

Pierre E Fournier

List of Publications by Year in descending order

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822
papers

37,597
citations

2963

93
h-index

5101

166
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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.	2.9	1,638
2	Update on Tick-Borne Rickettsioses around the World: a Geographic Approach. <i>Clinical Microbiology Reviews</i> , 2013, 26, 657-702.	5.7	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.	2.9	766
4	Culture of previously uncultured members of the human gut microbiota by culturomics. <i>Nature Microbiology</i> , 2016, 1, 16203.	5.9	735
5	Comparative Genomics of Multidrug Resistance in <i>Acinetobacter baumannii</i> . <i>PLoS Genetics</i> , 2006, 2, e7.	1.5	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.	5.7	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.	1.5	605
8	Diagnosis of Q Fever. <i>Journal of Clinical Microbiology</i> , 1998, 36, 1823-1834.	1.8	529
9	Culturing the human microbiota and culturomics. <i>Nature Reviews Microbiology</i> , 2018, 16, 540-550.	13.6	521
10	Q Fever 1985-1998: Clinical and Epidemiologic Features of 1,383 Infections. <i>Medicine (United States)</i> , 2000, 79, 109-123.	0.4	459
11	Cultivation of the <i>Bacillus</i> of Whipple's Disease. <i>New England Journal of Medicine</i> , 2000, 342, 620-625.	13.9	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.	2.9	418
13	Mechanisms of Evolution in <i>Rickettsia conorii</i> and <i>R. prowazekii</i> . <i>Science</i> , 2001, 293, 2093-2098.	6.0	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.	1.8	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.	1.5	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.	4.6	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.	1.8	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.	1.8	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.	0.8	330
20	Guidelines for the diagnosis of tick-borne bacterial diseases in Europe. <i>Clinical Microbiology and Infection</i> , 2004, 10, 1108-1132.	2.8	328
21	Diagnosis of 22 New Cases of Bartonella Endocarditis. <i>Annals of Internal Medicine</i> , 1996, 125, 646.	2.0	321
22	MALDI-TOF-mass spectrometry applications in clinical microbiology. <i>Future Microbiology</i> , 2010, 5, 1733-1754.	1.0	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.	1.5	286
24	Rickettsia africae, a Tick-Borne Pathogen in Travelers to Sub-Saharan Africa. <i>New England Journal of Medicine</i> , 2001, 344, 1504-1510.	13.9	282
25	Bacterial strain typing in the genomic era. <i>FEMS Microbiology Reviews</i> , 2009, 33, 892-916.	3.9	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.	1.2	271
28	Risks Factors and Prevention of Q Fever Endocarditis. <i>Clinical Infectious Diseases</i> , 2001, 33, 312-316.	2.9	264
29	Modern clinical microbiology: new challenges and solutions. <i>Nature Reviews Microbiology</i> , 2013, 11, 574-585.	13.6	264
30	Characterization of the Naturally Occurring Oxacillinase of Acinetobacter baumannii. <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 4174-4179.	1.4	254
31	A comprehensive repertoire of prokaryotic species identified in human beings. <i>Lancet Infectious Diseases</i> , The, 2015, 15, 1211-1219.	4.6	250
32	The Genome Sequence of Rickettsia felis Identifies the First Putative Conjugative Plasmid in an Obligate Intracellular Parasite. <i>PLoS Biology</i> , 2005, 3, e248.	2.6	242
33	The bacterial pangenome as a new tool for analysing pathogenic bacteria. <i>New Microbes and New Infections</i> , 2015, 7, 72-85.	0.8	241
34	Warmer Weather Linked to Tick Attack and Emergence of Severe Rickettsioses. <i>PLoS Neglected Tropical Diseases</i> , 2008, 2, e338.	1.3	228
35	Rickettsioses and the International Traveler. <i>Clinical Infectious Diseases</i> , 2004, 39, 1493-1499.	2.9	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.	1.5	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.	0.6	207
38	Epidemiologic and Clinical Characteristics of Bartonella quintana and Bartonella henselae Endocarditis. Medicine (United States), 2001, 80, 245-251.	0.4	207
39	A Flea-Associated Rickettsia Pathogenic for Humans. Emerging Infectious Diseases, 2001, 7, 73-81.	2.0	207
40	Significant Molecular and Systemic Adaptations after Repeated Sprint Training in Hypoxia. PLoS ONE, 2013, 8, e56522.	1.1	206
41	Outcome and Treatment of Bartonella Endocarditis. Archives of Internal Medicine, 2003, 163, 226.	4.3	202
42	African tick bite fever. Lancet Infectious Diseases, The, 2003, 3, 557-564.	4.6	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.	0.8	187
44	Evidence of Infection in Humans with Rickettsia helvetica in Eastern France. Emerging Infectious Diseases, 2000, 6, 389-392.	2.0	186
45	Comparison of PCR and Serology Assays for Early Diagnosis of Acute Q Fever. Journal of Clinical Microbiology, 2003, 41, 5094-5098.	1.8	182
46	Spotless Rickettsiosis Caused by Rickettsia slovaca and Associated with Dermacentor Ticks. Clinical Infectious Diseases, 2002, 34, 1331-1336.	2.9	177
47	African Tick Bite Fever in Travelers to Rural Sub-Equatorial Africa. Clinical Infectious Diseases, 2003, 36, 1411-1417.	2.9	177
48	Genotyping, Orientalis-like <i>Yersinia pestis</i> , and Plague Pandemics. Emerging Infectious Diseases, 2004, 10, 1585-1592.	2.0	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.	1.3	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.	0.8	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.	1.8	157
52	Mimivirus and the emerging concept of "giant" virus. Virus Research, 2006, 117, 133-144.	1.1	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.	26.7	157
54	Evidence for Louse-Transmitted Diseases in Soldiers of Napoleon's Grand Army in Vilnius. Journal of Infectious Diseases, 2006, 193, 112-120.	1.9	155

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55	First Documented Human <i>Rickettsia aeschlimannii</i> Infection. <i>Emerging Infectious Diseases</i> , 2002, 8, 748-749.	2.0	153
56	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. <i>FEMS Microbiology Reviews</i> , 2010, 34, 281-294.	3.9	153
57	HACEK Infective Endocarditis: Characteristics and Outcomes from a Large, Multi-National Cohort. <i>PLoS ONE</i> , 2013, 8, e63181.	1.1	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.	1.8	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.	0.8	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.	1.8	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, .	1.8	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.	2.0	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.	1.3	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.	2.9	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.	2.9	130
67	Asynchrony between the rates of standing height gain and bone mass accumulation during puberty. <i>Osteoporosis International</i> , 1997, 7, 525-532.	1.3	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.	1.3	127
69	Diagnosis of infectious endocarditis in patients undergoing valve surgery. <i>American Journal of Medicine</i> , 2005, 118, 230-238.	0.6	126
70	Genome sequence and description of <i>Anaerosalibacter massiliensis</i> sp. nov.. <i>New Microbes and New Infections</i> , 2016, 10, 66-76.	0.8	125
71	Human Pathogens in Body and Head Lice. <i>Emerging Infectious Diseases</i> , 2002, 8, 1515-1518.	2.0	123
72	Burden of emerging anaerobes in the MALDI-TOF and 16S rRNA gene sequencing era. <i>Anaerobe</i> , 2011, 17, 106-112.	1.0	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.	2.0	121
74	Molecular Evolution of Rickettsia Surface Antigens: Evidence of Positive Selection. <i>Molecular Biology and Evolution</i> , 2005, 22, 2073-2083.	3.5	119
75	From culturomics to taxonomogenomics: A need to change the taxonomy of prokaryotes in clinical microbiology. <i>Anaerobe</i> , 2015, 36, 73-78.	1.0	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.	1.3	115
78	Non-steroidal anti-inflammatory drugs for athletes: An update. <i>Annals of Physical and Rehabilitation Medicine</i> , 2010, 53, 278-288.	1.1	115
79	Genetic Identification of Rickettsiae Isolated from Ticks in Japan. <i>Journal of Clinical Microbiology</i> , 2002, 40, 2176-2181.	1.8	114
80	Selfish DNA in Protein-Coding Genes of Rickettsia. <i>Science</i> , 2000, 290, 347-350.	6.0	113
81	The human gut microbiome, a taxonomic conundrum. <i>Systematic and Applied Microbiology</i> , 2015, 38, 276-286.	1.2	113
82	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. <i>FEMS Microbiology Reviews</i> , 2010, 34, 281-94.	3.9	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.	1.8	111
84	Clinical detection and characterization of bacterial pathogens in the genomics era. <i>Genome Medicine</i> , 2014, 6, 114.	3.6	111
85	Blood culture-negative endocarditis. <i>Medicine (United States)</i> , 2017, 96, e8392.	0.4	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.	1.8	109
87	Analysis of the Rickettsia africae genome reveals that virulence acquisition in Rickettsia species may be explained by genome reduction. <i>BMC Genomics</i> , 2009, 10, 166.	1.2	107
88	Respiratory Viruses and Bacteria among Pilgrims during the 2013 Hajj. <i>Emerging Infectious Diseases</i> , 2014, 20, 1821-1827.	2.0	107
89	Non-contiguous finished genome sequence and description of Anaerococcus senegalensis 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.	5.1	106

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91	Mink, SARS-CoV-2, and the Human-Animal Interface. <i>Frontiers in Microbiology</i> , 2021, 12, 663815.	1.5	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.	2.9	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.	1.4	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, .	1.8	103
95	What does the future hold for clinical microbiology?. <i>Nature Reviews Microbiology</i> , 2004, 2, 151-159.	13.6	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.	2.9	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.	1.8	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.	1.8	93
99	Metagenomic Analysis of Brain Abscesses Identifies Specific Bacterial Associations. <i>Clinical Infectious Diseases</i> , 2012, 54, 202-210.	2.9	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.	1.8	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.	1.8	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.	2.8	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.	0.7	87
106	Suicide PCR on Skin Biopsy Specimens for Diagnosis of Rickettsioses. <i>Journal of Clinical Microbiology</i> , 2004, 42, 3428-3434.	1.8	86
107	Do we need new antibiotics?. <i>Clinical Microbiology and Infection</i> , 2016, 22, 408-415.	2.8	86
108	Real-time PCR for universal antibiotic susceptibility testing. <i>Journal of Antimicrobial Chemotherapy</i> , 2004, 54, 538-541.	1.3	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.	2.9	84
110	Identification of Novel Zoonotic Activity of Bartonella spp., France. Emerging Infectious Diseases, 2016, 22, 457-462.	2.0	84
111	Culture and Immunological Detection of <EMPH TYPE="ITAL">Tropheryma whippelii</EMPH> From the Duodenum of a Patient With Whipple Disease. JAMA - Journal of the American Medical Association, 2001, 285, 1039.	3.8	83
112	Use of Highly Variable Intergenic Spacer Sequences for Multispacer Typing of Rickettsia conorii Strains. Journal of Clinical Microbiology, 2004, 42, 5757-5766.	1.8	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.	1.8	83
114	Phylogenetic classification of Bartonella species by comparing groEL sequences.. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 165-171.	0.8	82
115	Genotyping reveals a wide heterogeneity of Tropheryma whippelii. Microbiology (United Kingdom), 2008, 154, 521-527.	0.7	81
116	Quantitative Detection of Tropheryma whippelii DNA by Real-Time PCR. Journal of Clinical Microbiology, 2002, 40, 1119-1120.	1.8	80
117	An Experimental Model of Human Body Louse Infection with Rickettsia prowazekii. Journal of Infectious Diseases, 2002, 186, 1639-1646.	1.9	80
118	Rickettsia tamurae sp. nov., isolated from Amblyomma testudinarium ticks. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1673-1675.	0.8	80
119	Bacterial genome sequencing and its use in infectious diseases. Lancet Infectious Diseases, The, 2007, 7, 711-723.	4.6	79
120	Genotyping of Human Lice Suggests Multiple Emergences of Body Lice from Local Head Louse Populations. PLoS Neglected Tropical Diseases, 2010, 4, e641.	1.3	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.	2.8	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.	2.9	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.	0.4	75
125	Systematic PCR Detection in Culture-negative Osteoarticular Infections. American Journal of Medicine, 2013, 126, 1143.e25-1143.e33.	0.6	75
126	<i>Tropheryma whippelii</i> Endocarditis. Emerging Infectious Diseases, 2013, 19, 1721-30.	2.0	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.	1.8	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.	1.3	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.	2.3	70
133	Genomics and metagenomics in medical microbiology. <i>Journal of Microbiological Methods</i> , 2013, 95, 415-424.	0.7	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.	1.3	69
135	Value of Microimmunofluorescence for Diagnosis and Follow-up of <i>Bartonella</i> Endocarditis. <i>Vaccine Journal</i> , 2002, 9, 795-801.	3.2	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.	1.0	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.	1.8	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.	1.5	63
144	Strikes, flooding, rats, and leptospirosis in Marseille, France. <i>International Journal of Infectious Diseases</i> , 2011, 15, e710-e715.	1.5	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.	1.8	61
146	Rapid identification of Legionella species by mass spectrometry. Journal of Medical Microbiology, 2010, 59, 273-284.	0.7	61
147	Genome sequence and description of Alistipes senegalensis sp. nov.. Standards in Genomic Sciences, 2012, 6, 304-314.	1.5	61
148	Genome and pan-genome analysis to classify emerging bacteria. Biology Direct, 2019, 14, 5.	1.9	61
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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.	1.3	13
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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.	1.8	12
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525	Genome Sequence of <i>Rickettsia sibirica</i> subsp. <i>mongolitimonae</i> . <i>Journal of Bacteriology</i> , 2012, 194, 2389-2390.	1.0	7
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543	Early mortality attributable to PICC-lines in 4 public hospitals of Marseille from 2010 to 2016 (Revised) <i>Tj ETQq1 1 0.784314, rjgBT /Over</i> 0.4	0.4	7
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545	Pangenome analysis and virulence profiling of <i>Streptococcus intermedius</i> . <i>BMC Genomics</i> , 2021, 22, 522.	1.2	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.	1.0	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.	0.8	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.	0.8	6
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813	NMNI editorial report, 2020. <i>New Microbes and New Infections</i> , 2021, 41, 100859.	0.8	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.	0.8	0
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818	Genomes of <i>Rickettsia</i> Species. , 2016, , 447-462.		0
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.	0.1	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.	0.3	0
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