov. and <i>Marinomonas atlantica</i> sp. nov., isolated from reared clams ( <i>Ruditapes decussatus</i> ). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 3183-3188.	0.8	14
77	Phylogeography of <i>Yersinia ruckeri</i> reveals effects of past evolutionary events on the current strain distribution and explains variations in the global transmission of enteric redmouth (ERM) disease. <i>Frontiers in Microbiology</i> , 2015, 6, 1198.	1.5	16
78	<i>Vibrio mexicanus</i> sp. nov., isolated from a cultured oyster <i>Crassostrea corteziensis</i> . <i>Antonie Van Leeuwenhoek</i> , 2015, 108, 355-364.	0.7	30
79	Seasonal variation of bacterial communities in shellfish harvesting waters: Preliminary study before applying phage therapy. <i>Marine Pollution Bulletin</i> , 2015, 90, 68-77.	2.3	17
80	Description of <i>Lacinutrix venerupis</i> sp. nov.: A novel bacterium associated with reared clams. <i>Systematic and Applied Microbiology</i> , 2015, 38, 115-119.	1.2	21
81	Molecular epidemiology of norovirus from patients with acute gastroenteritis in northwestern Spain. <i>Epidemiology and Infection</i> , 2015, 143, 316-324.	1.0	10
82	Isolation and identification of <i>Vibrio toranzoniae</i> associated with diseased red conger eel ( <i>Genypterus chilensis</i> ) farmed in Chile. <i>Veterinary Microbiology</i> , 2015, 179, 327-331.	0.8	23
83	Mathematical model for viral depuration kinetics in shellfish: An useful tool to estimate the risk for the consumers. <i>Food Microbiology</i> , 2015, 49, 220-225.	2.1	22
84	Efficiency of hepatitis A virus removal in six sewage treatment plants from central Tunisia. <i>Applied Microbiology and Biotechnology</i> , 2015, 99, 10759-10769.	1.7	20
85	<i>Sinobacterium norvegicum</i> sp. nov., isolated from great scallop ( <i>Pecten maximus</i> ) broodstock and emended description of <i>Sinobacterium caligoides</i> . <i>Antonie Van Leeuwenhoek</i> , 2015, 108, 983-991.	0.7	8
86	Detection and quantification of hepatitis A virus and norovirus in Spanish authorized shellfish harvesting areas. <i>International Journal of Food Microbiology</i> , 2015, 193, 43-50.	2.1	77
87	Solar water disinfection (SODIS): Impact on hepatitis A virus and on a human Norovirus surrogate under natural solar conditions. <i>International Microbiology</i> , 2015, 18, 41-9.	1.1	14
88	<i>Vibrio tapetis</i> isolated from vesicular skin lesions in Dover sole <i>Solea solea</i> . <i>Diseases of Aquatic Organisms</i> , 2015, 115, 81-86.	0.5	11
89	New <i>Vibrio</i> species associated to molluscan microbiota: a review. <i>Frontiers in Microbiology</i> , 2014, 4, 413.	1.5	118
90	Genome Sequence of <i>Streptococcus phocae</i> subsp. <i>salmonis</i> Strain C-4 T , Isolated from Atlantic Salmon ( <i>Salmo salar</i> ). <i>Genome Announcements</i> , 2014, 2, .	0.8	4

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91	<i>Vibrio ostreicida</i> sp. nov., a new pathogen of bivalve larvae. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 1641-1646.	0.8	26
92	Effectiveness of depuration for hepatitis A virus removal from mussels ( <i>Mytilus galloprovincialis</i> ). International Journal of Food Microbiology, 2014, 180, 24-29.	2.1	17
93	<i>Vibrio cortegadensis</i> sp. nov., isolated from clams. Antonie Van Leeuwenhoek, 2014, 105, 335-341.	0.7	12
94	Disentangling the Population Structure and Evolution of the Clam Pathogen <i>Vibrio tapetis</i> . Microbial Ecology, 2014, 67, 145-154.	1.4	4
95	Depuration kinetics of murine norovirus in shellfish. Food Research International, 2014, 64, 182-187.	2.9	23
96	<i>Vibrio crosai</i> sp. nov., isolated from a cultured oyster <i>Crassostrea gigas</i> . Antonie Van Leeuwenhoek, 2014, 106, 457-463.	0.7	11
97	Viral elimination during commercial depuration of shellfish. Food Control, 2014, 43, 206-212.	2.8	38
98	Depuration kinetics of hepatitis A virus in clams. Food Microbiology, 2014, 39, 103-107.	2.1	21
99	Comparative polyphasic characterization of <i>Streptococcus phocae</i> strains with different host origin and description of the subspecies <i>Streptococcus phocae</i> subsp. <i>salmonis</i> subsp. nov.. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 1775-1781.	0.8	19
100	Multilocus sequence analysis of <i>Vibrio tapetis</i> , the causative agent of Brown Ring Disease: Description of <i>Vibrio tapetis</i> subsp. <i>britannicus</i> subsp. nov. Systematic and Applied Microbiology, 2013, 36, 183-187.	1.2	29
101	Detection and Characterization of Hepatitis A Virus and Norovirus in Mussels from Galicia (NW) Tj ETQq1 1 0.784314 rgBT /Qverlock 10	1.5	31
102	<i>Vibrio toranzoniae</i> sp. nov., a new member of the Splendidus clade in the genus <i>Vibrio</i> . Systematic and Applied Microbiology, 2013, 36, 96-100.	1.2	30
103	Role of norovirus in acute gastroenteritis in the Northwest of Spain during 2010-2011. Journal of Medical Virology, 2013, 85, 2009-2015.	2.5	10
104	Identification and virulence of <i>Aeromonas dhakensis</i> , <i>Pseudomonas mosselii</i> and <i>Microbacterium paraoxydans</i> isolated from Nile tilapia, <i>Oreochromis niloticus</i> , cultivated in Mexico. Journal of Applied Microbiology, 2013, 115, 654-662.	1.4	55
105	A polyphasic approach to study the intraspecific diversity of <i>Yersinia ruckeri</i> strains isolated from recent outbreaks in salmonid culture. Veterinary Microbiology, 2012, 160, 176-182.	0.8	14
106	Virulence of <i>Vibrio harveyi</i> responsible for the "Bright-red" Syndrome in the Pacific white shrimp <i>Litopenaeus vannamei</i> . Journal of Invertebrate Pathology, 2012, 109, 307-317.	1.5	70
107	Evaluation of different culture media for the isolation and growth of the fastidious <i>Vibrio tapetis</i> , the causative agent of brown ring disease. Journal of Invertebrate Pathology, 2012, 111, 74-81.	1.5	4
108	Effectiveness of bivalent vaccines against <i>Aeromonas hydrophila</i> and <i>Lactococcus garvieae</i> infections in rainbow trout <i>Oncorhynchus mykiss</i> (Walbaum). Fish and Shellfish Immunology, 2012, 32, 756-761.	1.6	58

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109	<i>Pseudomonas baetica</i> sp. nov., a fish pathogen isolated from wedge sole, <i>Dicologlossa cuneata</i> (Moreau). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012, 62, 874-882.	0.8	56
110	Highly sensitive detection and quantification of the pathogen <i>Yersinia ruckeri</i> in fish tissues by using real-time PCR. <i>Applied Microbiology and Biotechnology</i> , 2012, 96, 511-520.	1.7	15
111	Comparative study on the antibiotic susceptibility and plasmid profiles of <i>Vibrio alginolyticus</i> strains isolated from four Tunisian marine biotopes. <i>World Journal of Microbiology and Biotechnology</i> , 2012, 28, 3345-3363.	1.7	23
112	Two-dimensional proteome reference map of <i>Vibrio tapetis</i> , the aetiological agent of brown ring disease in clams. <i>Journal of Applied Microbiology</i> , 2012, 112, 853-864.	1.4	2
113	<i>Arcobacter bivalviorum</i> sp. nov. and <i>Arcobacter venerupis</i> sp. nov., new species isolated from shellfish. <i>Systematic and Applied Microbiology</i> , 2012, 35, 133-138.	1.2	91
114	Multilocus sequence typing reveals high genetic diversity and epidemic population structure for the fish pathogen <i>Yersinia ruckeri</i> . <i>Environmental Microbiology</i> , 2012, 14, 1888-1897.	1.8	27
115	Phenotypical and genetic characterization of <i>Yersinia ruckeri</i> strains isolated from recent outbreaks in farmed rainbow trout <i>Oncorhynchus mykiss</i> (Walbaum) in Peru. <i>Aquaculture</i> , 2011, 317, 229-232.	1.7	17
116	Microbial contamination and purification of bivalve shellfish: Crucial aspects in monitoring and future perspectives – A mini-review. <i>Food Control</i> , 2011, 22, 805-816.	2.8	117
117	Evaluation of different species-specific PCR protocols for the detection of <i>Vibrio tapetis</i> . <i>Journal of Invertebrate Pathology</i> , 2011, 108, 85-91.	1.5	8
118	Characterization of <i>Vibrio tapetis</i> strains isolated from diseased cultured Wedge sole ( <i>Dicologlossa cuneata</i> Moreau). <i>Research in Veterinary Science</i> , 2011, 90, 189-195.	0.9	15
119	Serological and molecular heterogeneity among <i>Yersinia ruckeri</i> strains isolated from farmed Atlantic salmon <i>Salmo salar</i> in Chile. <i>Diseases of Aquatic Organisms</i> , 2011, 93, 207-214.	0.5	33
120	Proteomics and multilocus sequence analysis confirm intraspecific variability of <i>Vibrio tapetis</i> . <i>FEMS Microbiology Letters</i> , 2011, 324, 80-87.	0.7	5
121	The use of multiple typing methods allows a more accurate molecular characterization of <i>Vibrio parahaemolyticus</i> strains isolated from the Italian Adriatic Sea. <i>FEMS Microbiology Ecology</i> , 2011, 77, 611-622.	1.3	11
122	Pseudo-membranes on internal organs associated with <i>Rhodococcus qingshengii</i> infection in Atlantic salmon ( <i>Salmo salar</i> ). <i>Veterinary Microbiology</i> , 2011, 147, 200-204.	0.8	8
123	Evaluation of different RNA-extraction kits for sensitive detection of Hepatitis A virus in strawberry samples. <i>Food Microbiology</i> , 2011, 28, 38-42.	2.1	13
124	Norovirus, hepatitis A virus and enterovirus presence in shellfish from high quality harvesting areas in Portugal. <i>Food Microbiology</i> , 2011, 28, 936-941.	2.1	48
125	<i>Photobacterium swingsii</i> sp. nov., isolated from marine organisms. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 315-319.	0.8	43
126	<i>Vibrio atlanticus</i> sp. nov. and <i>Vibrio artabrorum</i> sp. nov., isolated from the clams <i>Ruditapes philippinarum</i> and <i>Ruditapes decussatus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 2406-2411.	0.8	34



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127	Comparison of phenotypical and genetic identification of <i>Aeromonas</i> strains isolated from diseased fish. <i>Systematic and Applied Microbiology</i> , 2010, 33, 149-153.	1.2	106
128	<i>Vibrio celticus</i> sp. nov., a new <i>Vibrio</i> species belonging to the Splendidus clade with pathogenic potential for clams. <i>Systematic and Applied Microbiology</i> , 2010, 33, 311-315.	1.2	37
129	Review of probiotics for use in bivalve hatcheries. <i>Veterinary Microbiology</i> , 2010, 145, 187-197.	0.8	95
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