

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	Ongoing Revolution in Bacteriology: Routine Identification of Bacteria by Matrix-Assisted Laser Desorption Ionization Time-of-Flight Mass Spectrometry. <i>Clinical Infectious Diseases</i> , 2009, 49, 543-551.	5.8	1,638
2	Update on Tick-Borne Rickettsioses around the World: a Geographic Approach. <i>Clinical Microbiology Reviews</i> , 2013, 26, 657-702.	13.6	1,033
3	The Epidemiology and Control of <i>Acinetobacter baumannii</i> in Health Care Facilities. <i>Clinical Infectious Diseases</i> , 2006, 42, 692-699.	5.8	766
4	Culture of previously uncultured members of the human gut microbiota by culturomics. <i>Nature Microbiology</i> , 2016, 1, 16203.	13.3	735
5	Comparative Genomics of Multidrug Resistance in <i>Acinetobacter baumannii</i> . <i>PLoS Genetics</i> , 2006, 2, e7.	3.5	677
6	The Rebirth of Culture in Microbiology through the Example of Culturomics To Study Human Gut Microbiota. <i>Clinical Microbiology Reviews</i> , 2015, 28, 237-264.	13.6	605
7	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
8	Diagnosis of Q Fever. <i>Journal of Clinical Microbiology</i> , 1998, 36, 1823-1834.	3.9	529
9	Culturing the human microbiota and culturomics. <i>Nature Reviews Microbiology</i> , 2018, 16, 540-550.	28.6	521
10	Q Fever 1985-1998: Clinical and Epidemiologic Features of 1,383 Infections. <i>Medicine (United States)</i> , 2000, 79, 109-123.	1.0	459
11	Cultivation of the <i>Bacillus</i> of Whipple's Disease. <i>New England Journal of Medicine</i> , 2000, 342, 620-625.	27.0	458
12	Comprehensive Diagnostic Strategy for Blood Culture-Negative Endocarditis: A Prospective Study of 819 New Cases. <i>Clinical Infectious Diseases</i> , 2010, 51, 131-140.	5.8	418
13	Mechanisms of Evolution in <i>Rickettsia conorii</i> and <i>R. prowazekii</i> . <i>Science</i> , 2001, 293, 2093-2098.	12.6	408
14	Differentiation of spotted fever group rickettsiae by sequencing and analysis of restriction fragment length polymorphism of PCR-amplified DNA of the gene encoding the protein rOmpA. <i>Journal of Clinical Microbiology</i> , 1996, 34, 2058-2065.	3.9	388
15	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
16	Clinical features and prognostic factors of listeriosis: the MONALISA national prospective cohort study. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 510-519.	9.1	366
17	Identification of Rare Pathogenic Bacteria in a Clinical Microbiology Laboratory: Impact of Matrix-Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2182-2194.	3.9	362
18	Gene Sequence-Based Criteria for Identification of New <i>Rickettsia</i> Isolates and Description of <i>Rickettsia heilongjiangensis</i> sp. nov. <i>Journal of Clinical Microbiology</i> , 2003, 41, 5456-5465.	3.9	347

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19	A polyphasic strategy incorporating genomic data for the taxonomic description of novel bacterial species. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 384-391.	1.7	330
20	Guidelines for the diagnosis of tick-borne bacterial diseases in Europe. <i>Clinical Microbiology and Infection</i> , 2004, 10, 1108-1132.	6.0	328
21	Diagnosis of 22 New Cases of Bartonella Endocarditis. <i>Annals of Internal Medicine</i> , 1996, 125, 646.	3.9	321
22	MALDI-TOF-mass spectrometry applications in clinical microbiology. <i>Future Microbiology</i> , 2010, 5, 1733-1754.	2.0	310
23	Genome Sequence of Rickettsia bellii Illuminates the Role of Amoebae in Gene Exchanges between Intracellular Pathogens. <i>PLoS Genetics</i> , 2006, 2, e76.	3.5	286
24	<i>Rickettsia africae</i> , a Tick-Borne Pathogen in Travelers to Sub-Saharan Africa. <i>New England Journal of Medicine</i> , 2001, 344, 1504-1510.	27.0	282
25	Bacterial strain typing in the genomic era. <i>FEMS Microbiology Reviews</i> , 2009, 33, 892-916.	8.6	278
26	Phylogenetic analysis of spotted fever group rickettsiae by study of the outer surface protein rOmpA. <i>International Journal of Systematic Bacteriology</i> , 1998, 48, 839-849.	2.8	274
27	Value and limitations of the duke criteria for the diagnosis of infective endocarditis. <i>Journal of the American College of Cardiology</i> , 1999, 33, 2023-2029.	2.8	271
28	Risks Factors and Prevention of Q Fever Endocarditis. <i>Clinical Infectious Diseases</i> , 2001, 33, 312-316.	5.8	264
29	Modern clinical microbiology: new challenges and solutions. <i>Nature Reviews Microbiology</i> , 2013, 11, 574-585.	28.6	264
30	Characterization of the Naturally Occurring Oxacillinase of <i>Acinetobacter baumannii</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 4174-4179.	3.2	254
31	A comprehensive repertoire of prokaryotic species identified in human beings. <i>Lancet Infectious Diseases</i> , The, 2015, 15, 1211-1219.	9.1	250
32	The Genome Sequence of <i>Rickettsia felis</i> Identifies the First Putative Conjugative Plasmid in an Obligate Intracellular Parasite. <i>PLoS Biology</i> , 2005, 3, e248.	5.6	242
33	The bacterial pangenome as a new tool for analysing pathogenic bacteria. <i>New Microbes and New Infections</i> , 2015, 7, 72-85.	1.6	241
34	Warmer Weather Linked to Tick Attack and Emergence of Severe Rickettsioses. <i>PLoS Neglected Tropical Diseases</i> , 2008, 2, e338.	3.0	228
35	Rickettsioses and the International Traveler. <i>Clinical Infectious Diseases</i> , 2004, 39, 1493-1499.	5.8	222
36	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

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37	Modification of the diagnostic criteria proposed by the duke endocarditis service to permit improved diagnosis of q fever endocarditis. American Journal of Medicine, 1996, 100, 629-633.	1.5	207
38	Epidemiologic and Clinical Characteristics of Bartonella quintana and Bartonella henselae Endocarditis. Medicine (United States), 2001, 80, 245-251.	1.0	207
39	A Flea-Associated Rickettsia Pathogenic for Humans. Emerging Infectious Diseases, 2001, 7, 73-81.	4.3	207
40	Significant Molecular and Systemic Adaptations after Repeated Sprint Training in Hypoxia. PLoS ONE, 2013, 8, e56522.	2.5	206
41	Outcome and Treatment of Bartonella Endocarditis. Archives of Internal Medicine, 2003, 163, 226.	3.8	202
42	African tick bite fever. Lancet Infectious Diseases, The, 2003, 3, 557-564.	9.1	199
43	Description of Tropheryma whipplei gen. nov., sp. nov., the Whipple's disease bacillus.. International Journal of Systematic and Evolutionary Microbiology, 2001, 51, 1471-1479.	1.7	187
44	Evidence of Infection in Humans with Rickettsia helvetica in Eastern France. Emerging Infectious Diseases, 2000, 6, 389-392.	4.3	186
45	Comparison of PCR and Serology Assays for Early Diagnosis of Acute Q Fever. Journal of Clinical Microbiology, 2003, 41, 5094-5098.	3.9	182
46	Spotless Rickettsiosis Caused by <i>Rickettsia slovaca</i> and Associated with <i>Dermacentor</i> Ticks. Clinical Infectious Diseases, 2002, 34, 1331-1336.	5.8	177
47	African Tick Bite Fever in Travelers to Rural Sub-Equatorial Africa. Clinical Infectious Diseases, 2003, 36, 1411-1417.	5.8	177
48	Genotyping, Orientalis-like <i>Yersinia pestis</i> , and Plague Pandemics. Emerging Infectious Diseases, 2004, 10, 1585-1592.	4.3	166
49	Complementarity between targeted real-time specific PCR and conventional broad-range 16S rDNA PCR in the syndrome-driven diagnosis of infectious diseases. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 561-570.	2.9	166
50	Cautionary tale of using 16S rRNA gene sequence similarity values in identification of human-associated bacterial species. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 1929-1934.	1.7	161
51	rpoB Gene Sequence-Based Identification of Aerobic Gram-Positive Cocci of the Genera Streptococcus , Enterococcus , Gemella , Abiotrophia , and Granulicatella. Journal of Clinical Microbiology, 2004, 42, 497-504.	3.9	157
52	Mimivirus and the emerging concept of "giant" virus. Virus Research, 2006, 117, 133-144.	2.2	157
53	Diagnosis and management of Q fever—United States, 2013: recommendations from CDC and the Q Fever Working Group. MMWR Recommendations and Reports, 2013, 62, 1-30.	61.1	157
54	Evidence for Louse-Transmitted Diseases in Soldiers of Napoleon's Grand Army in Vilnius. Journal of Infectious Diseases, 2006, 193, 112-120.	4.0	155

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55	First Documented Human <i>Rickettsia aeschlimannii</i> Infection. <i>Emerging Infectious Diseases</i> , 2002, 8, 748-749.	4.3	153
56	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. <i>FEMS Microbiology Reviews</i> , 2010, 34, 281-294.	8.6	153
57	HACEK Infective Endocarditis: Characteristics and Outcomes from a Large, Multi-National Cohort. <i>PLoS ONE</i> , 2013, 8, e63181.	2.5	148
58	Molecular Detection of <i>Coxiella burnetii</i> in the Sera of Patients with Q Fever Endocarditis or Vascular Infection. <i>Journal of Clinical Microbiology</i> , 2004, 42, 4919-4924.	3.9	147
59	<i>Rickettsia raoultii</i> sp. nov., a spotted fever group rickettsia associated with Dermacentor ticks in Europe and Russia. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 1635-1639.	1.7	146
60	Spotted fever rickettsioses in southern and eastern Europe. <i>FEMS Immunology and Medical Microbiology</i> , 2007, 49, 2-12.	2.7	141
61	Aneruptive Fever Associated with Antibodies to <i>Rickettsia helvetica</i> in Europe and Thailand. <i>Journal of Clinical Microbiology</i> , 2004, 42, 816-818.	3.9	136
62	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
63	<i>Rickettsia felis</i> Infection Acquired in Europe and Documented by Polymerase Chain Reaction. <i>Emerging Infectious Diseases</i> , 2002, 8, 207-208.	4.3	134
64	<i>Coxiella burnetii</i> infection of aortic aneurysms or vascular grafts: report of 30 new cases and evaluation of outcome. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2007, 26, 635-640.	2.9	132
65	Lymphangitis-Associated Rickettsiosis, a New Rickettsiosis Caused by <i>Rickettsia sibirica mongolotimonae</i> : Seven New Cases and Review of the Literature. <i>Clinical Infectious Diseases</i> , 2005, 40, 1435-1444.	5.8	131
66	Influence of the Timing of Cardiac Surgery on the Outcome of Patients With Infective Endocarditis and Stroke. <i>Clinical Infectious Diseases</i> , 2013, 56, 209-217.	5.8	130
67	Asynchrony between the rates of standing height gain and bone mass accumulation during puberty. <i>Osteoporosis International</i> , 1997, 7, 525-532.	3.1	128
68	Proposal to create subspecies of <i>Rickettsia conorii</i> based on multi-locus sequence typing and an emended description of <i>Rickettsia conorii</i> . <i>BMC Microbiology</i> , 2005, 5, 11.	3.3	127
69	Diagnosis of infectious endocarditis in patients undergoing valve surgery. <i>American Journal of Medicine</i> , 2005, 118, 230-238.	1.5	126
70	Genome sequence and description of <i>Anaerosalibacter massiliensis</i> sp. nov.. <i>New Microbes and New Infections</i> , 2016, 10, 66-76.	1.6	125
71	Human Pathogens in Body and Head Lice. <i>Emerging Infectious Diseases</i> , 2002, 8, 1515-1518.	4.3	123
72	Burden of emerging anaerobes in the MALDI-TOF and 16S rRNA gene sequencing era. <i>Anaerobe</i> , 2011, 17, 106-112.	2.1	122

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73	Acute Tick-borne Rickettsiosis Caused by <i>Rickettsia sibirica</i> in the Russian Far East. <i>Emerging Infectious Diseases</i> , 2004, 10, 810-817.	4.3	121
74	Molecular Evolution of Rickettsia Surface Antigens: Evidence of Positive Selection. <i>Molecular Biology and Evolution</i> , 2005, 22, 2073-2083.	8.9	119
75	From culturomics to taxonomogenomics: A need to change the taxonomy of prokaryotes in clinical microbiology. <i>Anaerobe</i> , 2015, 36, 73-78.	2.1	119
76	Prognosis and Treatment of Paraquat Poisoning: A Review of 28 Cases. <i>Journal of Toxicology: Clinical Toxicology</i> , 1982, 19, 461-474.	1.5	117
77	HIV-2 integrase gene polymorphism and phenotypic susceptibility of HIV-2 clinical isolates to the integrase inhibitors raltegravir and elvitegravir in vitro. <i>Journal of Antimicrobial Chemotherapy</i> , 2008, 62, 914-920.	3.0	115
78	Non-steroidal anti-inflammatory drugs for athletes: An update. <i>Annals of Physical and Rehabilitation Medicine</i> , 2010, 53, 278-288.	2.3	115
79	Genetic Identification of Rickettsiae Isolated from Ticks in Japan. <i>Journal of Clinical Microbiology</i> , 2002, 40, 2176-2181.	3.9	114
80	Selfish DNA in Protein-Coding Genes of <i>Rickettsia</i> . <i>Science</i> , 2000, 290, 347-350.	12.6	113
81	The human gut microbiome, a taxonomic conundrum. <i>Systematic and Applied Microbiology</i> , 2015, 38, 276-286.	2.8	113
82	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. <i>FEMS Microbiology Reviews</i> , 2010, 34, 281-94.	8.6	112
83	Current Knowledge on Phylogeny and Taxonomy of <i>Rickettsia</i> spp.. <i>Annals of the New York Academy of Sciences</i> , 2009, 1166, 1-11.	3.8	111
84	Clinical detection and characterization of bacterial pathogens in the genomics era. <i>Genome Medicine</i> , 2014, 6, 114.	8.2	111
85	Blood culture-negative endocarditis. <i>Medicine (United States)</i> , 2017, 96, e8392.	1.0	110
86	Bartonella, a Common Cause of Endocarditis: a Report on 106 Cases and Review. <i>Journal of Clinical Microbiology</i> , 2015, 53, 824-829.	3.9	109
87	Analysis of the <i>Rickettsia africae</i> genome reveals that virulence acquisition in Rickettsia species may be explained by genome reduction. <i>BMC Genomics</i> , 2009, 10, 166.	2.8	107
88	Respiratory Viruses and Bacteria among Pilgrims during the 2013 Hajj. <i>Emerging Infectious Diseases</i> , 2014, 20, 1821-1827.	4.3	107
89	Non-contiguous finished genome sequence and description of <i>Anaerococcus senegalensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 6, 116-125.	1.5	106
90	Update on blood culture-negative endocarditis. <i>Médecine Et Maladies Infectieuses</i> , 2015, 45, 1-8.	5.0	106

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91	Mink, SARS-CoV-2, and the Human-Animal Interface. <i>Frontiers in Microbiology</i> , 2021, 12, 663815.	3.5	106
92	Outbreak of <i>Rickettsia africae</i> Infections in Participants of an Adventure Race in South Africa. <i>Clinical Infectious Diseases</i> , 1998, 27, 316-323.	5.8	105
93	Isolation of <i>Rickettsia parkeri</i> and Identification of a Novel Spotted Fever Group <i>Rickettsia</i> sp. from Gulf Coast Ticks (<i>Amblyomma maculatum</i>) in the United States. <i>Applied and Environmental Microbiology</i> , 2010, 76, 2689-2696.	3.1	104
94	A Concise Review of the Epidemiology and Diagnostics of Rickettsioses: <i>Rickettsia</i> and <i>Orientia</i> spp. <i>Journal of Clinical Microbiology</i> , 2018, 56, .	3.9	103
95	What does the future hold for clinical microbiology?. <i>Nature Reviews Microbiology</i> , 2004, 2, 151-159.	28.6	97
96	<i>Coxiella burnetii</i> Infection of Aneurysms or Vascular Grafts: Report of Seven Cases and Review. <i>Clinical Infectious Diseases</i> , 1998, 26, 116-121.	5.8	95
97	Genomic Variation of <i>Bartonella henselae</i> Strains Detected in Lymph Nodes of Patients with Cat Scratch Disease. <i>Journal of Clinical Microbiology</i> , 2002, 40, 1023-1030.	3.9	95
98	Use of Genome Selected Repeated Sequences Increases the Sensitivity of PCR Detection of <i>Tropheryma whippelii</i> . <i>Journal of Clinical Microbiology</i> , 2004, 42, 401-403.	3.9	93
99	Metagenomic Analysis of Brain Abscesses Identifies Specific Bacterial Associations. <i>Clinical Infectious Diseases</i> , 2012, 54, 202-210.	5.8	93
100	Non contiguous-finished genome sequence and description of <i>Peptoniphilus timonensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 7, 1-11.	1.5	93
101	Naming of <i>Rickettsiae</i> and <i>Rickettsial</i> Diseases. <i>Annals of the New York Academy of Sciences</i> , 2005, 1063, 1-12.	3.8	90
102	Diagnosis of <i>Bartonella</i> Endocarditis by a Real-Time Nested PCR Assay Using Serum. <i>Journal of Clinical Microbiology</i> , 2003, 41, 919-925.	3.9	89
103	Non-contiguous finished genome sequence and description of <i>Alistipes timonensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 6, 315-324.	1.5	89
104	Advantages and limitations of genomics in prokaryotic taxonomy. <i>Clinical Microbiology and Infection</i> , 2013, 19, 790-795.	6.0	87
105	Description of <i>Mediterraneibacter massiliensis</i> , gen. nov., sp. nov., a new genus isolated from the gut microbiota of an obese patient and reclassification of <i>Ruminococcus faecis</i> , <i>Ruminococcus lactaris</i> , <i>Ruminococcus torques</i> , <i>Ruminococcus gnavus</i> and <i>Clostridium glycyrrhizinilyticum</i> as <i>Mediterraneibacter faecis</i> comb. nov., <i>Mediterraneibacter lactaris</i> comb. nov., <i>Mediterraneibacter torques</i> comb. nov., <i>Mediterraneibacter gnavus</i> comb. nov. and <i>Mediterraneibacter glycyrrhizinilyticus</i> comb. nov.. <i>Antonie Van Leeuwenhoek</i> , 2018, 111, 2107-2128.	1.7	87
106	Suicide PCR on Skin Biopsy Specimens for Diagnosis of Rickettsioses. <i>Journal of Clinical Microbiology</i> , 2004, 42, 3428-3434.	3.9	86
107	Do we need new antibiotics?. <i>Clinical Microbiology and Infection</i> , 2016, 22, 408-415.	6.0	86
108	Real-time PCR for universal antibiotic susceptibility testing. <i>Journal of Antimicrobial Chemotherapy</i> , 2004, 54, 538-541.	3.0	85

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109	Impact of Early Valve Surgery on Outcome of Staphylococcus aureus Prosthetic Valve Infective Endocarditis: Analysis in the International Collaboration of Endocarditisâ€“Prospective Cohort Study. Clinical Infectious Diseases, 2015, 60, 741-749.	5.8	84
110	Identification of Novel Zoonotic Activity of Bartonella spp., France. Emerging Infectious Diseases, 2016, 22, 457-462.	4.3	84
111	Culture and Immunological Detection of <i>Tropheryma whippelii</i> From the Duodenum of a Patient With Whipple Disease. JAMA - Journal of the American Medical Association, 2001, 285, 1039.	7.4	83
112	Use of Highly Variable Intergenic Spacer Sequences for Multispacer Typing of Rickettsia conorii Strains. Journal of Clinical Microbiology, 2004, 42, 5757-5766.	3.9	83
113	Detection of Members of the Genera Rickettsia, Anaplasma, and Ehrlichia in Ticks Collected in the Asiatic Part of Russia. Annals of the New York Academy of Sciences, 2006, 1078, 378-383.	3.8	83
114	Phylogenetic classification of Bartonella species by comparing groEL sequences.. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 165-171.	1.7	82
115	Genotyping reveals a wide heterogeneity of Tropheryma whipplei. Microbiology (United Kingdom), 2008, 154, 521-527.	1.8	81
116	Quantitative Detection of <i>Tropheryma whipplei</i> DNA by Real-Time PCR. Journal of Clinical Microbiology, 2002, 40, 1119-1120.	3.9	80
117	An Experimental Model of Human Body Louse Infection with Rickettsia prowazekii. Journal of Infectious Diseases, 2002, 186, 1639-1646.	4.0	80
118	Rickettsia tamurae sp. nov., isolated from Amblyomma testudinarium ticks. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1673-1675.	1.7	80
119	Bacterial genome sequencing and its use in infectious diseases. Lancet Infectious Diseases, The, 2007, 7, 711-723.	9.1	79
120	Genotyping of Human Lice Suggests Multiple Emergences of Body Lice from Local Head Louse Populations. PLoS Neglected Tropical Diseases, 2010, 4, e641.	3.0	79
121	Clinical Features and Complications of <i>Coxiella burnetii</i> Infections From the French National Reference Center for Q Fever. JAMA Network Open, 2018, 1, e181580.	5.9	77
122	Myocarditis, a Rare but Severe Manifestation of Q Fever: Report of 8 Cases and Review of the Literature. Clinical Infectious Diseases, 2001, 32, 1440-1447.	5.8	76
123	Genome sequence and description of Aeromicrobium massiliense sp. nov.. Standards in Genomic Sciences, 2012, 7, 246-257.	1.5	76
124	Quantitative Analysis of Valvular Lesions During Bartonella Endocarditis. American Journal of Clinical Pathology, 2000, 114, 880-889.	0.7	75
125	Systematic PCR Detection in Culture-negative Osteoarticular Infections. American Journal of Medicine, 2013, 126, 1143.e25-1143.e33.	1.5	75
126	<i>Tropheryma whipplei</i> Endocarditis. Emerging Infectious Diseases, 2013, 19, 1721-30.	4.3	75

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127	<i>Candidatus Rickettsia tarasevichiae</i> ™ in <i>Ixodes persulcatus</i> Ticks Collected in Russia. <i>Annals of the New York Academy of Sciences</i> , 2003, 990, 162-172.	3.8	72
128	Non-contiguous finished genome sequence and description of <i>Clostridium senegalense</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 6, 401-410.	1.5	70
129	Non contiguous-finished genome sequence and description of <i>Bacillus timonensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 6, 346-355.	1.5	70
130	Non-contiguous finished genome sequence and description of <i>Herbaspirillum massiliense</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 7, 1-14.	1.5	70
131	<i>Lactobacillus rhamnosus</i> bacteremia: an emerging clinical entity. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2012, 31, 2469-2480.	2.9	70
132	Diagnosis of blood culture-negative endocarditis and clinical comparison between blood culture-negative and blood culture-positive cases. <i>Infection</i> , 2016, 44, 459-466.	4.7	70
133	Genomics and metagenomics in medical microbiology. <i>Journal of Microbiological Methods</i> , 2013, 95, 415-424.	1.6	69
134	The role of <i>Streptococcus intermedius</i> in brain abscess. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2013, 32, 477-483.	2.9	69
135	Value of Microimmunofluorescence for Diagnosis and Follow-up of <i>Bartonella</i> Endocarditis. <i>Vaccine Journal</i> , 2002, 9, 795-801.	3.1	67
136	Nosocomial Infections Caused by <i>Acinetobacter baumannii</i> A Major Threat Worldwide. <i>Infection Control and Hospital Epidemiology</i> , 2006, 27, 645-646.	1.8	67
137	Non-contiguous finished genome sequence and description of <i>Paenibacillus senegalensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 7, 70-81.	1.5	66
138	Non contiguous-finished genome sequence and description of <i>Cellulomonas massiliensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 7, 258-270.	1.5	66
139	Non contiguous-finished genome sequence and description of <i>Senegalemassilia anaerobia</i> gen. nov., sp. nov.. <i>Standards in Genomic Sciences</i> , 2013, 7, 343-356.	1.5	66
140	Non contiguous-finished genome sequence and description of <i>Enterobacter massiliensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2013, 7, 399-412.	1.5	66
141	Detection of a <i>Rickettsia</i> Closely Related to <i>Rickettsia aeschlimannii</i> , <i>Rickettsia heilongjiangensis</i> , <i>Rickettsia</i> sp. Strain RpA4, and <i>Ehrlichia muris</i> in Ticks Collected in Russia and Kazakhstan. <i>Journal of Clinical Microbiology</i> , 2004, 42, 2221-2223.	3.9	65
142	Non-contiguous finished genome sequence and description of <i>Brevibacterium senegalense</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 7, 233-245.	1.5	64
143	Tick-borne rickettsioses in international travellers. <i>International Journal of Infectious Diseases</i> , 2004, 8, 139-146.	3.3	63
144	Strikes, flooding, rats, and leptospirosis in Marseille, France. <i>International Journal of Infectious Diseases</i> , 2011, 15, e710-e715.	3.3	63

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145	Detection and Culture of Bartonella quintana, Serratia marcescens , and Acinetobacter spp. from Decontaminated Human Body Lice. Journal of Clinical Microbiology, 2001, 39, 1707-1709.	3.9	61
146	Rapid identification of Legionella species by mass spectrometry. Journal of Medical Microbiology, 2010, 59, 273-284.	1.8	61
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290	Endocarditis Caused by <i>Cardiobacterium valvarum</i> . <i>Journal of Clinical Microbiology</i> , 2006, 44, 657-658.	3.9	24
291	<i>Bartonella australis</i> sp. nov. from Kangaroos, Australia. <i>Emerging Infectious Diseases</i> , 2007, 13, 1961-1963.	4.3	24
292	<i>Mycobacterium setense</i> Infection in Humans. <i>Emerging Infectious Diseases</i> , 2008, 14, 1330-1332.	4.3	24
293	Association of Soluble CD14 and Inflammatory Biomarkers With HIV-2 Disease Progression. <i>Clinical Infectious Diseases</i> , 2012, 55, 1417-1425.	5.8	24
294	Non-contiguous finished genome sequence and description of <i>Bacillus massilioanorexius</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2013, 8, 465-479.	1.5	24
295	Non contiguous-finished genome sequence and description of <i>Microbacterium gorillae</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2016, 11, 32.	1.5	24
296	Rickettsial genomics and the paradigm of genome reduction associated with increased virulence. <i>Microbes and Infection</i> , 2018, 20, 401-409.	1.9	24
297	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
298	Non contiguous-finished genome sequence and description of <i>Peptoniphilus obesi</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013, 7, 357-69.	1.5	24
299	First Isolation and Detection by Immunofluorescence Assay of <i>Bartonella koehlerae</i> in Erythrocytes from a French Cat. <i>Journal of Clinical Microbiology</i> , 2003, 41, 4001-4002.	3.9	23
300	A deadly aversion to pork. <i>Lancet</i> , The, 2011, 377, 1542.	13.7	23
301	Non-contiguous finished genome sequence and description of <i>Kallipyga massiliensis</i> gen. nov., sp. nov., a new member of the family <i>Clostridiales</i> Incertae Sedis XI. <i>Standards in Genomic Sciences</i> , 2013, 8, 500-515.	1.5	23
302	From Culturomics to Clinical Microbiology and Forward. <i>Emerging Infectious Diseases</i> , 2018, 24, 1683-1690.	4.3	23
303	Paradoxical evolution of rickettsial genomes. <i>Ticks and Tick-borne Diseases</i> , 2019, 10, 462-469.	2.7	23
304	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
305	Methyl iodide poisoning: Report of two cases. , 1996, 30, 759-764.		22
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308	Non contiguous-finished genome sequence and description of <i>Bacillus jeddahensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2015, 10, 47.	1.5	22
309	â€˜Enterococcus timonensisâ€™ TM sp. nov., â€˜Actinomyces marseillensisâ€™ TM sp. nov., â€˜Leptotrichia massiliensisâ€™ TM sp. nov., â€˜Actinomyces pacaensisâ€™ TM sp. nov., â€˜Actinomyces oralisâ€™ TM sp. nov., â€˜Actinomyces culturomiciâ€™ TM sp. nov. and â€˜Gemella massiliensisâ€™ TM sp. nov., new bacterial species isolated from the human respiratory microbiome. <i>New Microbes and New Infections</i> , 2018, 22, 37-43.	1.6	22
310	Nine Cases of Methanogenic Archaea in Refractory Sinusitis, an Emerging Clinical Entity. <i>Frontiers in Public Health</i> , 2019, 7, 38.	2.7	22
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312	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
313	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
314	Quantitative Histological Examination of Mechanical Heart Valves. <i>Clinical Infectious Diseases</i> , 2005, 40, 655-661.	5.8	21
315	Detection of <i>Rickettsia aeschlimannii</i> in <i>Hyalomma marginatum</i> ticks in western Russia. <i>Clinical Microbiology and Infection</i> , 2009, 15, 315-316.	6.0	21
316	Link between endocarditis on porcine bioprosthetic valves and allergy to pork. <i>International Journal of Cardiology</i> , 2013, 167, 600-602.	1.7	21
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319	A New Highly Sensitive and Specific Real-Time PCR Assay Targeting the Malate Dehydrogenase Gene of <i>Kingella kingae</i> and Application to 201 Pediatric Clinical Specimens. <i>Journal of Clinical Microbiology</i> , 2018, 56, .	3.9	21
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322	Viral Communities Associated with Human Pericardial Fluids in Idiopathic Pericarditis. <i>PLoS ONE</i> , 2014, 9, e93367.	2.5	21
323	First isolation of <i>Rickettsia conorii</i> from humans in the Trakya (European) region of Turkey. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2004, 23, 609-14.	2.9	20
324	Non-contiguous finished genome sequence and description of <i>Bartonella senegalensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2013, 8, 279-289.	1.5	20

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326	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
327	<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
328	Incidence and Outcome of Coinfections with SARS-CoV-2 and Rhinovirus. <i>Viruses</i> , 2021, 13, 2528.	3.3	20
329	Comparison of the Effect of Two Prazepam Formulations on Psychomotor and Cognitive Function in Healthy Volunteers. <i>International Clinical Psychopharmacology</i> , 1990, 5, 57-64.	1.7	19
330	Isolated Right-sided <i>Bartonella quintana</i> endocarditis in an Immunocompetent Adult. <i>Scandinavian Journal of Infectious Diseases</i> , 1998, 30, 424-425.	1.5	19
331	Sequential changes in hematologic and biochemical parameters in African tick bite fever. <i>Clinical Microbiology and Infection</i> , 2003, 9, 678-683.	6.0	19
332	Phylogenetic Study of <i>Rickettsia</i> Species Using Sequences of the Autotransporter Protein-Encoding Gene <i>sca2</i> . <i>Annals of the New York Academy of Sciences</i> , 2005, 1063, 94-99.	3.8	19
333	Transmission of <i>Aeromonas hydrophila</i> by leeches. <i>Lancet, The</i> , 2013, 381, 1686.	13.7	19
334	Hand, Foot and Mouth Disease and <i>Kingella kingae</i> Infections. <i>Pediatric Infectious Disease Journal</i> , 2015, 34, 547-548.	2.0	19
335	Noncontiguous finished genome sequence and description of <i>Raoultibacter massiliensis</i> gen. nov., sp. nov. and <i>Raoultibacter timonensis</i> sp. nov., two new bacterial species isolated from the human gut. <i>MicrobiologyOpen</i> , 2019, 8, e00758.	3.0	19
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337	Histamine food poisoning: a sudden, large outbreak linked to fresh yellowfin tuna from Reunion Island, France, April 2017. <i>Eurosurveillance</i> , 2019, 24, .	7.0	19
338	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
339	Enhancement of propoxyphene bioavailability by ethanol. <i>European Journal of Clinical Pharmacology</i> , 1991, 41, 147-52.	1.9	18
340	Evidence for the Presence of <i>Francisella</i> and Spotted Fever Group <i>Rickettsia</i> DNA in the Tick <i>Amblyomma fimbriatum</i> (Acari: Ixodidae), Northern Territory, Australia. <i>Journal of Medical Entomology</i> , 2009, 46, 926-933.	1.8	18
341	Presence or absence of plasmid in <i>Rickettsia felis</i> depending on the source of fleas. <i>Clinical Microbiology and Infection</i> , 2009, 15, 296-297.	6.0	18
342	Human rickettsioses in the Batna area, eastern Algeria. <i>Ticks and Tick-borne Diseases</i> , 2012, 3, 364-366.	2.7	18

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344	Absence of antibodies to <i>Rickettsia</i> spp., <i>Bartonella</i> spp., <i>Ehrlichia</i> spp. and <i>Coxiella burnetii</i> in Tahiti, French Polynesia. BMC Infectious Diseases, 2014, 14, 255.	2.9	18
345	Genome sequence and description of <i>Nesterenkonia massiliensis</i> sp. nov. strain NP1T. Standards in Genomic Sciences, 2014, 9, 866-882.	1.5	18
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347	Prospective Comparison of Infective Endocarditis in Khon Kaen, Thailand and Rennes, France. American Journal of Tropical Medicine and Hygiene, 2015, 92, 871-874.	1.4	18
348	High-quality genome sequence and description of <i>Bacillus dielmoensis</i> strain FF4T sp. nov.. Standards in Genomic Sciences, 2015, 10, 41.	1.5	18
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350	<i>Hugonella massiliensis</i> gen. nov., sp. nov., genome sequence, and description of a new strictly anaerobic bacterium isolated from the human gut. MicrobiologyOpen, 2017, 6, e00458.	3.0	18
351	<i>Anaerotruncus massiliensis</i> sp. nov., a succinate-producing bacterium isolated from human stool from an obese patient after bariatric surgery. New Microbes and New Infections, 2019, 29, 100508.	1.6	18
352	Utilization of Galectins by Pathogens for Infection. Frontiers in Immunology, 2020, 11, 1877.	4.8	18
353	The severity of the first 207 infections with the SARS-CoV-2 Omicron BA.2 variant, in Marseille, France, December 2021–February 2022. Journal of Medical Virology, 2022, 94, 3494-3497.	5.0	18
354	Acute tubulointerstitial nephritis complicating Legionnaires' disease: a case report. Journal of Medical Case Reports, 2012, 6, 100.	0.8	17
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356	Noncontiguous finished genome sequence and description of <i>Paenibacillus ihumii</i> sp. nov. strain AT5. New Microbes and New Infections, 2016, 10, 142-150.	1.6	17
357	<i>Libanicoccus massiliensis</i> gen. nov., sp. nov., a new bacterium isolated from human stool. New Microbes and New Infections, 2018, 21, 63-71.	1.6	17
358	<i>Corynebacterium urinaleomorphum</i> sp. nov., a new bacterial species isolated from human urine sample. New Microbes and New Infections, 2019, 31, 100576.	1.6	17
359	DNA Persistence and Relapses Questions on the Treatment Strategies of Enterococcus Infections of Prosthetic Valves. PLoS ONE, 2012, 7, e53335.	2.5	17
360	A SURVEY FOR SPOTTED FEVER GROUP RICKETTSIAE AND EHRLICHIAE IN AMBLYOMMA VARIEGATUM FROM ST. KITTS AND NEVIS. American Journal of Tropical Medicine and Hygiene, 2003, 69, 58-59.	1.4	17

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362	Tick-Transmitted Infections in Transvaal: Consider <i>Rickettsia africae</i> . <i>Emerging Infectious Diseases</i> , 1999, 5, 178-181.	4.3	16
363	African Tick Bite Fever in a Taiwanese Traveler Returning from South Africa: Molecular and Serologic Studies. <i>American Journal of Tropical Medicine and Hygiene</i> , 2009, 81, 735-739.	1.4	16
364	<i>Clostridium difficile</i> O27 Emerging Outbreak in Marseille, France. <i>Infection Control and Hospital Epidemiology</i> , 2013, 34, 1339-1341.	1.8	16
365	Genome sequence of <i>Oceanobacillus picturae</i> strain S1, an halophilic bacterium first isolated in human gut. <i>Standards in Genomic Sciences</i> , 2015, 10, 91.	1.5	16
366	<i>Christensenella massiliensis</i> , a new bacterial species isolated from the human gut. <i>New Microbes and New Infections</i> , 2016, 12, 69-70.	1.6	16
367	Outbreaks of Invasive <i>Kingella kingae</i> Infections in Daycare Facilities: Approach to Investigation and Management. <i>Journal of Pediatrics</i> , 2017, 182, 14-20.	1.8	16
368	Microbial Culturomics Broadens Human Vaginal Flora Diversity: Genome Sequence and Description of <i>Prevotella lascolaii</i> sp. nov. Isolated from a Patient with Bacterial Vaginosis. <i>OMICS A Journal of Integrative Biology</i> , 2018, 22, 210-222.	2.0	16
369	<i>Varibaculum massiliense</i> sp. nov., a new bacterium isolated from human urine with culturomics. <i>New Microbes and New Infections</i> , 2019, 32, 100591.	1.6	16
370	<i>Miniphocibacter massiliensis</i> gen. nov., sp. nov., a new species isolated from the human gut and its taxonomic description. <i>MicrobiologyOpen</i> , 2019, 8, e00735.	3.0	16
371	<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
372	Pericardial Effusion in a Homeless Man Due to <i>Bartonella quintana</i> . <i>Journal of Clinical Microbiology</i> , 2003, 41, 5291-5293.	3.9	15
373	Genome Sequence of <i>Rickettsia conorii</i> subsp. <i>indica</i> , the Agent of Indian Tick Typhus. <i>Journal of Bacteriology</i> , 2012, 194, 3288-3289.	2.2	15
374	The first reported cases of Q fever endocarditis in Thailand. <i>Gastroenterology Insights</i> , 2012, 4, e7.	1.2	15
375	<i>Bartonella henselae</i> Endocarditis in Laos – “The Unsought Will Go Undetected”. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3385.	3.0	15
376	Non-contiguous finished genome sequence and description of <i>Clostridium saudii</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2014, 9, 8.	1.5	15
377	Rise of Microbial Culturomics: Noncontiguous Finished Genome Sequence and Description of <i>Beduini massiliensis</i> gen. nov., sp. nov.. <i>OMICS A Journal of Integrative Biology</i> , 2015, 19, 766-776.	2.0	15
378	<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

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380	â€ˆCollinsella phocaeensisâ€™ sp. nov., â€ˆClostridium merdaeâ€™ sp. nov., â€ˆSutterella massiliensisâ€™ sp. nov., â€ˆSutturella timonensisâ€™ sp. nov., â€ˆEnorma phocaeensisâ€™ sp. nov., â€ˆMailhella massiliensisâ€™ gen. nov., sp. nov., â€ˆMordavella massiliensisâ€™ gen. nov., sp. nov. and â€ˆMassiliprevotella massiliensisâ€™ gen. nov., sp. nov., 9 new species isolated from fresh stool samples of healthy French patients. <i>New Microbes and New Infections</i> , 2017, 17, 89-95.	1.6	15
381	â€ˆKhelaifiella massiliensisâ€™, â€ˆNiameybacter massiliensisâ€™, â€ˆBrachybacterium massilienseâ€™, â€ˆEnterobacter timonensisâ€™, â€ˆMassilibacillus massiliensisâ€™, new bacterial species and genera isolated from the gut microbiota of healthy infants. <i>New Microbes and New Infections</i> , 2017, 19, 1-7.	1.6	15
382	Description and genomic characterization of <i>Massiliimalia massiliensis</i> gen. nov., sp. nov., and <i>Massiliimalia timonensis</i> gen. nov., sp. nov., two new members of the family Ruminococcaceae isolated from the human gut. <i>Antonie Van Leeuwenhoek</i> , 2019, 112, 905-918.	1.7	15
383	<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
384	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
385	Molecular identification of a collection of spotted Fever group rickettsiae obtained from patients and ticks from Russia. <i>American Journal of Tropical Medicine and Hygiene</i> , 2006, 74, 440-3.	1.4	15
386	Diversity of <i>Borrelia burgdorferi</i> Sensu Lato in Russian Far East. <i>Microbiology and Immunology</i> , 2005, 49, 191-197.	1.4	14
387	Tick Paralysis by <i>Ixodes holocyclus</i> in a Japanese Traveler Returning from Australia Associated with <i>Rickettsia helvetica</i> Infection. <i>Journal of Travel Medicine</i> , 2006, 10, 61-63.	3.0	14
388	Paleomicrobiology of <i>Bartonella</i> infections. <i>Microbes and Infection</i> , 2015, 17, 879-883.	1.9	14
389	Noncontiguous finished genome sequence and description of <i>Planococcus massiliensis</i> sp. nov., a moderately halophilic bacterium isolated from the human gut. <i>New Microbes and New Infections</i> , 2016, 10, 36-46.	1.6	14
390	<i>Anaerococcus rubiinfantis</i> sp. nov., isolated from the gut microbiota of a Senegalese infant with severe acute malnutrition. <i>Anaerobe</i> , 2016, 40, 85-94.	2.1	14
391	New Species Announcement: a new format to prompt the description of new human microbial species. <i>New Microbes and New Infections</i> , 2017, 15, 136-137.	1.6	14
392	<i>Olegusella massiliensis</i> gen. nov., sp. nov., strain KHD7 T, a new bacterial genus isolated from the female genital tract of a patient with bacterial vaginosis. <i>Anaerobe</i> , 2017, 44, 87-95.	2.1	14
393	â€ˆArcanobacterium urinimassilienseâ€™ sp. nov., a new bacterium isolated from the urogenital tract. <i>New Microbes and New Infections</i> , 2017, 18, 15-17.	1.6	14
394	â€ˆLachnoclostridium urinimassilienseâ€™ sp. nov. and â€ˆLachnoclostridium phocaeenseâ€™ sp. nov., two new bacterial species isolated from human urine after kidney transplantation. <i>New Microbes and New Infections</i> , 2017, 16, 73-75.	1.6	14
395	Paediatric Bone and Joint Infections in French Guiana: A 6 Year Retrospective Review. <i>Journal of Tropical Pediatrics</i> , 2017, 63, 380-388.	1.5	14
396	<i>Peptoniphilus lacydonensis</i> sp. nov., a new human-associated species isolated from a patient with chronic refractory sinusitis. <i>New Microbes and New Infections</i> , 2018, 23, 61-69.	1.6	14

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398	Characterization of a novel Gram-stain-positive anaerobic coccus isolated from the female genital tract: Genome sequence and description of <i>Murdochiella vaginalis</i> sp. nov.. <i>MicrobiologyOpen</i> , 2018, 7, e00570.	3.0	14
399	Emergence of <i>Clostridium difficile</i> tcdC variant 078 in Marseille, France. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2017, 36, 1971-1974.	2.9	14
400	Arterio-Venous Plasma Concentration Differences in Amitriptyline Overdose. <i>Journal of Toxicology: Clinical Toxicology</i> , 1985, 23, 391-406.	1.5	13
401	Molecular detection of <i>Bartonella quintana</i> DNA in the dental pulp of a homeless patient. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2004, 23, 920-2.	2.9	13
402	Deciphering Genomic Virulence Traits of a <i>Staphylococcus epidermidis</i> Strain Causing Native-Valve Endocarditis. <i>Journal of Clinical Microbiology</i> , 2013, 51, 1617-1621.	3.9	13
403	Diagnosis of human granulocytic anaplasmosis in Belgium by combining molecular and serological methods. <i>New Microbes and New Infections</i> , 2014, 2, 177-178.	1.6	13
404	Non-contiguous finished genome sequence and description of <i>Collinsella massiliensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2014, 9, 1144-1158.	1.5	13
405	Description of <i>Bacillus dakarensis</i> sp. nov., <i>Bacillus sinesaloumensis</i> sp. nov., <i>Gracilibacillus timonensis</i> sp. nov., <i>Halobacillus massiliensis</i> sp. nov., <i>Lentibacillus massiliensis</i> sp. nov., <i>Oceanobacillus senegalensis</i> sp. nov., <i>Oceanobacillus timonensis</i> sp. nov., <i>Virgibacillus dakarensis</i> sp. nov. and <i>Virgibacillus marseillensis</i> sp. nov., nine halophilic new species isolated from human stool. <i>New Microbes and New Infections</i> , 2017, 17, 45-51.	1.6	13
406	An Outbreak of <i>Kingella Kingae</i> Infections Complicating a Severe Hand, Foot, And Mouth Disease Outbreak in Nice, France, 2016. <i>Pediatric Infectious Disease Journal</i> , 2017, 36, 530-532.	2.0	13
407	<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
408	Characterization of a New <i>Ezakiella</i> Isolated from the Human Vagina: Genome Sequence and Description of <i>Ezakiella massiliensis</i> sp. nov.. <i>Current Microbiology</i> , 2018, 75, 456-463.	2.2	13
409	Draft genome and description of <i>Merdibacter massiliensis</i> gen.nov., sp. nov., a new bacterium genus isolated from the human ileum. <i>Scientific Reports</i> , 2019, 9, 7931.	3.3	13
410	<i>Yersinia pestis</i> Genotyping. <i>Emerging Infectious Diseases</i> , 2005, 11, 1318-1319.	4.3	13
411	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
412	VALPROATE POISONING. <i>Lancet, The</i> , 1982, 320, 97.	18.7	12
413	A Case of <i>Salmonella enteritidis</i> Septicemia Complicated by Disseminated Intravascular Coagulation, Severe Hepatitis, Rhabdomyolysis and Acute Renal Failure. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 1999, 18, 830-831.	2.9	12
414	Isolation of Mycobacteria from Clinical Samples Using the Centrifugation-Shell Vial Technique. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2000, 19, 69-70.	2.9	12

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415	Introduction of the Exotic Tick <i>Amblyomma hebraeum</i> into Florida on a Human Host. <i>Journal of Parasitology</i> , 2002, 88, 800-801.	0.7	12
416	Tick Paralysis by <i>Ixodes holocyclus</i> in a Japanese Traveler Returning from Australia. <i>Annals of the New York Academy of Sciences</i> , 2003, 990, 357-358.	3.8	12
417	Genomic Comparison of <i>Kingella kingae</i> Strains. <i>Journal of Bacteriology</i> , 2012, 194, 5972-5972.	2.2	12
418	The first reported cases of Q fever endocarditis in Thailand. <i>Gastroenterology Insights</i> , 2012, 4, 7.	1.2	12
419	Update on Rickettsioses in Slovakia. <i>Acta Virologica</i> , 2013, 57, 180-199.	0.8	12
420	Genome Sequence of <i>Bacillus simplex</i> Strain P558, Isolated from a Human Fecal Sample. <i>Genome Announcements</i> , 2014, 2, .	0.8	12
421	<i>Clostridium tetani</i> Osteitis without Tetanus. <i>Emerging Infectious Diseases</i> , 2014, 20, 1571-1573.	4.3	12
422	Noncontiguous finished genome sequence and description of <i>Weeksella massiliensis</i> sp. nov.. <i>New Microbes and New Infections</i> , 2015, 8, 89-98.	1.6	12
423	Meningitis caused by <i>Pasteurella multocida</i> in a dog owner without a dog bite: clonal lineage identification by MALDI-TOF mass spectrometry. <i>BMC Research Notes</i> , 2015, 8, 626.	1.4	12
424	<i>Colidextribacter massiliensis</i> gen. nov., sp. nov., isolated from human right colon. <i>New Microbes and New Infections</i> , 2017, 17, 27-29.	1.6	12
425	<i>Inediibacterium massiliense</i> gen. nov., sp. nov., a new bacterial species isolated from the gut microbiota of a severely malnourished infant. <i>Antonie Van Leeuwenhoek</i> , 2017, 110, 737-750.	1.7	12
426	Endocarditis in the Mediterranean Basin. <i>New Microbes and New Infections</i> , 2018, 26, S43-S51.	1.6	12
427	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
428	Interest of bacterial pangenome analyses in clinical microbiology. <i>Microbial Pathogenesis</i> , 2020, 149, 104275.	2.9	12
429	<i>Collinsella vaginalis</i> sp. nov. strain Marseille-P2666T, a new member of the <i>Collinsella</i> genus isolated from the genital tract of a patient suffering from bacterial vaginosis. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 949-956.	1.7	12
430	Simultaneous quantitation of glafenine and its major metabolites, glafenic and hydroxyglafenic acid, by high-performance liquid chromatography. <i>Biomedical Applications</i> , 1981, 224, 348-352.	1.7	11
431	Biological Evaluation of Hemoperfusion in Acute Poisoning. <i>Clinical Toxicology</i> , 1981, 18, 1213-1223.	0.5	11
432	<i>Grimontia hollisae</i> , a potential agent of gastroenteritis and bacteraemia in the Mediterranean area. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2009, 28, 705-707.	2.9	11

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434	Non-contiguous finished genome sequence and description of <i>Anaerococcus vaginalis</i> . <i>Standards in Genomic Sciences</i> , 2012, 6, 356-365.	1.5	11
435	Non-contiguous finished genome sequence and description of <i>Corynebacterium jeddahense</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2014, 9, 987-1002.	1.5	11
436	Non-contiguous finished genome sequence and description of <i>Clostridium thumii</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2015, 10, 63.	1.5	11
437	“ <i>Olsenella urinifantis</i> ”, a new bacterial species isolated from a urine sample of a 26-day-old boy suffering from gastroesophageal reflux. <i>Human Microbiome Journal</i> , 2016, 2, 17-18.	3.8	11
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439	Microbial Cultuomics to Map Halophilic Bacterium in Human Gut: Genome Sequence and Description of <i>Oceanobacillus jeddahense</i> sp. nov.. <i>OMICS A Journal of Integrative Biology</i> , 2016, 20, 248-258.	2.0	11
440	Noncontiguous finished genome sequence and description of <i>Gabonia massiliensis</i> gen. nov., sp. nov.. <i>New Microbes and New Infections</i> , 2016, 9, 35-44.	1.6	11
441	Description of “ <i>Gorbachella massiliensis</i> ”™ gen. nov., sp. nov., “ <i>Fenollaria timonensis</i> ”™ sp. nov., “ <i>Intestinimonas timonensis</i> ”™ sp. nov. and “ <i>Collinsella ihuae</i> ”™ sp. nov. isolated from healthy fresh stools with cultuomics. <i>New Microbes and New Infections</i> , 2017, 16, 60-62.	1.6	11
442	<i>Olsenella provencensis</i> sp. nov., <i>Olsenella phocaeensis</i> sp. nov., and <i>Olsenella mediterranea</i> sp. nov. isolated from the human colon. <i>Human Microbiome Journal</i> , 2017, 4, 22-23.	3.8	11
443	Description of “ <i>Blautia phocaeensis</i> ”™ sp. nov. and “ <i>Lachnoclostridium edouardi</i> ”™ sp. nov., isolated from healthy fresh stools of Saudi Arabia Bedouins by cultuomics. <i>New Microbes and New Infections</i> , 2017, 19, 129-131.	1.6	11
444	Multi-omics Analysis Sheds Light on the Evolution and the Intracellular Lifestyle Strategies of Spotted Fever Group <i>Rickettsia</i> spp.. <i>Frontiers in Microbiology</i> , 2017, 8, 1363.	3.5	11
445	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
446	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
447	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
448	<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
449	Searching for <i>Bacillus anthracis</i> in Suspect Powders: a French Experience. <i>Journal of Clinical Microbiology</i> , 2003, 41, 524-525.	3.9	10
450	Intraspecies Diversity of <i>Rickettsia conorii</i> . <i>Journal of Infectious Diseases</i> , 2009, 199, 1097-1098.	4.0	10

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452	A case of multifocal chronic Q fever osteomyelitis. <i>Infection</i> , 2011, 39, 167-169.	4.7	10
453	Non contiguous-finished genome sequence and description of <i>Enorma timonensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2014, 9, 970-986.	1.5	10
454	Genome sequence and description of <i>Corynebacterium ihumii</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2014, 9, 1128-1143.	1.5	10
455	Genome sequence and description of <i>Bacteroides timonensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2014, 9, 1181-1197.	1.5	10
456	Noncontiguous finished genome sequence and description of <i>Virgibacillus massiliensis</i> sp. nov., a moderately halophilic bacterium isolated from human gut. <i>New Microbes and New Infections</i> , 2015, 8, 78-88.	1.6	10
457	High-quality genome sequence and description of <i>Bacillus ndiopicus</i> strain FF3T sp. nov.. <i>New Microbes and New Infections</i> , 2015, 8, 154-163.	1.6	10
458	<i>Numidum massiliense</i> gen. nov., sp. nov., a new member of the Bacillaceae family isolated from the human gut. <i>New Microbes and New Infections</i> , 2016, 12, 76-85.	1.6	10
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460	Primary subacute epiphyseal osteomyelitis caused by <i>Mycobacterium</i> species in young children: a modern diagnostic approach. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2017, 36, 771-777.	2.9	10
461	A modified multilocus sequence typing protocol to genotype <i>Kingella kingae</i> from oropharyngeal swabs without bacterial isolation. <i>BMC Microbiology</i> , 2017, 17, 200.	3.3	10
462	â€œ <i>Cellulomonas timonensis</i> â€™™ sp. nov., taxonogenomics description of a new bacterial species isolated from human gut. <i>New Microbes and New Infections</i> , 2018, 23, 7-16.	1.6	10
463	Genomic analysis of a <i>Raoultella ornithinolytica</i> strain causing prosthetic joint infection in an immunocompetent patient. <i>Scientific Reports</i> , 2018, 8, 9462.	3.3	10
464	<i>Colibacter massiliensis</i> gen. nov. sp. nov., a novel Gram-stain-positive anaerobic diplococcal bacterium, isolated from the human left colon. <i>Scientific Reports</i> , 2019, 9, 17199.	3.3	10
465	Draft genome and description of <i>Cohnella massiliensis</i> sp. nov., a new bacterial species isolated from the blood culture of a hemodialysis patient. <i>Archives of Microbiology</i> , 2019, 201, 305-312.	2.2	10
466	<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
467	<i>Chryseobacterium schmidtea</i> sp. nov. a novel bacterial species isolated from planarian <i>Schmidtea mediterranea</i> . <i>Scientific Reports</i> , 2021, 11, 11002.	3.3	10
468	Cat-Scratch Disease and an Overview of Other <i>Bartonella henselae</i> -Related Infections. , 1998, 1, 32-62.		9

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470	Genome Sequence of <i>Rickettsia conorii</i> subsp. <i>israelensis</i> , the Agent of Israeli Spotted Fever. <i>Journal of Bacteriology</i> , 2012, 194, 5130-5131.	2.2	9
471	Genome Sequence of <i>Diplorickettsia massiliensis</i> , an Emerging <i>Ixodes ricinus</i> -Associated Human Pathogen. <i>Journal of Bacteriology</i> , 2012, 194, 3287-3287.	2.2	9
472	Leptospirosis in a French Traveler Returning From Mauritius: Table 1. <i>Journal of Travel Medicine</i> , 2012, 19, 69-71.	3.0	9
473	<i>Actinomyces gerencseriae</i> hip prosthesis infection: a case report. <i>Journal of Medical Case Reports</i> , 2015, 9, 223.	0.8	9
474	Increasing burden of urinary tract infections due to intrinsic colistin-resistant bacteria in hospitals in Marseille, France. <i>International Journal of Antimicrobial Agents</i> , 2015, 45, 144-150.	2.5	9
475	<i>Anaerococcus urinomassiliensis</i> sp. nov., isolated from a urine sample of a 17-year-old boy affected by autoimmune hepatitis and membranoproliferative glomerulonephritis. <i>New Microbes and New Infections</i> , 2016, 13, 56-58.	1.6	9
476	<i>Butyricimonas phoceensis</i> sp. nov., a new anaerobic species isolated from the human gut microbiota of a French morbidly obese patient. <i>New Microbes and New Infections</i> , 2016, 14, 38-48.	1.6	9
477	<i>Emergencia timonensis</i> , a new bacterium isolated from the stool of a healthy patient. <i>New Microbes and New Infections</i> , 2016, 12, 73-75.	1.6	9
478	<i>Ndongobacter massiliensis</i> gen. nov., sp. nov., a new bacterial genus isolated from a human urine sample after de novo kidney transplantation. <i>New Microbes and New Infections</i> , 2017, 16, 34-36.	1.6	9
479	<i>Actinotignum timonense</i> sp. nov., a new bacterial species isolated from a human urine sample. <i>New Microbes and New Infections</i> , 2017, 16, 47-48.	1.6	9
480	<i>Lactobacillus timonensis</i> sp. nov., a new bacterial species isolated from the human gut. <i>New Microbes and New Infections</i> , 2017, 19, 121-122.	1.6	9
481	Implementation and Initial Analysis of a Laboratory-Based Weekly Biosurveillance System, Provence-Alpes-Côte d'Azur, France. <i>Emerging Infectious Diseases</i> , 2017, 23, 582-589.	4.3	9
482	Naming microorganisms: the contribution of the IHU Méditerranée Infection, Marseille, France. <i>New Microbes and New Infections</i> , 2018, 26, S89-S95.	1.6	9
483	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
484	<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
485	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
486	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

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488	<i>Bartonella</i> infections diagnosed in the French reference center, 2014–2019, and focus on infections in the immunocompromised. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 2407-2410.	2.9	9
489	Draft Genome Sequence of <i>Salirhabdus euzebyi</i> Strain Q1438. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	9
490	Bacteriology, Taxonomy, and Phylogeny of <i>Rickettsia</i> . <i>Infectious Disease and Therapy</i> , 2007, , 1-14.	0.0	9
491	Diversity and distribution of ticks from domestic ruminants in Lebanon. <i>Veterinaria Italiana</i> , 2017, 53, 147-155.	0.5	9
492	First isolation of <i>Rickettsia conorii</i> from human blood in Croatia. <i>Croatian Medical Journal</i> , 2003, 44, 630-4.	0.7	9
493	Immunohistochemical detection of <i>Coxiella burnetii</i> in an aortic graft. <i>Clinical Microbiology and Infection</i> , 2009, 15, 171-172.	6.0	8
494	Identification of a new serotype of <i>Rickettsia heilongjiangensis</i> in wild rats from Guangdong Province, China. <i>Clinical Microbiology and Infection</i> , 2009, 15, 338-339.	6.0	8
495	Complete Genome Sequence of <i>Rickettsia slovaca</i> , the Agent of Tick-Borne Lymphadenitis. <i>Journal of Bacteriology</i> , 2012, 194, 1612-1612.	2.2	8
496	Genome Sequence of <i>Rickettsia australis</i> , the Agent of Queensland Tick Typhus. <i>Journal of Bacteriology</i> , 2012, 194, 5129-5129.	2.2	8
497	Diagnosis of <i>Bartonella henselae</i> Prosthetic Valve Endocarditis in Man, France. <i>Emerging Infectious Diseases</i> , 2014, 20, 1396-1397.	4.3	8
498	Noncontiguous finished genome sequence and description of <i>Necropsobacter massiliensis</i> sp. nov.. <i>New Microbes and New Infections</i> , 2015, 8, 41-50.	1.6	8
499	“ <i>Bacillus mediterraneensis</i> ,” a new bacterial species isolated from human gut microbiota. <i>New Microbes and New Infections</i> , 2016, 12, 86-87.	1.6	8
500	<i>Risungbinella massiliensis</i> sp. nov., a new member of Thermoactinomycetaceae isolated from human gut. <i>Antonie Van Leeuwenhoek</i> , 2016, 109, 773-784.	1.7	8
501	Genome Sequence of the Tick-Borne Pathogen <i>Rickettsia raoultii</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	8
502	“ <i>Raoultibacter massiliensis</i> ” gen. nov., sp. nov., a new bacterium isolated from the human gut of a Saudi Bedouin. <i>New Microbes and New Infections</i> , 2016, 14, 1-3.	1.6	8
503	“ <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
504	<i>Raoultella ornithinolytica</i> : An unusual pathogen for prosthetic joint infection. <i>IDCases</i> , 2016, 5, 46-48.	0.9	8

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506	“ <i>Negativibacillus massiliensis</i> ” gen. nov., sp. nov., isolated from human left colon. <i>New Microbes and New Infections</i> , 2017, 17, 36-38.	1.6	8
507	<i>Rubeoparvulum massiliense</i> gen. nov., sp. nov., a new bacterial genus isolated from the human gut of a Senegalese infant with severe acute malnutrition. <i>New Microbes and New Infections</i> , 2017, 15, 49-60.	1.6	8
508	Description of <i>Chryseobacterium timonianum</i> sp. nov., isolated from a patient with pneumonia. <i>Antonie Van Leeuwenhoek</i> , 2017, 110, 1121-1132.	1.7	8
509	“ <i>Angelakisella massiliensis</i> ”™ gen. nov., sp. nov., a new bacterial species isolated from human ileum. <i>New Microbes and New Infections</i> , 2017, 16, 51-53.	1.6	8
510	<i>Mycoplasma genitalium</i> , an agent of reemerging sexually transmitted infections. <i>Apmis</i> , 2017, 125, 916-920.	2.0	8
511	Hemodialysis vascular graft as a focus of persistent Q fever. <i>Infection</i> , 2018, 46, 881-884.	4.7	8
512	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
513	Whole-Genome Sequence of French Clinical <i>Olivibacter jilunii</i> Strain P8502. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	8
514	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
515	Statins potentiate the antibacterial effect of platelets on <i>Staphylococcus aureus</i> . <i>Platelets</i> , 2021, 32, 671-676.	2.3	8
516	Human galectin-1 and galectin-3 promote <i>Tropheryma whipplei</i> infection. <i>Gut Microbes</i> , 2021, 13, 1-15.	9.8	8
517	Planarians (Platyhelminthes) – An Emerging Model Organism for Investigating Innate Immune Mechanisms. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 619081.	3.9	8
518	Non-contiguous finished genome sequence and description of <i>Alistipes timonensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012, 6, 315-324.	1.5	8
519	Direct diagnosis of <i>Pasteurella multocida</i> meningitis using next-generation sequencing. <i>Lancet Microbe</i> , The, 2022, 3, e6.	7.3	8
520	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
521	Judicial Opinions 103–111. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	1.7	8
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524	Sequence and Annotation of <i>Rickettsia sibirica sibirica</i> Genome. Journal of Bacteriology, 2012, 194, 2377-2377.	2.2	7
525	Genome Sequence of <i>Rickettsia sibirica</i> subsp. <i>mongolitimonae</i> . Journal of Bacteriology, 2012, 194, 2389-2390.	2.2	7
526	Genome Sequence of <i>Rickettsia tamurae</i> , a Recently Detected Human Pathogen in Japan. Genome Announcements, 2014, 2, .	0.8	7
527	Genome of a chronic osteitis-causing <i>Clostridium tetani</i> . New Microbes and New Infections, 2014, 2, 25-26.	1.6	7
528	Update on Tick-Borne Rickettsioses around the World: a Geographic Approach. Clinical Microbiology Reviews, 2014, 27, 166-166.	13.6	7
529	Non-contiguous finished genome sequence and description of <i>Paenibacillus gorillae</i> sp. nov.. Standards in Genomic Sciences, 2014, 9, 1031-1045.	1.5	7
530	Two Cases of Leptospirosis in French Travelers Returning From Koh Samui, Thailand. Journal of Travel Medicine, 2015, 22, 419-421.	3.0	7
531	<i>Halomonas massiliensis</i> ™ sp. nov., a new halotolerant bacterium isolated from the human gut. New Microbes and New Infections, 2016, 14, 19-20.	1.6	7
532	Noncontiguous finished genome sequence and description of <i>Kallipyga gabonensis</i> sp. nov.. New Microbes and New Infections, 2016, 9, 15-23.	1.6	7
533	<i>Leibacterium massiliense</i> ™ gen. nov., sp. nov., a new bacterial species isolated from human ileum of a patient with Crohn disease. New Microbes and New Infections, 2017, 17, 25-26.	1.6	7
534	Description of <i>Arabia massiliensis</i> ™ gen. nov., sp. nov., <i>Gordonibacter massiliensis</i> ™ sp. nov., and <i>Bacilliculturomica massiliensis</i> ™ gen. nov., sp. nov., isolated from a faecal specimen of a 50-year-old Saudi Bedouin woman. New Microbes and New Infections, 2017, 19, 87-90.	1.6	7
535	Draft Genome Sequence of <i>Kingella negevensis</i> SW7208426, the First European Strain of <i>K. negevensis</i> Isolated from a Healthy Child in Switzerland. Genome Announcements, 2017, 5, .	0.8	7
536	Draft genome sequence of <i>Fermentimonas caenicola</i> strain SIT8, isolated from the human gut. Standards in Genomic Sciences, 2018, 13, 8.	1.5	7
537	Genome analysis and description of <i>Xanthomonas massiliensis</i> sp. nov., a new species isolated from human faeces. New Microbes and New Infections, 2018, 26, 63-72.	1.6	7
538	<i>Prevotella ihumii</i> sp. nov., a new bacterium isolated from a stool specimen of a healthy woman. New Microbes and New Infections, 2019, 32, 100607.	1.6	7
539	Western Immunoblotting for the Diagnosis of <i>Enterococcus faecalis</i> and <i>Streptococcus gallolyticus</i> Infective Endocarditis. Frontiers in Cellular and Infection Microbiology, 2019, 9, 314.	3.9	7
540	Diversity of Aerobic Bacilli Analysis Using Molecular and Culture-Based Approaches in Debagh Hot Spring. Geomicrobiology Journal, 2019, 36, 137-147.	2.0	7

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542	Genetic diversity of <i>Streptococcus pneumoniae</i> in Tunisia. <i>International Journal of Antimicrobial Agents</i> , 2019, 53, 63-69.	2.5	7
543	Early mortality attributable to PICC-lines in 4 public hospitals of Marseille from 2010 to 2016 (Revised) <i>Tj ETQq1 1 0,784314,rgBT /Over</i>	1.0	7
544	<i>Anaerococcus urinimassiliensis</i> sp. nov., a new bacterium isolated from human urine. <i>Scientific Reports</i> , 2021, 11, 2684.	3.3	7
545	Pangenome analysis and virulence profiling of <i>Streptococcus intermedius</i> . <i>BMC Genomics</i> , 2021, 22, 522.	2.8	7
546	Whole-Genome Sequence of <i>Haloimpatiens lingqiaonensis</i> Strain P8956. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	7
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554	Genomic Comparison of <i>Rickettsia honei</i> Strain RB T and Other <i>Rickettsia</i> Species. <i>Journal of Bacteriology</i> , 2012, 194, 4145-4145.	2.2	6
555	Genome Sequence of <i>Rickettsia gravesii</i> , Isolated from Western Australian Ticks. <i>Genome Announcements</i> , 2013, 1, .	0.8	6
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558	Noncontiguous finished genome sequence and description of <i>Nocardioides massiliensis</i> sp. nov. GD13T. <i>New Microbes and New Infections</i> , 2016, 10, 47-57.	1.6	6

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561	Genome sequence and description of <i>Anaeromassilibacillus senegalensis</i> gen. nov., sp. nov., isolated from the gut of patient with kwashiorkor. <i>New Microbes and New Infections</i> , 2017, 17, 54-64.	1.6	6
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580	Draft Genome Sequence of <i>Coxiella burnetii</i> Strain Cb196, an Agent of Endocarditis in Saudi Arabia. <i>Genome Announcements</i> , 2014, 2, .	0.8	5
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611	Draft Genome Sequence of <i>Rickettsia aeschlimannii</i> , Associated with <i>Hyalomma marginatum</i> Ticks. <i>Genome Announcements</i> , 2014, 2, .	0.8	4
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813	NMNI editorial report, 2020. <i>New Microbes and New Infections</i> , 2021, 41, 100859.	1.6	0
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815	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
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819	A concise overview on tick-borne human infections in Europe: a focus on Lyme borreliosis and tick-borne <i>Rickettsia</i> spp.. <i>Microbiology Australia</i> , 2018, 39, 207.	0.4	0
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821	Draft Genome Sequence of <i>Corynebacterium sanguinis</i> Strain Marseille-P8776. <i>Microbiology Resource Announcements</i> , 2022, 11, e0000822.	0.6	0
822	Culture-negative endocarditis caused by <i>Stenotrophomonas maltophilia</i> : a report of two cases. <i>Future Microbiology</i> , 0, , .	2.0	0