

# Pierre E Fournier

## List of Publications by Year in descending order

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822  
papers

37,597  
citations

2975

93  
h-index

5120

166  
g-index

866  
all docs

866  
docs citations

866  
times ranked

26380  
citing authors

#	ARTICLE	IF	CITATIONS
1	Direct diagnosis of <i>Pasteurella multocida</i> meningitis using next-generation sequencing. <i>Lancet Microbe</i> , The, 2022, 3, e6.	7.3	8
2	Occurrence of a substitution or deletion of SARS-CoV-2 spike amino acid 677 in various lineages in Marseille, France. <i>Virus Genes</i> , 2022, 58, 53-58.	1.6	8
3	<i>Buttiauxella massiliensis</i> sp. nov., Isolated from a Human Bone Infection. <i>Current Microbiology</i> , 2022, 79, 41.	2.2	5
4	Judicial Opinions 103â€“111. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	1.7	8
5	<i>Arabiibacter massiliensis</i> gen. nov. sp. nov., New Anaerobic Bacterium Isolated from the Human Gut. <i>Current Microbiology</i> , 2022, 79, 47.	2.2	6
6	Limited spread of a rare spike E484K-harboring SARS-CoV-2 in Marseille, France. <i>Archives of Virology</i> , 2022, 167, 583.	2.1	3
7	Characteristics of the first 1119 SARSâ€“CoVâ€“2 Omicron variant cases, in Marseille, France, Novemberâ€“December 2021. <i>Journal of Medical Virology</i> , 2022, 94, 2290-2295.	5.0	56
8	Introduction of the SARS-CoV-2 Beta variant from Comoros into the Marseille geographical area. <i>Travel Medicine and Infectious Disease</i> , 2022, 46, 102277.	3.0	3
9	Q fever endocarditis in India: A report of two cases. <i>Indian Journal of Medical Microbiology</i> , 2022, , .	0.8	1
10	SARS-CoV-2 reinfection and COVID-19 severity. <i>Emerging Microbes and Infections</i> , 2022, 11, 894-901.	6.5	46
11	<i>Peptostreptococcus faecalis</i> sp. nov., new bacterial species isolated from healthy indigenous congolese volunteer. <i>Heliyon</i> , 2022, 8, e09102.	3.2	4
12	Genomic evolution and adaptation of arthropod-associated <i>Rickettsia</i> . <i>Scientific Reports</i> , 2022, 12, 3807.	3.3	25
13	First cases of infection with the 21L/BA.2 Omicron variant in Marseille, France. <i>Journal of Medical Virology</i> , 2022, 94, 3421-3430.	5.0	19
14	Smart apps for self-reporting clinical information. <i>Lancet</i> , The, 2022, , .	13.7	1
15	The severity of the first 207 infections with the SARSâ€“CoVâ€“2 Omicron BA.2 variant, in Marseille, France, December 2021â€“February 2022. <i>Journal of Medical Virology</i> , 2022, 94, 3494-3497.	5.0	18
16	<i>Neglectibacter timonensis</i> gen. nov., sp. nov. and <i>Scatolibacter rhodanostii</i> gen. nov., sp. nov., two anaerobic bacteria isolated from human stool samples. <i>Archives of Microbiology</i> , 2022, 204, 45.	2.2	6
17	Culture and identification of a â€œDeltamicronâ€“SARSâ€“CoVâ€“2 in a three cases cluster in southern France. <i>Journal of Medical Virology</i> , 2022, 94, 3739-3749.	5.0	58
18	Description of <i>Acinetobacter ihumii</i> sp. nov., <i>Microbacterium ihumii</i> sp. nov., and <i>Gulosibacter massiliensis</i> sp. nov., three new bacteria isolated from human blood. <i>FEMS Microbiology Letters</i> , 2022, 369, .	1.8	5

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19	High rate of reinfection with the SARS-CoV-2 Omicron variant. <i>Journal of Infection</i> , 2022, 85, 174-211.	3.3	26
20	Draft Genome Sequence of <i>Corynebacterium sanguinis</i> Strain Marseille-P8776. <i>Microbiology Resource Announcements</i> , 2022, 11, e0000822.	0.6	0
21	Screening and Whole Genome Sequencing of SARS-CoV-2 Circulating During the First Three Waves of the COVID-19 Pandemic in Libreville and the Haut-Ogooué Province in Gabon. <i>Frontiers in Medicine</i> , 2022, 9, .	2.6	7
22	Sequential Appearance and Isolation of a SARS-CoV-2 Recombinant between Two Major SARS-CoV-2 Variants in a Chronically Infected Immunocompromised Patient. <i>Viruses</i> , 2022, 14, 1266.	3.3	13
23	Flying Fox Hemolytic Fever, Description of a New Zoonosis Caused by <i>Candidatus</i> <i>Mycoplasma haemohominis</i> . <i>Clinical Infectious Diseases</i> , 2021, 73, e1445-e1453.	5.8	24
24	Genomic Characterization of the Novel <i>Bartonella refiksaydamii</i> sp. Isolated from the Blood of a <i>Crocidura suaveolens</i> (Pallas, 1811). <i>Vector-Borne and Zoonotic Diseases</i> , 2021, 21, 432-440.	1.5	11
25	Evaluation of the Panbio COVID-19 Rapid Antigen Detection Test Device for the Screening of Patients with COVID-19. <i>Journal of Clinical Microbiology</i> , 2021, 59, .	3.9	136
26	Statins potentiate the antibacterial effect of platelets on <i>Staphylococcus aureus</i> . <i>Platelets</i> , 2021, 32, 671-676.	2.3	8
27	Human galectin-1 and galectin-3 promote <i>Tropheryma whipplei</i> infection. <i>Gut Microbes</i> , 2021, 13, 1-15.	9.8	8
28	<i>Anaerococcus urinimassiliensis</i> sp. nov., a new bacterium isolated from human urine. <i>Scientific Reports</i> , 2021, 11, 2684.	3.3	7
29	Early combination therapy with hydroxychloroquine and azithromycin reduces mortality in 10,429 COVID-19 outpatients. <i>Reviews in Cardiovascular Medicine</i> , 2021, 22, 1063.	1.4	21
30	Draft Genome Sequence of <i>Comamonas jiangduensis</i> Strain YW1 <sup>T</sup> . <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.6	0
31	Variations in respiratory pathogen carriage among a homeless population in a shelter for men in Marseille, France, March–July 2020: cross-sectional 1-day surveys. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 1579-1582.	2.9	9
32	Draft Genome Sequence of <i>Comamonas aquatilis</i> Strain LK (=CSUR P6418=CECT 9772), Isolated from the Planarian <i>Schmidtea mediterranea</i> . <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.6	1
33	Draft Genome Sequence of <i>Herminiimonas contaminans</i> Strain CCM 7991 <sup>T</sup> , a Biopharmaceutical Contaminant. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.6	3
34	Draft Genome Sequence of <i>Vogesella oryzae</i> L3B39 <sup>T</sup> , Isolated from the Rhizosphere of Saline-Tolerant Pokkali Rice. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.6	0
35	Automated Western immunoblotting detection of anti-SARS-CoV-2 serum antibodies. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 1309-1317.	2.9	23
36	Introduction into the Marseille geographical area of a mild SARS-CoV-2 variant originating from sub-Saharan Africa: An investigational study. <i>Travel Medicine and Infectious Disease</i> , 2021, 40, 101980.	3.0	31

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37	Planarians (Platyhelminthes)â€”An Emerging Model Organism for Investigating Innate Immune Mechanisms. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 619081.	3.9	8
38	Screening of SARS-CoV-2 among homeless people, asylum-seekers and other people living in precarious conditions in Marseille, France, Marchâ€”April 2020. <i>International Journal of Infectious Diseases</i> , 2021, 105, 1-6.	3.3	19
39	Mink, SARS-CoV-2, and the Human-Animal Interface. <i>Frontiers in Microbiology</i> , 2021, 12, 663815.	3.5	106
40	<i>Pedobacter ghigonii</i> sp. nov., Isolated from the Microbiota of the Planarian <i>Schmidtea mediterranea</i> . <i>Microbiology Research</i> , 2021, 12, 268-287.	1.9	5
41	<i>Metabacillus schmidteae</i> sp. nov., Cultivated from Planarian <i>Schmidtea mediterranea</i> Microbiota. <i>Microbiology Research</i> , 2021, 12, 299-316.	1.9	9
42	<i>Bartonella</i> infections diagnosed in the French reference center, 2014â€”2019, and focus on infections in the immunocompromised. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 2407-2410.	2.9	9
43	Genomic description and characterization of <i>Nigeribacterium massiliense</i> gen. nov., sp. nov., isolated from the human gut. <i>Microbes and Infection</i> , 2021, 23, 104842.	1.9	0
44	Evaluation of factors influencing tick bites and tick-borne infections: a longitudinal study. <i>Parasites and Vectors</i> , 2021, 14, 289.	2.5	3
45	<i>Chryseobacterium schmidteae</i> sp. nov. a novel bacterial species isolated from planarian <i>Schmidtea mediterranea</i> . <i>Scientific Reports</i> , 2021, 11, 11002.	3.3	10
46	Tick borne relapsing fever after travelling to a Greek island. <i>Journal of Travel Medicine</i> , 2021, , .	3.0	0
47	Is minor surgery safe during the COVID-19 pandemic? A multi-disciplinary study. <i>PLoS ONE</i> , 2021, 16, e0251122.	2.5	3
48	<i>Lagierella massiliensis</i> gen. nov., sp. nov., Isolated from a Stool Sample. <i>Current Microbiology</i> , 2021, 78, 2481-2487.	2.2	2
49	LamPORE SARS-CoV-2 diagnosis and genotyping: A preliminary report. <i>Journal of Clinical Virology</i> , 2021, 138, 104815.	3.1	5
50	NMNI editorial report, 2020. <i>New Microbes and New Infections</i> , 2021, 41, 100859.	1.6	0
51	Viral RNA in City Wastewater as a Key Indicator of COVID-19 Recrudescence and Containment Measures Effectiveness. <i>Frontiers in Microbiology</i> , 2021, 12, 664477.	3.5	22
52	New human-associated species of the family Atopobiaceae and proposal to reclassify members of the genus <i>Olsenella</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	1.7	46
53	Emergence and outcomes of the SARS-CoV-2 â€”Marseille-4â€” variant. <i>International Journal of Infectious Diseases</i> , 2021, 106, 228-236.	3.3	44
54	Implementation of an in-house real-time reverse transcription-PCR assay for the rapid detection of the SARS-CoV-2 Marseille-4 variant. <i>Journal of Clinical Virology</i> , 2021, 139, 104814.	3.1	27

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55	Contagion Management at the Mediterranean Infection University Hospital Institute. <i>Journal of Clinical Medicine</i> , 2021, 10, 2627.	2.4	1
56	Rapid Isothermal Amplification for the Buccal Detection SARS-CoV-2 in the Context of Out-Patient COVID-19 Screening. <i>Journal of Clinical Medicine</i> , 2021, 10, 2643.	2.4	0
57	<i>Vitreoscilla massiliensis</i> sp. nov., Isolated From the Stool of an Amazonian Patient. <i>Current Microbiology</i> , 2021, 78, 3313-3320.	2.2	4
58	Pangenome analysis and virulence profiling of <i>Streptococcus intermedius</i> . <i>BMC Genomics</i> , 2021, 22, 522.	2.8	7
59	Implementation of an in-house real-time reverse transcription-PCR assay to detect the emerging SARS-CoV-2 N501Y variants. <i>Journal of Clinical Virology</i> , 2021, 140, 104868.	3.1	15
60	Draft genome of <i>Metabacillus niabensis</i> strain 4T19T isolated from cotton-waste composts for mushroom cultivation. <i>New Microbes and New Infections</i> , 2021, 42, 100894.	1.6	0
61	SARS-CoV-2 variant from India to Marseille: The still active role of ports in the introduction of epidemics. <i>Travel Medicine and Infectious Disease</i> , 2021, 42, 102085.	3.0	11
62	Draft Genome Sequence of <i>Bacillus velezensis</i> Strain Marseille-Q1230, Isolated from a Stool Sample from a Severely Malnourished Child. <i>Microbiology Resource Announcements</i> , 2021, 10, e0051421.	0.6	3
63	Gut Microbiota in Military International Travelers with Doxycycline Malaria Prophylaxis: Towards the Risk of a Simpson Paradox in the Human Microbiome Field. <i>Pathogens</i> , 2021, 10, 1063.	2.8	2
64	<i>Gemella massiliensis</i> sp. nov., a new bacterium isolated from the human sputum. <i>Archives of Microbiology</i> , 2021, 203, 5817-5823.	2.2	2
65	Whole Genome Sequencing of SARS-CoV-2 Strains in COVID-19 Patients From Djibouti Shows Novel Mutations and Clades Replacing Over Time. <i>Frontiers in Medicine</i> , 2021, 8, 737602.	2.6	4
66	Spreading of a new SARS-CoV-2 N501Y spike variant in a new lineage. <i>Clinical Microbiology and Infection</i> , 2021, 27, 1352.e1-1352.e5.	6.0	30
67	<i>Limosilactobacillus caccae</i> sp. nov., a new bacterial species isolated from the human gut microbiota. <i>FEMS Microbiology Letters</i> , 2021, 368, .	1.8	1
68	Clinical outcomes in patients infected with different SARS-CoV-2 variants at one hospital during three phases of the COVID-19 epidemic in Marseille, France. <i>Infection, Genetics and Evolution</i> , 2021, 95, 105092.	2.3	22
69	<i>Collinsella ihumii</i> sp. nov., a new anaerobic bacterium isolated from human stool. <i>Archives of Microbiology</i> , 2021, 203, 6315-6322.	2.2	6
70	Real-time whole genome sequencing direct diagnosis of <i>Streptococcus pneumoniae</i> meningitis: A case report. <i>Journal of Infection</i> , 2021, , .	3.3	2
71	Pneumococcal pneumonia among shipyard workers: Inside the features of disease onset. <i>Travel Medicine and Infectious Disease</i> , 2021, 44, 102183.	3.0	1
72	Using MALDI-TOF spectra in epidemiological surveillance for the detection of bacterial subgroups with a possible epidemic potential. <i>BMC Infectious Diseases</i> , 2021, 21, 1109.	2.9	4

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73	SARS-CoV-2 Vaccination and Protection Against Clinical Disease: A Retrospective Study, Bouches-du-Rhône District, Southern France, 2021. <i>Frontiers in Microbiology</i> , 2021, 12, 796807.	3.5	5
74	Analysis of SARS-CoV-2 Variants From 24,181 Patients Exemplifies the Role of Globalization and Zoonosis in Pandemics. <i>Frontiers in Microbiology</i> , 2021, 12, 786233.	3.5	46
75	Culturomics revealed the bacterial constituents of the microbiota of a 10-year-old laboratory culture of planarian species <i>S. mediterranea</i> . <i>Scientific Reports</i> , 2021, 11, 24311.	3.3	1
76	Incidence and Outcome of Coinfections with SARS-CoV-2 and Rhinovirus. <i>Viruses</i> , 2021, 13, 2528.	3.3	20
77	Seroepidemiological and molecular investigation of spotted fever group rickettsiae and <i>Coxiella burnetii</i> in Sao Tome Island: A One Health approach. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 36-43.	3.0	12
78	Mediastinitis in the intensive care unit patient: a narrative review. <i>Clinical Microbiology and Infection</i> , 2020, 26, 26-34.	6.0	28
79	Tick-Borne Spotted Fever Rickettsioses. , 2020, , 587-593.		4
80	Early mortality attributable to PICC-lines in 4 public hospitals of Marseille from 2010 to 2016 (Revised) <i>TJ ETQq0 0 0 rgBT /Overlock 10 T</i>	1.0	7
81	Effect of antiplatelet agents on platelet antistaphylococcal capacity: An in vitro study. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105890.	2.5	9
82	Lack of <i>Vibrio cholerae</i> among French pilgrims during the 2017 and 2018 Hajj. <i>Travel Medicine and Infectious Disease</i> , 2020, 36, 101506.	3.0	3
83	Taxono-genomics and description of <i>Haloimpatiens massiliensis</i> sp. nov., a new bacterium isolated from the gut of a healthy infant. <i>New Microbes and New Infections</i> , 2020, 33, 100631.	1.6	1
84	Identification of Noncanonical Transcripts Produced by Systematic Nucleotide Exchanges in HIV-Associated Centroblastic Lymphoma. <i>DNA and Cell Biology</i> , 2020, 39, 1444-1448.	1.9	4
85	Culturing Ancient Bacteria Carrying Resistance Genes from Permafrost and Comparative Genomics with Modern Isolates. <i>Microorganisms</i> , 2020, 8, 1522.	3.6	6
86	Description of <i>Clostridium cagae</i> sp. nov., <i>Clostridium rectalis</i> sp. nov. and <i>Hathewayia massiliensis</i> sp. nov., new anaerobic bacteria isolated from human stool samples. <i>New Microbes and New Infections</i> , 2020, 37, 100719.	1.6	2
87	Genome sequence and description of <i>Urinicoccus timonensis</i> gen. nov., sp. nov., a new bacterium isolated from a human stool sample. <i>New Microbes and New Infections</i> , 2020, 37, 100720.	1.6	1
88	Tularemia: A Case Series of Patients Diagnosed at the National Reference Center for Rickettsioses From 2008 to 2017. <i>Open Forum Infectious Diseases</i> , 2020, 7, ofaa440.	0.9	11
89	Contribution of VitaPCR SARS-CoV-2 to the emergency diagnosis of COVID-19. <i>Journal of Clinical Virology</i> , 2020, 133, 104682.	3.1	31
90	Utilization of Galectins by Pathogens for Infection. <i>Frontiers in Immunology</i> , 2020, 11, 1877.	4.8	18

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91	Type 1 Diabetes in People Hospitalized for COVID-19: New Insights From the CORONADO Study. <i>Diabetes Care</i> , 2020, 43, e174-e177.	8.6	35
92	<i>Fenollaria timonensis</i> sp. nov., A New Bacterium Isolated from Healthy Human Fresh Stool. <i>Current Microbiology</i> , 2020, 77, 3780-3786.	2.2	3
93	Description of <i>Gracilibacillus phocaeensis</i> sp. nov., a new halophilic bacterium isolated from Senegalian human stool. <i>New Microbes and New Infections</i> , 2020, 38, 100799.	1.6	2
94	Investigation of <i>Ctenocephalides felis</i> on domestic dogs and <i>Rickettsia felis</i> infection in the Democratic Republic of Sao Tome and Principe. <i>Zoonoses and Public Health</i> , 2020, 67, 892-902.	2.2	3
95	Early treatment of COVID-19 patients with hydroxychloroquine and azithromycin: A retrospective analysis of 1061 cases in Marseille, France. <i>Travel Medicine and Infectious Disease</i> , 2020, 35, 101738.	3.0	372
96	Interest of bacterial pangenome analyses in clinical microbiology. <i>Microbial Pathogenesis</i> , 2020, 149, 104275.	2.9	12
97	Clinical efficacy of chloroquine derivatives in COVID-19 infection: comparative meta-analysis between the big data and the real world. <i>New Microbes and New Infections</i> , 2020, 38, 100709.	1.6	61
98	<i>Dialister massiliensis</i> sp. nov., a new bacterium isolated from the human gut. <i>New Microbes and New Infections</i> , 2020, 34, 100657.	1.6	3
99	<i>Anaerococcus marasmi</i> sp. nov., a new bacterium isolated from human gut microbiota. <i>New Microbes and New Infections</i> , 2020, 35, 100655.	1.6	2
100	COVID-19 virus and children: What do we know?. <i>Archives De Pediatrie</i> , 2020, 27, 117-118.	1.0	59
101	Systematic Nucleotide Exchange Analysis of ESTs From the Human Cancer Genome Project Report: Origins of 347 Unknown ESTs Indicate Putative Transcription of Non-Coding Genomic Regions. <i>Frontiers in Genetics</i> , 2020, 11, 42.	2.3	2
102	<i>Clostridium massiliamazoniense</i> sp. nov., New Bacterial Species Isolated from Stool Sample of a Volunteer Brazilian. <i>Current Microbiology</i> , 2020, 77, 2008-2015.	2.2	10
103	NMNI editorial report, 2019. <i>New Microbes and New Infections</i> , 2020, 35, 100662.	1.6	0
104	Detailed description of <i>Senegalia massiliensis</i> strain SIT17T, a bacterium isolated from the human gut. <i>New Microbes and New Infections</i> , 2020, 37, 100700.	1.6	0
105	Outcomes of 3,737 COVID-19 patients treated with hydroxychloroquine/azithromycin and other regimens in Marseille, France: A retrospective analysis. <i>Travel Medicine and Infectious Disease</i> , 2020, 36, 101791.	3.0	209
106	Description of <i>Prevotella rectalis</i> sp. nov., a new bacterium isolated from human rectum. <i>New Microbes and New Infections</i> , 2020, 36, 100703.	1.6	2
107	<i>Gorillibacterium timonense</i> sp. nov., isolated from an obese patient. <i>Archives of Microbiology</i> , 2020, 202, 1223-1229.	2.2	1
108	<i>Pseudoruminococcus massiliensis</i> gen. nov., sp. nov., a new bacterium isolated from the human gut microbiota. <i>New Microbes and New Infections</i> , 2020, 34, 100645.	1.6	2



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109	Actinomycetoma Caused by <i>Actinomadura mexicana</i> , A Neglected Entity in the Caribbean. <i>Emerging Infectious Diseases</i> , 2020, 26, 379-380.	4.3	5
110	Epidemiological serosurvey of vector-borne and zoonotic pathogens among homeless people living in shelters in Marseille: cross-sectional one-day surveys (2005–2015). <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020, 39, 1663-1672.	2.9	8
111	<i>Massilistercora timonensis</i> gen. nov., sp. nov., a new bacterium isolated from the human microbiota. <i>New Microbes and New Infections</i> , 2020, 35, 100664.	1.6	3
112	Clinical and microbiological effect of a combination of hydroxychloroquine and azithromycin in 80 COVID-19 patients with at least a six-day follow up: A pilot observational study. <i>Travel Medicine and Infectious Disease</i> , 2020, 34, 101663.	3.0	605
113	<i>Pedobacter schmidtea</i> sp. nov., a new bacterium isolated from the microbiota of the planarian <i>Schmidtea mediterranea</i> . <i>Scientific Reports</i> , 2020, 10, 6113.	3.3	15
114	High-Content Screening, a Reliable System for <i>Coxiella burnetii</i> Isolation from Clinical Samples. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	3.9	3
115	Survival of <i>Rickettsia conorii</i> in artificially contaminated whole and leukoreduced canine blood units during the storage period. <i>Parasites and Vectors</i> , 2020, 13, 118.	2.5	4
116	Draft genome and description of <i>Negativicoccus massiliensis</i> strain Marseille-P2082, a new species isolated from the gut microbiota of an obese patient. <i>Antonie Van Leeuwenhoek</i> , 2020, 113, 997-1008.	1.7	0
117	Dynamics and genetic diversity of <i>Haemophilus influenzae</i> carriage among French pilgrims during the 2018 Hajj: A prospective cohort survey. <i>Travel Medicine and Infectious Disease</i> , 2020, 38, 101883.	3.0	4
118	Genome sequence-based criteria for demarcation and definition of species in the genus <i>Rickettsia</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 1738-1750.	1.7	22
119	<i>Chitinophaga vietnamensis</i> sp. nov., a multi-drug resistant bacterium infecting humans. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 1758-1768.	1.7	16
120	Phenotypic and molecular characterization of macrolide resistance mechanisms among <i>Streptococcus pneumoniae</i> isolated in Tunisia. <i>Journal of Medical Microbiology</i> , 2020, 69, 505-520.	1.8	6
121	Draft Genome Sequence of <i>Salirhabdus euzebyi</i> Strain Q1438. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	9
122	Case Report: <i>Vibrio cholerae</i> Biliary Tract Infections in Two North Africans in France. <i>American Journal of Tropical Medicine and Hygiene</i> , 2020, 102, 1306-1308.	1.4	3
123	<i>Vaginimicrobium propionicum</i> gen. nov., sp. nov., a novel propionic acid bacterium derived from human vaginal discharge. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 4091-4097.	1.7	11
124	Multispacer sequence typing of <i>Coxiella burnetii</i> from milk and hard tick samples from ruminant farms in Lebanon. <i>Veterinaria Italiana</i> , 2020, 56, 289-296.	0.5	2
125	Genome sequence and description of <i>Gracilibacillus timonensis</i> sp. nov. strain Marseille-P2481T, a moderate halophilic bacterium isolated from the human gut microflora. <i>MicrobiologyOpen</i> , 2019, 8, e00638.	3.0	8
126	Description of three new <i>Peptoniphilus</i> species cultured in the vaginal fluid of a woman diagnosed with bacterial vaginosis: <i>Peptoniphilus pacaensis</i> sp. nov., <i>Peptoniphilus raoultii</i> sp. nov., and <i>Peptoniphilus vaginalis</i> sp. nov.. <i>MicrobiologyOpen</i> , 2019, 8, e00661.	3.0	20



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127	Human pegivirus isolates characterized by deep sequencing from hepatitis C virus RNA and human immunodeficiency virus RNA positive blood donations, France. <i>Journal of Medical Virology</i> , 2019, 91, 38-44.	5.0	9
128	<i>Arcanobacterium ihumii</i> sp. nov., <i>Varibaculum vaginae</i> sp. nov. and <i>Tessaracoccus timonensis</i> sp. nov., isolated from vaginal swabs from healthy Senegalese women. <i>New Microbes and New Infections</i> , 2019, 31, 100585.	1.6	20
129	Draft genome and description of <i>Chryseobacterium phocaeense</i> sp. nov.: a new bacterial species isolated from the sputum of a cystic fibrosis patient. <i>Archives of Microbiology</i> , 2019, 201, 1361-1368.	2.2	0
130	<i>Varibaculum timonense</i> sp. nov., a new bacterial species isolated from human stool sample. <i>New Microbes and New Infections</i> , 2019, 31, 100575.	1.6	9
131	<i>Corynebacterium urinapleomorphum</i> sp. nov., a new bacterial species isolated from human urine sample. <i>New Microbes and New Infections</i> , 2019, 31, 100576.	1.6	17
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281	â€ˆ <i>Bacteroides ihuae</i> â€™ sp. nov., a new bacterial species isolated from the human respiratory microbiome. <i>New Microbes and New Infections</i> , 2017, 17, 52-53.	1.6	3
282	â€ˆ <i>Actinomyces provencensis</i> â€™ sp. nov., â€ˆ <i>Corynebacterium bouchesdurhonense</i> â€™ sp. nov., â€ˆ <i>Corynebacterium provencense</i> â€™ sp. nov. and â€ˆ <i>Xanthomonas massiliensis</i> â€™ sp. nov., 4 new species isolated from fresh stools of obese French patients. <i>New Microbes and New Infections</i> , 2017, 18, 24-27.	1.6	4
283	â€ˆ <i>Urmitella timonensis</i> â€™ gen. nov., sp. nov., â€ˆ <i>Blautia marasmii</i> â€™ sp. nov., â€ˆ <i>Lachnoclostridium pacaense</i> â€™ sp. nov., â€ˆ <i>Bacillus marasmii</i> â€™ sp. nov. and â€ˆ <i>Anaerotruncus rubiinfantis</i> â€™ sp. nov., isolated from stool samples of undernourished African children. <i>New Microbes and New Infections</i> , 2017, 17, 84-88.	1.6	40
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298	New Laboratory Tools for Emerging Bacterial Challenges. <i>Clinical Infectious Diseases</i> , 2017, 65, S39-S49.	5.8	3
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301	â€œ <i>Veillonella massiliensis</i> â€™, a new anaerobic species isolated from human colostrum. <i>Human Microbiome Journal</i> , 2017, 4, 20-21.	3.8	5
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317	<i>Bartonella henselae</i> is usually not viable in lymph nodes of patients with cat scratch disease. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2017, 36, 2207-2213.	2.9	13
318	Discovery of a new species within <i>Bacillus</i> genus in a stool sample from Gabon: <i>Bacillus massilogabonensis</i> sp. nov.. <i>New Microbes and New Infections</i> , 2017, 15, 3-5.	1.6	2
319	<i>Advances in Diagnostic Microbiology</i> , 2017, , 1386-1389.e1.		0
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323	Blood Culture–Negative Endocarditis, Morocco. <i>Emerging Infectious Diseases</i> , 2017, 23, 1908-1909.	4.3	5
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340	<i>Collinsella vaginalis</i> sp. nov., a new bacterial species cultivated from human female genital tract. <i>Human Microbiome Journal</i> , 2016, 2, 19-20.	3.8	1
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345	Noncontiguous finished genome sequence and description of <i>Bacillus andreraoultii</i> strain SIT1T sp. nov.. <i>New Microbes and New Infections</i> , 2016, 10, 25-35.	1.6	2
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363	<i>Corynebacterium phoceense</i> sp. nov., strain MC1T a new bacterial species isolated from human urine. <i>New Microbes and New Infections</i> , 2016, 14, 73-82.	1.6	15
364	<i>Clostridium massiliodiemoense</i> sp. nov., a new species isolated from the human gut microbiota. <i>New Microbes and New Infections</i> , 2016, 13, 27-28.	1.6	1
365	<i>Neglecta timonensis</i> gen. nov., sp. nov., a new human-associated species. <i>New Microbes and New Infections</i> , 2016, 13, 13-14.	1.6	8
366	<i>Peptoniphilus raoultii</i> sp. nov., a new species isolated from human female genital tract. <i>New Microbes and New Infections</i> , 2016, 13, 29-31.	1.6	2
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368	<i>Peptoniphilus pacaensis</i> sp. nov., a new species isolated from human female genital tract. <i>New Microbes and New Infections</i> , 2016, 13, 37-39.	1.6	2
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370	<i>Culturomica massiliensis</i> , a new bacterium isolated from the human gut. <i>New Microbes and New Infections</i> , 2016, 13, 60-61.	1.6	4
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