## Didier Raoult

List of Publications by Year in descending order

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2,428 papers

147,088 citations

159 h-index 280 g-index

2515 all docs

2515 docs citations

times ranked

2515

86421 citing authors

#	Article	IF	Citations
1	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial. International Journal of Antimicrobial Agents, 2020, 56, 105949.	2.5	3,955
2	Gut microbiome influences efficacy of PD-1–based immunotherapy against epithelial tumors. Science, 2018, 359, 91-97.	12.6	3,689
3	Anticancer immunotherapy by CTLA-4 blockade relies on the gut microbiota. Science, 2015, 350, 1079-1084.	12.6	2,539
4	Ongoing Revolution in Bacteriology: Routine Identification of Bacteria by Matrixâ€Assisted Laser Desorption Ionization Timeâ€ofâ€Flight Mass Spectrometry. Clinical Infectious Diseases, 2009, 49, 543-551.	5.8	1,638
5	The abundance and variety of carbohydrate-active enzymes in the human gut microbiota. Nature Reviews Microbiology, 2013, 11, 497-504.	28.6	1,240
6	Update on Tick-Borne Rickettsioses around the World: a Geographic Approach. Clinical Microbiology Reviews, 2013, 26, 657-702.	13.6	1,033
7	The 1.2-Megabase Genome Sequence of Mimivirus. Science, 2004, 306, 1344-1350.	12.6	959
8	Tick-Borne Rickettsioses around the World: Emerging Diseases Challenging Old Concepts. Clinical Microbiology Reviews, 2005, 18, 719-756.	13.6	920
9	Microbial culturomics: paradigm shift in the human gut microbiome study. Clinical Microbiology and Infection, 2012, 18, 1185-1193.	6.0	905
10	Microorganisms Resistant to Free-Living Amoebae. Clinical Microbiology Reviews, 2004, 17, 413-433.	13.6	895
11	16S Ribosomal DNA Sequence Analysis of a Large Collection of Environmental and Clinical Unidentifiable Bacterial Isolates. Journal of Clinical Microbiology, 2000, 38, 3623-3630.	3.9	873
12	New insights on the antiviral effects of chloroquine against coronavirus: what to expect for COVID-19?. International Journal of Antimicrobial Agents, 2020, 55, 105938.	2.5	842
13	Viral RNA load as determined by cell culture as a management tool for discharge of SARS-CoV-2 patients from infectious disease wards. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1059-1061.	2.9	767
14	A Giant Virus in Amoebae. Science, 2003, 299, 2033-2033.	12.6	742
15	Monitoring Bacterial Community of Human Gut Microbiota Reveals an Increase in Lactobacillus in Obese Patients and Methanogens in Anorexic Patients. PLoS ONE, 2009, 4, e7125.	2.5	735
16	Culture of previously uncultured members of the human gut microbiota by culturomics. Nature Microbiology, 2016, 1, 16203.	13.3	735
17	Chloroquine and hydroxychloroquine as available weapons to fight COVID-19. International Journal of Antimicrobial Agents, 2020, 55, 105932.	2.5	724
18	Comparative Genomics of Multidrug Resistance in Acinetobacter baumannii. PLoS Genetics, 2006, 2, e7.	3.5	677

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19	Endocarditis Due to Rare and Fastidious Bacteria. Clinical Microbiology Reviews, 2001, 14, 177-207.	13.6	668
20	Enterococcus hirae and Barnesiella intestinihominis Facilitate Cyclophosphamide-Induced Therapeutic Immunomodulatory Effects. Immunity, 2016, 45, 931-943.	14.3	645
21	From Q Fever to Coxiella burnetii Infection: a Paradigm Change. Clinical Microbiology Reviews, 2017, 30, 115-190.	13.6	616
22	The Rebirth of Culture in Microbiology through the Example of Culturomics To Study Human Gut Microbiota. Clinical Microbiology Reviews, 2015, 28, 237-264.	13.6	605
23	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. Travel Medicine and Infectious Disease, 2020, 34, 101663.	3.0	605
24	Risk of Embolism and Death in Infective Endocarditis: Prognostic Value of Echocardiography. Circulation, 2005, 112, 69-75.	1.6	600
25	Whipple's Disease. New England Journal of Medicine, 2007, 356, 55-66.	27.0	574
26	Pig Liver Sausage as a Source of Hepatitis E Virus Transmission to Humans. Journal of Infectious Diseases, 2010, 202, 825-834.	4.0	571
27	Obesity-associated gut microbiota is enriched in Lactobacillus reuteri and depleted in Bifidobacterium animalis and Methanobrevibacter smithii. International Journal of Obesity, 2012, 36, 817-825.	3.4	567
28	Diagnosis of Q Fever. Journal of Clinical Microbiology, 1998, 36, 1823-1834.	3.9	529
29	Q fever. Veterinary Microbiology, 2010, 140, 297-309.	1.9	525
30	The microbiome in cancer immunotherapy: Diagnostic tools and therapeutic strategies. Science, 2018, 359, 1366-1370.	12.6	525
31	Culturing the human microbiota and culturomics. Nature Reviews Microbiology, 2018, 16, 540-550.	28.6	521
32	Methanomassiliicoccus luminyensis gen. nov., sp. nov., a methanogenic archaeon isolated from human faeces. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 1902-1907.	1.7	510
33	The virophage as a unique parasite of the giant mimivirus. Nature, 2008, 455, 100-104.	27.8	505
34	Genome sequences of the human body louse and its primary endosymbiont provide insights into the permanent parasitic lifestyle. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12168-12173.	7.1	482
35	Q Fever 1985-1998: Clinical and Epidemiologic Features of 1,383 Infections. Medicine (United States), 2000, 79, 109-123.	1.0	459
36	Cultivation of the Bacillus of Whipple's Disease. New England Journal of Medicine, 2000, 342, 620-625.	27.0	458

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37	Recommendations for Treatment of Human Infections Caused by Bartonella Species. Antimicrobial Agents and Chemotherapy, 2004, 48, 1921-1933.	3.2	456
38	rpoB sequence analysis as a novel basis for bacterial identification. Molecular Microbiology, 1997, 26, 1005-1011.	2.5	450
39	Positron Emission Tomography/Computed Tomography forÂDiagnosis of Prosthetic Valve Endocarditis. Journal of the American College of Cardiology, 2013, 61, 2374-2382.	2.8	440
40	Comprehensive Diagnostic Strategy for Blood Culture–Negative Endocarditis: A Prospective Study of 819 New Cases. Clinical Infectious Diseases, 2010, 51, 131-140.	5.8	418
41	ACE2 receptor polymorphism: Susceptibility to SARS-CoV-2, hypertension, multi-organ failure, and COVID-19 disease outcome. Journal of Microbiology, Immunology and Infection, 2020, 53, 425-435.	3.1	410
42	Mechanisms of Evolution in <i>Rickettsia conorii</i> and <i>R. prowazekii</i> Science, 2001, 293, 2093-2098.	12.6	408
43	Laboratory diagnosis of rickettsioses: current approaches to diagnosis of old and new rickettsial diseases. Journal of Clinical Microbiology, 1997, 35, 2715-2727.	3.9	397
44	Chikungunya Outbreaks â€" The Globalization of Vectorborne Diseases. New England Journal of Medicine, 2007, 356, 769-771.	27.0	394
45	Echocardiography predicts embolic events in infective endocarditis. Journal of the American College of Cardiology, 2001, 37, 1069-1076.	2.8	388
46	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. Journal of Clinical Microbiology, 1996, 34, 2058-2065.	3.9	388
47	Giant Marseillevirus highlights the role of amoebae as a melting pot in emergence of chimeric microorganisms. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 21848-21853.	7.1	385
48	Blood Culture-Negative Endocarditis in a Reference Center. Medicine (United States), 2005, 84, 162-173.	1.0	377
49	Early treatment of COVID-19 patients with hydroxychloroquine and azithromycin: A retrospective analysis of 1061 cases in Marseille, France. Travel Medicine and Infectious Disease, 2020, 35, 101738.	3.0	372
50	Comparative meta-analysis of the effect of Lactobacillus species on weight gain in humans and animals. Microbial Pathogenesis, 2012, 53, 100-108.	2.9	364
51	Identification of Rare Pathogenic Bacteria in a Clinical Microbiology Laboratory: Impact of Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry. Journal of Clinical Microbiology, 2013, 51, 2182-2194.	3.9	362
52	Management of infective endocarditis: challenges and perspectives. Lancet, The, 2012, 379, 965-975.	13.7	359
53	Redefining viruses: lessons from Mimivirus. Nature Reviews Microbiology, 2008, 6, 315-319.	28.6	358
54	Current and Past Strategies for Bacterial Culture in Clinical Microbiology. Clinical Microbiology Reviews, 2015, 28, 208-236.	13.6	358

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55	Autoantibodies neutralizing type I IFNs are present in ~4% of uninfected individuals over 70 years old and account for ~20% of COVID-19 deaths. Science Immunology, 2021, 6, .	11.9	357
56	<i>Bartonella (Rochalimaea) quintana</i> Endocarditis in Three Homeless Men. New England Journal of Medicine, 1995, 332, 419-423.	27.0	355
57	Chloroquine for the 2019 novel coronavirus SARS-CoV-2. International Journal of Antimicrobial Agents, 2020, 55, 105923.	2.5	354
58	Gene Sequence-Based Criteria for Identification of New <i>Rickettsia</i> Isolates and Description of <i>Rickettsia heilongjiangensis</i> Sp. nov. Journal of Clinical Microbiology, 2003, 41, 5456-5465.	3.9	347
59	Recycling of chloroquine and its hydroxyl analogue to face bacterial, fungal and viral infections in the 21st century. International Journal of Antimicrobial Agents, 2007, 30, 297-308.	2.5	332
60	Direct Identification of Bacteria in Positive Blood Culture Bottles by Matrix-Assisted Laser Desorption Ionisation Time-of-Flight Mass Spectrometry. PLoS ONE, 2009, 4, e8041.	2.5	331
61	A polyphasic strategy incorporating genomic data for the taxonomic description of novel bacterial species. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 384-391.	1.7	330
62	rpoB gene sequence-based characterization of emerging non-tuberculous mycobacteria with descriptions of Mycobacterium bolletii sp. nov., Mycobacterium phocaicum sp. nov. and Mycobacterium aubagnense sp. nov International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 133-143.	1.7	329
63	Epidemiologic features and clinical presentation of acute Q fever in hospitalized patients: 323 French cases. American Journal of Medicine, 1992, 93, 427-434.	1.5	328
64	Guidelines for the diagnosis of tick-borne bacterial diseases in Europe. Clinical Microbiology and Infection, 2004, 10, 1108-1132.	6.0	328
65	Whipple's disease: new aspects of pathogenesis and treatment. Lancet Infectious Diseases, The, 2008, 8, 179-190.	9.1	328
66	Q Fever. Clinical Infectious Diseases, 1995, 20, 489-496.	5.8	322
67	Diagnosis of 22 New Cases of Bartonella Endocarditis. Annals of Internal Medicine, 1996, 125, 646.	3.9	321
68	Sequencing of the rpoB Gene and Flanking Spacers for Molecular Identification of Acinetobacter Species. Journal of Clinical Microbiology, 2006, 44, 827-832.	3.9	321
69	Eukaryotic large nucleo-cytoplasmic DNA viruses: Clusters of orthologous genes and reconstruction of viral genome evolution. Virology Journal, 2009, 6, 223.	3.4	321
70	Impact of cerebrovascular complications on mortality and neurologic outcome during infective endocarditis: a prospective multicentre study. European Heart Journal, 2007, 28, 1155-1161.	2,2	320
71	The Body Louse as a Vector of Reemerging Human Diseases. Clinical Infectious Diseases, 1999, 29, 888-911.	5.8	316
72	Comparative Analysis of Acinetobacters: Three Genomes for Three Lifestyles. PLoS ONE, 2008, 3, e1805.	2.5	315

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73	High Prevalence of Methanobrevibacter smithii and Methanosphaera stadtmanae Detected in the Human Gut Using an Improved DNA Detection Protocol. PLoS ONE, 2009, 4, e7063.	2.5	312
74	Fleas and flea-borne diseases. International Journal of Infectious Diseases, 2010, 14, e667-e676.	3.3	312
75	MALDI-TOF-mass spectrometry applications in clinical microbiology. Future Microbiology, 2010, 5, 1733-1754.	2.0	310
76	Q fever serology: cutoff determination for microimmunofluorescence. Vaccine Journal, 1994, 1, 189-196.	2.6	308
77	Treatment of Q Fever Endocarditis. Archives of Internal Medicine, 1999, 159, 167.	3.8	306
78	The relationship between gut microbiota and weight gain in humans. Future Microbiology, 2012, 7, 91-109.	2.0	306
79	Detection of 400-year-old Yersinia pestis DNA in human dental pulp: An approach to the diagnosis of ancient septicemia. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 12637-12640.	7.1	301
80	The Q fever epidemic in The Netherlands: history, onset, response and reflection. Epidemiology and Infection, 2011, 139, 1-12.	2.1	298
81	Genome of Acanthamoeba castellanii highlights extensive lateral gene transfer and early evolution of tyrosine kinase signaling. Genome Biology, 2013, 14, R11.	9.6	296
82	Novel Chikungunya Virus Variant in Travelers Returning from Indian Ocean Islands. Emerging Infectious Diseases, 2006, 12, 1493-1499.	4.3	295
83	Staphylococcus aureus Native Valve Infective Endocarditis: Report of 566 Episodes from the International Collaboration on Endocarditis Merged Database. Clinical Infectious Diseases, 2005, 41, 507-514.	5.8	289
84	Molecular identification by "suicide PCR―of <i>Yersinia pestis</i> as the agent of Medieval Black Death. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 12800-12803.	7.1	288
85	Genome Sequence of Rickettsia bellii Illuminates the Role of Amoebae in Gene Exchanges between Intracellular Pathogens. PLoS Genetics, 2006, 2, e76.	3.5	286
86	Chronic <i>Bartonella quintana</i> Bacteremia in Homeless Patients. New England Journal of Medicine, 1999, 340, 184-189.	27.0	285
87	<i>Rickettsia africae,</i> a Tick-Borne Pathogen in Travelers to Sub-Saharan Africa. New England Journal of Medicine, 2001, 344, 1504-1510.	27.0	282
88	Malaria morbidity and pyrethroid resistance after the introduction of insecticide-treated bednets and artemisinin-based combination therapies: a longitudinal study. Lancet Infectious Diseases, The, 2011, 11, 925-932.	9.1	282
89	Bacterial strain typing in the genomic era. FEMS Microbiology Reviews, 2009, 33, 892-916.	8.6	278
90	Molecular Detection of <i>Bartonella quintana </i> , <i>B. koehlerae, B. henselae, B. clarridgeiae, Rickettsia felis </i> , and <i>Wolbachia pipientis </i> in Cat Fleas, France. Emerging Infectious Diseases, 2003, 9, 338-342.	4.3	275

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91	Value and limitations of the duke criteria for the diagnosis of infective endocarditis. Journal of the American College of Cardiology, 1999, 33, 2023-2029.	2.8	271
92	Amoebal Coculture of " Mycobacterium massiliense ―sp. nov. from the Sputum of a Patient with Hemoptoic Pneumonia. Journal of Clinical Microbiology, 2004, 42, 5493-5501.	3.9	271
93	Coronavirus infections: Epidemiological, clinical and immunological features and hypotheses. Cell Stress, 2020, 4, 66-75.	3.2	271
94	Molecular Quantification of <i>Gardnerella vaginalis </i> Predict Bacterial Vaginosis. Clinical Infectious Diseases, 2008, 47, 33-43.	5.8	267
95	Risks Factors and Prevention of Q Fever Endocarditis. Clinical Infectious Diseases, 2001, 33, 312-316.	5.8	264
96	Modern clinical microbiology: new challenges and solutions. Nature Reviews Microbiology, 2013, 11, 574-585.	28.6	264
97	Whipple's disease. Lancet, The, 2003, 361, 239-246.	13.7	263
98	Gene-sequence-based criteria for species definition in bacteriology: the Bartonella paradigm. Trends in Microbiology, 2003, 11, 318-321.	7.7	259
99	The rpoB gene as a tool for clinical microbiologists. Trends in Microbiology, 2009, 17, 37-45.	7.7	257
100	"Megaviralesâ€, a proposed new order for eukaryotic nucleocytoplasmic large DNA viruses. Archives of Virology, 2013, 158, 2517-2521.	2.1	256
101	rpoB Gene Sequence-Based Identification of Staphylococcus Species. Journal of Clinical Microbiology, 2002, 40, 1333-1338.	3.9	255
102	Dramatic Reduction in Infective Endocarditis–Related Mortality With a Management-Based Approach. Archives of Internal Medicine, 2009, 169, 1290.	3.8	255
103	Characterization of the Naturally Occurring Oxacillinase of Acinetobacter baumannii. Antimicrobial Agents and Chemotherapy, 2005, 49, 4174-4179.	3.2	254
104	Understanding the Cholera Epidemic, Haiti. Emerging Infectious Diseases, 2011, 17, 1161-1168.	4.3	252
105	Human Gut Microbiota: Repertoire and Variations. Frontiers in Cellular and Infection Microbiology, 2012, 2, 136.	3.9	252
106	Analysis of 525 Samples To Determine the Usefulness of PCR Amplification and Sequencing of the 16S rRNA Gene for Diagnosis of Bone and Joint Infections. Journal of Clinical Microbiology, 2006, 44, 1018-1028.	3.9	251
107	A comprehensive repertoire of prokaryotic species identified in human beings. Lancet Infectious Diseases, The, 2015, 15, 1211-1219.	9.1	250
108	Tick- and flea-borne rickettsial emerging zoonoses. Veterinary Research, 2005, 36, 469-492.	3.0	248

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109	Tailed giant Tupanvirus possesses the most complete translational apparatus of the known virosphere. Nature Communications, 2018, 9, 749.	12.8	247
110	Worldwide emergence of colistin resistance in Klebsiella pneumoniae from healthy humans and patients in Lao PDR, Thailand, Israel, Nigeria and France owing to inactivation of the PhoP/PhoQ regulator mgrB: an epidemiological and molecular study. International Journal of Antimicrobial Agents, 2014, 44, 500-507.	2.5	246
111	In vitro testing of combined hydroxychloroquine and azithromycin on SARS-CoV-2 shows synergistic effect. Microbial Pathogenesis, 2020, 145, 104228.	2.9	246
112	Wind in November, Q Fever in December. Emerging Infectious Diseases, 2004, 10, 1264-1269.	4.3	244
113	rpoB Gene Sequencing for Identification of Corynebacterium Species. Journal of Clinical Microbiology, 2004, 42, 3925-3931.	3.9	243
114	The Genome Sequence of Rickettsia felis Identifies the First Putative Conjugative Plasmid in an Obligate Intracellular Parasite. PLoS Biology, 2005, 3, e248.	5.6	242
115	Detection of ehrlichiae in African ticks by polymerase chain reaction. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2000, 94, 707-708.	1.8	238
116	Culture of <i>Bartonella quintana</i> and <i>Bartonella henselae</i> from Human Samples: a 5-Year Experience (1993 to 1998). Journal of Clinical Microbiology, 1999, 37, 1899-1905.	3.9	232
117	Warmer Weather Linked to Tick Attack and Emergence of Severe Rickettsioses. PLoS Neglected Tropical Diseases, 2008, 2, e338.	3.0	228
118	Antibiotic discovery: history, methods and perspectives. International Journal of Antimicrobial Agents, 2019, 53, 371-382.	2.5	223
119	Rickettsioses and the International Traveler. Clinical Infectious Diseases, 2004, 39, 1493-1499.	5.8	222
120	Rickettsial evolution in the light of comparative genomics. Biological Reviews, 2011, 86, 379-405.	10.4	219
121	The two faces of interleukin 10 in human infectious diseases. Lancet Infectious Diseases, The, 2006, 6, 557-569.	9.1	218
122	Cross-reactivity between tumor MHC class l–restricted antigens and an enterococcal bacteriophage. Science, 2020, 369, 936-942.	12.6	217
123	Gut microbiota signatures are associated with toxicity to combined CTLA-4 and PD-1 blockade. Nature Medicine, 2021, 27, 1432-1441.	30.7	216
124	Use of rpoB Gene Analysis for Detection and Identification of Bartonella Species. Journal of Clinical Microbiology, 2001, 39, 430-437.	3.9	211
125	<i>Coxiella burnetii</i> Genotyping. Emerging Infectious Diseases, 2005, 11, 1211-1217.	4.3	210
126	Structural Studies of the Giant Mimivirus. PLoS Biology, 2009, 7, e1000092.	5.6	209

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127	Outcomes of 3,737 COVID-19 patients treated with hydroxychloroquine/azithromycin and other regimens in Marseille, France: A retrospective analysis. Travel Medicine and Infectious Disease, 2020, 36, 101791.	3.0	209
128	Epidemiologic and Clinical Characteristics of Bartonella quintana and Bartonella henselae Endocarditis. Medicine (United States), 2001, 80, 245-251.	1.0	207
129	A Flea-Associated Rickettsia Pathogenic for Humans. Emerging Infectious Diseases, 2001, 7, 73-81.	4.3	207
130	Teicoplanin: an alternative drug for the treatment of COVID-19?. International Journal of Antimicrobial Agents, 2020, 55, 105944.	2.5	205
131	Outcome and Treatment of Bartonella Endocarditis. Archives of Internal Medicine, 2003, 163, 226.	3.8	202
132	<i>Propionibacterium acnes</i> Postoperative Shoulder Arthritis: An Emerging Clinical Entity. Clinical Infectious Diseases, 2008, 46, 1884-1886.	5.8	202
133	Molecular Detection of Multiple Emerging Pathogens in Sputa from Cystic Fibrosis Patients. PLoS ONE, 2008, 3, e2908.	2.5	201
134	African tick bite fever. Lancet Infectious Diseases, The, 2003, 3, 557-564.	9.1	199
135	Rickettsial Infections and Fever, Vientiane, Laos. Emerging Infectious Diseases, 2006, 12, 256-262.	4.3	197
136	Long-term outcome of Q fever endocarditis: a 26-year personal survey. Lancet Infectious Diseases, The, 2010, 10, 527-535.	9.1	197
137	Systemic Tropheryma whipplei. Medicine (United States), 2010, 89, 337-345.	1.0	195
138	Infections in the homeless. Lancet Infectious Diseases, The, 2001, 1, 77-84.	9.1	194
139	Provirophages and transpovirons as the diverse mobilome of giant viruses. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 18078-18083.	7.1	194
140	Gut Bacteria Composition Drives Primary Resistance to Cancer Immunotherapy in Renal Cell Carcinoma Patients. European Urology, 2020, 78, 195-206.	1.9	192
141	<i>Rickettsia slovaca</i> and <i>R. raoultii</i> ii> Tick <i></i> borne Rickettsioses. Emerging Infectious Diseases, 2009, 15, 1105-1108.	4.3	191
142	Faustovirus, an Asfarvirus-Related New Lineage of Giant Viruses Infecting Amoebae. Journal of Virology, 2015, 89, 6585-6594.	3.4	191
143	Epidemic typhus. Lancet Infectious Diseases, The, 2008, 8, 417-426.	9.1	189
144	Massive comparative genomic analysis reveals convergent evolution of specialized bacteria. Biology Direct, 2009, 4, 13.	4.6	187

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145	<i>Rickettsia massiliae</i> Human Isolation. Emerging Infectious Diseases, 2006, 12, 174-175.	4.3	186
146	High-level colonisation of the human gut by Verrucomicrobia following broad-spectrum antibiotic treatment. International Journal of Antimicrobial Agents, 2013, 41, 149-155.	2.5	186
147	A Cluster of Coxiella Burnetii Infections Associated with Exposure to Vaccinated Goats and their Unpasteurized Dairy Products. American Journal of Tropical Medicine and Hygiene, 1992, 47, 35-40.	1.4	186
148	The timing of surgery influences mortality and morbidity in adults with severe complicated infective endocarditis: a propensity analysis. European Heart Journal, 2011, 32, 2027-2033.	2.2	184
149	<i>Bartonella vinsonii</i> subsp. <i>berkhoffii</i> as an Agent of Afebrile Blood Culture-Negative Endocarditis in a Human. Journal of Clinical Microbiology, 2000, 38, 1698-1700.	3.9	183
150	Comparison of PCR and Serology Assays for Early Diagnosis of Acute Q Fever. Journal of Clinical Microbiology, 2003, 41, 5094-5098.	3.9	182
151	Validation of partial rpoB gene sequence analysis for the identification of clinically important and emerging Acinetobacter species. Microbiology (United Kingdom), 2009, 155, 2333-2341.	1.8	182
152	Coxiella burnetii in Humans and Ticks in Rural Senegal. PLoS Neglected Tropical Diseases, 2010, 4, e654.	3.0	181
153	Genome-based design of a cell-free culture medium for Tropheryma whipplei. Lancet, The, 2003, 362, 447-449.	13.7	180
154	Mediterranean Spotted Fever: Clinical, Laboratory and Epidemiological Features of 199 Cases. American Journal of Tropical Medicine and Hygiene, 1986, 35, 845-850.	1.4	179
155	Spotless Rickettsiosis Caused by <i>Rickettsia slovaca</i> and Associated with <i>Dermacentor</i> Ticks. Clinical Infectious Diseases, 2002, 34, 1331-1336.	5 <b>.</b> 8	177
156	African Tick Bite Fever in Travelers to Rural Sub-Equatorial Africa. Clinical Infectious Diseases, 2003, 36, 1411-1417.	5.8	177
157	Molecular diagnosis of bloodstream infections caused by non-cultivable bacteria. International Journal of Antimicrobial Agents, 2007, 30, 7-15.	2.5	176
158	Gut microbiota and malnutrition. Microbial Pathogenesis, 2017, 106, 127-138.	2.9	173
159	Etiologic Diagnosis of 204 Pericardial Effusions. Medicine (United States), 2003, 82, 385-391.	1.0	170
160	Value of <i>Tropheryma whipplei</i> Quantitative Polymerase Chain Reaction Assay for the Diagnosis of Whipple Disease: Usefulness of Saliva and Stool Specimens for Firstâ€Line Screening. Clinical Infectious Diseases, 2008, 47, 659-667.	5.8	170
161	Detection of Ehrlichia spp., Anaplasma spp., Rickettsia spp., and Other Eubacteria in Ticks from the Thai-Myanmar Border and Vietnam. Journal of Clinical Microbiology, 2003, 41, 1600-1608.	3.9	167
162	Reductive Genome Evolution from the Mother of Rickettsia. PLoS Genetics, 2007, 3, e14.	3.5	167

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163	Pathogenicity and treatment of Bartonella infections. International Journal of Antimicrobial Agents, 2014, 44, 16-25.	2.5	167
164	Genotyping, Orientalis-like <i>Yersinia pestis </i> , and Plague Pandemics. Emerging Infectious Diseases, 2004, 10, 1585-1592.	4.3	166
165	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
166	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
167	<i>Tropheryma whipplei</i> Twist: A Human Pathogenic Actinobacteria With a Reduced Genome. Genome Research, 2003, 13, 1800-1809.	<b>5.</b> 5	161
168	Correlation Between 3790 Quantitative Polymerase Chain Reaction–Positives Samples and Positive Cell Cultures, Including 1941 Severe Acute Respiratory Syndrome Coronavirus 2 Isolates. Clinical Infectious Diseases, 2021, 72, e921-e921.	5 <b>.</b> 8	158
169	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
170	Prevalence of Asymptomatic <i>Tropheryma whipplei</i> Primates. Journal of Infectious Diseases, 2008, 197, 880-887.	4.0	157
171	Q fever in children. Lancet Infectious Diseases, The, 2002, 2, 686-691.	9.1	155
172	Evidence for Louseâ€Transmitted Diseases in Soldiers of Napoleon's Grand Army in Vilnius. Journal of Infectious Diseases, 2006, 193, 112-120.	4.0	155
173	Related actions of probiotics and antibiotics on gut microbiota and weight modification. Lancet Infectious Diseases, The, 2013, 13, 889-899.	9.1	154
174	First Documented Human <i>Rickettsia aeschlimannii</i> li>Infection. Emerging Infectious Diseases, 2002, 8, 748-749.	4.3	153
175	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. FEMS Microbiology Reviews, 2010, 34, 281-294.	8.6	153
176	Pepper Mild Mottle Virus, a Plant Virus Associated with Specific Immune Responses, Fever, Abdominal Pains, and Pruritus in Humans. PLoS ONE, 2010, 5, e10041.	2.5	153
177	<i>Bartonella quintana</i> Characteristics and Clinical Management. Emerging Infectious Diseases, 2006, 12, 217-223.	4.3	152
178	Archaea as emerging organisms in complex human microbiomes. Anaerobe, 2011, 17, 56-63.	2.1	152
179	Matrix-Assisted Laser Desorption/Ionization Time-of-Flight Mass Spectrometry Identification of Mycobacteria in Routine Clinical Practice. PLoS ONE, 2011, 6, e24720.	2.5	152
180	Emerging Rickettsioses of the Thai-Myanmar Border1. Emerging Infectious Diseases, 2003, 9, 592-595.	4.3	151

#	Article	IF	Citations
181	Culturomics identified 11 new bacterial species from a single anorexia nervosa stool sample. European Journal of Clinical Microbiology and Infectious Diseases, 2013, 32, 1471-1481.	2.9	150
182	Bacterial culture through selective and non-selective conditions: the evolution of culture media in clinical microbiology. New Microbes and New Infections, 2020, 34, 100622.	1.6	150
183	<i>Massilia timonae</i> gen. nov., sp. nov., Isolated from Blood of an Immunocompromised Patient with Cerebellar Lesions. Journal of Clinical Microbiology, 1998, 36, 2847-2852.	3.9	150
184	Mimivirus in Pneumonia Patients. Emerging Infectious Diseases, 2005, 11, 449-452.	4.3	149
185	Reclassification of Giant Viruses Composing a Fourth Domain of Life in the New Order & lt;b> <i>Megavirales</i> . Intervirology, 2012, 55, 321-332.	2.8	149
186	Comparison between rpoB and 16S rRNA Gene Sequencing for Molecular Identification of 168 Clinical Isolates of Corynebacterium. Journal of Clinical Microbiology, 2005, 43, 1934-1936.	3.9	148
187	Palaeomicrobiology: current issues and perspectives. Nature Reviews Microbiology, 2005, 3, 23-35.	28.6	147
188	The Genome of Borrelia recurrentis, the Agent of Deadly Louse-Borne Relapsing Fever, Is a Degraded Subset of Tick-Borne Borrelia duttonii. PLoS Genetics, 2008, 4, e1000185.	3.5	146
189	PCR Detection of Bacteria on Cardiac Valves of Patients with Treated Bacterial Endocarditis. Journal of Clinical Microbiology, 2005, 43, 163-167.	3.9	145
190	Effect of Sex onCoxiella burnetiiInfection: Protective Role of 17βâ€Estradiol. Journal of Infectious Diseases, 2004, 189, 339-345.	4.0	143
191	Ketogenic diet and ketone bodies enhance the anticancer effects of PD-1 blockade. JCI Insight, 2021, 6, .	5.0	143
192	Collateral damage: insights into bacterial mechanisms that predispose host cells to cancer. Nature Reviews Microbiology, 2017, 15, 109-128.	28.6	142
193	Detection of <i>Ehrlichia platys</i> DNA in Brown Dog Ticks ( <i>Rhipicephalus sanguineus</i> ) in Okinawa Island, Japan. Journal of Clinical Microbiology, 2000, 38, 4219-4221.	3.9	142
194	Spotted fever rickettsioses in southern and eastern Europe. FEMS Immunology and Medical Microbiology, 2007, 49, 2-12.	2.7	141
195	Q Fever. Infectious Disease Clinics of North America, 2008, 22, 505-514.	5.1	141
196	The relationship between spotted fever group <i>Rickettsiae</i> and Ixodid ticks. Veterinary Research, 2009, 40, 34.	3.0	141
197	Monocytes and Macrophages, Targets of Severe Acute Respiratory Syndrome Coronavirus 2: The Clue for Coronavirus Disease 2019 Immunoparalysis. Journal of Infectious Diseases, 2021, 224, 395-406.	4.0	141
198	The contribution of culturomics to the repertoire of isolated human bacterial and archaeal species. Microbiome, 2018, 6, 94.	11.1	139

#	Article	IF	Citations
199	Ultrastructural Characterization of the Giant Volcano-like Virus Factory of Acanthamoeba polyphaga Mimivirus. PLoS ONE, 2007, 2, e328.	2.5	139
200	<i>Rickettsia felis</i> à€"associated Uneruptive Fever, Senegal. Emerging Infectious Diseases, 2010, 16, 1140-1142.	4.3	138
201	Mimivirus shows dramatic genome reduction after intraamoebal culture. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 10296-10301.	7.1	138
202	Tentative Characterization of New Environmental Giant Viruses by MALDI-TOF Mass Spectrometry. Intervirology, 2010, 53, 344-353.	2.8	137
203	Prediction of Symptomatic Embolism in Infective Endocarditis. Journal of the American College of Cardiology, 2013, 62, 1384-1392.	2.8	137
204	Aneruptive Fever Associated with Antibodies to <i>Rickettsia helvetica</i> in Europe and Thailand. Journal of Clinical Microbiology, 2004, 42, 816-818.	3.9	136
205	Survival of Environmental Mycobacteria in Acanthamoeba polyphaga. Applied and Environmental Microbiology, 2006, 72, 5974-5981.	3.1	136
206	Evaluation of the Panbio COVID-19 Rapid Antigen Detection Test Device for the Screening of Patients with COVID-19. Journal of Clinical Microbiology, 2021, 59, .	3.9	136
207	Ameba-associated Microorganisms and Diagnosis of Nosocomial Pneumonia. Emerging Infectious Diseases, 2006, 12, 248-255.	4.3	135
208	Phylogenetic and Phyletic Studies of Informational Genes in Genomes Highlight Existence of a 4th Domain of Life Including Giant Viruses. PLoS ONE, 2010, 5, e15530.	2.5	135
209	Rickettsia felis Infection Acquired in Europe and Documented by Polymerase Chain Reaction. Emerging Infectious Diseases, 2002, 8, 207-208.	4.3	134
210	Yersinia pestis as a telluric, human ectoparasite-borne organism. Lancet Infectious Diseases, The, 2006, 6, 234-241.	9.1	134
211	Ameobal Pathogen Mimivirus Infects Macrophages through Phagocytosis. PLoS Pathogens, 2008, 4, e1000087.	4.7	134
212	<i>Coxiella</i> â€^ <i>burnetii</i> Survival in THP-1 Monocytes Involves the Impairment of Phagosome Maturation: IFN-γ Mediates its Restoration and Bacterial Killing. Journal of Immunology, 2002, 169, 4488-4495.	0.8	133
213	Q Fever during Pregnancy A Risk for Women, Fetuses, and Obstetricians. New England Journal of Medicine, 1994, 330, 371-371.	27.0	132
214	Scalp Eschar and Neck Lymphadenopathy Caused by <i>Bartonella henselae</i> after Tick Bite. Clinical Infectious Diseases, 2010, 50, 549-551.	5.8	132
215	Mimivirus: leading the way in the discovery of giant viruses of amoebae. Nature Reviews Microbiology, 2017, 15, 243-254.	28.6	132
216	Ambulatory Treatment of Multidrug-Resistant <i>Staphylococcus</i> li>lnfected Orthopedic Implants with High-Dose Oral Co-trimoxazole (Trimethoprim-Sulfamethoxazole). Antimicrobial Agents and Chemotherapy, 1998, 42, 3086-3091.	3.2	132

#	Article	IF	Citations
217	Lymphangitis-Associated Rickettsiosis, a New Rickettsiosis Caused by Rickettsia sibirica mongolotimonae: Seven New Cases and Review of the Literature. Clinical Infectious Diseases, 2005, 40, 1435-1444.	5.8	131
218	Managing Q Fever during Pregnancy: The Benefits of Long-Term Cotrimoxazole Therapy. Clinical Infectious Diseases, 2007, 45, 548-555.	5.8	131
219	The Expansion of the Microbiological Spectrum of Brain Abscesses with Use of Multiple 16S Ribosomal DNA Sequencing. Clinical Infectious Diseases, 2009, 48, 1169-1178.	5.8	131
220	First Isolation of Mimivirus in a Patient With Pneumonia. Clinical Infectious Diseases, 2013, 57, e127-e134.	5.8	131
221	History of the ADP/ATP-Translocase-Encoding Gene, a Parasitism Gene Transferred from a <i>Chlamydiales</i> Ancestor to Plants 1 Billion Years Ago. Applied and Environmental Microbiology, 2003, 69, 5530-5535.	3.1	130
222	Antibiotic Susceptibility of <i>Tropheryma whipplei</i> in MRC5 Cells. Antimicrobial Agents and Chemotherapy, 2004, 48, 747-752.	3.2	130
223	Cryo-electron Microscopy of the Giant Mimivirus. Journal of Molecular Biology, 2005, 353, 493-496.	4.2	130
224	Scrub Typhus in Himalayas. Emerging Infectious Diseases, 2006, 12, 1590-1592.	4.3	130
225	Questions on Mediterranean Spotted Fever a Century after Its Discovery. Emerging Infectious Diseases, 2008, 14, 1360-1367.	4.3	130
226	Body Lice as Tools for Diagnosis and Surveillance of Reemerging Diseases. Journal of Clinical Microbiology, 1999, 37, 596-599.	3.9	130
227	Chronic Q fever: Expert opinion versus literature analysis and consensus. Journal of Infection, 2012, 65, 102-108.	3.3	128
228	Proposal to create subspecies of Rickettsia conorii based on multi-locus sequence typing and an emended description of Rickettsia conorii. BMC Microbiology, 2005, 5, 11.	3.3	127
229	A simple method for amplification of DNA from paraffin-embedded tissues. Nucleic Acids Research, 1992, 20, 5237-5238.	14.5	126
230	Diagnosis of infectious endocarditis in patients undergoing valve surgery. American Journal of Medicine, 2005, 118, 230-238.	1.5	126
231	The Point-of-Care Laboratory in Clinical Microbiology. Clinical Microbiology Reviews, 2016, 29, 429-447.	13.6	126
232	The Increase of Lactobacillus Species in the Gut Flora of Newborn Broiler Chicks and Ducks Is Associated with Weight Gain. PLoS ONE, 2010, 5, e10463.	2.5	125
233	The Discovery and Characterization of Mimivirus, the Largest Known Virus and Putative Pneumonia Agent. Clinical Infectious Diseases, 2007, 45, 95-102.	5.8	124
234	Tick-Borne Rickettsioses, Neglected Emerging Diseases in Rural Senegal. PLoS Neglected Tropical Diseases, 2010, 4, e821.	3.0	124

#	Article	IF	CITATIONS
235	Vancomycin Treatment of Infective Endocarditis Is Linked with Recently Acquired Obesity. PLoS ONE, 2010, 5, e9074.	2.5	124
236	Host Factors in the Severity of Q Fever. Annals of the New York Academy of Sciences, 1990, 590, 33-38.	3.8	123
237	Human Pathogens in Body and Head Lice. Emerging Infectious Diseases, 2002, 8, 1515-1518.	4.3	123
238	Lateral gene transfer between obligate intracellular bacteria: Evidence from the <i>Rickettsia massiliae</i> genome. Genome Research, 2007, 17, 1657-1664.	5.5	123
239	Isolation and identification of amoebaâ€resisting bacteria from water in human environment by using an <i>Acanthamoeba polyphaga</i> coâ€culture procedure. Environmental Microbiology, 2008, 10, 1135-1144.	3.8	123
240	The gut microbiota of a patient with resistant tuberculosis is more comprehensively studied by culturomics than by metagenomics. European Journal of Clinical Microbiology and Infectious Diseases, 2013, 32, 637-645.	2.9	123
241	Outbreak of epidemic typhus in Russia. Lancet, The, 1998, 352, 1151.	13.7	122
242	Whipple's Endocarditis: Review of the Literature and Comparisons with Q Fever, <i>Bartonella</i> Infection, and Blood Culture–Positive Endocarditis. Clinical Infectious Diseases, 2001, 33, 1309-1316.	5.8	122
243	Genomes of the Most Dangerous Epidemic Bacteria Have a Virulence Repertoire Characterized by Fewer Genes but More Toxin-Antitoxin Modules. PLoS ONE, 2011, 6, e17962.	2.5	122
244	Burden of emerging anaerobes in the MALDI-TOF and 16S rRNA gene sequencing era. Anaerobe, 2011, 17, 106-112.	2.1	122
245	Genotyping, evolution and epidemiological findings of Rickettsia species. Infection, Genetics and Evolution, 2014, 25, 122-137.	2.3	122
246	Natural History of <i>Bartonella</i> Infections (an Exception to Koch's Postulate). Vaccine Journal, 2002, 9, 8-18.	3.1	121
247	Q Fever During Pregnancy. Archives of Internal Medicine, 2002, 162, 701.	3.8	121
248	Acute Tick-borne Rickettsiosis Caused by <i>Rickettsia sibirica </i> in the Russian Far East. Emerging Infectious Diseases, 2004, 10, 810-817.	4.3	121
249	Ectoparasitism and Vector-Borne Diseases in 930 Homeless People From Marseilles. Medicine (United) Tj ETQq1 1	0.784314 1.0	l IggT /Ove
250	High Frequency of Tropheryma whipplei in Culture-Negative Endocarditis. Journal of Clinical Microbiology, 2012, 50, 216-222.	3.9	121
251	Crescent Bodies of Parachlamydia acanthamoeba and Its Life Cycle within Acanthamoeba polyphaga: an Electron Micrograph Study. Applied and Environmental Microbiology, 2002, 68, 3076-3084.	3.1	120
252	Emerging tools for identification of arthropod vectors. Future Microbiology, 2016, 11, 549-566.	2.0	120

#	Article	IF	Citations
253	Cardiac Valves in Patients with Q Fever Endocarditis: Microbiological, Molecular, and Histologic Studies. Journal of Infectious Diseases, 2003, 187, 1097-1106.	4.0	119
254	Molecular Evolution of Rickettsia Surface Antigens: Evidence of Positive Selection. Molecular Biology and Evolution, 2005, 22, 2073-2083.	8.9	119
255	<i>Yersinia pestis</i> Orientalis in Remains of Ancient Plague Patients. Emerging Infectious Diseases, 2007, 13, 332-333.	4.3	119
256	Long-term outcomes following infection of cardiac implantable electronic devices: a prospective matched cohort study. Heart, 2012, 98, 724-731.	2.9	119
257	Transmission potential of <i>Rickettsia felis</i> infection by <i>Anopheles gambiae</i> mosquitoes. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 8088-8093.	7.1	119
258	From culturomics to taxonomogenomics: A need to change the taxonomy of prokaryotes in clinical microbiology. Anaerobe, 2015, 36, 73-78.	2.1	119
259	Diet influence on the gut microbiota and dysbiosis related to nutritional disorders. Human Microbiome Journal, 2016, 1, 3-11.	3.8	119
260	Q feverâ€"a review and issues for the next century. International Journal of Antimicrobial Agents, 1997, 8, 145-161.	2.5	118
261	Amoebae-resisting Bacteria Isolated from Human Nasal Swabs by Amoebal Coculture. Emerging Infectious Diseases, 2004, 10, 470-477.	4.3	118
262	Complete rpoB gene sequencing as a suitable supplement to DNA-DNA hybridization for bacterial species and genus delineation. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 1807-1814.	1.7	118
263	Related Giant Viruses in Distant Locations and Different Habitats: Acanthamoeba polyphaga moumouvirus Represents a Third Lineage of the Mimiviridae That Is Close to the Megavirus Lineage. Genome Biology and Evolution, 2012, 4, 1324-1330.	2.5	118
264	Chemotherapy-induced ileal crypt apoptosis and the ileal microbiome shape immunosurveillance and prognosis of proximal colon cancer. Nature Medicine, 2020, 26, 919-931.	30.7	118
265	Preventing and Controlling Emerging and Reemerging Transmissible Diseases in the Homeless. Emerging Infectious Diseases, 2008, 14, 1353-1359.	4.3	116
266	Cowpox Virus Transmission from Pet Rats to Humans, France. Emerging Infectious Diseases, 2009, 15, 781-784.	<b>4.</b> 3	115
267	Genetic Identification of Rickettsiae Isolated from Ticks in Japan. Journal of Clinical Microbiology, 2002, 40, 2176-2181.	3.9	114
268	New <i>Rickettsiae</i> in Ticks Collected in Territories of the Former Soviet Union. Emerging Infectious Diseases, 1999, 5, 811-814.	4.3	113
269	Selfish DNA in Protein-Coding Genes of <i>Rickettsia</i> . Science, 2000, 290, 347-350.	12.6	113
270	Genomic analysis of an emerging multiresistant Staphylococcus aureus strain rapidly spreading in cystic fibrosis patients revealed the presence of an antibiotic inducible bacteriophage. Biology Direct, 2009, 4, 1.	4.6	113

#	Article	IF	CITATIONS
271	The Rhizome of the Multidrug-Resistant Enterobacter aerogenes Genome Reveals How New "Killer Bugs―Are Created because of a Sympatric Lifestyle. Molecular Biology and Evolution, 2013, 30, 369-383.	8.9	113
272	The human gut microbiome, a taxonomic conundrum. Systematic and Applied Microbiology, 2015, 38, 276-286.	2.8	113
273	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. FEMS Microbiology Reviews, 2010, 34, 281-94.	8.6	112
274	Citrate Synthase Gene Sequence: a New Tool for Phylogenetic Analysis and Identification of Ehrlichia. Journal of Clinical Microbiology, 2001, 39, 3031-3039.	3.9	111
275	First Isolation of Bartonella alsatica from a Valve of a Patient with Endocarditis. Journal of Clinical Microbiology, 2006, 44, 278-279.	3.9	111
276	Current Knowledge on Phylogeny and Taxonomy of <i>Rickettsia</i> spp Annals of the New York Academy of Sciences, 2009, 1166, 1-11.	3.8	111
277	Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry for Rapid Identification of Tick Vectors. Journal of Clinical Microbiology, 2013, 51, 522-528.	3.9	111
278	A Toxin-Antitoxin Module of Salmonella Promotes Virulence in Mice. PLoS Pathogens, 2013, 9, e1003827.	4.7	111
279	Viruses in the desert: a metagenomic survey of viral communities in four perennial ponds of the Mauritanian Sahara. ISME Journal, 2013, 7, 359-369.	9.8	111
280	Clinical detection and characterization of bacterial pathogens in the genomics era. Genome Medicine, 2014, 6, 114.	8.2	111
281	Treatment of classic Whipple's disease: from in vitro results to clinical outcome. Journal of Antimicrobial Chemotherapy, 2014, 69, 219-227.	3.0	111
282	Lipopolysaccharide from i>Coxiella burnetii li>Is Involved in Bacterial Phagocytosis, Filamentous Actin Reorganization, and Inflammatory Responses through Toll-Like Receptor 4. Journal of Immunology, 2004, 172, 3695-3703.	0.8	110
283	Giant Viruses of Amoebas: An Update. Frontiers in Microbiology, 2016, 7, 349.	3.5	110
284	Blood culture-negative endocarditis. Medicine (United States), 2017, 96, e8392.	1.0	110
285	The risk of COVID-19 death is much greater and age dependent with type I IFN autoantibodies. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2200413119.	7.1	110
286	African tick-bite fever: a new spotted fever group rickettsiosis under an old name. Lancet, The, 1992, 340, 982-983.	13.7	109
287	Evaluation of Antibiotic Susceptibilities of Three Rickettsial Species Including Rickettsia felis by a Quantitative PCR DNA Assay. Antimicrobial Agents and Chemotherapy, 2002, 46, 2747-2751.	3.2	109
288	Bartonella, a Common Cause of Endocarditis: a Report on 106 Cases and Review. Journal of Clinical Microbiology, 2015, 53, 824-829.	3.9	109

#	Article	IF	CITATIONS
289	<i>Clostridium butyricum</i> Strains and Dysbiosis Linked to Necrotizing Enterocolitis in Preterm Neonates. Clinical Infectious Diseases, 2015, 61, 1107-1115.	5.8	109
290	Rapid viral diagnosis and ambulatory management of suspected COVID-19 cases presenting at the infectious diseases referral hospital in Marseille, France, - January 31st to March 1st, 2020: A respiratory virus snapshot. Travel Medicine and Infectious Disease, 2020, 36, 101632.	3.0	109
291	Identification and Characterization of a Phospholipase D–Superfamily Gene in Rickettsiae. Journal of Infectious Diseases, 2003, 188, 1276-1283.	4.0	108
292	Role of Sex, Age, Previous Valve Lesion, and Pregnancy in the Clinical Expression and Outcome of Q Fever after a Large Outbreak. Clinical Infectious Diseases, 2007, 44, 232-237.	5.8	108
293	<i>Parachlamydiaceae</i> : Potential Emerging Pathogens. Emerging Infectious Diseases, 2002, 8, 626-630.	4.3	107
294	Analysis of the Rickettsia africae genome reveals that virulence acquisition in Rickettsia species may be explained by genome reduction. BMC Genomics, 2009, 10, 166.	2.8	107
295	<i>Borrelia</i> , <i>Rickettsia</i> , and <i>Ehrlichia</i> Species in Bat Ticks, France, 2010. Emerging Infectious Diseases, 2012, 18, 1966-1975.	4.3	107
296	Respiratory Viruses and Bacteria among Pilgrims during the 2013 Hajj. Emerging Infectious Diseases, 2014, 20, 1821-1827.	4.3	107
297	Potential for Tick-borne Bartonelloses. Emerging Infectious Diseases, 2010, 16, 385-391.	4.3	107
298	Q Fever During Pregnancy: A Public Health Problem in Southern France. Clinical Infectious Diseases, 1998, 27, 592-596.	5.8	106
299	Tick-Borne Relapsing Fever Borreliosis, Rural Senegal. Emerging Infectious Diseases, 2011, 17, 883-885.	4.3	106
300	Non-contiguous finished genome sequence and description of Anaerococcus senegalensis sp. nov Standards in Genomic Sciences, 2012, 6, 116-125.	1.5	106
301	"Marseilleviridaeâ€, a new family of giant viruses infecting amoebae. Archives of Virology, 2013, 158, 915-920.	2.1	106
302	Early Fecal Microbiota Transplantation Improves Survival in Severe Clostridium difficile Infections. Clinical Infectious Diseases, 2018, 66, 645-650.	5.8	106
303	Mink, SARS-CoV-2, and the Human-Animal Interface. Frontiers in Microbiology, 2021, 12, 663815.	3.5	106
304	Outbreak of <i>Rickettsia africae </i> Infections in Participants of an Adventure Race in South Africa. Clinical Infectious Diseases, 1998, 27, 316-323.	5.8	105
305	The history of epidemic typhus. Infectious Disease Clinics of North America, 2004, 18, 127-140.	5.1	105
306	Mediterranean Spotted Fever. Infectious Disease Clinics of North America, 2008, 22, 515-530.	5.1	105

#	Article	IF	Citations
307	Q Fever Pneumonia in French Guiana: Prevalence, Risk Factors, and Prognostic Score. Clinical Infectious Diseases, 2012, 55, 67-74.	5.8	105
308	Matrix-Assisted Laser Desorption Ionization - Time of Flight Mass Spectrometry: An Emerging Tool for the Rapid Identification of Mosquito Vectors. PLoS ONE, 2013, 8, e72380.	2.5	105
309	Diagnosis of Whipple Disease by Immunohistochemical Analysis. American Journal of Clinical Pathology, 2002, 118, 742-748.	0.7	104
310	<i>Bartonella henselae</i> in <i>Ixodes ricinus</i> Ticks (Acari: Ixodida) Removed from Humans, Belluno Province, Italy. Emerging Infectious Diseases, 2003, 9, 329-332.	4.3	104
311	<i>Tropheryma whipplei</i> in Children with Gastroenteritis. Emerging Infectious Diseases, 2010, 16, 776-782.	4.3	104
312	16S/23S rRNA Intergenic Spacer Regions for Phylogenetic Analysis, Identification, and Subtyping of Bartonella Species. Journal of Clinical Microbiology, 2001, 39, 2768-2778.	3.9	103
313	Parachlamydia acanthamoeba Enters and Multiplies within Human Macrophages and Induces Their Apoptosis. Infection and Immunity, 2003, 71, 5979-5985.	2.2	103
314	Common Epidemiology of <i>Rickettsia felis </i> li>Infection and Malaria, Africa. Emerging Infectious Diseases, 2013, 19, 1775-1783.	4.3	103
315	Immune responses during COVID-19 infection. Oncolmmunology, 2020, 9, 1807836.	4.6	103
316	Molecular Identification of <i>Gemella</i> Species from Three Patients with Endocarditis. Journal of Clinical Microbiology, 1998, 36, 866-871.	3.9	103
317	Arthropod-Borne Diseases in Homeless. Annals of the New York Academy of Sciences, 2006, 1078, 223-235.	3.8	102
318	Immuno-PCR: a promising ultrasensitive diagnostic method to detect antigens and antibodies. Trends in Microbiology, 2011, 19, 295-302.	7.7	102
319	Tropheryma whipplei and Whipple's disease. Journal of Infection, 2014, 69, 103-112.	3.3	102
320	Rickettsia felis: The Complex Journey of an Emergent Human Pathogen. Trends in Parasitology, 2016, 32, 554-564.	3.3	102
321	Comparison of clinical and echocardiographic characteristics of Streptococcus bovis endocarditis with that caused by other pathogens. American Journal of Cardiology, 2001, 88, 871-875.	1.6	101
322	Mycobacterium marseillense sp. nov., Mycobacterium timonense sp. nov. and Mycobacterium bouchedurhonense sp. nov., members of the Mycobacterium avium complex. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 2803-2808.	1.7	101
323	Zamilon, a Novel Virophage with Mimiviridae Host Specificity. PLoS ONE, 2014, 9, e94923.	2.5	101
324	A Metagenomic Investigation of the Duodenal Microbiota Reveals Links with Obesity. PLoS ONE, 2015, 10, e0137784.	2.5	101

#	Article	IF	CITATIONS
325	Emerging role of Raoultella ornithinolytica in human infections: a series of cases and review of the literature. International Journal of Infectious Diseases, 2016, 45, 65-71.	3.3	101
326	Natural history of COVID-19 and therapeutic options. Expert Review of Clinical Immunology, 2020, 16, 1159-1184.	3.0	101
327	Evidence of the megavirome in humans. Journal of Clinical Virology, 2013, 57, 191-200.	3.1	100
328	Point-of-Care Laboratory of Pathogen Diagnosis in Rural Senegal. PLoS Neglected Tropical Diseases, 2013, 7, e1999.	3.0	100
329	Goat-Associated Q Fever: A New Disease in Newfoundland. Emerging Infectious Diseases, 2001, 7, 413-419.	4.3	99
330	Rapid and Accurate Bacterial Identification in Probiotics and Yoghurts by MALDIâ€₹OF Mass Spectrometry. Journal of Food Science, 2011, 76, M568-72.	3.1	99
331	Virophages, polintons, and transpovirons: a complex evolutionary network of diverse selfish genetic elements with different reproduction strategies. Virology Journal, 2013, 10, 158.	3.4	99
332	Recent advances in the study of Q fever epidemiology, diagnosis and management. Journal of Infection, 2015, 71, S2-S9.	3.3	99
333	Pacmanvirus, a New Giant Icosahedral Virus at the Crossroads between Asfarviridae and Faustoviruses. Journal of Virology, 2017, 91, .	3.4	99
334	Excess mortality and morbidity in patients surviving infective endocarditis. American Heart Journal, 2012, 164, 94-101.	2.7	98
335	Cedratvirus, a Double-Cork Structured Giant Virus, is a Distant Relative of Pithoviruses. Viruses, 2016, 8, 300.	3.3	98
336	Validated Risk Score for Predicting 6â€Month Mortality in Infective Endocarditis. Journal of the American Heart Association, 2016, 5, e003016.	3.7	98
337	MIMIVIRE is a defence system in mimivirus that confers resistance to virophage. Nature, 2016, 531, 249-252.	27.8	98
338	Genetic Classification and Differentiation of Bartonella Species Based on Comparison of Partial ftsZ Gene Sequences. Journal of Clinical Microbiology, 2002, 40, 3641-3647.	3.9	97
339	What does the future hold for clinical microbiology?. Nature Reviews Microbiology, 2004, 2, 151-159.	28.6	97
340	New insight into the diagnosis of fastidious bacterial endocarditis. FEMS Immunology and Medical Microbiology, 2006, 47, 1-13.	2.7	97
341	Evaluation of the LightCycler $\hat{A}^{\otimes}$ SeptiFast test in the rapid etiologic diagnostic of infectious endocarditis. European Journal of Clinical Microbiology and Infectious Diseases, 2009, 28, 569-573.	2.9	97
342	Impaired Immune Functions of Monocytes and Macrophages in Whipple's Disease. Gastroenterology, 2010, 138, 210-220.	1.3	97

#	Article	IF	Citations
343	Q Fever in France, 1985–2009. Emerging Infectious Diseases, 2011, 17, 350-356.	4.3	97
344	Repertoire of Intensive Care Unit Pneumonia Microbiota. PLoS ONE, 2012, 7, e32486.	2.5	97
345	Mimiviridae: clusters of orthologous genes, reconstruction of gene repertoire evolution and proposed expansion of the giant virus family. Virology Journal, 2013, 10, 106.	3.4	97
346	Inhaled Lactonase Reduces Pseudomonas aeruginosa Quorum Sensing and Mortality in Rat Pneumonia. PLoS ONE, 2014, 9, e107125.	2.5	97
347	Molecular genetic methods for the diagnosis of fastidious microorganisms. Apmis, 2004, 112, 785-807.	2.0	96
348	rpoB sequence-based identification of Mycobacterium avium complex species. Microbiology (United) Tj ETQq0 C	0 rgBT /0	Overlock 10 Tf
349	Partial sequence comparison of the rpoB, sodA, groEL and gyrB genes within the genus Streptococcus. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 2317-2322.	1.7	96
350	Amoebae as Genitors and Reservoirs of Giant Viruses. Intervirology, 2010, 53, 321-329.	2.8	96
351	Biology and genetics of human head and body lice. Trends in Parasitology, 2012, 28, 563-571.	3.3	96
352	Coxiella burnetii Infection of Aneurysms or Vascular Grafts: Report of Seven Cases and Review. Clinical Infectious Diseases, 1998, 26, 116-121.	5.8	95
353	Genomic Variation of Bartonella henselae Strains Detected in Lymph Nodes of Patients with Cat Scratch Disease. Journal of Clinical Microbiology, 2002, 40, 1023-1030.	3.9	95
354	<i>Acinetobacter baumannii</i> in Human Body Louse. Emerging Infectious Diseases, 2004, 10, 1671-1673.	4.3	95
355	Experimental Model to Evaluate the Human Body Louse as a Vector of Plague. Journal of Infectious Diseases, 2006, 194, 1589-1596.	4.0	95
356	From Acute Q Fever to Endocarditis: Serological Follow-Up Strategy. Clinical Infectious Diseases, 2007, 44, 1337-1340.	5.8	95
357	Pigeon Pneumonia in Provence: A Bird-Borne Q Fever Outbreak. Clinical Infectious Diseases, 1999, 29, 617-620.	5.8	94
358	Molecular insights into the history of plague. Microbes and Infection, 2002, 4, 105-109.	1.9	94
359	Q Fever Pneumonia: Virulence of Coxiella burnetii Pathovars in a Murine Model of Aerosol Infection. Infection and Immunity, 2005, 73, 2469-2477.	2.2	94
360	Long-term persistence of virulent Yersinia pestis in soil. Microbiology (United Kingdom), 2008, 154, 2865-2871.	1.8	94

#	Article	IF	Citations
361	High Vaginal Concentrations of Atopobium vaginae and Gardnerella vaginalis in Women Undergoing Preterm Labor. Obstetrics and Gynecology, 2010, 115, 134-140.	2.4	94
362	Detection of Bartonella tamiae, Coxiella burnetii and rickettsiae in arthropods and tissues from wild and domestic animals in northeastern Algeria. Parasites and Vectors, 2016, 9, 27.	2.5	94
363	Asymptomatic hypoxia in COVID-19 is associated with poor outcome. International Journal of Infectious Diseases, 2021, 102, 233-238.	3.3	94
364	Whipple's disease: immunospecific and quantitative immunohistochemical study of intestinal biopsy specimens. Human Pathology, 2003, 34, 589-596.	2.0	93
365	Use of Genome Selected Repeated Sequences Increases the Sensitivity of PCR Detection of Tropheryma whipplei. Journal of Clinical Microbiology, 2004, 42, 401-403.	3.9	93
366	Isolation of Bartonella rattimassiliensis sp. nov. and Bartonella phoceensis sp. nov. from European Rattus norvegicus. Journal of Clinical Microbiology, 2004, 42, 3816-3818.	3.9	93
367	MALDI-TOF-MS for rapid detection of staphylococcal Panton–Valentine leukocidin. International Journal of Antimicrobial Agents, 2009, 34, 467-470.	2.5	93
368	The antimicrobial resistance pattern of cultured human methanogens reflects the unique phylogenetic position of archaea. Journal of Antimicrobial Chemotherapy, 2011, 66, 2038-2044.	3.0	93
369	Metagenomic Analysis of Brain Abscesses Identifies Specific Bacterial Associations. Clinical Infectious Diseases, 2012, 54, 202-210.	5 <b>.</b> 8	93
370	Non contiguous-finished genome sequence and description of Peptoniphilus timonensis sp. nov Standards in Genomic Sciences, 2012, 7, 1-11.	1.5	93
371	Marseillevirus-Like Virus Recovered From Blood Donated by Asymptomatic Humans. Journal of Infectious Diseases, 2013, 208, 1042-1050.	4.0	93
372	Gut Bacteria Missing in Severe Acute Malnutrition, Can We Identify Potential Probiotics by Culturomics?. Frontiers in Microbiology, 2017, 8, 899.	<b>3.</b> 5	93
373	Molecular and MALDI-TOF identification of ticks and tick-associated bacteria in Mali. PLoS Neglected Tropical Diseases, 2017, 11, e0005762.	3.0	93
374	Serological Differentiation of Murine Typhus and Epidemic Typhus Using Cross-Adsorption and Western Blotting. Vaccine Journal, 2000, 7, 612-616.	2.6	92
375	Diagnostic methods. Infectious Disease Clinics of North America, 2002, 16, 339-361.	5.1	92
376	Dysregulation of Cytokines in Acute Q Fever: Role of Interleukinâ€10 and Tumor Necrosis Factor in Chronic Evolution of Q Fever. Journal of Infectious Diseases, 2003, 187, 956-962.	4.0	92
377	Paenibacillus massiliensis sp. nov., Paenibacillus sanguinis sp. nov. and Paenibacillus timonensis sp. nov., isolated from blood cultures. International Journal of Systematic and Evolutionary Microbiology, 2004, 54, 1049-1054.	1.7	91
378	Seasonal Variation in <i>Klebsiella pneumoniae </i> Infectious Diseases, 2008, 197, 752-756.	4.0	91

#	Article	IF	CITATIONS
379	<i>Tropheryma whipplei</i> in Patients with Pneumonia. Emerging Infectious Diseases, 2010, 16, 258-263.	4.3	91
380	A Versatile Medium for Cultivating Methanogenic Archaea. PLoS ONE, 2013, 8, e61563.	2.5	91
381	Immunohistologic demonstration of coxiella burnetii in the valves of patients with Q fever endocarditis. American Journal of Medicine, 1994, 97, 451-458.	1.5	90
382	Bartonella (Rochalimaea) quintana isolation In patient with chronic adenopathy, lymphopenia, and a cat. Lancet, The, 1994, 343, 977.	13.7	90
383	Brill-Zinsser disease in France. Lancet, The, 1999, 353, 1936.	13.7	90
384	Naming of Rickettsiae and Rickettsial Diseases. Annals of the New York Academy of Sciences, 2005, 1063, 1-12.	3.8	90
385	Defining Pathogenic Bacterial Species in the Genomic Era. Frontiers in Microbiology, 2010, 1, 151.	3.5	90
386	Plague: History and contemporary analysis. Journal of Infection, 2013, 66, 18-26.	3.3	90
387	Identification of flea species using MALDI-TOF/MS. Comparative Immunology, Microbiology and Infectious Diseases, 2014, 37, 153-157.	1.6	90
388	Yersinia pestis: the Natural History of Plague. Clinical Microbiology Reviews, 2020, 34, .	13.6	90
389	Diagnosis of <i>Bartonella</i> Endocarditis by a Real-Time Nested PCR Assay Using Serum. Journal of Clinical Microbiology, 2003, 41, 919-925.	3.9	89
390	Bacteriophages and diffusion of genes encoding antimicrobial resistance in cystic fibrosis sputum microbiota. Journal of Antimicrobial Chemotherapy, 2011, 66, 2448-2454.	3.0	89
391	Non-contiguous finished genome sequence and description of Alistipes timonensis sp. nov Standards in Genomic Sciences, 2012, 6, 315-324.	1.5	89
392	Colistin: An Antimicrobial for the 21st Century?. Clinical Infectious Diseases, 2002, 35, 901-902.	5.8	88
393	Bartonella quintana in human erythrocytes. Lancet, The, 2002, 360, 226-228.	13.7	88
394	Nanobacteria Are Mineralo Fetuin Complexes. PLoS Pathogens, 2008, 4, e41.	4.7	88
395	A Decade of Improvements in Mimiviridae and Marseilleviridae Isolation from Amoeba. Intervirology, 2013, 56, 354-363.	2.8	88
396	Abnormal Weight Gain and Gut Microbiota Modifications Are Side Effects of Long-Term Doxycycline and Hydroxychloroquine Treatment. Antimicrobial Agents and Chemotherapy, 2014, 58, 3342-3347.	3.2	88

#	Article	IF	Citations
397	High Atopobium vaginae and Gardnerella vaginalis Vaginal Loads Are Associated With Preterm Birth. Clinical Infectious Diseases, 2015, 60, 860-867.	5.8	88
398	Diagnosis of Mediterranean Spotted Fever by Indirect Immunofluorescence of Rickettsia conorii in Circulating Endothelial Cells Isolated with Monoclonal Antibody-Coated Immunomagnetic Beads. Journal of Infectious Diseases, 1992, 166, 660-663.	4.0	87
399	Interleukin-10 Stimulates Coxiella burnetiiReplication in Human Monocytes through Tumor Necrosis Factor Down-Modulation: Role in Microbicidal Defect of Q Fever. Infection and Immunity, 2001, 69, 2345-2352.	2.2	87
400	Legionella drancourtii sp. nov., a strictly intracellular amoebal pathogen. International Journal of Systematic and Evolutionary Microbiology, 2004, 54, 699-703.	1.7	87
401	Molecular Identification of Lice from Preâ€Columbian Mummies. Journal of Infectious Diseases, 2008, 197, 535-543.	4.0	87
402	Samba virus: a novel mimivirus from a giant rain forest, the Brazilian Amazon. Virology Journal, 2014, 11, 95.	3.4	87
403	microbiota of an obese patient and reclassification of Ruminococcus faecis, Ruminococcus lactaris, Ruminococcus torques, Ruminococcus gnavus and Clostridium glycyrrhizinilyticum as Mediterraneibacter faecis comb. nov., Mediterraneibacter lactaris comb. nov., Mediterraneibacter torques comb. nov., Mediterraneibacter gnavus comb. nov. and Mediterraneibacter	1.7	87
404	glycyrrhizinllyticus comb. nov. Antonie Van Leeuwenhoek, 2018, 111, 2107-2128. Suicide PCR on Skin Biopsy Specimens for Diagnosis of Rickettsioses. Journal of Clinical Microbiology, 2004, 42, 3428-3434.	3.9	86
405	Cytomegalovirus and Herpes Simplex Virus Effect on the Prognosis of Mechanically Ventilated Patients Suspected to Have Ventilator-Associated Pneumonia. PLoS ONE, 2012, 7, e51340.	2.5	86
406	Do we need new antibiotics?. Clinical Microbiology and Infection, 2016, 22, 408-415.	6.0	86
407	Molecular Detection of Eukaryotes in a Single Human Stool Sample from Senegal. PLoS ONE, 2012, 7, e40888.	2.5	86
408	A Natural Polyphenol Exerts Antitumor Activity and Circumvents Anti–PD-1 Resistance through Effects on the Gut Microbiota. Cancer Discovery, 2022, 12, 1070-1087.	9.4	86
409	Survival of Tropheryma whipplei, the Agent of Whipple's Disease, Requires Phagosome Acidification. Infection and Immunity, 2002, 70, 1501-1506.	2.2	85
410	<i>Tropheryma whipplei</i> Bacteremia during Fever in Rural West Africa. Clinical Infectious Diseases, 2010, 51, 515-521.	5.8	85
411	Metabolic role of lactobacilli in weight modification in humans and animals. Microbial Pathogenesis, 2017, 106, 182-194.	2.9	85
412	Infectious Disease Risk Across the Growing Human-Non Human Primate Interface: A Review of the Evidence. Frontiers in Public Health, 2019, 7, 305.	2.7	85
413	Repertoire of human breast and milk microbiota: a systematic review. Future Microbiology, 2019, 14, 623-641.	2.0	85
414	Q fever and HIV infection. Aids, 1993, 7, 81-86.	2,2	84

#	Article	IF	CITATIONS
415	Surgical Treatment of Active Aortic Endocarditis: Homografts Are Not the Cornerstone of Outcome. Annals of Thoracic Surgery, 2007, 84, 1935-1942.	1.3	84
416	Rapid identification and typing of Yersinia pestis and other Yersinia species by matrix-assisted laser desorption/ionization time-of-flight (MALDI-TOF) mass spectrometry. BMC Microbiology, 2010, 10, 285.	3.3	84
417	Diagnostic yield of FDG positron-emission tomography/computed tomography in patients with CEID infection: a pilot study. Europace, 2013, 15, 252-257.	1.7	84
418	Culturomics and pyrosequencing evidence of the reduction in gut microbiota diversity in patients with broad-spectrum antibiotics. International Journal of Antimicrobial Agents, 2014, 44, 117-124.	2.5	84
419	Impact of Early Valve Surgery on Outcome of Staphylococcus aureus Prosthetic Valve Infective Endocarditis: Analysis in the International Collaboration of Endocarditis–Prospective Cohort Study. Clinical Infectious Diseases, 2015, 60, 741-749.	5.8	84
420	Identification of Novel Zoonotic Activity of Bartonella spp., France. Emerging Infectious Diseases, 2016, 22, 457-462.	4.3	84
421	Antimicrobial susceptibility of Rochalimaea quintana, Rocholimaea vinsonii, and the newly recognized Rochalimaea henselae. Journal of Antimicrobial Chemotherapy, 1993, 32, 587-594.	3.0	83
422	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.	7.4	83
423	Neurological Involvement in Acute Q Fever. Archives of Internal Medicine, 2002, 162, 693.	3.8	83
424	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
425	Bacterial Zoonoses and Infective Endocarditis, Algeria. Emerging Infectious Diseases, 2005, 11, 216-224.	4.3	83
426	Bartonella quintanain a 4000‥earâ€Old Human Tooth. Journal of Infectious Diseases, 2005, 191, 607-611.	4.0	83
427	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
428	Viruses with More Than 1,000 Genes: Mamavirus, a New Acanthamoeba polyphagamimivirus Strain, and Reannotation of Mimivirus Genes. Genome Biology and Evolution, 2011, 3, 737-742.	2.5	83
429	Identification of Rickettsial Infections by Using Cutaneous Swab Specimens and PCR. Emerging Infectious Diseases, 2011, 17, 83-86.	4.3	83
430	Central Nervous System Involvement in Whipple Disease. Medicine (United States), 2013, 92, 324-330.	1.0	83
431	Bacillary angiomatosis in immunocompromised patients. Aids, 1998, 12, 1793-1803.	2.2	82
432	Phylogenetic classification of Bartonella species by comparing groEL sequences International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 165-171.	1.7	82

#	Article	IF	CITATIONS
433	IL-16 Is Critical for <i>Tropheryma whipplei</i> Replication in Whipple's Disease. Journal of Immunology, 2005, 175, 4575-4582.	0.8	82
434	Bartonella vinsonii subsp. arupensis as an Agent of Blood Culture-Negative Endocarditis in a Human. Journal of Clinical Microbiology, 2005, 43, 945-947.	3.9	82
435	Paper money and coins as potential vectors of transmissible disease. Future Microbiology, 2014, 9, 249-261.	2.0	82
436	Multiple Pathogens Including Potential New Species in Tick Vectors in Côte d'Ivoire. PLoS Neglected Tropical Diseases, 2016, 10, e0004367.	3.0	82
437	Survey of Three Bacterial Louseâ€Associated Diseases Among Rural Andean Communities in Peru: Prevalence of Epidemic Typhus, Trench Fever, and Relapsing Fever. Clinical Infectious Diseases, 1999, 29, 434-436.	5.8	81
438	Genotyping reveals a wide heterogeneity of Tropheryma whipplei. Microbiology (United Kingdom), 2008, 154, 521-527.	1.8	81
439	Tick-borne rickettsioses in America: Unanswered questions and emerging diseases. Current Infectious Disease Reports, 2009, 11, 40-50.	3.0	81
440	The origin and distribution of human lice in the world. Infection, Genetics and Evolution, 2014, 23, 209-217.	2.3	81
441	Quantitative Detection of <i>Tropheryma whipplei</i> DNA by Real-Time PCR. Journal of Clinical Microbiology, 2002, 40, 1119-1120.	3.9	80
442	An Experimental Model of Human Body Louse Infection with Rickettsia prowazekii. Journal of Infectious Diseases, 2002, 186, 1639-1646.	4.0	80
443	Rickettsia tamurae sp. nov., isolated from Amblyomma testudinarium ticks. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1673-1675.	1.7	80
444	Usefulness of broad-range PCR for the diagnosis of osteoarticular infections. Current Opinion in Rheumatology, 2008, 20, 463-470.	4.3	80
445	High Prevalence of Fastidious Bacteria in 1520 Cases of Uveitis of Unknown Etiology. Medicine (United) Tj ETQq1	1,0.78431 1.0	14 rgBT /CIV 80
446	Q Fever during Pregnancy. Annals of the New York Academy of Sciences, 2009, 1166, 79-89.	3.8	80
447	Lymph Node Biopsy Specimens and Diagnosis of Cat-scratch Disease. Emerging Infectious Diseases, 2006, 12, 1338-1344.	4.3	79
448	Bacterial genome sequencing and its use in infectious diseases. Lancet Infectious Diseases, The, 2007, 7, 711-723.	9.1	79
449	A bioinformatic approach to understanding antibiotic resistance in intracellular bacteria through whole genome analysis. International Journal of Antimicrobial Agents, 2008, 32, 207-220.	2.5	79
450	Persistent Coxiella burnetii Infection in Mice Overexpressing IL-10: An Efficient Model for Chronic Q Fever Pathogenesis. PLoS Pathogens, 2008, 4, e23.	4.7	79

#	Article	IF	CITATIONS
451	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
452	Murine Typhus in Returned Travelers: A Report of Thirty-Two Cases. American Journal of Tropical Medicine and Hygiene, 2012, 86, 1049-1053.	1.4	79
453	Treatment of <i>Rickettsia </i> spp. infections: a review. Expert Review of Anti-Infective Therapy, 2012, 10, 1425-1437.	4.4	79
454	Identification of European mosquito species by MALDI-TOF MS. Parasitology Research, 2014, 113, 2375-2378.	1.6	79
455	Rhodococcus equi infection in patients with AIDS. Journal of Infection, 1992, 24, 123-131.	3.3	78
456	Use of Shell-Vial Cell Culture Assay for Isolation of Bacteria from Clinical Specimens: 13 Years of Experience. Journal of Clinical Microbiology, 2005, 43, 4993-5002.	3.9	78
457	Bartonella rattaustraliani sp. nov., Bartonella queenslandensis sp. nov. and Bartonella coopersplainsensis sp. nov., identified in Australian rats. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 2956-2961.	1.7	78
458	Probiotics and obesity: a link?. Nature Reviews Microbiology, 2009, 7, 616-616.	28.6	78
459	<i>Candidatus</i> Bartonella mayotimonensis and Endocarditis. Emerging Infectious Diseases, 2010, 16, 500-503.	4.3	78
460	Complete Genome Sequence of Methanomassiliicoccus luminyensis, the Largest Genome of a Human-Associated Archaea Species. Journal of Bacteriology, 2012, 194, 4745-4745.	2.2	78
461	B-cell non-Hodgkin lymphoma linked to Coxiella burnetii. Blood, 2016, 127, 113-121.	1.4	78
462	Culturomics and Amplicon-based Metagenomic Approaches for the Study of Fungal Population in Human Gut Microbiota. Scientific Reports, 2017, 7, 16788.	3.3	78
463	Azithromycin Inhibits the Replication of Zika Virus. Journal of Antivirals & Antiretrovirals, 2018, 10, .	0.1	78
464	The nasopharyngeal microbiota in patients with viral respiratory tract infections is enriched in bacterial pathogens. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 1725-1733.	2.9	78
465	Bosea eneae sp. nov., Bosea massiliensis sp. nov. and Bosea vestrisii sp. nov., isolated from hospital water supplies, and emendation of the genus Bosea (Das et al. 1996). International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 15-20.	1.7	77
466	Whipple Disease: Intestinal Infiltrating Cells Exhibit a Transcriptional Pattern of M2/Alternatively Activated Macrophages. Journal of Infectious Diseases, 2005, 192, 1642-1646.	4.0	77
467	Tropical rickettsioses. Clinics in Dermatology, 2006, 24, 191-200.	1.6	77
468	The post-Darwinist rhizome of life. Lancet, The, 2010, 375, 104-105.	13.7	77

#	Article	IF	CITATIONS
469	Evidence of lifetime susceptibility to Tropheryma whipplei in patients with Whipple's disease. Journal of Antimicrobial Chemotherapy, 2011, 66, 1188-1189.	3.0	77
470	Evolution From Acute Q Fever to Endocarditis Is Associated With Underlying Valvulopathy and Age and Can Be Prevented by Prolonged Antibiotic Treatment. Clinical Infectious Diseases, 2013, 57, 836-844.	5.8	77
471	Development of a new PCR-based assay to detect Anaplasmataceae and the first report of Anaplasma phagocytophilum and Anaplasma platys in cattle from Algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2015, 39, 39-45.	1.6	77
472	Clinical Features and Complications of <i>Coxiella burnetii </i> li>Infections From the French National Reference Center for Q Fever. JAMA Network Open, 2018, 1, e181580.	5.9	77
473	Genome sequence and description of Aeromicrobium massiliense sp. nov Standards in Genomic Sciences, 2012, 7, 246-257.	1.5	76
474	Endosymbiotic bacteria associated with nematodes, ticks and amoebae. FEMS Immunology and Medical Microbiology, 2012, 64, 21-31.	2.7	76
475	Detection of Rickettsia felis, Rickettsia typhi, Bartonella Species and Yersinia pestis in Fleas (Siphonaptera) from Africa. PLoS Neglected Tropical Diseases, 2014, 8, e3152.	3.0	76
476	Quantitative Analysis of Valvular Lesions During Bartonella Endocarditis. American Journal of Clinical Pathology, 2000, 114, 880-889.	0.7	75
477	Some lessons fromRickettsiagenomics. FEMS Microbiology Reviews, 2005, 29, 99-117.	8.6	75
478	Obesity pandemics and the modification of digestive bacterial flora. European Journal of Clinical Microbiology and Infectious Diseases, 2008, 27, 631-634.	2.9	75
479	Systematic PCR Detection in Culture-negative Osteoarticular Infections. American Journal of Medicine, 2013, 126, 1143.e25-1143.e33.	1.5	75
480	<i>Tropheryma whipplei</i> Endocarditis. Emerging Infectious Diseases, 2013, 19, 1721-30.	4.3	75
481	Kaumoebavirus, a New Virus That Clusters with Faustoviruses and Asfarviridae. Viruses, 2016, 8, 278.	3.3	75
482	Survey of laboratory-acquired infections around the world in biosafety level 3 and 4 laboratories. European Journal of Clinical Microbiology and Infectious Diseases, 2016, 35, 1247-1258.	2.9	75
483	Plant and Fungal Diversity in Gut Microbiota as Revealed by Molecular and Culture Investigations. PLoS ONE, 2013, 8, e59474.	2.5	75
484	Shan Virus: A New Mimivirus Isolated from the Stool of a Tunisian Patient with Pneumonia. Intervirology, 2013, 56, 424-429.	2.8	74
485	Common occurrence of antibacterial agents in human intestinal microbiota. Frontiers in Microbiology, 2015, 6, 441.	3.5	74
486	Atomic force microscopy investigation of the giant mimivirus. Virology, 2010, 404, 127-137.	2.4	73

#	Article	IF	CITATIONS
487	The Rhizome of Life: The Sympatric Rickettsia felis Paradigm Demonstrates the Random Transfer of DNA Sequences. Molecular Biology and Evolution, 2011, 28, 3213-3223.	8.9	73
488	Structure of Sputnik, a virophage, at 3.5-Ã resolution. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 18431-18436.	7.1	73
489	Accurate identification of Culicidae at aquatic developmental stages by MALDI-TOF MS profiling. Parasites and Vectors, 2014, 7, 544.	2.5	73
490	Competence of Cimex lectularius Bed Bugs for the Transmission of Bartonella quintana, the Agent of Trench Fever. PLoS Neglected Tropical Diseases, 2015, 9, e0003789.	3.0	73
491	â€~ <i>Candidatus Rickettsia tarasevichiae</i> ' in <i>Ixodes persulcatus</i> Ticks Collected in Russia. Annals of the New York Academy of Sciences, 2003, 990, 162-172.	3.8	72
492	Laboratory Infection of a Technician by Mimivirus. Annals of Internal Medicine, 2006, 144, 702.	3.9	72
493	Use of pyrosequencing and DNA barcodes to monitor variations in Firmicutes and Bacteroidetes communities in the gut microbiota of obese humans. BMC Genomics, 2008, 9, 576.	2.8	72
494	Altitude-dependent i>Bartonella quintana / i>Genotype C in Head Lice, Ethiopia. Emerging Infectious Diseases, 2011, 17, 2357-2359.	4.3	72
495	Lack of MERS Coronavirus but Prevalence of Influenza Virus in French Pilgrims after 2013 Hajj. Emerging Infectious Diseases, 2014, 20, 726-728.	4.3	72
496	Prognostic Value of 18F-Fluorodeoxyglucose Positron Emission Tomography/Computed Tomography in Infective Endocarditis. Journal of the American College of Cardiology, 2019, 74, 1031-1040.	2.8	72
497	FIRST MOLECULAR DETECTION OF RICKETTSIA FELIS IN FLEAS FROM ALGERIA. American Journal of Tropical Medicine and Hygiene, 2006, 74, 532-535.	1.4	72
498	Prevalence and clinical significance of IgG isotype anti- $\hat{l}^2$ 2-glycoprotein I antibodies in antiphospholipid syndrome: A comparative study w. Translational Research, 1997, 129, 499-506.	2.3	71
499	Spotted Fever Group Rickettsiae in Ticks, Morocco. Emerging Infectious Diseases, 2008, 14, 1067-1073.	4.3	71
500	Babela massiliensis, a representative of a widespread bacterial phylum with unusual adaptations to parasitism in amoebae. Biology Direct, 2015, 10, 13.	4.6	71
501	Traditional and syndromic surveillance of infectious diseases and pathogens. International Journal of Infectious Diseases, 2016, 48, 22-28.	3.3	71
502	Coxiella burnetii seropositivity in parturient women is associated with adverse pregnancy outcomes. American Journal of Obstetrics and Gynecology, 2003, 189, 228-232.	1.3	70
503	Successive Emergence of Enterobacter aerogenes Strains Resistant to Imipenem and Colistin in a Patient. Antimicrobial Agents and Chemotherapy, 2005, 49, 1354-1358.	3.2	70
504	Whipple's Disease: a Macrophage Disease. Vaccine Journal, 2006, 13, 170-178.	3.1	70

#	Article	IF	Citations
505	High Throughput Sequencing and Proteomics to Identify Immunogenic Proteins of a New Pathogen: The Dirty Genome Approach. PLoS ONE, 2009, 4, e8423.	2.5	70
506	Failure and relapse after treatment with trimethoprim/sulfamethoxazole in classic Whipple's disease. Journal of Antimicrobial Chemotherapy, 2010, 65, 2005-2012.	3.0	70
507	Acinetobacter baumannii Resistant to Colistin With Impaired Virulence: A Case Report From France. Journal of Infectious Diseases, 2011, 204, 1146-1147.	4.0	70
508	Non-contiguous finished genome sequence and description of Clostridium senegalense sp. nov Standards in Genomic Sciences, 2012, 6, 401-410.	1.5	70
509	Non contiguous-finished genome sequence and description of Bacillus timonensis sp. nov Standards in Genomic Sciences, 2012, 6, 346-355.	1.5	70
510	Non-contiguous finished genome sequence and description of Herbaspirillum massiliense sp. nov Standards in Genomic Sciences, 2012, 7, 1-14.	1.5	70
511	Lactobacillus rhamnosus bacteremia: an emerging clinical entity. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 2469-2480.	2.9	70
512	Real-Time Sequencing To Decipher the Molecular Mechanism of Resistance of a Clinical Pan-Drug-Resistant Acinetobacter baumannii Isolate from Marseille, France. Antimicrobial Agents and Chemotherapy, 2013, 57, 592-596.	3.2	70
513	Diagnosis of blood culture-negative endocarditis and clinical comparison between blood culture-negative and blood culture-positive cases. Infection, 2016, 44, 459-466.	4.7	70
514	Repertoire of human gut microbes. Microbial Pathogenesis, 2017, 106, 103-112.	2.9	70
515	Orpheovirus IHUMI-LCC2: A New Virus among the Giant Viruses. Frontiers in Microbiology, 2017, 8, 2643.	3.5	70
516	Expression of ACE2, Soluble ACE2, Angiotensin I, Angiotensin II and Angiotensin-(1-7) Is Modulated in COVID-19 Patients. Frontiers in Immunology, 2021, 12, 625732.	4.8	70
517	A Novel Obligate Intracellular Gamma-Proteobacterium Associated with Ixodid Ticks, Diplorickettsia massiliensis, Gen. Nov., Sp. Nov. PLoS ONE, 2010, 5, e11478.	2.5	70
518	Genomics and metagenomics in medical microbiology. Journal of Microbiological Methods, 2013, 95, 415-424.	1.6	69
519	Complete genome sequence of Tunisvirus, a new member of the proposed family Marseilleviridae. Archives of Virology, 2014, 159, 2349-2358.	2.1	69
520	Etiology of Pericarditis in a Prospective Cohort of 1162 Cases. American Journal of Medicine, 2015, 128, 784.e1-784.e8.	1.5	69
521	Eukaryote Culturomics of the Gut Reveals New Species. PLoS ONE, 2014, 9, e106994.	2.5	69
522	Cardiac Valves in Patients with Whipple Endocarditis: Microbiological, Molecular, Quantitative Histologic, and Immunohistochemical Studies of 5 Patients. Journal of Infectious Diseases, 2004, 190, 935-945.	4.0	68

#	Article	IF	CITATIONS
523	Rickettsia parkeriInfection and Other Spotted Fevers in the United States. New England Journal of Medicine, 2005, 353, 626-627.	27.0	68
524	PartialrpoBgene sequencing for identification of Leptospiraspecies. FEMS Microbiology Letters, 2006, 263, 142-147.	1.8	68
525	Description of Mycobacterium conceptionense sp. nov., a Mycobacterium fortuitum Group Organism Isolated from a Posttraumatic Osteitis Inflammation. Journal of Clinical Microbiology, 2006, 44, 1268-1273.	3.9	68
526	Oral Ivermectin in the Treatment of Body Lice. Journal of Infectious Diseases, 2006, 193, 474-476.	4.0	68
527	Ageâ€related prevalence of <i><scp>M</scp>ethanomassiliicoccus luminyensis</i> in the human gut microbiome. Apmis, 2012, 120, 773-777.	2.0	68
528	Postgenomic analysis of bacterial pathogens repertoire reveals genome reduction rather than virulence factors. Briefings in Functional Genomics, 2013, 12, 291-304.	2.7	68
529	Can Plant Viruses Cross the Kingdom Border and Be Pathogenic to Humans?. Viruses, 2015, 7, 2074-2098.	3.3	68
530	Review of European and American guidelines for the diagnosis of Lyme borreliosis. Médecine Et Maladies Infectieuses, 2019, 49, 121-132.	5.0	68
531	Whipple's Disease. Vaccine Journal, 2001, 8, 1-8.	2.6	67
532	Value of Microimmunofluorescence for Diagnosis and Follow-up of Bartonella Endocarditis. Vaccine Journal, 2002, 9, 795-801.	3.1	67
533	Detection of Mycobacterium tuberculosis complex organisms in the stools of patients with pulmonary tuberculosis. Microbiology (United Kingdom), 2009, 155, 2384-2389.	1.8	67
534	Orientia tsutsugamushi Stimulates an Original Gene Expression Program in Monocytes: Relationship with Gene Expression in Patients with Scrub Typhus. PLoS Neglected Tropical Diseases, 2011, 5, e1028.	3.0	67
535	Highâ€throughput isolation of giant viruses of the <i><scp>M</scp>imiviridae</i> and <i><scp>M</scp>arseilleviridae</i> families in the <scp>T</scp> unisian environment. Environmental Microbiology, 2013, 15, 2000-2007.	3.8	67
536	Coxiella burnetii DNA, But Not Viable Bacteria, in Dairy Products in France. American Journal of Tropical Medicine and Hygiene, 2013, 88, 765-769.	1.4	67
537	New insights in gut microbiota and mucosal immunity of the small intestine. Human Microbiome Journal, 2018, 7-8, 23-32.	3.8	67
538	Identification of Rickettsiae from Ticks Collected in the Central African Republic Using the Polymerase Chain Reaction. American Journal of Tropical Medicine and Hygiene, 1994, 50, 373-380.	1.4	67
539	Traditional and Molecular Techniques for the Study of Emerging Bacterial Diseases: One Laboratory's Perspective. Emerging Infectious Diseases, 2002, 8, 122-131.	4.3	66
540	Non-contiguous finished genome sequence and description of Paenibacillus senegalensis sp. nov Standards in Genomic Sciences, 2012, 7, 70-81.	1.5	66

#	Article	IF	CITATIONS
541	Non contiguous-finished genome sequence and description of Cellulomonas massiliensis sp. nov Standards in Genomic Sciences, 2012, 7, 258-270.	1.5	66
542	Multiple tick-associated bacteria in Ixodes ricinus from Slovakia. Ticks and Tick-borne Diseases, 2012, 3, 406-410.	2.7	66
543	Molecular Detection of Spotted Fever Group Rickettsiae Associated with Ixodid Ticks in Egypt. Vector-Borne and Zoonotic Diseases, 2012, 12, 346-359.	1.5	66
544	Non contiguous-finished genome sequence and description of Senegalemassilia anaerobia gen. nov., sp. nov Standards in Genomic Sciences, 2013, 7, 343-356.	1.5	66
545	Non contiguous-finished genome sequence and description of Enterobacter massiliensis sp. nov Standards in Genomic Sciences, 2013, 7, 399-412.	1.5	66
546	Infective Endocarditis: Prevention, Diagnosis, and Management. Canadian Journal of Cardiology, 2014, 30, 1046-1057.	1.7	66
547	Isolation of new Brazilian giant viruses from environmental samples using a panel of protozoa. Frontiers in Microbiology, 2015, 6, 1086.	3.5	66
548	Gut microbiota associated with HIV infection is significantly enriched in bacteria tolerant to oxygen. BMJ Open Gastroenterology, 2016, 3, e000080.	2.7	66
549	Analysis of the 16S rRNA Gene Sequence of Anaplasma centrale and Its Phylogenetic Relatedness to Other Ehrlichiae. Vaccine Journal, 2001, 8, 241-244.	2.6	65
550	Serological Hint Suggesting That Parachlamydiaceae Are Agents of Pneumonia in Polytraumatized Intensive Care Patients. Annals of the New York Academy of Sciences, 2003, 990, 311-319.	3.8	65
551	Attempted Isolation of Nanobacterium sp. Microorganisms from Upper Urinary Tract Stones. Journal of Clinical Microbiology, 2003, 41, 368-372.	3.9	65
552	Detection of a Rickettsia Closely Related to Rickettsia aeschlimannii , " Rickettsia heilongjiangensis ,― Rickettsia sp. Strain RpA4, and Ehrlichia muris in Ticks Collected in Russia and Kazakhstan. Journal of Clinical Microbiology, 2004, 42, 2221-2223.	3.9	65
553	Endocarditis After Acute Q Fever in Patients with Previously Undiagnosed Valvulopathies. Clinical Infectious Diseases, 2006, 42, 818-821.	5.8	65
554	Pneumonia in mice inoculated experimentally with Acanthamoeba polyphaga mimivirus. Microbial Pathogenesis, 2007, 42, 56-61.	2.9	65
555	False positive PCR detection of Tropheryma whipplei in the saliva of healthy people. BMC Microbiology, 2007, 7, 48.	3.3	65
556	Revolutionizing Clinical Microbiology Laboratory Organization in Hospitals with In Situ Point-of-Care. PLoS ONE, 2011, 6, e22403.	2.5	65
557	Isolation of Vermamoeba vermiformis and associated bacteria in hospital water. Microbial Pathogenesis, 2015, 80, 14-20.	2.9	65
558	Diagnosis of Infective Endocarditis AfterÂTAVR. JACC: Cardiovascular Imaging, 2018, 11, 143-146.	5 <b>.</b> 3	65

#	Article	IF	Citations
559	The Impact of Human Immunodeficiency Virus Infection on Gut Microbiota α-Diversity: An Individual-level Meta-analysis. Clinical Infectious Diseases, 2020, 70, 615-627.	5.8	65
560	" Actinobaculum massiliae ,―a New Species Causing Chronic Urinary Tract Infection. Journal of Clinical Microbiology, 2002, 40, 3938-3941.	3.9	64
561	<i>Parachlamydia acanthamoeba</i> Is Endosymbiotic or Lytic for <i>Acanthamoeba polyphaga</i> Depending on the Incubation Temperature. Annals of the New York Academy of Sciences, 2003, 990, 628-634.	3.8	64
562	Link between Impaired Maturation of Phagosomes and DefectiveCoxiella burnetiiKilling in Patients with Chronic Q Fever. Journal of Infectious Diseases, 2004, 190, 1767-1772.	4.0	64
563	Genome Analysis of Minibacterium massiliensis Highlights the Convergent Evolution of Water-Living Bacteria. PLoS Genetics, 2007, 3, e138.	3.5	64
564	Non-contiguous finished genome sequence and description of Brevibacterium senegalense sp. nov Standards in Genomic Sciences, 2012, 7, 233-245.	1.5	64
565	Epidemic Genotype of <i>Coxiella burnetii </i> li>among Goats, Sheep, and Humans in the Netherlands. Emerging Infectious Diseases, 2012, 18, 887-889.	4.3	64
566	Survey of Anaplasmataceae bacteria in sheep from Senegal. Tropical Animal Health and Production, 2013, 45, 1557-1561.	1.4	64
567	Where Are We With Human Lice? A Review of the Current State of Knowledge. Frontiers in Cellular and Infection Microbiology, 2019, 9, 474.	3.9	64
568	Ehrlichial DNA Amplified from Ixodes ricinus (Acari: Ixodidae) in France. Journal of Medical Entomology, 1998, 35, 180-183.	1.8	63
569	Tick-borne rickettsioses in international travellers. International Journal of Infectious Diseases, 2004, 8, 139-146.	3.3	63
570	First Molecular Detection of R. conorii, R. aeschlimannii, and R. massiliae in Ticks from Algeria. Annals of the New York Academy of Sciences, 2006, 1078, 368-372.	3.8	63
571	Bacteriophages as vehicles of the resistome in cystic fibrosis. Journal of Antimicrobial Chemotherapy, 2011, 66, 2444-2447.	3.0	63
572	Strikes, flooding, rats, and leptospirosis in Marseille, France. International Journal of Infectious Diseases, 2011, 15, e710-e715.	3.3	63
573	Dramatic reduction in Clostridium difficile ribotype 027-associated mortality with early fecal transplantation by the nasogastric route: a preliminary report. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 1597-1601.	2.9	63
574	Metagenomic Analysis of Antibiotic-Induced Changes in Gut Microbiota in a Pregnant Rat Model. Frontiers in Pharmacology, 2016, 7, 104.	3.5	63
575	The puzzling mutational landscape of the SARSâ€2â€variant Omicron. Journal of Medical Virology, 2022, 94, 2019-2025.	5.0	63
576	Isolation of spotted fever group rickettsias from triturated ticks using a modification of the centrifugation-shell vial technique. Transactions of the Royal Society of Tropical Medicine and Hygiene, 1991, 85, 397-398.	1.8	62

#	Article	IF	CITATIONS
577	Correlation between Ratio of Serum Doxycycline Concentration to MIC and Rapid Decline of Antibody Levels during Treatment of Q Fever Endocarditis. Antimicrobial Agents and Chemotherapy, 2005, 49, 2673-2676.	3.2	62
578	Yersinia massiliensis sp. nov., isolated from fresh water. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 779-784.	1.7	62
579	Genomic, proteomic, and transcriptomic analysis of virulent and avirulent <i>Rickettsia prowazekii</i> reveals its adaptive mutation capabilities. Genome Research, 2010, 20, 655-663.	5.5	62
580	Mycobacterium tuberculosis Complex Mycobacteria as Amoeba-Resistant Organisms. PLoS ONE, 2011, 6, e20499.	2.5	62
581	No evidence of Bartonella quintana but detection of Acinetobacter baumannii in head lice from elementary schoolchildren in Paris. Comparative Immunology, Microbiology and Infectious Diseases, 2011, 34, 475-477.	1.6	62
582	Detection of Ehrlichial Infection by PCR in Dogs from Yamaguchi and Okinawa Prefectures, Japan Journal of Veterinary Medical Science, 2001, 63, 815-817.	0.9	61
583	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
584	<i>Bartonella quintana</i> and <i>Rickettsia felis</i> in Gabon. Emerging Infectious Diseases, 2005, 11, 1742-1744.	4.3	61
585	Migrated Foreign Body Liver Abscess. Medicine (United States), 2010, 89, 85-95.	1.0	61
586	Genome sequence and description of Alistipes senegalensis sp. nov Standards in Genomic Sciences, 2012, 6, 304-314.	1.5	61
587	Profile: The Niakhar Health and Demographic Surveillance System. International Journal of Epidemiology, 2013, 42, 1002-1011.	1.9	61
588	Fighting viruses with antibiotics: an overlooked path. International Journal of Antimicrobial Agents, 2016, 48, 349-352.	2.5	61
589	Giant Viruses of Amoebae: A Journey Through Innovative Research and Paradigm Changes. Annual Review of Virology, 2017, 4, 61-85.	6.7	61
590	Genome and pan-genome analysis to classify emerging bacteria. Biology Direct, 2019, 14, 5.	4.6	61
591	Clinical efficacy of chloroquine derivatives in COVID-19 infection: comparative meta-analysis between the big data and the real world. New Microbes and New Infections, 2020, 38, 100709.	1.6	61
592	SARS-CoV-2: fear versus data. International Journal of Antimicrobial Agents, 2020, 55, 105947.	2.5	61
593	Wolbachia pipientis Growth Kinetics and Susceptibilities to 13 Antibiotics Determined by Immunofluorescence Staining and Real-Time PCR. Antimicrobial Agents and Chemotherapy, 2003, 47, 1665-1671.	3.2	60
594	Western Immunoblotting for Bartonella Endocarditis. Vaccine Journal, 2003, 10, 95-102.	3.1	60

#	Article	IF	CITATIONS
595	Identification of two putative rickettsial adhesins by proteomic analysis. Research in Microbiology, 2006, 157, 605-612.	2.1	60
596	<i>Rickettsia felis</i> Infection, Tunisia. Emerging Infectious Diseases, 2006, 12, 138-140.	4.3	60
597	Body Lice, <i>Yersinia pestis</i> Orientalis, and Black Death. Emerging Infectious Diseases, 2010, 16, 892-893.	4.3	60
598	Real-time PCR assay allows detection of the New Delhi metallo- $\hat{l}^2$ -lactamase (NDM-1)-encoding gene in France. International Journal of Antimicrobial Agents, 2011, 37, 544-546.	2.5	60
599	Crystal Structure of the VgrG1 Actin Cross-linking Domain of the Vibrio cholerae Type VI Secretion System. Journal of Biological Chemistry, 2012, 287, 38190-38199.	3.4	60
600	Relapsing Fever Borreliae in Africa. American Journal of Tropical Medicine and Hygiene, 2013, 89, 288-292.	1.4	60
601	Microbiota of Demodex mites from rosacea patients and controls. Microbial Pathogenesis, 2014, 71-72, 37-40.	2.9	60
602	<i>Coxiella burnetii</i> Induces Reorganization of the Actin Cytoskeleton in Human Monocytes. Infection and Immunity, 1998, 66, 5527-5533.	2.2	60
603	Identification of Algerian Field-Caught Phlebotomine Sand Fly Vectors by MALDI-TOF MS. PLoS Neglected Tropical Diseases, 2016, 10, e0004351.	3.0	60
604	HIGH PREVALENCE OF BARTONELLA QUINTANA ENDOCARDITIS IN SFAX, TUNISIA. American Journal of Tropical Medicine and Hygiene, 2005, 72, 503-507.	1.4	60
605	Characterization of Mutations in the <i>rpoB</i> Gene in Naturally Rifampin-Resistant <i>Rickettsia</i> Species. Antimicrobial Agents and Chemotherapy, 1999, 43, 2400-2403.	3.2	59
606	<i>Parachlamydiaceae</i> as Rare Agents of Pneumonia. Emerging Infectious Diseases, 2003, 9, 755-756.	4.3	59
607	Bartonella quintanain head louse nits. FEMS Immunology and Medical Microbiology, 2011, 62, 244-246.	2.7	59
608	Tropheryma whipplei: A Common Bacterium in Rural Senegal. PLoS Neglected Tropical Diseases, 2011, 5, e1403.	3.0	59
609	An evaluation of the effects of Lactobacillus ingluviei on body weight, the intestinal microbiome and metabolism in mice. Microbial Pathogenesis, 2012, 52, 61-68.	2.9	59
610	Rickettsia Species in African Anopheles Mosquitoes. PLoS ONE, 2012, 7, e48254.	2.5	59
611	Q fever and pregnancy: disease, prevention, and strain specificity. European Journal of Clinical Microbiology and Infectious Diseases, 2013, 32, 361-368.	2.9	59
612	Broad Spectrum of Mimiviridae Virophage Allows Its Isolation Using a Mimivirus Reporter. PLoS ONE, 2013, 8, e61912.	2.5	59

#	Article	lF	CITATIONS
613	Determination of the animal origin of meat and gelatin by MALDI-TOF-MS. Journal of Food Composition and Analysis, 2015, 41, 104-112.	3.9	59
614	Molecular evidence of tick-borne hemoprotozoan-parasites (Theileria ovis and Babesia ovis) and bacteria in ticks and blood from small ruminants in Northern Algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2017, 50, 34-39.	1.6	59
615	Evidence of Archaeal Methanogens in Brain Abscess. Clinical Infectious Diseases, 2017, 65, 1-5.	5.8	59
616	Q FEVER IN CHILDREN IN GREECE. American Journal of Tropical Medicine and Hygiene, 2004, 70, 540-544.	1.4	59
617	MOLECULAR DETECTION OF BARTONELLA QUINTANA, B. ELIZABETHAE, B. KOEHLERAE, B. DOSHIAE, B. TAYLORII, AND RICKETTSIA FELIS IN RODENT FLEAS COLLECTED IN KABUL, AFGHANISTAN. American Journal of Tropical Medicine and Hygiene, 2006, 74, 436-439.	1.4	59
618	<i>Bartonella quintana</i> i>in Domestic Cat. Emerging Infectious Diseases, 2005, 11, 1287-1289.	4.3	58
619	Excretion of LivingBorrelia recurrentisin Feces of Infected Human Body Lice. Journal of Infectious Diseases, 2005, 191, 1898-1906.	4.0	58
620	Molecular analysis of pericardial fluid: a 7-year experience. European Heart Journal, 2006, 27, 1942-1946.	2.2	58
621	African Tick Bite Fever in Elderly Patients: 8 Cases in French Tourists Returning from South Africa. Clinical Infectious Diseases, 2008, 47, e28-e35.	5.8	58
622	Partial recN gene sequencing: a new tool for identification and phylogeny within the genus Streptococcus. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 2140-2148.	1.7	58
623	Viruses in a 14th-Century Coprolite. Applied and Environmental Microbiology, 2014, 80, 2648-2655.	3.1	58
624	Aerobic culture of methanogenic archaea without an external source of hydrogen. European Journal of Clinical Microbiology and Infectious Diseases, 2016, 35, 985-991.	2.9	58
625	Linking gut redox to human microbiome. Human Microbiome Journal, 2018, 10, 27-32.	3.8	58
626	Identification of Spotted Fever Group Rickettsiae Isolated from Dermacentor Marginatus and Ixodes Ricinus Ticks Collected in Switzerland. American Journal of Tropical Medicine and Hygiene, 1994, 51, 138-148.	1.4	58
627	Culture and identification of a "Deltamicron―SARSâ€CoVâ€2 in a three cases cluster in southern France. Journal of Medical Virology, 2022, 94, 3739-3749.	5.0	58
628	Serological and molecular evidence of Rickettsia helvetica in Denmark. Scandinavian Journal of Infectious Diseases, 2004, 36, 559-563.	1.5	57
629	Culture of < i>T. whipplei < /i>from the Stool of a Patient with Whipple's Disease. New England Journal of Medicine, 2006, 355, 1503-1505.	27.0	57
630	Geographic Distributions and Origins of Human Head Lice (Pediculus humanus capitis) Based on Mitochondrial Data. Journal of Parasitology, 2008, 94, 1275-1281.	0.7	57

#	Article	IF	CITATIONS
631	Resistance to trimethoprim/sulfamethoxazole and Tropheryma whipplei. International Journal of Antimicrobial Agents, 2009, 34, 255-259.	2.5	57
632	High Throughput, Multiplexed Pathogen Detection Authenticates Plague Waves in Medieval Venice, Italy. PLoS ONE, 2011, 6, e16735.	2.5	57
633	Viral Metagenomics on Animals as a Tool for the Detection of Zoonoses Prior to Human Infection?. International Journal of Molecular Sciences, 2014, 15, 10377-10397.	4.1	57
634	Dramatic reduction of culture time of Mycobacterium tuberculosis. Scientific Reports, 2014, 4, 4236.	3.3	57
635	A quasi-universal medium to break the aerobic/anaerobic bacterial culture dichotomy in clinical microbiology. Clinical Microbiology and Infection, 2016, 22, 53-58.	6.0	57
636	A gene transfer event suggests a long-term partnership between eustigmatophyte algae and a novel lineage of endosymbiotic bacteria. ISME Journal, 2018, 12, 2163-2175.	9.8	57
637	Salt in stools is associated with obesity, gut halophilic microbiota and Akkermansia muciniphila depletion in humans. International Journal of Obesity, 2019, 43, 862-871.	3.4	57
638	Sexual Dimorphism and Gender in Infectious Diseases. Frontiers in Immunology, 2021, 12, 698121.	4.8	57
639	Astrakhan Fever Rickettsiae: Antigenic and Genotypic Analysis of Isolates Obtained from Human and Rhipicephalus pumilio Ticks. American Journal of Tropical Medicine and Hygiene, 1994, 51, 697-706.	1.4	57
640	In Vitro Activities of Telithromycin (HMR 3647) against Rickettsia rickettsii, Rickettsia conorii, Rickettsia africae, Rickettsia typhi, Rickettsia prowazekii, Coxiella burnetii, Bartonella henselae, Bartonella quintana, Bartonella bacilliformis, and Ehrlichia chaffeensis. Antimicrobial Agents and Chemotherapy, 2000, 44, 1391-1393.	3.2	56
641	The geographical segregation of human lice preceded that of Pediculus humanus capitis and Pediculus humanus humanus. Comptes Rendus - Biologies, 2003, 326, 565-574.	0.2	56
642	Mycoplasma Endocarditis: Two Case Reports and a Review. Clinical Infectious Diseases, 2004, 38, e21-e24.	5.8	56
643	Pediatric Scrub typhus in Indian Himalayas. Indian Journal of Pediatrics, 2008, 75, 947-949.	0.8	56
644	Intrafamilial Circulation of <i>Tropheryma whipplei </i> , France. Emerging Infectious Diseases, 2012, 18, 949-55.	4.3	56
645	Acquisition of a High Diversity of Bacteria during the Hajj Pilgrimage, Including Acinetobacter baumannii with <i>bla</i> <sub>OXA-72</sub> and Escherichia coli with <i>bla</i> <sub>NDM-5</sub> Carbapenemase Genes. Antimicrobial Agents and Chemotherapy, 2016, 60, 5942-5948.	3.2	56
646	The Giant Cafeteria roenbergensis Virus That Infects a Widespread Marine Phagocytic Protist Is a New Member of the Fourth Domain of Life. PLoS ONE, 2011, 6, e18935.	2.5	56
647	Characteristics of the first 1119 SARSâ€CoVâ€2 Omicron variant cases, in Marseille, France, Novemberâ^'December 2021. Journal of Medical Virology, 2022, 94, 2290-2295.	5.0	56
648	Rickettsia conoriiInfection Enhances Vascular Cell Adhesion Moleculeâ€1 ―and Intercellular Adhesion Moleculeâ€1â€Dependent Mononuclear Cell Adherence to Endothelial Cells. Journal of Infectious Diseases, 1997, 175, 1142-1152.	4.0	55

#	Article	IF	CITATIONS
649	Experimental Model of Human Body Louse Infection Using Green Fluorescent Protein-Expressing Bartonella quintana. Infection and Immunity, 2001, 69, 1876-1879.	2.2	55
650	<i>Rickettsia felis, Bartonella henselae,</i> and <i>B. clarridgeiae,</i> New Zealand. Emerging Infectious Diseases, 2004, 10, 967-968.	4.3	55
651	<i>Rickettsia africae</i> , Western Africa. Emerging Infectious Diseases, 2010, 16, 571-573.	4.3	55
652	Non-contiguous finished genome sequence and description of Kurthia massiliensis sp. nov Standards in Genomic Sciences, 2012, 7, 221-232.	1.5	55
653	Plague Epidemics and Lice, Democratic Republic of the Congo. Emerging Infectious Diseases, 2013, 19, 505-6.	4.3	55
654	First Isolation of a Marseillevirus in the Diptera SyrphidaeEristalis tenax. Intervirology, 2013, 56, 386-394.	2.8	55
655	Are stool samples suitable for studying the link between gut microbiota and obesity?. European Journal of Epidemiology, 2014, 29, 307-309.	5.7	55
656	Faustoviruses: Comparative Genomics of New Megavirales Family Members. Frontiers in Microbiology, 2016, 7, 3.	3.5	55
657	Emerging Tick-Borne Bacterial Pathogens. Microbiology Spectrum, 2016, 4, .	3.0	55
658	Gut microbiome and dietary patterns in different Saudi populations and monkeys. Scientific Reports, 2016, 6, 32191.	3.3	55
659	Unifying view of stem–loop hairpin RNA as origin of current and ancient parasitic and non-parasitic RNAs, including in giant viruses. Current Opinion in Microbiology, 2016, 31, 1-8.	5.1	55
660	Changing Clinical Presentation of Q Fever Endocarditis. Clinical Infectious Diseases, 2002, 34, e28-e31.	5.8	54
661	Morphology of Legionella pneumophila according to their location within Hartmanella vermiformis. Research in Microbiology, 2003, 154, 619-621.	2.1	54
662	Culture of Tropheryma whipplei from Human Samples: a 3-Year Experience (1999 to 2002). Journal of Clinical Microbiology, 2003, 41, 3816-3822.	3.9	54
663	Rapid and cost-effective identification of Bartonella species using mass spectrometry. Journal of Medical Microbiology, 2009, 58, 1154-1159.	1.8	54
664	Clinical significance of a positive serology for mimivirus in patients presenting a suspicion of ventilator-associated pneumonia. Critical Care Medicine, 2009, 37, 111-118.	0.9	54
665	Imaging investigations in infective endocarditis: Current approach and perspectives. Archives of Cardiovascular Diseases, 2013, 106, 52-62.	1.6	54
666	Non contiguous-finished genome sequence and description of Alistipes obesi sp. nov. Standards in Genomic Sciences, 2013, 7, 427-439.	1.5	54

#	Article	IF	CITATIONS
667	Microbial culturomics unravels the halophilic microbiota repertoire of table salt: description of <i>Gracilibacillus massiliensis</i> sp. nov Microbial Ecology in Health and Disease, 2016, 27, 32049.	3.5	54
668	Long-term persistence of olfactory and gustatory disorders in COVID-19 patients. Clinical Microbiology and Infection, 2021, 27, 931-932.	6.0	54
669	MOLECULAR DETECTION OF RICKETTSIA FELIS, RICKETTSIA TYPHI AND TWO GENOTYPES CLOSELY RELATED TO BARTONELLA ELIZABETHAE. American Journal of Tropical Medicine and Hygiene, 2006, 75, 727-731.	1.4	54
670	Evaluation of Serological Response to Bartonella henselae, Bartonella quintana and Afipia felis Antigens in 64 Patients with Suspected Cat-scratch Disease. Scandinavian Journal of Infectious Diseases, 1996, 28, 361-366.	1.5	53
671	Amoeba-Resisting Bacteria and Ventilator-Associated Pneumonia. Emerging Infectious Diseases, 2003, 9, 815-821.	4.3	53
672	Prospects for the Future Using Genomics and Proteomics in Clinical Microbiology. Annual Review of Microbiology, 2011, 65, 169-188.	7.3	53
673	Gene gain and loss events in Rickettsia and Orientiaspecies. Biology Direct, 2011, 6, 6.	4.6	53
674	Widespread use of real-time PCR for rickettsial diagnosis: Figure 1. FEMS Immunology and Medical Microbiology, 2012, 64, 126-129.	2.7	53
675	Gorilla gorilla gorilla gut: a potential reservoir of pathogenic bacteria as revealed using culturomics and molecular tools. Scientific Reports, 2014, 4, 7174.	3.3	53
676	Comparison of a Modern and FossilPithovirusReveals Its Genetic Conservation and Evolution. Genome Biology and Evolution, 2016, 8, 2333-2339.	2.5	53
677	Whole-exome sequencing to analyze population structure, parental inbreeding, and familial linkage. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 6713-6718.	7.1	53
678	Emended description of Rickettsia felis (Bouyer et al. 2001), a temperature-dependent cultured bacterium International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 2035-2041.	1.7	53
679	$\hat{l}$ ± v $\hat{l}^2$ 3 Integrin and Bacterial Lipopolysaccharide Are Involved in Coxiella burnetii -Stimulated Production of Tumor Necrosis Factor by Human Monocytes. Infection and Immunity, 2000, 68, 5673-5678.	2.2	52
680	Response to Comment on "The 1.2-Megabase Genome Sequence of Mimivirus". Science, 2005, 308, 1114-1114.	12.6	52
681	Autoimmunohistochemistry: A New Method for the Histologic Diagnosis of Infective Endocarditis. Journal of Infectious Diseases, 2006, 193, 1711-1717.	4.0	52
682	Paenibacillus provencensis sp. nov., isolated from human cerebrospinal fluid, and Paenibacillus urinalis sp. nov., isolated from human urine. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 682-687.	1.7	52
683	A Paradoxical <i>Tropheryma whipplei </i> Western Blot Differentiates Patients with Whipple Disease from Asymptomatic Carriers. Clinical Infectious Diseases, 2009, 49, 717-723.	5.8	52
684	Likely Correlation between Sources of Information and Acceptability of A/H1N1 Swine-Origin Influenza Virus Vaccine in Marseille, France. PLoS ONE, 2010, 5, e11292.	2.5	52

#	Article	IF	CITATIONS
685	Sputnik, a Virophage Infecting the Viral Domain of Life. Advances in Virus Research, 2012, 82, 63-89.	2.1	52
686	Non contiguous-finished genome sequence and description of Dielma fastidiosa gen. nov., sp. nov., a new member of the Family Erysipelotrichaceae. Standards in Genomic Sciences, 2013, 8, 336-351.	1.5	52
687	Acquisition of extended-spectrum cephalosporin- and colistin-resistant Salmonella enterica subsp. enterica serotype Newport by pilgrims during Hajj. International Journal of Antimicrobial Agents, 2015, 45, 600-604.	2.5	52
688	Faecal microbiota transplantation for stool decolonization of OXA-48 carbapenemase-producing Klebsiella pneumoniae. Journal of Hospital Infection, 2015, 90, 173-174.	2.9	52
689	New Rickettsia species in soft ticks Ornithodoros hasei collected from bats in French Guiana. Ticks and Tick-borne Diseases, 2016, 7, 1089-1096.	2.7	52
690	Natural Anaplasmataceae infection in Rhipicephalus bursa ticks collected from sheep in the French Basque Country. Ticks and Tick-borne Diseases, 2017, 8, 18-24.	2.7	52
691	Ultrarapid diagnosis, microscope imaging, genome sequencing, and culture isolation of SARS-CoV-2. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1601-1603.	2.9	52
692	Bartonella quintanain Body Lice Collected from Homeless Persons in Russia. Emerging Infectious Diseases, 1999, 5, 176-178.	4.3	52
693	Bartonella quintanaEndocarditis in Dogs. Emerging Infectious Diseases, 2006, 12, 1869-1872.	4.3	52
694	Seroepidemiological survey of rickettsial infections among blood donors in central Tunisia. Transactions of the Royal Society of Tropical Medicine and Hygiene, 1995, 89, 266-268.	1.8	51
695	Use of Amplification and Sequencing of the 16S rRNA Gene to Diagnose <i>Mycoplasma pneumoniae</i> Osteomyelitis in a Patient with Hypogammaglobulinemia. Clinical Infectious Diseases, 1997, 24, 1161-1163.	5.8	51
696	A Focus of Tickâ€Borne Relapsing Fever in Southern Zaire. Clinical Infectious Diseases, 1997, 25, 139-144.	5.8	51
697	Intraspecies diversity of Coxiella burnetii as revealed by com1 and mucZ sequence comparison. FEMS Microbiology Letters, 1999, 180, 61-67.	1.8	51
698	Molecular phylogeny of the genusBartonella: what is the current knowledge?. FEMS Microbiology Letters, 2001, 200, 1-7.	1.8	51
699	IFN-Î <sup>3</sup> -Induced Apoptosis and Microbicidal Activity in Monocytes Harboring the Intracellular Bacterium <i>Coxiella burnetii</i> Require Membrane TNF and Homotypic Cell Adherence. Journal of Immunology, 2002, 169, 6309-6315.	0.8	51
700	Crystal structure of the DNA-bound VapBC2 antitoxin/toxin pair from Rickettsia felis. Nucleic Acids Research, 2012, 40, 3245-3258.	14.5	51
701	Detection of Acinetobacter baumannii in human head and body lice from Ethiopia and identification of new genotypes. International Journal of Infectious Diseases, 2012, 16, e680-e683.	3.3	51
702	<i>Bartonella quintana</i> in Head Lice from Sénégal. Vector-Borne and Zoonotic Diseases, 2012, 12, 564-567.	1.5	51

#	Article	IF	CITATIONS
703	Rickettsiae in arthropods collected from the North African Hedgehog (Atelerix algirus) and the desert hedgehog (Paraechinus aethiopicus) in Algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 117-122.	1.6	51
704	Tropheryma whipplei prevalence strongly suggests human transmission in homeless shelters. International Journal of Infectious Diseases, 2013, 17, e67-e68.	3.3	51
705	Describing the Silent Human Virome with an Emphasis on Giant Viruses. Intervirology, 2013, 56, 395-412.	2.8	51
706	MALDI-TOF Mass Spectrometry: A Powerful Tool for Clinical Microbiology at Hôpital Principal de Dakar, Senegal (West Africa). PLoS ONE, 2015, 10, e0145889.	2.5	51
707	Identification of tick species and disseminate pathogen using hemolymph by MALDI-TOF MS. Ticks and Tick-borne Diseases, 2015, 6, 579-586.	2.7	51
708	The Strengths of Scanning Electron Microscopy in Deciphering SARS-CoV-2 Infectious Cycle. Frontiers in Microbiology, 2020, 11, 2014.	3.5	51
709	COVIDâ€19 reâ€infection. European Journal of Clinical Investigation, 2021, 51, e13537.	3.4	51
710	Effect of Rickettsial Toxin VapC on Its Eukaryotic Host. PLoS ONE, 2011, 6, e26528.	2.5	51
711	<i>Rickettsia felis</i> in <i>Aedes albopictus</i> Mosquitoes, Libreville, Gabon. Emerging Infectious Diseases, 2012, 18, 1687-1689.	4.3	51
712	Prevalence of Rickettsia-Like Organisms and Spotted Fever Group Rickettsiae in Ticks (Acari: Ixodidae) from Zimbabwe. Journal of Medical Entomology, 1995, 32, 787-792.	1.8	50
713	<i>Coxiella burnetii</i> Survives in Monocytes from Patients with Q Fever Endocarditis: Involvement of Tumor Necrosis Factor. Infection and Immunity, 2000, 68, 160-164.	2.2	50
714	Whipple Disease Associated with Giardiasis. Journal of Infectious Diseases, 2003, 188, 828-834.	4.0	50
715	Corynebacterium Species Isolated from Bone and Joint Infections Identified by 16S rRNA Gene Sequence Analysis. Journal of Clinical Microbiology, 2004, 42, 2231-2233.	3.9	50
716	Acquired Resistance to Trimethoprim‧ulfamethoxazole during Whipple Disease and Expression of the Causative Target Gene. Journal of Infectious Diseases, 2008, 198, 101-108.	4.0	50
717	Frequent transmission of hepatitis E virus among piglets in farms in Southern France. Journal of Medical Virology, 2009, 81, 1750-1759.	5.0	50
718	Phylogenomic Analysis of Odyssella thessalonicensis Fortifies the Common Origin of Rickettsiales, Pelagibacter ubique and Reclimonas americana Mitochondrion. PLoS ONE, 2011, 6, e24857.	2.5	50
719	Analysis of risk factors for malignant Mediterranean spotted fever indicates that fluoroquinolone treatment has a deleterious effect. Journal of Antimicrobial Chemotherapy, 2011, 66, 1821-1830.	3.0	50
720	Carbapenem Resistance and Acinetobacter baumannii in Senegal: The Paradigm of a Common Phenomenon in Natural Reservoirs. PLoS ONE, 2012, 7, e39495.	2.5	50

#	Article	IF	CITATIONS
721	Aerobic culture of anaerobic bacteria using antioxidants: a preliminary report. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 1781-1783.	2.9	50
722	Glycans affect DNA extraction and induce substantial differences in gut metagenomic studies. Scientific Reports, 2016, 6, 26276.	3.3	50
723	Mimivirus inaugurated in the 21st century the beginning of a reclassification of viruses. Current Opinion in Microbiology, 2016, 31, 16-24.	5.1	50
724	MALDI-TOF MS as an innovative tool for detection of Plasmodium parasites in Anopheles mosquitoes. Malaria Journal, 2017, 16, 5.	2.3	50
725	MALDI-TOF Identification of the Human Gut Microbiome in People with and without Diarrhea in Senegal. PLoS ONE, 2014, 9, e87419.	2.5	50
726	Diagnostic methods. Cardiology Clinics, 2003, 21, 207-217.	2.2	49
727	<i>Coxiella burnetii</i> Avoids Macrophage Phagocytosis by Interfering with Spatial Distribution of Complement Receptor 3. Journal of Immunology, 2003, 170, 4217-4225.	0.8	49
728	Intracellular Rickettsiales: Insights into manipulators of eukaryotic cells. Trends in Molecular Medicine, 2011, 17, 573-583.	6.7	49
729	Marseillevirus Adenitis in an 11-Month-Old Child. Journal of Clinical Microbiology, 2013, 51, 4102-4105.	3.9	49
730	Detection of Rickettsia spp in Ticks by MALDI-TOF MS. PLoS Neglected Tropical Diseases, 2015, 9, e0003473.	3.0	49
731	Activation of Protein Tyrosine Kinases by <i>Coxiella burnetii </i> Role in Actin Cytoskeleton Reorganization and Bacterial Phagocytosis. Infection and Immunity, 2001, 69, 2520-2526.	2.2	48
732	Molecular techniques in Whipple's disease. Expert Review of Molecular Diagnostics, 2001, 1, 299-309.	3.1	48
733	Genotypic Characteristics of Two Serotypes of Bartonella henselae. Journal of Clinical Microbiology, 2002, 40, 2002-2008.	3.9	48
734	Proteome analysis of Rickettsia conorii by two-dimensional gel electrophoresis coupled with mass spectrometry. FEMS Microbiology Letters, 2005, 245, 231-238.	1.8	48
735	Threats to international travellers posed by tick-borne diseases. Travel Medicine and Infectious Disease, 2006, 4, 4-13.	3.0	48
736	<i>Tropheryma whipplei</i> in Fecal Samples from Children, Senegal. Emerging Infectious Diseases, 2009, 15, 922-924.	4.3	48
737	Axenic culture of fastidious and intracellular bacteria. Trends in Microbiology, 2013, 21, 92-99.	7.7	48
738	Molecular Studies Neglect Apparently Gram-Negative Populations in the Human Gut Microbiota. Journal of Clinical Microbiology, 2013, 51, 3286-3293.	3.9	48

#	Article	IF	Citations
739	Non-contiguous finished genome sequence and description of Alistipes ihumii sp. nov Standards in Genomic Sciences, 2014, 9, 1221-1235.	1.5	48
740	Management and Treatment of Human Lice. BioMed Research International, 2016, 2016, 1-12.	1.9	48
741	Plasmid-Mediated <i>mcr-1</i> Gene in Colistin-Resistant Clinical Isolates of Klebsiella pneumoniae in France and Laos. Antimicrobial Agents and Chemotherapy, 2016, 60, 6994-6995.	3.2	48
742	Molecular evidence of vector-borne pathogens in dogs and cats and their ectoparasites in Algiers, Algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2016, 45, 23-28.	1.6	48
<b>7</b> 43	Molecular investigation and phylogeny of Anaplasmataceae species infecting domestic animals and ticks in Corsica, France. Parasites and Vectors, 2017, 10, 302.	2.5	48
744	Detection of bacterial pathogens including potential new species in human head lice from Mali. PLoS ONE, 2017, 12, e0184621.	2.5	48
745	A Large Open Pangenome and a Small Core Genome for Giant Pandoraviruses. Frontiers in Microbiology, 2018, 9, 1486.	3.5	48
746	Massive analysis of 64,628 bacterial genomes to decipher water reservoir and origin of mobile colistin resistance genes: is there another role for these enzymes?. Scientific Reports, 2020, 10, 5970.	3.3	48
747	Diagnosis of Rickettsial Diseases Using Samples Dried on Blotting Paper. Vaccine Journal, 1999, 6, 483-488.	2.6	48
748	Monoclonal Antibodies to <i>Coxiella burnetii</i> for Antigenic Detection in Cell Cultures and in Paraffin-embedded Tissues. American Journal of Clinical Pathology, 1994, 101, 318-320.	0.7	47
749	Return of trench fever. Lancet, The, 1995, 345, 450-451.	13.7	47
750	Afipia felis in hospital water supply in association with free-living amoebae. Lancet, The, 1999, 353, 1330.	13.7	47
751	Kinetics of Antibody Responses in Rickettsia africae and Rickettsia conorii Infections. Vaccine Journal, 2002, 9, 324-328.	3.1	47
752	Increased Levels of Soluble CD40L in African Tick Bite Fever: Possible Involvement of TLRs in the Pathogenic Interaction between <i>Rickettsia africae</i> , Endothelial Cells, and Platelets. Journal of Immunology, 2006, 177, 2699-2706.	0.8	47
753	Cost-Effectiveness of Blood Agar for Isolation of Mycobacteria. PLoS Neglected Tropical Diseases, 2007, 1, e83.	3.0	47
754	The Uptake of Apoptotic Cells Drives Coxiella burnetii Replication and Macrophage Polarization: A Model for Q Fever Endocarditis. PLoS Pathogens, 2008, 4, e1000066.	4.7	47
755	Contribution of a shelter-based survey for screening respiratory diseases in the homeless. European Journal of Public Health, 2009, 19, 157-160.	0.3	47
756	Mediterranean spotted fever in Algeria â€" new trends. International Journal of Infectious Diseases, 2009, 13, 227-235.	3.3	47

#	Article	IF	Citations
757	Insight into cross-talk between intra-amoebal pathogens. BMC Genomics, 2011, 12, 542.	2.8	47
758	The Rise of Tropheryma whipplei: a 12-Year Retrospective Study of PCR Diagnoses in Our Reference Center. Journal of Clinical Microbiology, 2012, 50, 3917-3920.	3.9	47
759	Rickettsiae of spotted fever group, Borrelia valaisiana, and Coxiella burnetii in ticks on passerine birds and mammals from the Camargue in the south of France. Ticks and Tick-borne Diseases, 2012, 3, 355-360.	2.7	47
760	<i>Borrelia recurrentis</i> in Head Lice, Ethiopia. Emerging Infectious Diseases, 2013, 19, 796-8.	4.3	47
761	Non-contiguous finished genome sequence and description of Brevibacillus massiliensis sp. nov Standards in Genomic Sciences, 2013, 8, 1-14.	1.5	47
762	A New Clade of African Body and Head Lice Infected by Bartonella quintana and Yersinia pestis—Democratic Republic of the Congo. American Journal of Tropical Medicine and Hygiene, 2015, 93, 990-993.	1.4	47
763	<i>Candidatus</i> Coxiella massiliensis Infection. Emerging Infectious Diseases, 2016, 22, 285-288.	4.3	47
764	Morphological, molecular and MALDI-TOF mass spectrometry identification of ixodid tick species collected in Oromia, Ethiopia. Parasitology Research, 2016, 115, 4199-4210.	1.6	47
765	Standardization of sample homogenization for mosquito identification using an innovative proteomic tool based on protein profiling. Proteomics, 2016, 16, 3148-3160.	2.2	47
766	High-quality draft genome sequence and description of Haemophilus massiliensis sp. nov Standards in Genomic Sciences, 2016, 11, 31.	1.5	47
767	Identification of blood meal sources in the main African malaria mosquito vector by MALDI-TOF MS. Malaria Journal, 2016, 15, 87.	2.3	47
768	Emerging methodologies for pathogen identification in positive blood culture testing. Expert Review of Molecular Diagnostics, 2016, 16, 97-111.	3.1	47
769	Infective endocarditis in octogenarians. Heart, 2017, 103, 1602-1609.	2.9	47
770	Fluorescence in situ hybridization, a complementary molecular tool for the clinical diagnosis of infectious diseases by intracellular and fastidious bacteria. FEMS Microbiology Reviews, 2019, 43, 88-107.	8.6	47
771	Evidence of SARS-CoV-2 re-infection with a different genotype. Journal of Infection, 2021, 82, 84-123.	3.3	47
772	Head Lice of Pygmies Reveal the Presence of Relapsing Fever Borreliae in the Republic of Congo. PLoS Neglected Tropical Diseases, 2016, 10, e0005142.	3.0	47
773	Non-contiguous finished genome sequence and description of Paenibacillus senegalensis sp. nov Standards in Genomic Sciences, 2012, 7, 70-81.	1.5	47
774	Mediterranean Spotted Fever in Marseille, France: Correlation between Prevalence of Hospitalized Patients, Seroepidemiology, and Prevalence of Infected Ticks in Three Different Areas. American Journal of Tropical Medicine and Hygiene, 1993, 48, 249-256.	1.4	47

#	Article	IF	Citations
775	First molecular evidence of new Bartonella spp. in fleas and a tick from Peru American Journal of Tropical Medicine and Hygiene, 2002, 67, 135-136.	1.4	47
776	Optimum Treatment of Intracellular Infection. Drugs, 1996, 52, 45-59.	10.9	46
777	Coxiella burnetii: the â€~query' fever bacterium: A model of immune subversion by a strictly intracellular microorganism. FEMS Microbiology Reviews, 1997, 19, 209-217.	8.6	46
778	<i>Tropheryma whipplei</i> Circulating in Blood Monocytes. New England Journal of Medicine, 2001, 345, 548-548.	27.0	46
779	Intracellular trafficking of Parachlamydia acanthamoebae. Cellular Microbiology, 2005, 7, 581-589.	2.1	46
780	Transcriptional response of Rickettsia conorii exposed to temperature variation and stress starvation. Research in Microbiology, 2005, 156, 211-218.	2.1	46
781	Sca1, a previously undescribed paralog from autotransporter protein-encoding genes in Rickettsia species. BMC Microbiology, 2006, 6, 12.	3.3	46
782	Rickettsia asiatica sp. nov., isolated in Japan. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 2365-2368.	1.7	46
783	Phocaeicola abscessus gen. nov., sp. nov., an anaerobic bacterium isolated from a human brain abscess sample. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 2232-2237.	1.7	46
784	Outbreak of <i>Corynebacterium pseudodiphtheriticum </i> Infection in Cystic Fibrosis Patients, France. Emerging Infectious Diseases, 2010, 16, 1231-1236.	4.3	46
785	Intrinsic fluoroquinolone resistance in Orientia tsutsugamushi. International Journal of Antimicrobial Agents, 2010, 35, 338-341.	2.5	46
786	Tick-borne rickettsiae in Guinea and Liberia. Ticks and Tick-borne Diseases, 2012, 3, 43-48.	2.7	46
787	The Gene Expression Analysis of Blood Reveals S100A11 and AQP9 as Potential Biomarkers of Infective Endocarditis. PLoS ONE, 2012, 7, e31490.	2.5	46
788	Detection of Bartonella quintana in African Body and Head Lice. American Journal of Tropical Medicine and Hygiene, 2014, 91, 294-301.	1.4	46
789	Scalp eschar and neck lymphadenopathy after tick bite: an emerging syndrome with multiple causes. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 1449-1456.	2.9	46
790	Acquisition of <i>mcr-1</i> Plasmid-Mediated Colistin Resistance in Escherichia coli and Klebsiella pneumoniae during Hajj 2013 and 2014. Antimicrobial Agents and Chemotherapy, 2016, 60, 6998-6999.	3.2	46
791	MALDI-TOF MS identification of ticks of domestic and wild animals in Algeria and molecular detection of associated microorganisms. Comparative Immunology, Microbiology and Infectious Diseases, 2018, 57, 39-49.	1.6	46
792	Metagenomic and culturomic analysis of gut microbiota dysbiosis during Clostridium difficile infection. Scientific Reports, 2019, 9, 12807.	3.3	46

#	Article	IF	CITATIONS
<b>7</b> 93	New human-associated species of the family Atopobiaceae and proposal to reclassify members of the genus Olsenella. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	46
794	MOLECULAR IDENTIFICATION OF A COLLECTION OF SPOTTED FEVER GROUP RICKETTSIAE OBTAINED FROM PATIENTS AND TICKS FROM RUSSIA. American Journal of Tropical Medicine and Hygiene, 2006, 74, 440-443.	1.4	46
795	Analysis of SARS-CoV-2 Variants From 24,181 Patients Exemplifies the Role of Globalization and Zoonosis in Pandemics. Frontiers in Microbiology, 2021, 12, 786233.	3.5	46
796	SARS-CoV-2 reinfection and COVID-19 severity. Emerging Microbes and Infections, 2022, 11, 894-901.	<b>6.</b> 5	46
797	Repeated Pregnancies in BALB/c Mice Infected withCoxiella burnetiiCause Disseminated Infection, Resulting in Stillbirth and Endocarditis. Journal of Infectious Diseases, 2000, 181, 188-194.	4.0	45
798	Improved Culture from Lymph Nodes of Patients with Cat Scratch Disease and Genotypic Characterization of Bartonella henselae Isolates in Australia. Journal of Clinical Microbiology, 2002, 40, 3620-3624.	3.9	45
799	Diagnostic methods. Infectious Disease Clinics of North America, 2002, 16, 377-392.	5.1	45
800	Natural History of Q Fever in Goats. Vector-Borne and Zoonotic Diseases, 2003, 3, 11-15.	1.5	45
801	Magnitude and Features of Scrub Typhus and Spotted Fever in Children in India. Journal of Tropical Pediatrics, 2006, 52, 228-229.	1.5	45
802	First isolation of two colistin-resistant emerging pathogens, Brevundimonas diminuta and Ochrobactrum anthropi, in a woman with cystic fibrosis: a case report. Journal of Medical Case Reports, 2008, 2, 373.	0.8	45
803	Rapid comparative genomic analysis for clinical microbiology: The <i>Francisella tularensis</i> paradigm. Genome Research, 2008, 18, 742-750.	5.5	45
804	Human Case of <i>Bartonella alsatica </i> Lymphadenitis. Emerging Infectious Diseases, 2008, 14, 1951-1953.	4.3	45
805	Amoeba co-culture of soil specimens recovered 33 different bacteria, including four new species and Streptococcus pneumoniae. Microbiology (United Kingdom), 2009, 155, 657-664.	1.8	45
806	Successful treatment of immune reconstitution inflammatory syndrome in Whipple's disease using thalidomide. Journal of Infection, 2010, 60, 79-82.	3.3	45
807	Diagnosis of Cardiac Device–Related Infective Endocarditis After Device Removal. JACC: Cardiovascular Imaging, 2010, 3, 673-681.	5.3	45
808	Non-contiguous finished genome sequence and description of Clostridium dakarense sp. nov Standards in Genomic Sciences, 2013, 9, 14-27.	1.5	45
809	Cardiac multidetector computed tomography in infective endocarditis: a pictorial essay. Insights Into Imaging, 2014, 5, 559-570.	3.4	45
810	Monitoring human tick-borne disease risk and tick bite exposure in Europe: Available tools and promising future methods. Ticks and Tick-borne Diseases, 2014, 5, 607-619.	2.7	45

#	Article	IF	Citations
811	Culture-negative Prosthetic Joint Arthritis RelatedÂtoÂCoxiella burnetii. American Journal of Medicine, 2014, 127, 786.e7-786.e10.	1.5	45
812	Co-culture of Methanobrevibacter smithii with enterobacteria during urinary infection. EBioMedicine, 2019, 43, 333-337.	6.1	45
813	New Insights Into the Physiopathology of COVID-19: SARS-CoV-2-Associated Gastrointestinal Illness. Frontiers in Medicine, 2021, 8, 640073.	2.6	45
814	Description of Afipia birgiae sp. nov. and Afipia massiliensis sp. nov. and recognition of Afipia felis genospecies A International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 1773-1782.	1.7	45
815	Immunodetection of Tropheryma whippleiin Intestinal Tissues from Dr. Whipple's 1907 Patient. New England Journal of Medicine, 2003, 348, 1411-1412.	27.0	44
816	Multispacer Typing of Rickettsia prowazekii Enabling Epidemiological Studies of Epidemic Typhus. Journal of Clinical Microbiology, 2005, 43, 4708-4712.	3.9	44
817	Multispacer Typing To Study the Genotypic Distribution of Bartonella henselae Populations. Journal of Clinical Microbiology, 2006, 44, 2499-2506.	3.9	44
818	Propionibacterium endocarditis: A case series from the International Collaboration on Endocarditis Merged Database and Prospective Cohort Study. Scandinavian Journal of Infectious Diseases, 2007, 39, 840-848.	1.5	44
819	Regulation of whole bacterial pathogen transcription within infected hosts. FEMS Microbiology Reviews, 2008, 32, 440-460.	8.6	44
820	Rocky Mountain spotted fever in the USA: a benign disease or a common diagnostic error?. Lancet Infectious Diseases, The, 2008, 8, 587-589.	9.1	44
821	Structural Studies of the Sputnik Virophage. Journal of Virology, 2010, 84, 894-897.	3.4	44
822	Adipose Tissue Serves as a Reservoir for Recrudescent Rickettsia prowazekii Infection in a Mouse Model. PLoS ONE, 2010, 5, e8547.	2.5	44
823	Evidence of a Louse-Borne Outbreak Involving Typhus in Douai, 1710-1712 during the War of Spanish Succession. PLoS ONE, 2010, 5, e15405.	2.5	44
824	Palaeogenomics of Mycobacterium tuberculosis: epidemic bursts with a degrading genome. Lancet Infectious Diseases, The, 2011, 11, 641-650.	9.1	44
825	Diagnosis of Rickettsioses from Eschar Swab Samples, Algeria. Emerging Infectious Diseases, 2011, 17, 1968-1969.	4.3	44
826	Complete genome sequence of Cannes 8 virus, a new member of the proposed family "Marseilleviridae― Virus Genes, 2013, 47, 550-555.	1.6	44
827	Marseillevirus prevalence in multitransfused patients suggests blood transmission. Journal of Clinical Virology, 2013, 58, 722-725.	3.1	44
828	Acquisition and excretion of <i><scp>B</scp>artonella quintana</i> by the cat flea, <i><scp>C</scp>tenocephalides felis felis</i> Molecular Ecology, 2014, 23, 1204-1212.	3.9	44

#	Article	IF	CITATIONS
829	Molecular detection of Anaplasma platys and Ehrlichia canis in dogs from Kabylie, Algeria. Ticks and Tick-borne Diseases, 2015, 6, 198-203.	2.7	44
830	Ancestrality and Mosaicism of Giant Viruses Supporting the Definition of the Fourth TRUC of Microbes. Frontiers in Microbiology, 2018, 9, 2668.	3.5	44
831	Among Live and Dead Bacteria, the Optimization of Sample Collection and Processing Remains Essential in Recovering Gut Microbiota Components. Frontiers in Microbiology, 2019, 10, 1606.	3.5	44
832	Extensive culturomics of 8 healthy samples enhances metagenomics efficiency. PLoS ONE, 2019, 14, e0223543.	2.5	44
833	Metaproteomics of the human gut microbiota: Challenges and contributions to other OMICS. Clinical Mass Spectrometry, 2019, 14, 18-30.	1.9	44
834	Human Bacterial Repertoire of the Urinary Tract: a Potential Paradigm Shift. Journal of Clinical Microbiology, 2019, 57, .	3.9	44
835	Emergence and outcomes of the SARS-CoV-2 â€~Marseille-4' variant. International Journal of Infectious Diseases, 2021, 106, 228-236.	3.3	44
836	A Nosocomial Outbreak of <i>Legionella pneumophila </i> Caused by Contaminated Transesophageal Echocardiography Probes. Infection Control and Hospital Epidemiology, 2003, 24, 619-622.	1.8	43
837	Experimental Infection Models of Ticks of theRhipicephalus sanguineusGroup withRickettsia conorii. Vector-Borne and Zoonotic Diseases, 2005, 5, 363-372.	1.5	43
838	A fourth dose of DTPa-IPV vaccine given to 4â€"6 year old children in Italy and Sweden following primary vaccination at 3, 5 and 11â€"12 months of age. Scandinavian Journal of Infectious Diseases, 2005, 37, 221-229.	1.5	43
839	Estimation of prokaryote genomic DNA G+C content by sequencing universally conserved genes. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1025-1029.	1.7	43
840	<i>Finegoldia magna: </i> A Forgotten Pathogen in Prosthetic Joint Infection Rediscovered by Molecular Biology. Clinical Infectious Diseases, 2009, 49, 1244-1247.	5 <b>.</b> 8	43
841	Evidence that the intra-amoebal Legionella drancourtii acquired a sterol reductase gene from eukaryotes. BMC Research Notes, 2009, 2, 51.	1.4	43
842	Evidence of Transfer by Conjugation of Type IV Secretion System Genes between Bartonella Species and Rhizobium radiobacter in Amoeba. PLoS ONE, 2010, 5, e12666.	2.5	43
843	Molecular diversity of the <i>Planctomycetes</i> in the human gut microbiota in France and Senegal. Apmis, 2013, 121, 1082-1090.	2.0	43
844	Skin microbiota: overview and role in the skin diseases acne vulgaris and rosacea. Future Microbiology, 2013, 8, 209-222.	2.0	43
845	Coxiella burnetii-positive PCR in febrile patients in rural and urban Africa. International Journal of Infectious Diseases, 2014, 28, 107-110.	3.3	43
846	Gut microeukaryotes during anorexia nervosa: a case report. BMC Research Notes, 2014, 7, 33.	1.4	43

#	Article	IF	CITATIONS
847	The Ongoing Revolution of MALDI-TOF Mass Spectrometry for Microbiology Reaches Tropical Africa. American Journal of Tropical Medicine and Hygiene, 2015, 92, 641-647.	1.4	43
848	IRF4 haploinsufficiency in a family with Whippleâ $\in$ <sup>™</sup> s disease. ELife, 2018, 7, .	6.0	43
849	Relationship between nasopharyngeal microbiota and patient's susceptibility to viral infection. Expert Review of Anti-Infective Therapy, 2019, 17, 437-447.	4.4	43
850	Culture of Methanogenic Archaea from Human Colostrum and Milk. Scientific Reports, 2019, 9, 18653.	3.3	43
851	Tick-Borne Infection Caused byRickettsia africaein the West Indies. New England Journal of Medicine, 1998, 338, 1391-1392.	27.0	42
852	Molecular detection of spotted fever group rickettsiae in ticks from Ethiopia and Chad. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2008, 102, 945-949.	1.8	42
853	Human microbiome: take-home lesson on growth promoters?. Nature, 2008, 454, 690-691.	27.8	42
854	Chronic Q Fever: Different Serological Results in 3 Countries-Results of a Follow-up Study 6 Years After a Point Source Outbreak. Clinical Infectious Diseases, 2011, 52, 1013-1019.	5.8	42
855	Mimivirus Collagen Is Modified by Bifunctional Lysyl Hydroxylase and Glycosyltransferase Enzyme. Journal of Biological Chemistry, 2011, 286, 43701-43709.	3.4	42
856	Matrixâ€assisted laser desorption/ionization timeâ€ofâ€flight mass spectrometry identification of <i>Archaea</i> : towards the universal identification of living organisms. Apmis, 2012, 120, 85-91.	2.0	42
857	Unique Clone of <i>Coxiella burnetii </i> Causing Severe Q Fever, French Guiana. Emerging Infectious Diseases, 2013, 19, 1102-1104.	4.3	42
858	Three-Toed Sloth as Putative Reservoir of (i) Coxiella burnetii (i), Cayenne, French Guiana. Emerging Infectious Diseases, 2014, 20, 1760-1761.	4.3	42
859	High-Throughput Isolation of Giant Viruses in Liquid Medium Using Automated Flow Cytometry and Fluorescence Staining. Frontiers in Microbiology, 2016, 7, 26.	3.5	42
860	Use of MALDI-TOF MS and culturomics to identify mosquitoes and their midgut microbiota. Parasites and Vectors, 2016, 9, 495.	2.5	42
861	Evaluation of two DNA extraction methods for the PCR-based detection of eukaryotic enteric pathogens in fecal samples. BMC Research Notes, 2018, 11, 206.	1.4	42
862	First Documentation of <i>Rickettsia conorii </i> Infection (Strain Indian Tick Typhus) in a Traveler. Emerging Infectious Diseases, 2001, 7, 909-910.	4.3	41
863	Editorial Commentary: A New Rickettsial Disease in the United States. Clinical Infectious Diseases, 2004, 38, 812-813.	5.8	41
864	Measurement of the antibiotic susceptibility of Coxiella burnetii using real time PCR. International Journal of Antimicrobial Agents, 2004, 23, 169-174.	2.5	41

#	Article	IF	CITATIONS
865	Bacteroides massiliensis sp. nov., isolated from blood culture of a newborn. International Journal of Systematic and Evolutionary Microbiology, 2005, 55, 1335-1337.	1.7	41
866	Far Eastern Tick-Borne Rickettsiosis: Identification of Two New Cases and Tick Vector. Annals of the New York Academy of Sciences, 2006, 1078, 80-88.	3.8	41
867	Characterization of Mimivirus DNA Topoisomerase IB Suggests Horizontal Gene Transfer between Eukaryal Viruses and Bacteria. Journal of Virology, 2006, 80, 314-321.	3.4	41
868	Cultivation of Tropheryma whipplei from the synovial fluid in Whipple's arthritis. Arthritis and Rheumatism, 2007, 56, 1713-1718.	6.7	41
869	Proteome analysis ofRickettsia felis highlights the expression profile of intracellular bacteria. Proteomics, 2007, 7, 1232-1248.	2.2	41
870	Gene Repertoire of Amoeba-Associated Giant Viruses. Intervirology, 2010, 53, 330-343.	2.8	41
871	Comparison of Real-Time Quantitative PCR and Culture for the Diagnosis of Emerging Rickettsioses. PLoS Neglected Tropical Diseases, 2012, 6, e1540.	3.0	41
872	Immunopathology of Immune Reconstitution Inflammatory Syndrome in Whipple's Disease. Journal of Immunology, 2013, 190, 2354-2361.	0.8	41
873	DNA-Dependent RNA Polymerase Detects Hidden Giant Viruses in Published Databanks. Genome Biology and Evolution, 2014, 6, 1603-1610.	2.5	41
874	Patterns of Kingella kingae Disease Outbreaks. Pediatric Infectious Disease Journal, 2016, 35, 340-346.	2.0	41
875	Identification of virulence factors and antibiotic resistance markers using bacterial genomics. Future Microbiology, 2016, 11, 455-466.	2.0	41
876	Nonhuman primates across sub-Saharan Africa are infected with the yaws bacterium <i>Treponema pallidum</i> subsp. <i>pertenue</i> Emerging Microbes and Infections, 2018, 7, 1-4.	6.5	41
877	Deciphering the Urinary Microbiota Repertoire by Culturomics Reveals Mostly Anaerobic Bacteria From the Gut. Frontiers in Microbiology, 2020, 11, 513305.	3.5	41
878	Optimization and standardization of the culturomics technique for human microbiome exploration. Scientific Reports, 2020, 10, 9674.	3.3	41
879	In Vitro Susceptibilities of Four <i>Bartonella bacilliformis</i> Strains to 30 Antibiotic Compounds. Antimicrobial Agents and Chemotherapy, 1999, 43, 2090-2092.	3.2	40
880	RNA polymerase $\hat{I}^2$ -subunit-based phylogeny of Ehrlichia spp., Anaplasma spp., Neorickettsia spp. and Wolbachia pipientis. International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 455-458.	1.7	40
881	A new example of viral intein in Mimivirus. Virology Journal, 2005, 2, 8.	3.4	40
882	Streptococcus massiliensis sp. nov., isolated from a patient blood culture. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1127-1131.	1.7	40

#	Article	IF	CITATIONS
883	Deciphering the Relationships between <i>Rickettsia conorii conorii</i> and <i>Rhipicephalus sanguineus</i> in the Ecology and Epidemiology of Mediterranean Spotted Fever. Annals of the New York Academy of Sciences, 2009, 1166, 49-54.	3.8	40
884	Ameba-associated Keratitis, France. Emerging Infectious Diseases, 2011, 17, 1306-1308.	4.3	40
885	Emergence of human granulocytic anaplasmosis in France. Ticks and Tick-borne Diseases, 2012, 3, 403-405.	2.7	40
886	Synergistic activity of sulbactam combined with colistin against colistin-resistant Acinetobacter baumannii. International Journal of Antimicrobial Agents, 2012, 39, 180-181.	2.5	40
887	TRUC or the Need for a New Microbial Classification. Intervirology, 2013, 56, 349-353.	2.8	40
888	Evaluating the Clinical Burden and Mortality Attributable to Antibiotic Resistance: The Disparity of Empirical Data and Simple Model Estimations. Clinical Infectious Diseases, 2017, 65, S58-S63.	5.8	40
889	Children account for a small proportion of diagnoses of SARS-CoV-2 infection and do not exhibit greater viral loads than adults. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1983-1987.	2.9	40
890	<i>Alistipes finegoldii</i> in Blood Cultures from Colon Cancer Patients. Emerging Infectious Diseases, 2007, 13, 1260-1262.	4.3	40
891	Coxiella burnetii Pericarditis: Report of 15 Cases and Review. Clinical Infectious Diseases, 1999, 29, 393-397.	5.8	39
892	Serosurvey among Mediterranean spotted fever patients of a new spotted fever group rickettsial strain (Bar29). European Journal of Epidemiology, 2002, 18, 351-354.	5.7	39
893	Q Fever in Thailand. Emerging Infectious Diseases, 2003, 9, 1186-1187.	4.3	39
894	Finegoldia magna, an early post-operative cause of infectious endocarditis: Report of two cases and review of the literature. Anaerobe, 2008, 14, 310-312.	2.1	39
895	Intracellular Life of <i>Coxiella burnetii</i> in Macrophages. Annals of the New York Academy of Sciences, 2009, 1166, 55-66.	3.8	39
896	The infective causes of hepatitis and jaundice amongst hospitalised patients in Vientiane, Laos. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2010, 104, 475-483.	1.8	39
897	Scrub Typhus Involving Central Nervous System, India, 2004–2006. Emerging Infectious Diseases, 2010, 16, 1641-1643.	4.3	39
898	African tick-bite fever: a new entity in the differential diagnosis of multiple eschars in travelers. Description of five cases imported from South Africa to Switzerland. International Journal of Infectious Diseases, 2010, 14, e274-e276.	3.3	39
899	Whipple Disease Revealed by Lung Involvement. Chest, 2012, 141, 1595-1598.	0.8	39
900	Molecular Detection of Acinetobacter Species in Lice and Keds of Domestic Animals in Oromia Regional State, Ethiopia. PLoS ONE, 2012, 7, e52377.	2.5	39

#	Article	IF	CITATIONS
901	Bartonella and Rickettsia in arthropods from the Lao PDR and from Borneo, Malaysia. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 51-57.	1.6	39
902	Description of "yaafâ€, the vesicular fever caused byÂacute Rickettsia felis infection in Senegal. Journal of Infection, 2013, 66, 536-540.	3.3	39
903	Impaired Virulence and Fitness of a Colistin-Resistant Clinical Isolate of Acinetobacter baumannii in a Rat Model of Pneumonia. Antimicrobial Agents and Chemotherapy, 2013, 57, 5120-5121.	3.2	39
904	<i>Tropheryma whipplei</i> as a commensal bacterium. Future Microbiology, 2013, 8, 57-71.	2.0	39
905	Whipple's disease and Tropheryma whipplei infections: when to suspect them and how to diagnose and treat them. Current Opinion in Infectious Diseases, 2018, 31, 463-470.	3.1	39
906	Molecular investigation and phylogeny of species of the Anaplasmataceae infecting animals and ticks in Senegal. Parasites and Vectors, 2019, 12, 495.	2.5	39
907	Testing the repatriated for SARS-Cov2: Should laboratory-based quarantine replace traditional quarantine?. Travel Medicine and Infectious Disease, 2020, 34, 101624.	3.0	39
908	Emergence in southern France of a new SARS-CoV-2 variant harbouring both N501Y and E484K substitutions in the spike protein. Archives of Virology, 2022, 167, 1185-1190.	2.1	39
909	Nonculture laboratory methods for the diagnosis of infectious endocarditis. Current Infectious Disease Reports, 1999, 1, 136-141.	3.0	38
910	HumanRickettsia felisInfection, Canary Islands, Spain. Emerging Infectious Diseases, 2005, 11, 1961-1964.	4.3	38
911	Histologic Features and Immunodetection of African Tick-bite Fever Eschar. Emerging Infectious Diseases, 2006, 12, 1332-1337.	4.3	38
912	Infection of Endothelial Cells with VirulentRickettsia prowazekiilncreases the Transmigration of Leukocytes. Journal of Infectious Diseases, 2008, 197, 142-147.	4.0	38
913	Classification and Determination of Possible Origins of ORFans through Analysis of Nucleocytoplasmic Large DNA Viruses. Intervirology, 2010, 53, 310-320.	2.8	38
914	Immunoglobulin G Anticardiolipin Antibodies and Progression to Q Fever Endocarditis. Clinical Infectious Diseases, 2013, 57, 57-64.	5.8	38
915	Non contiguous-finished genome sequence and description of Peptoniphilus senegalensis sp. nov Standards in Genomic Sciences, 2013, 7, 370-381.	1.5	38
916	Real-Time Microbiology Laboratory Surveillance System to Detect Abnormal Events and Emerging Infections, Marseille, France. Emerging Infectious Diseases, 2015, 21, 1302-1310.	4.3	38
917	How a multidisciplinary â€~One Health' approach can combat the tick-borne pathogen threat in Europe. Future Microbiology, 2015, 10, 809-818.	2.0	38
918	Comparison of Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry and Molecular Biology Techniques for Identification of Culicoides (Diptera: Ceratopogonidae) Biting Midges in Senegal. Journal of Clinical Microbiology, 2015, 53, 410-418.	3.9	38

#	Article	IF	CITATIONS
919	Arsenophonus nasoniae and Rickettsiae Infection of Ixodes ricinus Due to Parasitic Wasp Ixodiphagus hookeri. PLoS ONE, 2016, 11, e0149950.	2.5	38
920	18F-FDG PET/CT as a central tool in the shift from chronic Q fever to Coxiella burnetii persistent focalized infection. Medicine (United States), 2016, 95, e4287.	1.0	38
921	Current Status of Putative Animal Sources of SARS-CoV-2 Infection in Humans: Wildlife, Domestic Animals and Pets. Microorganisms, 2021, 9, 868.	3.6	38
922	Isolation and characterization of Reyranella massiliensis gen. nov., sp. nov. from freshwater samples by using an amoeba co-culture procedure. International Journal of Systematic and Evolutionary Microbiology, 2011, 61, 2151-2154.	1.7	38
923	Variations of Plasmid Content in Rickettsia felis. PLoS ONE, 2008, 3, e2289.	2.5	38
924	Distinguishing Body Lice from Head Lice by Multiplex Real-Time PCR Analysis of the Phum_PHUM540560 Gene. PLoS ONE, 2013, 8, e58088.	2.5	38
925	Prevalence of skin infections in sheltered homeless. European Journal of Dermatology, 2005, 15, 382-6.	0.6	38
926	Rickettsial Infections of the Central Nervous System. Seminars in Neurology, 1992, 12, 213-224.	1.4	37
927	Survey of seroprevalence of Q fever in dogs in the southeast of France, French Guyana, Martinique, Senegal and the Ivory Coast. Veterinary Microbiology, 1998, 64, 1-5.	1.9	37
928	Afebrile Blood Culture–Negative Endocarditis. Annals of Internal Medicine, 1999, 131, 144.	3.9	37
929	Determination of the Nucleotide Sequences of Heat Shock Operon groESL and the Citrate Synthase Gene ( gltA ) of Anaplasma ( Ehrlichia ) platys for Phylogenetic and Diagnostic Studies. Vaccine Journal, 2002, 9, 1132-1136.	3.1	37
930	Molecular Detection of Bartonella henselae DNA in the Dental Pulp of 800-Year-Old French Cats. Clinical Infectious Diseases, 2004, 39, 1391-1394.	5.8	37
931	Identification of candidate antigen in Whipple's disease using a serological proteomic approach. Proteomics, 2006, 6, 3294-3305.	2.2	37
932	<b>Infection in Man, France</b> . Emerging Infectious Diseases, 2009, 15, 1126-1127.	4.3	37
933	Brief communication: Coâ€detection of <i>Bartonella quintana</i> and <i>Yersinia pestis</i> in an 11th–15th burial site in Bondy, France. American Journal of Physical Anthropology, 2011, 145, 489-494.	2.1	37
934	Non contiguous-finished genome sequence and description of Enorma massiliensis gen. nov., sp. nov., a new member of the Family Coriobacteriaceae. Standards in Genomic Sciences, 2013, 8, 290-305.	1.5	37
935	Reevaluation of the Risk of Fetal Death and Malformation After Q Fever. Clinical Infectious Diseases, 2014, 59, 256-260.	5.8	37
936	Characterization of Viral Communities of Biting Midges and Identification of Novel Thogotovirus Species and Rhabdovirus Genus. Viruses, 2016, 8, 77.	3.3	37

#	Article	lF	Citations
937	Pattern of SARS-CoV-2 infection among dependant elderly residents living in long-term care facilities in Marseille, France, March–June 2020. International Journal of Antimicrobial Agents, 2020, 56, 106219.	2.5	37
938	Bartonella: new explanations for old diseases. Journal of Medical Microbiology, 2002, 51, 915-923.	1.8	37
939	Positive Predictive Value of Rochalimaea henselae Antibodies in the Diagnosis of Cat-Scratch Disease. Clinical Infectious Diseases, 1994, 19, 355-355.	5.8	36
940	Detection of Astrakhan Fever Rickettsia from Ticks in Kosovo. Annals of the New York Academy of Sciences, 2003, 990, 158-161.	3.8	36
941	Systemic Inflammatory Responses in African Tickâ€Bite Fever. Journal of Infectious Diseases, 2003, 187, 1332-1336.	4.0	36
942	Usefulness of rpoB Gene Sequencing for Identification of Afipia and Bosea Species, Including a Strategy for Choosing Discriminative Partial Sequences. Applied and Environmental Microbiology, 2003, 69, 6740-6749.	3.1	36
943	Proposal to Create Subspecies of Rickettsia sibirica and an Emended Description of Rickettsia sibirica. Annals of the New York Academy of Sciences, 2006, 1078, 597-606.	3.8	36
944	Global Analysis of Circulating Immune Cells by Matrix-Assisted Laser Desorption Ionization Time-of-Flight Mass Spectrometry. PLoS ONE, 2010, 5, e13691.	2.5	36
945	Beyond ancient microbial DNA: nonnucleotidic biomolecules for paleomicrobiology. BioTechniques, 2011, 50, 370-380.	1.8	36
946	A New <i>Rickettsia </i> Species Found in Fleas Collected from Human Dwellings and from Domestic Cats and Dogs in Senegal. Vector-Borne and Zoonotic Diseases, 2012, 12, 360-365.	1.5	36
947	Urban family cluster of spotted fever rickettsiosis linked to Rhipicephalus sanguineus infected with Rickettsia conorii subsp. caspia and Rickettsia massiliae. Ticks and Tick-borne Diseases, 2012, 3, 389-392.	2.7	36
948	Serologic Prevalence of Amoeba-Associated Microorganisms in Intensive Care Unit Pneumonia Patients. PLoS ONE, 2013, 8, e58111.	2.5	36
949	Marseillevirus in lymphoma: a giant in the lymph node. Lancet Infectious Diseases, The, 2016, 16, e225-e234.	9.1	36
950	The impact of culturomics on taxonomy in clinical microbiology. Antonie Van Leeuwenhoek, 2017, 110, 1327-1337.	1.7	36
951	Extremely and pandrug-resistant bacteria extra-deaths: myth or reality?. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 1687-1697.	2.9	36
952	SARS-CoV-2 Infectivity and Severity of COVID-19 According to SARS-CoV-2 Variants: Current Evidence. Journal of Clinical Medicine, 2021, 10, 2635.	2.4	36
953	Isolation of <i>Rickettsia prowazekii</i> from Blood by Shell Vial Cell Culture. Journal of Clinical Microbiology, 1999, 37, 3722-3724.	3.9	36
954	Isolation of <i>Legionella pneumophila</i> by Centrifugation of Shell Vial Cell Cultures from Multiple Liver and Lung Abscesses. Journal of Clinical Microbiology, 1999, 37, 785-787.	3.9	36

#	Article	IF	CITATIONS
955	Sex-Related Differences in Gene Expression Following Coxiella burnetii Infection in Mice: Potential Role of Circadian Rhythm. PLoS ONE, 2010, 5, e12190.	2.5	36
956	Development and Validation of a Microarray for the Investigation of the CAZymes Encoded by the Human Gut Microbiome. PLoS ONE, 2013, 8, e84033.	2.5	36
957	Culture of <i>Tropheryma whippelii</i> from the Vitreous Fluid of a Patient Presenting with Unilateral Uveitis. Annals of Internal Medicine, 2003, 139, 1046.	3.9	36
958	Zoonotic Focus of Plague, Algeria. Emerging Infectious Diseases, 2006, 12, 1975-1977.	4.3	35
959	Antiangiogenic Effect of Erythromycin: An In Vitro Model ofBartonella quintanaInfection. Journal of Infectious Diseases, 2006, 193, 380-386.	4.0	35
960	Bacillus massiliensis sp. nov., isolated from cerebrospinal fluid. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1485-1488.	1.7	35
961	Identification of rickettsial isolates at the species level using multi-spacer typing. BMC Microbiology, 2007, 7, 72.	3.3	35
962	Rickettsial diseases: from <i>Rickettsia</i> àê"arthropod relationships to pathophysiology and animal models. Future Microbiology, 2008, 3, 223-236.	2.0	35
963	A Single-Step Sequencing Method for the Identification of Mycobacterium tuberculosis Complex Species. PLoS Neglected Tropical Diseases, 2008, 2, e253.	3.0	35
964	Murine Typhus and Leptospirosis as Causes of Acute Undifferentiated Fever, Indonesia. Emerging Infectious Diseases, 2009, 15, 975-977.	4.3	35
965	Detection of Termites and Other Insects Consumed by African Great Apes using Molecular Fecal Analysis. Scientific Reports, 2014, 4, 4478.	3.3	35
966	Draft genome of Gemmata massiliana sp. nov, a water-borne Planctomycetes species exhibiting two variants. Standards in Genomic Sciences, 2015, 10, 120.	1.5	35
967	Developmental Cycle and Genome Analysis of "Rubidus massiliensis,―a New Vermamoeba vermiformis Pathogen. Frontiers in Cellular and Infection Microbiology, 2016, 6, 31.	3.9	35
968	Microbiome of HIV-infected people. Microbial Pathogenesis, 2017, 106, 85-93.	2.9	35
969	Blastocystis Colonization Is Associated with Increased Diversity and Altered Gut Bacterial Communities in Healthy Malian Children. Microorganisms, 2019, 7, 649.	3 <b>.</b> 6	35
970	Comparison Between ESC and Duke Criteria for the Diagnosis of Prosthetic Valve Infective Endocarditis. JACC: Cardiovascular Imaging, 2020, 13, 2605-2615.	<b>5.</b> 3	35
971	Can ACE2 Receptor Polymorphism Predict Species Susceptibility to SARS-CoV-2?. Frontiers in Public Health, 2020, 8, 608765.	2.7	35
972	Spread of Mink SARS-CoV-2 Variants in Humans: A Model of Sarbecovirus Interspecies Evolution. Frontiers in Microbiology, 2021, 12, 675528.	3 <b>.</b> 5	35

#	Article	IF	Citations
973	Isolation of Francisella tularensis by Centrifugation of Shell Vial Cell Culture from an Inoculation Eschar. Journal of Clinical Microbiology, 1998, 36, 2782-2783.	3.9	35
974	Mansonella, including a Potential New Species, as Common Parasites in Children in Gabon. PLoS Neglected Tropical Diseases, 2015, 9, e0004155.	3.0	35
975	SerotypingCoxiella burnetiiisolates from acute and chronic Q fever patients by using monoclonal antibodies. FEMS Microbiology Letters, 1994, 117, 15-19.	1.8	34
976	Isolation of new fastidious α Proteobacteria and Afipia felis from hospital water supplies by direct plating and amoebal co-culture procedures. FEMS Microbiology Ecology, 2000, 34, 129-137.	2.7	34
977	Molecular survey of Ehrlichia infection in ticks from animals in Yamaguchi Prefecture, Japan. Veterinary Parasitology, 2001, 99, 335-339.	1.8	34
978	Are clinical isolates of Pseudomonas aeruginosa more virulent than hospital environmental isolates in amebal co-culture test?. Critical Care Medicine, 2006, 34, 823-828.	0.9	34
979	Murine Typhus Poorly Responsive to Ciprofloxacin: A Case Report. Journal of Travel Medicine, 2002, 9, 103-104.	3.0	34
980	Genotyping of <i>Orientia tsutsugamushi </i> Ii>from Humans with Scrub Typhus, Laos. Emerging Infectious Diseases, 2008, 14, 1483-1485.	4.3	34
981	<i>Tropheryma whipplei</i> Glycosylation in the Pathophysiologic Profile of Whipple's Disease. Journal of Infectious Diseases, 2009, 199, 1043-1052.	4.0	34
982	There is no such thing as a tree of life (and of course viruses are out!). Nature Reviews Microbiology, 2009, 7, 615-615.	28.6	34
983	Relative chemokine and adhesion molecule expression in Mediterranean spotted fever and African tick bite fever. Journal of Infection, 2009, 58, 68-75.	3.3	34
984	<i>Rickettsia sibirica mongolitimonae</i> i>in Traveler from Egypt. Emerging Infectious Diseases, 2010, 16, 1495-1496.	4.3	34
985	Rickettsia africae in Hyalomma dromedarii ticks from sub-Saharan Algeria. Ticks and Tick-borne Diseases, 2012, 3, 377-379.	2.7	34
986	Investigation of <i><scp>A</scp>cinetobacter baumannii</i> resistance to carbapenems in <scp>M</scp> arseille hospitals, south of <scp>F</scp> rance: a transition from an epidemic to an endemic situation. Apmis, 2013, 121, 64-71.	2.0	34
987	Bartonella quintana in Cimex hemipterus, Rwanda. American Journal of Tropical Medicine and Hygiene, 2013, 89, 986-987.	1.4	34
988	Non contiguous-finished genome sequence and description of Peptoniphilus grossensis sp. nov Standards in Genomic Sciences, 2012, 7, 320-330.	1.5	34
989	Multiplex Real-Time PCR Diagnostic of Relapsing Fevers in Africa. PLoS Neglected Tropical Diseases, 2013, 7, e2042.	3.0	34
990	Bacterial Lymphadenitis at a Major Referral Hospital in France from 2008 to 2012. Journal of Clinical Microbiology, 2014, 52, 1161-1167.	3.9	34

#	Article	IF	Citations
991	Q Fever in French Guiana. American Journal of Tropical Medicine and Hygiene, 2014, 91, 771-776.	1.4	34
992	High quality draft genome sequence and description of Occidentia massiliensis gen. nov., sp. nov., a new member of the family Rickettsiaceae. Standards in Genomic Sciences, 2014, 9, 9.	1.5	34
993	Causes of Fever in Rural Southern Laos. American Journal of Tropical Medicine and Hygiene, 2015, 93, 517-520.	1.4	34
994	An Outbreak of Kingella kingae Infections Associated with Hand, Foot and Mouth Disease/Herpangina Virus Outbreak in Marseille, France, 2013. Pediatric Infectious Disease Journal, 2015, 34, 246-250.	2.0	34
995	MG-Digger: An Automated Pipeline to Search for Giant Virus-Related Sequences in Metagenomes. Frontiers in Microbiology, 2016, 7, 428.	3 <b>.</b> 5	34
996	Communicable and non-communicable disease risks at the Grand Magal of Touba: The largest mass gathering in Senegal. Travel Medicine and Infectious Disease, 2017, 19, 56-60.	3.0	34
997	Stem-Loop RNA Hairpins in Giant Viruses: Invading rRNA-Like Repeats and a Template Free RNA. Frontiers in Microbiology, 2018, 9, 101.	3.5	34
998	The Microbiological Memory, an Epigenetic Regulator Governing the Balance Between Good Health and Metabolic Disorders. Frontiers in Microbiology, 2018, 9, 1379.	3.5	34
999	Human metallo- $\hat{l}^2$ -lactamase enzymes degrade penicillin. Scientific Reports, 2019, 9, 12173.	3.3	34
1000	Isolation and characterization of Kingella negevensis sp. nov., a novel Kingella species detected in a healthy paediatric population. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2370-2376.	1.7	34
1001	Rickettsia conorii in Greece: Comparison of a Microimmunofluorescence Assay and Western Blotting for Seroepidemiology. American Journal of Tropical Medicine and Hygiene, 1993, 48, 784-792.	1.4	34
1002	The release in vitro of vancomycin and tobramycin from acrylic bone cement. Journal of Antimicrobial Chemotherapy, 1994, 33, 337-339.	3.0	33
1003	An Experimental Model of Human Body Louse Infection with <i>Rickettsia typhi</i> i>. Annals of the New York Academy of Sciences, 2003, 990, 617-627.	3.8	33
1004	Prosthetic valve endocarditis caused by Pseudomonas luteola. BMC Infectious Diseases, 2005, 5, 82.	2.9	33
1005	Prevotella timonensis sp. nov., isolated from a human breast abscess. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 883-886.	1.7	33
1006	Development of a method for recovering rickettsial RNA from infected cells to analyze gene expression profiling of obligate intracellular bacteria. Journal of Microbiological Methods, 2007, 71, 292-297.	1.6	33
1007	Molecular mechanisms of resistance to antibiotics in Bartonella bacilliformis. Journal of Antimicrobial Chemotherapy, 2007, 59, 1065-1070.	3.0	33
1008	Seasonality of mosquitoes and chikungunya in Italy. Lancet Infectious Diseases, The, 2008, 8, 5-6.	9.1	33

#	Article	IF	CITATIONS
1009	Murine Typhus as a Cause of Fever in Travelers From Tunisia and Mediterranean Areas. Journal of Travel Medicine, 2010, 17, 310-315.	3.0	33
1010	Investigation of blood culture-negative early prosthetic valve endocarditis reveals high prevalence of fungi. Heart, 2010, 96, 743-747.	2.9	33
1011	Progressive dementia associated with ataxia or obesity in patients with Tropheryma whipplei encephalitis. BMC Infectious Diseases, 2011, 11, 171.	2.9	33
1012	First detection of Rickettsia aeschlimannii in Hyalomma dromedarii ticks from Tunisia. Ticks and Tick-borne Diseases, 2012, 3, 398-402.	2.7	33
1013	First isolation of Tropheryma whipplei from bronchoalveolar fluid and clinical implications. Journal of Infection, 2012, 65, 275-278.	3.3	33
1014	Giant Viruses of Amoebae as Potential Human Pathogens. Intervirology, 2013, 56, 376-385.	2.8	33
1015	Looking for Tropheryma whipplei Source and Reservoir in Rural Senegal. American Journal of Tropical Medicine and Hygiene, 2013, 88, 339-343.	1.4	33
1016	Non contiguous-finished genome sequence and description of Bacillus massiliosenegalensis sp. nov Standards in Genomic Sciences, 2013, 8, 264-278.	1.5	33
1017	Comparison between Emerging Q Fever in French Guiana and Endemic Q fever in Marseille, France. American Journal of Tropical Medicine and Hygiene, 2014, 90, 915-919.	1.4	33
1018	Pan-genomic analysis to redefine species and subspecies based on quantum discontinuous variation: the Klebsiella paradigm. Biology Direct, 2015, 10, 55.	4.6	33
1019	The aerobic activity of metronidazole against anaerobic bacteria. International Journal of Antimicrobial Agents, 2015, 45, 537-540.	2.5	33
1020	High Prevalence of Tropheryma whipplei in Lao Kindergarten Children. PLoS Neglected Tropical Diseases, 2015, 9, e0003538.	3.0	33
1021	Occurrence and Genotyping of Coxiella burnetii in Ixodid Ticks in Oromia, Ethiopia. American Journal of Tropical Medicine and Hygiene, 2015, 93, 1074-1081.	1.4	33
1022	Antiphospholipid Antibody Syndrome With Valvular Vegetations in Acute Q Fever. Clinical Infectious Diseases, 2016, 62, 537-544.	5.8	33
1023	Intracranial haemorrhage in infective endocarditis. Archives of Cardiovascular Diseases, 2018, 111, 712-721.	1.6	33
1024	State of the Art in the Culture of the Human Microbiota: New Interests and Strategies. Clinical Microbiology Reviews, 2020, 34, .	13.6	33
1025	A new protectant medium preserving bacterial viability after freeze drying. Microbiological Research, 2020, 236, 126454.	5.3	33
1026	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open label non-randomized clinical trial revisited. International Journal of Antimicrobial Agents, 2021, 57, 106243.	2.5	33

#	Article	IF	CITATIONS
1027	Clinical evidence of the role of Methanobrevibacter smithii in severe acute malnutrition. Scientific Reports, 2021, 11, 5426.	3.3	33
1028	Fournierella massiliensis gen. nov., sp. nov., a new human-associated member of the family Ruminococcaceae. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 1393-1399.	1.7	33
1029	RickA Expression Is Not Sufficient to Promote Actin-Based Motility of Rickettsia raoultii. PLoS ONE, 2008, 3, e2582.	2.5	33
1030	Molecular Evidence for the Presence of Rickettsia Felis in the Feces of Wild-living African Apes. PLoS ONE, 2013, 8, e54679.	2.5	33
1031	Sennetsu Neorickettsiosis: A Probable Fish-Borne Cause of Fever Rediscovered in Laos. American Journal of Tropical Medicine and Hygiene, 2009, 81, 190-194.	1.4	33
1032	The line blot: an immunoassay for monoclonal and other antibodies. Journal of Immunological Methods, 1989, 125, 57-65.	1.4	32
1033	Coxiella burnetii infection of pseudoaneurysm of an aortic bypass graft with contiguous vertebral osteomyelitis. Journal of Vascular Surgery, 1994, 19, 165-168.	1.1	32
1034	Detection of Coxiella burnetii DNA in dental pulp during experimental bacteremia. Microbial Pathogenesis, 2000, 28, 249-254.	2.9	32
1035	Patients in the Intensive Care Unit are Exposed to Amoeba-Associated Pathogens. Infection Control and Hospital Epidemiology, 2002, 23, 462-465.	1.8	32
1036	Culture and Phenotypic Characterization of a Wolbachia pipientis Isolate. Journal of Clinical Microbiology, 2003, 41, 5434-5441.	3.9	32
1037	Aeromonas popoffii Urinary Tract Infection. Journal of Clinical Microbiology, 2004, 42, 5427-5428.	3.9	32
1038	Identification of <i>Rickettsia africae </i> and <i>Wolbachia </i> p. in <i>Ceratophyllus garei </i> Fleas from Passerine Birds Migrated from Africa. Vector-Borne and Zoonotic Diseases, 2012, 12, 539-543.	1.5	32
1039	Non-contiguous finished genome sequence and description of Bacillus massiliogorillae sp. nov Standards in Genomic Sciences, 2013, 9, 93-105.	1.5	32
1040	Genome sequence and description of Timonella senegalensis gen. nov., sp. nov., a new member of the suborder Micrococcinae. Standards in Genomic Sciences, 2013, 8, 318-335.	1.5	32
1041	<i>Borrelia crocidurae</i> Infection in Acutely Febrile Patients, Senegal. Emerging Infectious Diseases, 2014, 20, 1335-1338.	<b>4.</b> 3	32
1042	Infective Endocarditis in Northeastern Thailand. Emerging Infectious Diseases, 2014, 20, 473-6.	4.3	32
1043	Two Human Cases of <i>Rickettsia felis </i> Infection, Thailand. Emerging Infectious Diseases, 2014, 20, 1780-1781.	4.3	32
1044	Whole-genome assembly of Akkermansia muciniphila sequenced directly from human stool. Biology Direct, 2015, 10, 5.	4.6	32

#	Article	IF	Citations
1045	Louse-borne relapsing fever among East African refugees in Europe. Travel Medicine and Infectious Disease, 2016, 14, 110-114.	3.0	32
1046	Human Polyomavirus-6 Infecting Lymph Nodes of a Patient With an Angiolymphoid Hyperplasia With Eosinophilia or Kimura Disease. Clinical Infectious Diseases, 2016, 62, 1419-1421.	5.8	32
1047	Using MALDI-TOF MS to identify mosquitoes collected in Mali and their blood meals. Parasitology, 2018, 145, 1170-1182.	1.5	32
1048	Body lice of homeless people reveal the presence of several emerging bacterial pathogens in northern Algeria. PLoS Neglected Tropical Diseases, 2018, 12, e0006397.	3.0	32
1049	High-dose trimethoprim-sulfamethoxazole and clindamycin for Staphylococcus aureus endocarditis. International Journal of Antimicrobial Agents, 2019, 54, 143-148.	2.5	32
1050	Faecal microbiota transplantation shortens the colonisation period and allows re-entry of patients carrying carbapenamase-producing bacteria into medical care facilities. International Journal of Antimicrobial Agents, 2019, 53, 355-361.	2.5	32
1051	New insights in <i>Coxiella burnetii</i> infection: diagnosis and therapeutic update. Expert Review of Anti-Infective Therapy, 2020, 18, 75-86.	4.4	32
1052	<i>rpoB</i> Gene Analysis as a Novel Strategy for Identification of Spirochetes from the Genera <i>Borrelia</i> , <i>Treponema</i> , and <i>Leptospira</i> . Journal of Clinical Microbiology, 2000, 38, 2200-2203.	3.9	32
1053	Heteroplasmy in the Mitochondrial Genomes of Human Lice and Ticks Revealed by High Throughput Sequencing. PLoS ONE, 2013, 8, e73329.	2.5	32
1054	EXPERIMENTAL INFECTION OF HUMAN BODY LICE WITH ACINETOBACTER BAUMANNII. American Journal of Tropical Medicine and Hygiene, 2006, 74, 526-531.	1.4	32
1055	Identification of Bartonella bacilliformis Genotypes and Their Relevance to Epidemiological Investigations of Human Bartonellosis. Journal of Clinical Microbiology, 2002, 40, 3606-3612.	3.9	31
1056	Bartonella clarridgeiaeandB. henselaein Dogs, Gabon. Emerging Infectious Diseases, 2004, 10, 2257-2258.	4.3	31
1057	Increased Levels of Circulating IL-16 and Apoptosis Markers Are Related to the Activity of Whipple's Disease. PLoS ONE, 2007, 2, e494.	2.5	31
1058	Heterogeneity of susceptibility to fluoroquinolones in Bartonella isolates from Australia reveals a natural mutation in gyrA. Journal of Antimicrobial Chemotherapy, 2008, 61, 1252-1255.	3.0	31
1059	Systemic Immune Presentations of Coxiella burnetii Infection (Q Fever). Seminars in Arthritis and Rheumatism, 2010, 39, 405-409.	3.4	31
1060	Multi-Locus Sequence Typing of a Geographically and Temporally Diverse Sample of the Highly Clonal Human Pathogen Bartonella quintana. PLoS ONE, 2010, 5, e9765.	2.5	31
1061	Genome Sequence of Coxiella burnetii 109, a Doxycycline-Resistant Clinical Isolate. Journal of Bacteriology, 2012, 194, 6939-6939.	2.2	31
1062	Deleterious effect of ciprofloxacin on Rickettsia conorii-infected cells is linked to toxin-antitoxin module up-regulation. Journal of Antimicrobial Chemotherapy, 2012, 67, 1677-1682.	3.0	31

#	Article	IF	CITATIONS
1063	Bartonella and Coxiella infective endocarditis in Brazil: molecular evidence from excised valves from a cardiac surgery referral center in Rio de Janeiro, Brazil, 1998 to 2009. International Journal of Infectious Diseases, 2013, 17, e65-e66.	3.3	31
1064	Open membranes are the precursors for assembly of large DNA viruses. Cellular Microbiology, 2013, 15, n/a-n/a.	2.1	31
1065	Non contiguous-finished genome sequence and description of Peptoniphilus obesi sp. nov Standards in Genomic Sciences, 2013, 7, 357-369.	1.5	31
1066	Worldwide decrease in methicillin-resistant Staphylococcus aureus: do we understand something?. Clinical Microbiology and Infection, 2015, 21, 515-517.	6.0	31
1067	Malaria in urban, semi-urban and rural areas of southern of Gabon: comparison of the Pfmdr 1 and Pfcrt genotypes from symptomatic children. Malaria Journal, 2016, 15, 420.	2.3	31
1068	Molecular history of plague. Clinical Microbiology and Infection, 2016, 22, 911-915.	6.0	31
1069	Blautia massiliensis sp. nov., isolated from a fresh human fecal sample and emended description of the genus Blautia. Anaerobe, 2017, 43, 47-55.	2.1	31
1070	Child with liver transplant recovers from COVID-19 infection. A case report. Archives De Pediatrie, 2020, 27, 275-276.	1.0	31
1071	Introduction into the Marseille geographical area of a mild SARS-CoV-2 variant originating from sub-Saharan Africa: An investigational study. Travel Medicine and Infectious Disease, 2021, 40, 101980.	3.0	31
1072	Prolonged SARS-CoV-2 RNA virus shedding and lymphopenia are hallmarks of COVID-19 in cancer patients with poor prognosis. Cell Death and Differentiation, 2021, 28, 3297-3315.	11.2	31
1073	Astrakhan Fever Rickettsia Is Identical to Israel Tick Typhus Rickettsia, a Genotype of the Rickettsia conorii Complex. Journal of Infectious Diseases, 1992, 165, 1167-1168.	4.0	30
1074	Scrub Typhus after a Trip to Vietnam. New England Journal of Medicine, 1997, 336, 1613-1614.	27.0	30
1075	A Guinea Pig Model for Q Fever Endocarditis. Journal of Infectious Diseases, 1998, 178, 278-281.	4.0	30
1076	Immunohistological detection of Tropheryma whipplei (Whipple bacillus) in lymph nodes. American Journal of Medicine, 2002, 113, 334-336.	1.5	30
1077	Isolation of a Rickettsia Related to Astrakhan Fever Rickettsia from a Patient in Chad. Annals of the New York Academy of Sciences, 2003, 990, 152-157.	3.8	30
1078	Biocides Currently Used for Bronchoscope Decontamination Are Poorly Effective Against Free-Living Amoebae. Infection Control and Hospital Epidemiology, 2003, 24, 784-786.	1.8	30
1079	Culture and Antibiotic Susceptibility of Bartonella quintana in Human Erythrocytes. Antimicrobial Agents and Chemotherapy, 2003, 47, 614-619.	3.2	30
1080	Antigenic Classification of Rickettsia felis by Using Monoclonal and Polyclonal Antibodies. Vaccine Journal, 2003, 10, 221-228.	3.1	30

#	Article	IF	CITATIONS
1081	<i>Bartonella quintana</i> Bacteremia and Overproduction of Interleukinâ€10: Model of Bacterial Persistence in Homeless People. Journal of Infectious Diseases, 2003, 187, 837-844.	4.0	30
1082	Molecular detection of Yersinia pestis in dental pulp. Microbiology (United Kingdom), 2004, 150, 263-264.	1.8	30
1083	TLR2 Is Necessary to Inflammatory Response in Coxiella burnetii Infection. Annals of the New York Academy of Sciences, 2005, 1063, 161-166.	3.8	30
1084	Global Transcriptome Analysis of Tropheryma whipplei in Response to Temperature Stresses. Journal of Bacteriology, 2006, 188, 5228-5239.	2.2	30
1085	Role for the CD28 Molecule in the Control of Coxiella burnetii Infection. Infection and Immunity, 2006, 74, 1800-1808.	2.2	30
1086	Genetic Diversity of <i>Bartonella henselae </i> <in 1178-1183.<="" 13,="" 2007,="" detected="" diseases,="" emerging="" human="" infection="" infectious="" multispacer="" td="" typing.="" with=""><td>4.3</td><td>30</td></in>	4.3	30
1087	Structural, Biochemical, and in Vivo Characterization of the First Virally Encoded Cyclophilin from the Mimivirus. Journal of Molecular Biology, 2008, 378, 71-86.	4.2	30
1088	The effect of a single dose of oral ivermectin on pruritus in the homeless. Journal of Antimicrobial Chemotherapy, 2008, 62, 404-409.	3.0	30
1089	Rickettsia aeschlimanniiInfection, Algeria. Emerging Infectious Diseases, 2008, 14, 1814-1815.	4.3	30
1090	Brevibacterium massiliense sp. nov., isolated from a human ankle discharge. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 1960-1964.	1.7	30
1091	Host, pathogen and treatment-related prognostic factors in rickettsioses. European Journal of Clinical Microbiology and Infectious Diseases, 2011, 30, 1139-1150.	2.9	30
1092	Multispacer Sequence Typing Relapsing Fever Borreliae in Africa. PLoS Neglected Tropical Diseases, 2012, 6, e1652.	3.0	30
1093	Prevalence and Genetic Diversity of Bartonella spp. in Small Mammals from Southeastern Asia. Applied and Environmental Microbiology, 2012, 78, 8463-8466.	3.1	30
1094	Why Are There So Few Rickettsia conorii conorii-Infected Rhipicephalus sanguineus Ticks in the Wild?. PLoS Neglected Tropical Diseases, 2012, 6, e1697.	3.0	30
1095	Antibacterial efficacy of inhaled squalamine in a rat model of chronic Pseudomonas aeruginosa pneumonia. Journal of Antimicrobial Chemotherapy, 2012, 67, 2452-2458.	3.0	30
1096	Virophages question the existence of satellites. Nature Reviews Microbiology, 2012, 10, 234-234.	28.6	30
1097	Comparison of the performance of IFA, CFA, and ELISA assays for the serodiagnosis of acute Q fever by quality assessment. Diagnostic Microbiology and Infectious Disease, 2013, 75, 16-21.	1.8	30
1098	The expanding family Marseilleviridae. Virology, 2014, 466-467, 27-37.	2.4	30

#	Article	IF	CITATIONS
1099	Bartonella melophagi in Melophagus ovinus (sheep ked) collected from sheep in northern Oromia, Ethiopia. Comparative Immunology, Microbiology and Infectious Diseases, 2014, 37, 69-76.	1.6	30
1100	Prevalence of Bartonella quintana in Patients with Fever and Head Lice from Rural Areas of Sine-Saloum, Senegal. American Journal of Tropical Medicine and Hygiene, 2014, 91, 291-293.	1.4	30
1101	Detection of Bartonella spp. in fleas by MALDI-TOF MS. PLoS Neglected Tropical Diseases, 2018, 12, e0006189.	3.0	30
1102	Isolation of Yasminevirus, the First Member of Klosneuvirinae Isolated in Coculture with Vermamoeba vermiformis, Demonstrates an Extended Arsenal of Translational Apparatus Components. Journal of Virology, 2019, 94, .	3.4	30
1103	Molecular identification of protozoal and bacterial organisms in domestic animals and their infesting ticks from north-eastern Algeria. Ticks and Tick-borne Diseases, 2020, 11, 101330.	2.7	30
1104	Spreading of a new SARS-CoV-2 N501Y spike variant in a new lineage. Clinical Microbiology and Infection, 2021, 27, 1352.e1-1352.e5.	6.0	30
1105	High Ancient Genetic Diversity of Human Lice, Pediculus humanus, from Israel Reveals New Insights into the Origin of Clade B Lice. PLoS ONE, 2016, 11, e0164659.	2.5	30
1106	EXPERIMENTALLY INFECTED HUMAN BODY LICE (PEDICULUS HUMANUS HUMANUS) AS VECTORS OF RICKETTSIA RICKETTSII AND RICKETTSIA CONORII IN A RABBIT MODEL. American Journal of Tropical Medicine and Hygiene, 2006, 74, 521-525.	1.4	30
1107	Bacteriostatic and Bactericidal Activities of Moxifloxacin against Coxiella burnetii. Antimicrobial Agents and Chemotherapy, 2001, 45, 301-302.	3.2	29
1108	Human Infection Caused by <i>Leptospira fainei </i> . Emerging Infectious Diseases, 2002, 8, 865-868.	4.3	29
1109	Q Fever Outbreak in Homeless Shelter. Emerging Infectious Diseases, 2004, 10, 1297-1299.	4.3	29
1110	Multi-pathogens sequence containing plasmids as positive controls for universal detection of potential agents of bioterrorism. BMC Microbiology, 2004, 4, 21.	3.3	29
1111	Genetic Differentiation of Chinese Isolates of Rickettsia sibirica by Partial ompA Gene Sequencing and Multispacer Typing. Journal of Clinical Microbiology, 2006, 44, 2465-2467.	3.9	29
1112	Evaluation of clinical specimens for <i>Rickettsia</i> , <i>Bartonella</i> , <i>Borrelia</i> , <i>Coxiella</i> , <i>Anaplasma</i> , <i>Franciscella</i> and <i>using serological and molecular biology methods. FEMS Immunology and Medical Microbiology, 2012, 64, 82-91.</i>	Diploricket 2.7	ttsigposi
1113	Relevance of the positron emission tomography in the diagnosis of vascular graft infection with Coxiella burnetii. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 45-49.	1.6	29
1114	Rapid MALDI-TOF mass spectrometry identification of Leptospira organisms. Veterinary Microbiology, 2012, 158, 142-146.	1.9	29
1115	First Isolation of a Giant Virus from Wild Hirudo medicinalis Leech: Mimiviridae isolation in Hirudo medicinalis. Viruses, 2013, 5, 2920-2930.	3.3	29
1116	MALDI-TOF mass spectrometry and identification of new bacteria species in air samples from Makkah, Saudi Arabia. BMC Research Notes, 2014, 7, 892.	1.4	29

#	Article	IF	Citations
1117	Loss of TSS1 in hypervirulent Coxiella burnetii 175, the causative agent of Q fever in French Guiana. Comparative Immunology, Microbiology and Infectious Diseases, 2015, 41, 35-41.	1.6	29
1118	Enterococcus faecalis urinary-tract infections: Do they have a zoonotic origin?. Journal of Infection, 2016, 73, 305-313.	3.3	29
1119	Rickettsia massiliae infection and SENLAT syndrome in Romania. Ticks and Tick-borne Diseases, 2016, 7, 759-762.	2.7	29
1120	Bartonella bovis and Candidatus Bartonella davousti in cattle from Senegal. Comparative Immunology, Microbiology and Infectious Diseases, 2017, 50, 63-69.	1.6	29
1121	International experts' practice in the antibiotic therapy of infective endocarditis is not following the guidelines. Clinical Microbiology and Infection, 2017, 23, 736-739.	6.0	29
1122	No Such Thing as Chronic Q Fever. Emerging Infectious Diseases, 2017, 23, 856-857.	4.3	29
1123	<i>Delftia tsuruhatensis,</i> an Emergent Opportunistic Healthcare-Associated Pathogen. Emerging Infectious Diseases, 2018, 24, 594-596.	4.3	29
1124	Louse-Borne Relapsing Fever (Borrelia recurrentis) in a Somali Refugee Arriving in Italy: A Re-emerging Infection in Europe?. PLoS Neglected Tropical Diseases, 2016, 10, e0004522.	3.0	29
1125	The Transcriptional Programme of Human Heart Valves Reveals the Natural History of Infective Endocarditis. PLoS ONE, 2010, 5, e8939.	2.5	29
1126	First molecular detection of Rickettsia felis in fleas from Algeria. American Journal of Tropical Medicine and Hygiene, 2006, 74, 532-5.	1.4	29
1127	Seroprevalence of antibodies to Coxiella burnetti among pregnant women in South Eastern France. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2000, 93, 151-156.	1.1	28
1128	rpoB Sequence Analysis of Cultured Tropheryma whippelii. Journal of Clinical Microbiology, 2001, 39, 2425-2430.	3.9	28
1129	Prevalence of Bartonella clarridgeiae and Bartonella henselae in Domestic Cats from France and Detection of the Organisms in Erythrocytes by Immunofluorescence. Vaccine Journal, 2004, 11, 423-425.	2.6	28
1130	Sub-acute neuropathy in patients with African tick bite fever. Scandinavian Journal of Infectious Diseases, 2006, 38, 114-118.	1.5	28
1131	A Serum Protein Signature with High Diagnostic Value in Bacterial Endocarditis: Results from a Study Based on Surfaceâ€Enhanced Laser Desorption/Ionization Timeâ€ofâ€Flight Mass Spectrometry. Journal of Infectious Diseases, 2006, 194, 1356-1366.	4.0	28
1132	Outcome after surgical treatment performed within the first week of antimicrobial therapy during infective endocarditis: A prospective study. Archives of Cardiovascular Diseases, 2008, 101, 687-695.	1.6	28
1133	<i>Bartonella quintana</i> and <i>Coxiella burnetii</i> as Causes of Endocarditis, India. Emerging Infectious Diseases, 2008, 14, 1168-1169.	4.3	28
1134	Legionella tunisiensis sp. nov. and Legionella massiliensis sp. nov., isolated from environmental water samples. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 3003-3006.	1.7	28

#	Article	IF	Citations
1135	Complete Genome Sequence of Borrelia crocidurae. Journal of Bacteriology, 2012, 194, 3723-3724.	2.2	28
1136	Tobacco mosaic virus in cigarettes and saliva of smokers. Journal of Clinical Virology, 2012, 55, 374-376.	3.1	28
1137	Treatment of Staphylococcus aureus endocarditis with high doses of trimethoprim/sulfamethoxazole and clindamycinâ€"Preliminary report. International Journal of Antimicrobial Agents, 2013, 42, 190-191.	2.5	28
1138	Non-contiguous finished genome sequence and description of Megasphaera massiliensis sp. nov Standards in Genomic Sciences, 2013, 8, 525-538.	1.5	28
1139	Emergence of Q Fever Arthritis in France. Journal of Clinical Microbiology, 2014, 52, 1064-1067.	3.9	28
1140	Molecular Detection of Rickettsia felis and Bartonella henselae in Dog and Cat Fleas in Central Oromia, Ethiopia. American Journal of Tropical Medicine and Hygiene, 2014, 90, 457-462.	1.4	28
1141	The genome of Coxiella burnetii Z3055, a clone linked to the Netherlands Q fever outbreaks, provides evidence for the role of drift in the emergence of epidemic clones. Comparative Immunology, Microbiology and Infectious Diseases, 2014, 37, 281-288.	1.6	28
1142	High Prevalence of Mansonella perstans Filariasis in Rural Senegal. American Journal of Tropical Medicine and Hygiene, 2015, 93, 601-606.	1.4	28
1143	Novel Single-Stranded DNA Circular Viruses in Pericardial Fluid of Patient with Recurrent Pericarditis. Emerging Infectious Diseases, 2016, 22, 1839-1841.	4.3	28
1144	The History of Epidemic Typhus. Microbiology Spectrum, 2016, 4, .	3.0	28
1145	Simultaneous UHPLC-UV analysis of hydroxychloroquine, minocycline and doxycycline from serum samples for the therapeutic drug monitoring of Q fever and Whipple's disease. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2017, 1060, 166-172.	2.3	28
1146	Coronary events complicating infective endocarditis. Heart, 2017, 103, 1906-1910.	2.9	28
1147	Coxiella burnetii in Infertile Dairy Cattle With Chronic Endometritis. Veterinary Pathology, 2018, 55, 539-542.	1.7	28
1148	Repertoire of bacterial species cultured from the human oral cavity and respiratory tract. Future Microbiology, 2018, 13, 1611-1624.	2.0	28
1149	Respiratory tract infections among French Hajj pilgrims from 2014 to 2017. Scientific Reports, 2019, 9, 17771.	3.3	28
1150	Taxonomic Relationships among Spotted Fever Group Rickettsiae as Revealed by Antigenic Analysis with Monoclonal Antibodies. Journal of Clinical Microbiology, 1998, 36, 887-896.	3.9	28
1151	Third Human Isolate of a Desulfovibrio sp. Identical to the Provisionally Named Desulfovibrio fairfieldensis. Journal of Clinical Microbiology, 1999, 37, 3076-3077.	3.9	28
1152	Multispacer Sequence Typing for Mycobacterium tuberculosis Genotyping. PLoS ONE, 2008, 3, e2433.	2.5	28

#	Article	IF	Citations
1153	Point of Care Strategy for Rapid Diagnosis of Novel A/H1N1 Influenza Virus. PLoS ONE, 2010, 5, e9215.	2.5	28
1154	Classification of Ancient Mammal Individuals Using Dental Pulp MALDI-TOF MS Peptide Profiling. PLoS ONE, 2011, 6, e17319.	2.5	28
1155	<i>Inquilinus limosus</i> and Cystic Fibrosis. Emerging Infectious Diseases, 2008, 14, 993-995.	4.3	28
1156	Seroepidemiology of Rickettsia africae Infection in Norwegian Travellers to Rural Africa. Scandinavian Journal of Infectious Diseases, 2002, 34, 93-96.	1.5	27
1157	Comparison of Immunofluorescence, Western Blotting, and Cross-Adsorption Assays for Diagnosis of African Tick Bite Fever. Vaccine Journal, 2004, 11, 786-788.	3.1	27
1158	Lack of microbicidal response in human macrophages infected with Parachlamydia acanthamoebae. Microbes and Infection, 2005, 7, 714-719.	1.9	27
1159	Activity of Telithromycin against Thirteen New Isolates of C. burnetii Including Three Resistant to Doxycycline. Annals of the New York Academy of Sciences, 2005, 1063, 252-256.	3.8	27
1160	Molecular Screening of Bartonella Species in Rodents from the Russian Far East. Annals of the New York Academy of Sciences, 2005, 1063, 308-311.	3.8	27
1161	Bartonella bovis in cattle in Africa. Veterinary Microbiology, 2005, 105, 155-156.	1.9	27
1162	" <i>Candidatus Rickettsia kellyi</i> ," India. Emerging Infectious Diseases, 2006, 12, 483-485.	4.3	27
1163	From cat scratch disease to endocarditis, the possible natural history of Bartonella henselaeinfection. BMC Infectious Diseases, 2007, 7, 30.	2.9	27
1164	Actinomyces massiliensis sp. nov., isolated from a patient blood culture. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 540-544.	1.7	27
1165	Revised Mimivirus major capsid protein sequence reveals intron-containing gene structure and extra domain. BMC Molecular Biology, 2009, 10, 39.	3.0	27
1166	Rickettsia slovaca from Dermacentor marginatus ticks in Sardinia, Italy. Ticks and Tick-borne Diseases, 2012, 3, 393-395.	2.7	27
1167	The Genealogic Tree of Mycobacteria Reveals a Long-Standing Sympatric Life into Free-Living Protozoa. PLoS ONE, 2012, 7, e34754.	2.5	27
1168	Monocyte Responses in the Context of Q Fever: From a Static Polarized Model to a Kinetic Model of Activation. Journal of Infectious Diseases, 2013, 208, 942-951.	4.0	27
1169	Molecular Identification of Pathogenic Bacteria in Eschars from Acute Febrile Patients, Senegal. American Journal of Tropical Medicine and Hygiene, 2014, 91, 1015-1019.	1.4	27
1170	Alphaproteobacteria species as a source and target of lateral sequence transfers. Trends in Microbiology, 2014, 22, 147-156.	7.7	27

#	Article	IF	Citations
1171	Divergent Gemycircularvirus in HIV-Positive Blood, France. Emerging Infectious Diseases, 2015, 21, 2096-2098.	4.3	27
1172	Correlation between Sputum and Bronchoalveolar Lavage Fluid Cultures. Journal of Clinical Microbiology, 2015, 53, 994-996.	3.9	27
1173	Real-time video imaging as a new and rapid tool for antibiotic susceptibility testing by the disc diffusion method: A paradigm for evaluating resistance to imipenem and identifying extended-spectrum $\hat{l}^2$ -lactamases. International Journal of Antimicrobial Agents, 2015, 45, 61-65.	2.5	27
1174	Microvirga massiliensis sp. nov., the human commensal with the largest genome. MicrobiologyOpen, 2016, 5, 307-322.	3.0	27
1175	Molecular Tests That Target the RTX Locus Do Not Distinguish between Kingella kingae and the Recently Described Kingella negevensis Species. Journal of Clinical Microbiology, 2017, 55, 3113-3122.	3.9	27
1176	Flow Cytometry Sorting to Separate Viable Giant Viruses from Amoeba Co-culture Supernatants. Frontiers in Cellular and Infection Microbiology, 2016, 6, 202.	3.9	27
1177	Detection of novel RNA viruses from free-living gorillas, Republic of the Congo: genetic diversity of picobirnaviruses. Virus Genes, 2018, 54, 256-271.	1.6	27
1178	A Phylogenomic Study of Acanthamoeba polyphaga Draft Genome Sequences Suggests Genetic Exchanges With Giant Viruses. Frontiers in Microbiology, 2018, 9, 2098.	3.5	27
1179	Implementation of an in-house real-time reverse transcription-PCR assay for the rapid detection of the SARS-CoV-2 Marseille-4 variant. Journal of Clinical Virology, 2021, 139, 104814.	3.1	27
1180	SARS-CoV-2 antibodies seroprevalence in dogs from France using ELISA and an automated western blotting assay. One Health, 2021, 13, 100293.	3.4	27
1181	Host-Associated Metagenomics: A Guide to Generating Infectious RNA Viromes. PLoS ONE, 2015, 10, e0139810.	2.5	27
1182	High prevalence of Bartonella quintana endocarditis in Sfax, Tunisia. American Journal of Tropical Medicine and Hygiene, 2005, 72, 503-7.	1.4	27
1183	Molecular detection of Bartonella quintana, B. Elizabethae, B. Koehlerae, B. Doshiae, B. Taylorii, and Rickettsia felis in rodent fleas collected in Kabul, Afghanistan. American Journal of Tropical Medicine and Hygiene, 2006, 74, 436-9.	1.4	27
1184	Bartonella infections. Current Opinion in Infectious Diseases, 1998, 11, 189-194.	3.1	26
1185	Bartonella and Coxiella Antibodies in 334 Prospectively Studied Episodes of Infective Endocarditis in Sweden. Scandinavian Journal of Infectious Diseases, 2003, 35, 724-727.	1.5	26
1186	Reemergence of Rickettsiosis in Oran, Algeria. Annals of the New York Academy of Sciences, 2006, 1078, 180-184.	3.8	26
1187	SVARAP and aSVARAP: simple tools for quantitative analysis of nucleotide and amino acid variability and primer selection for clinical microbiology. BMC Microbiology, 2006, 6, 21.	3.3	26
1188	Serological microarray for a paradoxical diagnostic of Whipple's disease. European Journal of Clinical Microbiology and Infectious Diseases, 2008, 27, 959-968.	2.9	26

#	Article	IF	CITATIONS
1189	Fatal Case of Israeli Spotted Fever after Mediterranean Cruise. Emerging Infectious Diseases, 2008, 14, 1944-1946.	4.3	26
1190	Gordonia sputiBacteremia. Emerging Infectious Diseases, 2009, 15, 1535-1537.	4.3	26
1191	"Candidatus Bartonella thailandensisâ€! A new genotype of Bartonella identified from rodents. Veterinary Microbiology, 2009, 139, 197-201.	1.9	26
1192	Old and new tick-borne rickettsioses. International Health, 2009, 1, 17-25.	2.0	26
1193	Seasonality of Cat-Scratch Disease, France, 1999–2009. Emerging Infectious Diseases, 2011, 17, 705-707.	4.3	26
1194	Spotted fever group rickettsiae in ticks and fleas from the Democratic Republic of the Congo. Ticks and Tick-borne Diseases, 2012, 3, 371-373.	2.7	26
1195	The use of eschar swabs for the diagnosis of African tick-bite fever. Ticks and Tick-borne Diseases, 2012, 3, 361-363.	2.7	26
1196	Diplorickettsia massiliensis as a human pathogen. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 365-369.	2.9	26
1197	Non-contiguous finished genome sequence and description of Bartonella florenciae sp. nov Standards in Genomic Sciences, 2013, 9, 185-196.	1.5	26
1198	Non contiguous-finished genome sequence and description of Clostridium jeddahense sp. nov Standards in Genomic Sciences, 2014, 9, 1003-1019.	1.5	26
1199	Faustovirus-Like Asfarvirus in Hematophagous Biting Midges and Their Vertebrate Hosts. Frontiers in Microbiology, 2015, 6, 1406.	3 <b>.</b> 5	26
1200	Mouse Model of Coxiella burnetii Aerosolization. Infection and Immunity, 2016, 84, 2116-2123.	2.2	26
1201	Evidence of Bartonella spp. in Blood and Ticks (Ornithodoros hasei) of Bats, in French Guiana. Vector-Borne and Zoonotic Diseases, 2016, 16, 516-519.	1.5	26
1202	Synergistic activity of antibiotics combined with ivermectin to kill body lice. International Journal of Antimicrobial Agents, 2016, 47, 217-223.	2.5	26
1203	Malaria in Dielmo, a Senegal village: Is its elimination possible after seven years of implementation of long-lasting insecticide-treated nets?. PLoS ONE, 2017, 12, e0179528.	2.5	26
1204	Detection of a Potential New Bartonella Species "Candidatus Bartonella rondoniensis―in Human Biting Kissing Bugs (Reduviidae; Triatominae). PLoS Neglected Tropical Diseases, 2017, 11, e0005297.	3.0	26
1205	Detection of bacterial pathogens in clade E head lice collected from Niger's refugees in Algeria. Parasites and Vectors, 2018, 11, 348.	2.5	26
1206	Isolation and culture of Methanobrevibacter smithii by co-culture with hydrogen-producing bacteria on agar plates. Clinical Microbiology and Infection, 2019, 25, 1561.e1-1561.e5.	6.0	26

#	Article	IF	CITATIONS
1207	Detection of Methanobrevibacter smithii in vaginal samples collected from women diagnosed with bacterial vaginosis. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 1643-1649.	2.9	26
1208	Prospective case-control analysis of the aetiologies of acute undifferentiated fever in Vietnam. Emerging Microbes and Infections, 2019, 8, 339-352.	6.5	26
1209	Emerging methodologies for pathogen identification in bloodstream infections: an update. Expert Review of Molecular Diagnostics, 2019, 19, 161-173.	3.1	26
1210	The Antioxidants Glutathione, Ascorbic Acid and Uric Acid Maintain Butyrate Production by Human Gut Clostridia in The Presence of Oxygen In Vitro. Scientific Reports, 2020, 10, 7705.	3.3	26
1211	Role of plants in the transmission of Asaia sp., which potentially inhibit the Plasmodium sporogenic cycle in Anopheles mosquitoes. Scientific Reports, 2020, 10, 7144.	3.3	26
1212	Detection of relapsing fever Borrelia spp., Bartonella spp. and Anaplasmataceae bacteria in argasid ticks in Algeria. PLoS Neglected Tropical Diseases, 2017, 11, e0006064.	3.0	26
1213	Origin and Evolution of Rickettsial Plasmids. PLoS ONE, 2016, 11, e0147492.	2.5	26
1214	High rate of reinfection with the SARS-CoV-2 Omicron variant. Journal of Infection, 2022, 85, 174-211.	3.3	26
1215	Rickettsial infections - a threat to travellers?. Current Opinion in Infectious Diseases, 2004, 17, 433-437.	3.1	25
1216	Advances in Rickettsia Pathogenicity. Annals of the New York Academy of Sciences, 2009, 1166, 94-105.	3.8	25
1217	Novel Virus Influenza A (H1N1sw) in South-Eastern France, April-August 2009. PLoS ONE, 2010, 5, e9214.	2.5	25
1218	The rhizome of Reclinomonas americana, Homo sapiens, Pediculus humanus and Saccharomyces cerevisiae mitochondria. Biology Direct, 2011, 6, 55.	4.6	25
1219	Relapsing fever Borrelia inOrnithodorosticks from Bolivia. Annals of Tropical Medicine and Parasitology, 2011, 105, 407-411.	1.6	25
1220	Detection of Rickettsioses and Q fever in Sri Lanka. American Journal of Tropical Medicine and Hygiene, 2012, 86, 711-712.	1.4	25
1221	The first molecular detection of Rickettsia aeschlimannii in the ticks of camels from southern Algeria. Ticks and Tick-borne Diseases, 2012, 3, 374-376.	2.7	25
1222	Immuno-PCR - A New Tool for Paleomicrobiology: The Plague Paradigm. PLoS ONE, 2012, 7, e31744.	2.5	25
1223	Molecular detection of rickettsial agents in ticks and fleas collected from a European hedgehog (Erinaceus europaeus) in Marseilles, France. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 77-79.	1.6	25
1224	Q fever osteomyelitis: A case report and literature review. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 169-172.	1.6	25

#	Article	IF	Citations
1225	Amoebae as Battlefields for Bacteria, Giant Viruses, and Virophages. Journal of Virology, 2013, 87, 4783-4785.	3.4	25
1226	A Case of Q Fever Prosthetic Joint Infection and Description of an Assay for Detection of Coxiella burnetii. Journal of Clinical Microbiology, 2013, 51, 66-69.	3.9	25
1227	Non-contiguous finished genome sequence and description of Holdemania massiliensis sp. nov Standards in Genomic Sciences, 2013, 9, 395-409.	1.5	25
1228	Possible Role of <i>Rickettsia felis</i> in Acute Febrile Illness among Children in Gabon. Emerging Infectious Diseases, 2015, 21, 1808-1815.	4.3	25
1229	Spotted fever group rickettsiae in ixodid ticks in Oromia, Ethiopia. Ticks and Tick-borne Diseases, 2015, 6, 8-15.	2.7	25
1230	Wild Gorillas as a Potential Reservoir of <i>Leishmania major </i> . Journal of Infectious Diseases, 2015, 211, 267-273.	4.0	25
1231	Treatment and Prophylactic Strategy for Coxiella burnetii Infection of Aneurysms and Vascular Grafts. Medicine (United States), 2016, 95, e2810.	1.0	25
1232	Sennetsu Neorickettsiosis, Spotted Fever Group, and Typhus Group Rickettsioses in Three Provinces in Thailand. American Journal of Tropical Medicine and Hygiene, 2016, 95, 43-49.	1.4	25
1233	Risk factors for acquisition of CTX-M genes in pilgrims during Hajj 2013 and 2014. Journal of Antimicrobial Chemotherapy, 2017, 72, 2627-2635.	3.0	25
1234	Use of eschar swabbing for the molecular diagnosis and genotyping of Orientia tsutsugamushi causing scrub typhus in Quang Nam province, Vietnam. PLoS Neglected Tropical Diseases, 2017, 11, e0005397.	3.0	25
1235	Molecular Survey of Head and Body Lice, <i>Pediculus humanus </i> , in France. Vector-Borne and Zoonotic Diseases, 2018, 18, 243-251.	1.5	25
1236	Field application of MALDI-TOF MS on mosquito larvae identification. Parasitology, 2018, 145, 677-687.	1.5	25
1237	The dynamics and interactions of respiratory pathogen carriage among French pilgrims during the 2018 Hajj. Emerging Microbes and Infections, 2019, 8, 1701-1710.	6.5	25
1238	COVID-19 pandemic more than a century after the Spanish flu. Lancet Infectious Diseases, The, 2021, 21, e78.	9.1	25
1239	Rickettsia gravesii sp. nov.: a novel spotted fever group rickettsia in Western Australian Amblyomma triguttatum triguttatum ticks. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3156-3161.	1.7	25
1240	Q Fever in French Guiana: Tip of the Iceberg or Epidemiological Exception?. PLoS Neglected Tropical Diseases, 2016, 10, e0004598.	3.0	25
1241	Drug repurposing against SARS-CoV-1, SARS-CoV-2Âand MERS-CoV. Future Microbiology, 2021, 16, 1341-1370.	2.0	25
1242	Update on Q fever, including Q fever endocarditis. Current Clinical Topics in Infectious Diseases, 2002, 22, 97-124.	0.3	25

#	Article	IF	CITATIONS
1243	Genomic evolution and adaptation of arthropod-associated Rickettsia. Scientific Reports, 2022, 12, 3807.	3.3	25
1244	Psoriasis is a risk factor for hip-prosthesis infection. European Journal of Epidemiology, 1997, 13, 205-207.	5.7	24
1245	Coxiella burnetii vascular graft infection. BMC Infectious Diseases, 2005, 5, 109.	2.9	24
1246	<i>Rickettsia slovaca</i> li>Infection, France. Emerging Infectious Diseases, 2006, 12, 521-523.	4.3	24
1247	Molecular Characterization of Resistance to Macrolides in Bartonella henselae. Antimicrobial Agents and Chemotherapy, 2006, 50, 3192-3193.	3.2	24
1248	Quantitative Histological Examination of Bioprosthetic Heart Valves. Clinical Infectious Diseases, 2006, 42, 590-596.	5.8	24
1249	Endocarditis Caused by Cardiobacterium valvarum. Journal of Clinical Microbiology, 2006, 44, 657-658.	3.9	24
1250	<i>Bartonella australis</i> sp. nov. from Kangaroos, Australia. Emerging Infectious Diseases, 2007, 13, 1961-1963.	4.3	24
1251	Actinomyces timonensis sp. nov., isolated from a human clinical osteo-articular sample. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 1516-1521.	1.7	24
1252	New Rural Focus of Plague, Algeria. Emerging Infectious Diseases, 2010, 16, 1639-1640.	4.3	24
1253	Emergence of <i>Rickettsia africae </i> , Oceania. Emerging Infectious Diseases, 2011, 17, 100-102.	4.3	24
1254	Totally resistant tuberculosis: will antileprosy drugs be helpful?. International Journal of Antimicrobial Agents, 2013, 42, 584-585.	2.5	24
1255	Real-Time PCR Systems Targeting Giant Viruses of Amoebae and Their Virophages. Intervirology, 2013, 56, 413-423.	2.8	24
1256	Non-contiguous finished genome sequence and description of Bacillus massilioanorexius sp. nov Standards in Genomic Sciences, 2013, 8, 465-479.	1.5	24
1257	Clustered Cases of <i>Rickettsia sibirica mongolitimonae </i> Diseases, 2013, 19, 337-338.	4.3	24
1258	Imbalance of Circulating Monocyte Subsets and PD-1 Dysregulation in Q Fever Endocarditis: The Role of IL-10 in PD-1 Modulation. PLoS ONE, 2014, 9, e107533.	2.5	24
1259	Occurrence of Tropheryma whipplei during diarrhea in Hajj pilgrims: A PCR analysis of paired rectal swabs. Travel Medicine and Infectious Disease, 2014, 12, 481-484.	3.0	24
1260	Welcome to pandoraviruses at the ââ,¬ËœFourth TRUCââ,¬â,,¢ club. Frontiers in Microbiology, 2015, 6, 423.	3.5	24

#	Article	IF	Citations
1261	Non contiguous-finished genome sequence and description of Microbacterium gorillae sp. nov Standards in Genomic Sciences, 2016, 11, 32.	1.5	24
1262	Tropheryma whipplei DNA in bronchoalveolar lavage samples: a case control study. Clinical Microbiology and Infection, 2016, 22, 875-879.	6.0	24
1263	Molecular survey of Dirofilaria immitis and Dirofilaria repens by new real-time TaqMan® PCR assay in dogs and mosquitoes (Diptera: Culicidae) in Corsica (France). Veterinary Parasitology, 2017, 235, 1-7.	1.8	24
1264	Acute infections caused by <i>Tropheryma whipplei</i> . Future Microbiology, 2017, 12, 247-254.	2.0	24
1265	Rickettsial genomics and the paradigm of genome reduction associated with increased virulence. Microbes and Infection, 2018, 20, 401-409.	1.9	24
1266	Attributable deaths caused by infections with antibiotic-resistant bacteria in France. Lancet Infectious Diseases, The, 2019, 19, 128-129.	9.1	24
1267	Respiratory and gastrointestinal infections at the 2017 Grand Magal de Touba, Senegal: A prospective cohort survey. Travel Medicine and Infectious Disease, 2019, 32, 101410.	3.0	24
1268	Mitochondrial diversity and phylogeographic analysis of Pediculus humanus reveals a new Amazonian clade "F― Infection, Genetics and Evolution, 2019, 70, 1-8.	2.3	24
1269	Flying Fox Hemolytic Fever, Description of a New Zoonosis Caused by <i>Candidatus</i> Mycoplasma haemohominis. Clinical Infectious Diseases, 2021, 73, e1445-e1453.	5.8	24
1270	Evidence for an African Cluster of Human Head and Body Lice with Variable Colors and Interbreeding of Lice between Continents. PLoS ONE, 2012, 7, e37804.	2.5	24
1271	Evidence That Head and Body Lice on Homeless Persons Have the Same Genotype. PLoS ONE, 2012, 7, e45903.	2.5	24
1272	Tobacco Mosaic Virus in the Lungs of Mice following Intra-Tracheal Inoculation. PLoS ONE, 2013, 8, e54993.	2.5	24
1273	Rhizome of life, catastrophes, sequence exchanges, gene creations, and giant viruses: how microbial genomics challenges Darwin. Frontiers in Cellular and Infection Microbiology, 2012, 2, 113.	3.9	24
1274	Coxiella burnetii in Dromedary Camels (Camelus dromedarius): A Possible Threat for Humans and Livestock in North Africa and the Near and Middle East?. Frontiers in Veterinary Science, 2020, 7, 558481.	2.2	24
1275	Seroepidemiology of Rickettsia typhi, Spotted Fever Group Rickettsiae, and Coxiella burnetti Infection in Pregnant Women from Urban Tanzania. American Journal of Tropical Medicine and Hygiene, 1997, 57, 187-189.	1.4	24
1276	Non contiguous-finished genome sequence and description of Peptoniphilus obesi sp. nov. Standards in Genomic Sciences, 2013, 7, 357-69.	1.5	24
1277	Lyme disease presenting as isolated acute urinary retention caused by transverse myelitis: An electrophysiological and urodynamical study. Archives of Physical Medicine and Rehabilitation, 1995, 76, 1171-1172.	0.9	23
1278	Domestic cats as indicators of the presence of spotted fever and typhus group rickettsiae. European Journal of Epidemiology, 1997, 13, 109-111.	5.7	23

#	Article	IF	Citations
1279	Determination of Coxiella burnetii rpoB sequence and its use for phylogenetic analysis. Gene, 1998, 207, 97-103.	2.2	23
1280	Doxycycline and Eradication of Microfilaremia in Patients with Loiasis. Emerging Infectious Diseases, 2001, 7, 604-605.	4.3	23
1281	Rhodobacter massiliensis sp. nov., a new amoebae-resistant species isolated from the nose of a patient. Research in Microbiology, 2003, 154, 631-635.	2.1	23
1282	First Isolation and Detection by Immunofluorescence Assay of Bartonella koehlerae in Erythrocytes from a French Cat. Journal of Clinical Microbiology, 2003, 41, 4001-4002.	3.9	23
1283	Mycobacterium barrassiae sp. nov., a Mycobacterium moriokaense Group Species Associated with Chronic Pneumonia. Journal of Clinical Microbiology, 2006, 44, 3493-3498.	3.9	23
1284	Identification of Rickettsiae, Uganda and Djibouti. Emerging Infectious Diseases, 2007, 13, 1508-1509.	4.3	23
1285	Corynebacterium timonense sp. nov. and Corynebacterium massiliense sp. nov., isolated from human blood and human articular hip fluid. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 1953-1959.	1.7	23
1286	Helcobacillus massiliensis gen. nov., sp. nov., a novel representative of the family Dermabacteraceae isolated from a patient with a cutaneous discharge. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 2346-2351.	1.7	23
1287	Characterization of rickettsial adhesin Adr2 belonging to a new group of adhesins in $\hat{l}\pm$ -proteobacteria. Microbial Pathogenesis, 2011, 50, 233-242.	2.9	23
1288	A deadly aversion to pork. Lancet, The, 2011, 377, 1542.	13.7	23
1289	Bartonellae in animals and vectors in New Caledonia. Comparative Immunology, Microbiology and Infectious Diseases, 2011, 34, 497-501.	1.6	23
1290	A case of rickettsialpox in Northern Europe. International Journal of Infectious Diseases, 2012, 16, e221-e222.	3.3	23
1291	Vector-Borne Rickettsioses in North Africa. Infectious Disease Clinics of North America, 2012, 26, 455-478.	5.1	23
1292	Non-contiguous finished genome sequence and description of Kallipyga massiliensis gen. nov., sp. nov., a new member of the family Clostridiales Incertae Sedis XI. Standards in Genomic Sciences, 2013, 8, 500-515.	1.5	23
1293	Non-contiguous finished genome sequence and description of Oceanobacillus massiliensis sp. nov Standards in Genomic Sciences, 2013, 9, 370-384.	1.5	23
1294	<i>Bartonella quintana</i> in Body Lice from Scalp Hair of Homeless Persons, France. Emerging Infectious Diseases, 2014, 20, 907-908.	4.3	23
1295	Identification of giant Mimivirus protein functions using RNA interference. Frontiers in Microbiology, 2015, 6, 345.	3.5	23
1296	Cost-Effective Pooling of DNA from Nasopharyngeal Swab Samples for Large-Scale Detection of Bacteria by Real-Time PCR. Journal of Clinical Microbiology, 2015, 53, 1002-1004.	3.9	23

#	Article	IF	CITATIONS
1297	Screening for Viral Pathogens in African Simian Bushmeat Seized at A French Airport. Transboundary and Emerging Diseases, 2017, 64, 1159-1167.	3.0	23
1298	From Culturomics to Clinical Microbiology and Forward. Emerging Infectious Diseases, 2018, 24, 1683-1690.	4.3	23
1299	Paradoxical evolution of rickettsial genomes. Ticks and Tick-borne Diseases, 2019, 10, 462-469.	2.7	23
1300	The Butyrogenic and Lactic Bacteria of the Gut Microbiota Determine the Outcome of Allogenic Hematopoietic Cell Transplant. Frontiers in Microbiology, 2020, 11, 1642.	3.5	23
1301	An Earliest Endosymbiont, Wolbachia massiliensis sp. nov., Strain PL13 from the Bed Bug (Cimex) Tj ETQq1 1 (8064.	0.784314 rgB <sup>-</sup> 4.1	T /Overlock 23
1302	Development of MALDI-TOF mass spectrometry for the identification of lice isolated from farm animals. Parasite, 2020, 27, 28.	2.0	23
1303	Automated Western immunoblotting detection of anti-SARS-CoV-2 serum antibodies. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 1309-1317.	2.9	23
1304	Infective Endocarditis in Patients on Chronic Hemodialysis. Journal of the American College of Cardiology, 2021, 77, 1629-1640.	2.8	23
1305	Rickettsia conorii Transcriptional Response within Inoculation Eschar. PLoS ONE, 2008, 3, e3681.	2.5	23
1306	Metabarcoding analysis of eukaryotic microbiota in the gut of HIV-infected patients. PLoS ONE, 2018, 13, e0191913.	2.5	23
1307	Performance of Real-Time Polymerase Chain Reaction Assays for the Detection of 20 Gastrointestinal Parasites in Clinical Samples from Senegal. American Journal of Tropical Medicine and Hygiene, 2017, 97, 173-182.	1.4	23
1308	Intracellular organisms. International Journal of Antimicrobial Agents, 1997, 9, 61-70.	2.5	22
1309	Expression of green fluorescent protein in Rickettsia conorii. Microbial Pathogenesis, 2002, 33, 17-21.	2.9	22
1310	Can Whipple's Disease Be Transmitted by Gastroscopes?. Infection Control and Hospital Epidemiology, 2003, 24, 191-194.	1.8	22
1311	Shell Vial Culture as a Tool for Isolation of Brucella melitensis in Chronic Hepatic Abscess. Journal of Clinical Microbiology, 2003, 41, 4460-4461.	3.9	22
1312	Fever of Unknown Origin Due toÂRickettsioses. Infectious Disease Clinics of North America, 2007, 21, 997-1011.	5.1	22
1313	Detection of "Rickettsia sp. strain Uilenbergi" and "Rickettsia sp. strain Davousti" in Amblyomma tholloni ticks from elephants in Africa. BMC Microbiology, 2007, 7, 74.	3.3	22
1314	A murine model of infection with Rickettsia prowazekii: implications for pathogenesis of epidemic typhus. Microbes and Infection, 2007, 9, 898-906.	1.9	22

#	Article	lF	CITATIONS
1315	<i>Sphingomonas mucosissima</i> Bacteremia in Patient with Sickle Cell Disease. Emerging Infectious Diseases, 2009, 15, 133-134.	4.3	22
1316	Hepatitis E Virus Infection in Sheltered Homeless Persons, France. Emerging Infectious Diseases, 2010, 16, 1761-1763.	4.3	22
1317	<i>Rickettsia honei</i> lnfection in Human, Nepal, 2009. Emerging Infectious Diseases, 2011, 17, 1865-1867.	4.3	22
1318	Defective Monocyte Dynamics in Q Fever Granuloma Deficiency. Journal of Infectious Diseases, 2012, 205, 1086-1094.	4.0	22
1319	<i>Legionella longbeachae</i> and Endocarditis. Emerging Infectious Diseases, 2012, 18, 95-97.	4.3	22
1320	Immuno-PCR for the early serological diagnosis of acute infectious diseases: the Q fever paradigm. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 1951-1960.	2.9	22
1321	The Role of the Manipulation of the Gut Microbiota in Obesity. Current Infectious Disease Reports, 2013, 15, 25-30.	3.0	22
1322	Persistence of DNA in a Cured Patient and Positive Culture in Cases with Low Antibody Levels Bring into Question Diagnosis of Q Fever Endocarditis. Journal of Clinical Microbiology, 2013, 51, 3012-3017.	3.9	22
1323	<i>Borrelia garinii</i> and <i>Rickettsia monacensis</i> in <i>lxodes ricinus</i> Ticks, Algeria. Emerging Infectious Diseases, 2014, 20, 1776-1777.	4.3	22
1324	Pointâ€ofâ€Care Syndromeâ€Based, Rapid Diagnosis of Infections on Commercial Ships. Journal of Travel Medicine, 2014, 21, 12-16.	3.0	22
1325	Tropheryma whipplei natural resistance to trimethoprim and sulphonamides in vitro. International Journal of Antimicrobial Agents, 2014, 43, 388-390.	2.5	22
1326	Reduction in incidence of Q fever endocarditis: 27 years of experience of a national reference center. Journal of Infection, 2014, 68, 141-148.	3.3	22
1327	Draft genome sequences of Terra1 and Terra2 viruses, new members of the family Mimiviridae isolated from soil. Virology, 2014, 452-453, 125-132.	2.4	22
1328	A severe Whipple disease with an immune reconstitution inflammatory syndrome: An additional case of thalidomide efficiency. Joint Bone Spine, 2014, 81, 260-262.	1.6	22
1329	Rickettsia conorii israelensis in Rhipicephalus sanguineus ticks, Sardinia, Italy. Ticks and Tick-borne Diseases, 2014, 5, 446-448.	2.7	22
1330	Non contiguous-finished genome sequence and description of Bacillus jeddahensis sp. nov Standards in Genomic Sciences, 2015, 10, 47.	1.5	22
1331	Assessment of the in vitro antimicrobial activity of Lactobacillus species for identifying new potential antibiotics. International Journal of Antimicrobial Agents, 2015, 46, 590-593.	2,5	22
1332	Protochlamydia phocaeensis sp. nov., a new Chlamydiales species with host dependent replication cycle. Microbes and Infection, 2017, 19, 343-350.	1.9	22

#	Article	IF	CITATIONS
1333	Medical Entomology: A Reemerging Field of Research to Better Understand Vector-Borne Infectious Diseases. Clinical Infectious Diseases, 2017, 65, S30-S38.	5.8	22
1334	Q fever epidemic in Cayenne, French Guiana, epidemiologically linked to three-toed sloth. Comparative Immunology, Microbiology and Infectious Diseases, 2018, 56, 34-38.	1.6	22
1335	Bacterial infection and non-Hodgkin's lymphoma. Critical Reviews in Microbiology, 2020, 46, 270-287.	6.1	22
1336	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. Infection, Genetics and Evolution, 2021, 95, 105092.	2.3	22
1337	Genome sequence-based criteria for demarcation and definition of species in the genus Rickettsia. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 1738-1750.	1.7	22
1338	Prevalence of antibodies to Rickettsia conorii Ricketsia africae, Rickettsia typhi and Coxiella burnetii in Mauritania. European Journal of Epidemiology, 1998, 14, 817-818.	5.7	21
1339	Whipple's disease. Current Gastroenterology Reports, 2003, 5, 379-385.	2.5	21
1340	Quantitative Histological Examination of Mechanical Heart Valves. Clinical Infectious Diseases, 2005, 40, 655-661.	5.8	21
1341	Development and assessment of a new early scoring system using non-specific clinical signs and biological results to identify children and adult patients with a high probability of infective endocarditis on admission. Journal of Antimicrobial Chemotherapy, 2008, 62, 1434-1440.	3.0	21
1342	Emerging <i>Mycobacteria </i> spp. in Cooling Towers. Emerging Infectious Diseases, 2009, 15, 121-122.	4.3	21
1343	Prediction of rickettsial skin eschars in humans using an experimental guinea pig model. Microbial Pathogenesis, 2009, 47, 128-133.	2.9	21
1344	Tropheryma whipplei in the skin of patients with classic Whipple's disease. Journal of Infection, 2010, 61, 266-269.	3.3	21
1345	Epidemiologic Implications of the First Isolation and Cultivation of <i>Tropheryma whipplei</i> From a Saliva Sample. Annals of Internal Medicine, 2011, 154, 443.	3.9	21
1346	Brill-Zinsser Disease in Moroccan Man, France, 2011. Emerging Infectious Diseases, 2012, 18, 171-172.	4.3	21
1347	The combination of chloroquine and minocycline, a therapeutic option in cerebrospinal infection of Whipple's disease refractory to treatment with ceftriaxone, meropenem and co-trimoxazole. Journal of Antimicrobial Chemotherapy, 2012, 67, 1295-1296.	3.0	21
1348	Ehrlichia canis in Rhipicephalus sanguineus ticks in the Ivory Coast. Ticks and Tick-borne Diseases, 2012, 3, 411-413.	2.7	21
1349	<i>Acanthamoeba polyphaga mimivirus</i> Virophage Seroconversion in Travelers Returning from Laos. Emerging Infectious Diseases, 2012, 18, 1500-1502.	4.3	21
1350	Rickettsiae in arthropods collected from red foxes (Vulpes vulpes) in France. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 59-62.	1.6	21

#	Article	IF	Citations
1351	A multi-gene analysis of diversity of bartonella detected in fleas from algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 71-76.	1.6	21
1352	Species and strain specificity of Lactobacillus probiotics effect on weight regulation. Microbial Pathogenesis, 2013, 55, 52-54.	2.9	21
1353	Link between endocarditis on porcine bioprosthetic valves and allergy to pork. International Journal of Cardiology, 2013, 167, 600-602.	1.7	21
1354	<i>Bordetella hinzii</i> in Rodents, Southeast Asia. Emerging Infectious Diseases, 2013, 19, 502-503.	4.3	21
1355	Cell Extract-Containing Medium for Culture of Intracellular Fastidious Bacteria. Journal of Clinical Microbiology, 2013, 51, 2599-2607.	3.9	21
1356	To Tree or Not to Tree? Genome-Wide Quantification of Recombination and Reticulate Evolution during the Diversification of Strict Intracellular Bacteria. Genome Biology and Evolution, 2013, 5, 2305-2317.	2.5	21
1357	Complete genome sequence of Courdo11 virus, a member of the family Mimiviridae. Virus Genes, 2014, 48, 218-223.	1.6	21
1358	Identification of rickettsial pathogens in ixodid ticks in northern Senegal. Ticks and Tick-borne Diseases, 2014, 5, 552-556.	2.7	21
1359	Non-contiguous finished genome sequence and description of Fenollaria massiliensis gen. nov., sp. nov., a new genus of anaerobic bacterium. Standards in Genomic Sciences, 2014, 9, 704-717.	1.5	21
1360	EPIMIC: A Simple Homemade Computer Program for Real-Time EPIdemiological Surveillance and Alert Based on MICrobiological Data. PLoS ONE, 2015, 10, e0144178.	2.5	21
1361	Evidence that clade <scp>A</scp> and clade <scp>B</scp> head lice live in sympatry and recombine in <scp>A</scp> lgeria. Medical and Veterinary Entomology, 2015, 29, 94-98.	1.5	21
1362	The Rhizome of Lokiarchaeota Illustrates the Mosaicity of Archaeal Genomes. Genome Biology and Evolution, 2017, 9, 2635-2639.	2.5	21
1363	Molecular strategy for the diagnosis of infectious lymphadenitis. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 1179-1186.	2.9	21
1364	Acquisition of enteric pathogens by pilgrims during the 2016 Hajj pilgrimage: A prospective cohort study. Travel Medicine and Infectious Disease, 2018, 25, 26-30.	3.0	21
1365	Fluorescence <i>In Situ</i> Hybridization (FISH) and Peptide Nucleic Acid Probe-Based FISH for Diagnosis of Q Fever Endocarditis and Vascular Infections. Journal of Clinical Microbiology, 2018, 56, .	3.9	21
1366	A New Highly Sensitive and Specific Real-Time PCR Assay Targeting the Malate Dehydrogenase Gene of Kingella kingae and Application to 201 Pediatric Clinical Specimens. Journal of Clinical Microbiology, 2018, 56, .	3.9	21
1367	Antibiotic use for respiratory infections among Hajj pilgrims: A cohort survey and review of the literature. Travel Medicine and Infectious Disease, 2019, 30, 39-45.	3.0	21
1368	Genetic diversity of human head lice and molecular detection of associated bacterial pathogens in Democratic Republic of Congo. Parasites and Vectors, 2019, 12, 290.	2.5	21

#	Article	IF	Citations
1369	Rapid MALDI-TOF MS identification of commercial truffles. Scientific Reports, 2019, 9, 17686.	3.3	21
1370	COVID-19 Therapeutic and Prevention. International Journal of Antimicrobial Agents, 2020, 55, 105937.	2.5	21
1371	Early combination therapy with hydroxychloroquine and azithromycin reduces mortality in 10,429 COVID-19 outpatients. Reviews in Cardiovascular Medicine, 2021, 22, 1063.	1.4	21
1372	Cat Scratch Disease Due to Bartonella henselae Serotype Marseille (Swiss Cat) in a Seronegative Patient. Journal of Clinical Microbiology, 1998, 36, 2800-2800.	3.9	21
1373	Fulminant Myocardial Failure in a Previously Healthy Young Man. Circulation, 1997, 95, 1654-1657.	1.6	21
1374	Evidence of Sympatry of Clade A and Clade B Head Lice in a Pre-Columbian Chilean Mummy from Camarones. PLoS ONE, 2013, 8, e76818.	2.5	21
1375	Viral Communities Associated with Human Pericardial Fluids in Idiopathic Pericarditis. PLoS ONE, 2014, 9, e93367.	2.5	21
1376	AUTOCHTHONOUS EPIDEMIC TYPHUS ASSOCIATED WITH BARTONELLA QUINTANA BACTEREMIA IN A HOMELESS PERSON. American Journal of Tropical Medicine and Hygiene, 2005, 72, 638-639.	1.4	21
1377	High Individual Heterogeneity of Neutralizing Activities against the Original Strain and Nine Different Variants of SARS-CoV-2. Viruses, 2021, 13, 2177.	3.3	21
1378	Emerging Bartonella in humans and animals in Asia and Australia. Journal of the Medical Association of Thailand = Chotmaihet Thangphaet, 2009, 92, 707-31.	0.1	21
1379	Exposure of cats in southern Africa to Coxiella burnetii, the agent of Q fever. European Journal of Epidemiology, 1997, 13, 477-479.	5.7	20
1380	A novel alpha-Proteobacterium, Nordella oligomobilis gen. nov., sp. nov., isolated by using amoebal co-cultures. Research in Microbiology, 2004, 155, 47-51.	2.1	20
1381	<i>Rickettsia prowazekii</i> and Real-time Polymerase Chain Reaction. Emerging Infectious Diseases, 2006, 12, 428-432.	4.3	20
1382	Reemergence of Q Fever after 11 September 2001. Clinical Infectious Diseases, 2009, 48, 558-559.	5.8	20
1383	In vitro antibacterial activity of aminosterols against multidrug-resistant bacteria from patients with cystic fibrosis. Journal of Antimicrobial Chemotherapy, 2009, 64, 810-814.	3.0	20
1384	Squalamine ointment for Staphylococcus aureus skin decolonization in a mouse model. Journal of Antimicrobial Chemotherapy, 2011, 66, 1306-1310.	3.0	20
1385	An Experimental Mouse Model to Establish Tropheryma whipplei as a Diarrheal Agent. Journal of Infectious Diseases, 2011, 204, 44-50.	4.0	20
1386	First detection of Ehrlichia canis in Rhipicephalus bursa ticks in Sardinia, Italy. Ticks and Tick-borne Diseases, 2012, 3, 396-397.	2.7	20

#	Article	IF	CITATIONS
1387	The rhizome of life: what about metazoa?. Frontiers in Cellular and Infection Microbiology, 2012, 2, 50.	3.9	20
1388	Pneumonia pathogen detection and microbial interactions in polymicrobial episodes. Future Microbiology, 2013, 8, 633-660.	2.0	20
1389	Non-contiguous finished genome sequence and description of Bartonella senegalensis sp. nov Standards in Genomic Sciences, 2013, 8, 279-289.	1.5	20
1390	Granulomatous response to Coxiella burnetii, the agent of Q fever: the lessons from gene expression analysis. Frontiers in Cellular and Infection Microbiology, 2014, 4, 172.	3.9	20
1391	Pithovirus sibericum, a new bona fide member of the "Fourth TRUC―club. Frontiers in Microbiology, 2015, 6, 722.	3.5	20
1392	A New Zamilon-like Virophage Partial Genome Assembled from a Bioreactor Metagenome. Frontiers in Microbiology, 2015, 6, 1308.	3.5	20
1393	Malaria Risk Factors in Dielmo, A Senegalese Malaria-Endemic Village, Between October and November of 2013: A Case-Control Study. American Journal of Tropical Medicine and Hygiene, 2015, 92, 565-568.	1.4	20
1394	The role of giant viruses of amoebas in humans. Current Opinion in Microbiology, 2016, 31, 199-208.	5.1	20
1395	Bacterial agents in 248 ticks removed from people from 2002 to 2013. Ticks and Tick-borne Diseases, 2016, 7, 475-481.	2.7	20
1396	Classic Whipple's disease diagnosed by 18 F-fluorodeoxyglucose PET. Lancet Infectious Diseases, The, 2016, 16, 130.	9.1	20
1397	Many More Microbes in Humans: Enlarging the Microbiome Repertoire. Clinical Infectious Diseases, 2017, 65, S20-S29.	5.8	20
1398	Unexpected invasion of miniature inverted-repeat transposable elements in viral genomes. Mobile DNA, 2018, 9, 19.	3.6	20
1399	Successful Fecal Microbiota Transplantation in a Patient Suffering From Irritable Bowel Syndrome and Recurrent Urinary Tract Infections. Open Forum Infectious Diseases, 2019, 6, ofz398.	0.9	20
1400	Adenovirus Infections in African Humans and Wild Non-Human Primates: Great Diversity and Cross-Species Transmission. Viruses, 2020, 12, 657.	3.3	20
1401	Effect of hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial, an update with an intention-to-treat analysis and clinical outcomes. International Journal of Antimicrobial Agents, 2021, 57, 106239.	2.5	20
1402	Running after ghosts: are dead bacteria the dark matter of the human gut microbiota?. Gut Microbes, 2021, 13, 1-12.	9.8	20
1403	Multiple vector-borne pathogens of domestic animals in Egypt. PLoS Neglected Tropical Diseases, 2021, 15, e0009767.	3.0	20
1404	Species-specific BALB/c mouse antibodies to rickettsiae studied by Western blotting. FEMS Microbiology Letters, 1994, 119, 339-344.	1.8	19

#	Article	IF	Citations
1405	Phylogenic Analysis of Rickettsial Patatin-like Protein with Conserved Phospholipase A2 Active Sites. Annals of the New York Academy of Sciences, 2005, 1063, 83-86.	3.8	19
1406	Phylogenetic Study of Rickettsia Species Using Sequences of the Autotransporter Protein-Encoding Gene sca2. Annals of the New York Academy of Sciences, 2005, 1063, 94-99.	3.8	19
1407	Deficient Transendothelial Migration of Leukocytes in Q Fever: The Role Played by Interleukin‶0. Journal of Infectious Diseases, 2006, 194, 365-369.	4.0	19
1408	Comment on: Therapy for Whipple's disease. Journal of Antimicrobial Chemotherapy, 2008, 61, 968-969.	3.0	19
1409	<i>Acetobacter indonesiensis</i> Pneumonia after Lung Transplant. Emerging Infectious Diseases, 2008, 14, 997-998.	4.3	19
1410	Human Case of <i>Atopobium rimae </i> Bacteremia. Emerging Infectious Diseases, 2009, 15, 354-355.	4.3	19
1411	<i>Bartonella henselae</i> in Skin Biopsy Specimens of Patients with Cat-Scratch Disease. Emerging Infectious Diseases, 2010, 16, 1963-1965.	4.3	19
1412	Staphylococcus massiliensis sp. nov., isolated from a human brain abscess. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 1066-1072.	1.7	19
1413	Rickettsia felis, West Indies. Emerging Infectious Diseases, 2010, 16, 570-571.	4.3	19
1414	Review of microarray studies for host–intracellular pathogen interactions. Journal of Microbiological Methods, 2010, 81, 81-95.	1.6	19
1415	Airways colonizations in patients undergoing lung cancer surgeryâ*†. European Journal of Cardio-thoracic Surgery, 2011, 40, 309-19.	1.4	19
1416	Vectorborne diseases in West Africa: geographic distribution and geospatial characteristics. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2013, 107, 273-284.	1.8	19
1417	Detection of a New <i>Borrelia</i> Species in Ticks Taken from Cattle in Southwest Ethiopia. Vector-Borne and Zoonotic Diseases, 2013, 13, 266-269.	1.5	19
1418	Sudden death in patients with infective endocarditis: Findings from a large cohort study. International Journal of Cardiology, 2013, 162, 129-132.	1.7	19
1419	Detection of <i>Rickettsia sibirica mongolitimonae </i> by Using Cutaneous Swab Samples and Quantitative PCR. Emerging Infectious Diseases, 2014, 20, 716-718.	4.3	19
1420	Hand, Foot and Mouth Disease and Kingella kingae Infections. Pediatric Infectious Disease Journal, 2015, 34, 547-548.	2.0	19
1421	New Borrelia species detected in ixodid ticks in Oromia, Ethiopia. Ticks and Tick-borne Diseases, 2015, 6, 401-407.	2.7	19
1422	In vitro activity of â€~old antibiotics' against highly resistant Gram-negative bacteria. International Journal of Antimicrobial Agents, 2015, 46, 718-720.	2.5	19

#	Article	IF	Citations
1423	Blood-Borne Candidatus Borrelia algerica in a Patient with Prolonged Fever in Oran, Algeria. American Journal of Tropical Medicine and Hygiene, 2015, 93, 1070-1073.	1.4	19
1424	Evaluation of the PREVI® Isola automated seeder system compared to reference manual inoculation for antibiotic susceptibility testing by the disk diffusion method. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 1859-1869.	2.9	19
1425	Marseillevirus in the Pharynx of a Patient with Neurologic Disorders. Emerging Infectious Diseases, 2016, 22, 2008-2010.	4.3	19
1426	Rickettsia felis: the next mosquito-borne outbreak?. Lancet Infectious Diseases, The, 2016, 16, 1112-1113.	9.1	19
1427	Thrombosis and antiphospholipid antibody syndrome during acute Q fever. Medicine (United States), 2017, 96, e7578.	1.0	19
1428	Meta-analysis on efficacy of amoxicillin in uncomplicated severe acute malnutrition. Microbial Pathogenesis, 2017, 106, 76-77.	2.9	19
1429	Phylogenomic Analysis of $\hat{l}^2$ -Lactamase in Archaea and Bacteria Enables the Identification of Putative New Members. Genome Biology and Evolution, 2018, 10, 1106-1114.	2.5	19
1430	Metagenomics, culturomics, and the human gut microbiota. Expert Review of Anti-Infective Therapy, 2018, 16, 373-375.	4.4	19
1431	Deciphering viral presences: two novel partial giant viruses detected in marine metagenome and in a mine drainage metagenome. Virology Journal, 2018, 15, 66.	3.4	19
1432	Noncontiguous finished genome sequence and description of <i>Intestinimonas massilien</i> sis sp. nov strain <scp>GD</scp> 2 <sup>T</sup> , the second <i>Intestinimonas</i> species cultured from the human gut. MicrobiologyOpen, 2019, 8, e00621.	3.0	19
1433	Measles, the need for a paradigm shift. European Journal of Epidemiology, 2019, 34, 897-915.	5.7	19
1434	Limitations of diagnostic tests for bacterial infections. Médecine Et Maladies Infectieuses, 2019, 49, 98-101.	5.0	19
1435	<i>Methanobrevibacter smithii</i> Archaemia in Febrile Patients With Bacteremia, Including Those With Endocarditis. Clinical Infectious Diseases, 2021, 73, e2571-e2579.	5 <b>.</b> 8	19
1436	Point-of-care multiplexed diagnosis of meningitis using the FilmArray® ME panel technology. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1573-1580.	2.9	19
1437	Difference between kwashiorkor and marasmus: Comparative meta-analysis of pathogenic characteristics and implications for treatment. Microbial Pathogenesis, 2021, 150, 104702.	2.9	19
1438	Screening of SARS-CoV-2 among homeless people, asylum-seekers and other people living in precarious conditions in Marseille, France, March–April 2020. International Journal of Infectious Diseases, 2021, 105, 1-6.	3.3	19
1439	Q FEVER AND PLASMODIUM FALCIPARUM MALARIA CO-INFECTION IN A PATIENT RETURNING FROM THE COMOROS ARCHIPELAGO. American Journal of Tropical Medicine and Hygiene, 2005, 73, 1028-1030.	1.4	19
1440	First cases of infection with the 21L/BA.2 Omicron variant in Marseille, France. Journal of Medical Virology, 2022, 94, 3421-3430.	5.0	19

#	Article	IF	CITATIONS
1441	Candidate Phyla Radiation, an Underappreciated Division of the Human Microbiome, and Its Impact on Health and Disease. Clinical Microbiology Reviews, 2022, 35, .	13.6	19
1442	Study of the 16S-23S ribosomal DNA internal spacer of Coxiella burnetii. European Journal of Epidemiology, 1997, 13, 471-475.	5.7	18
1443	Species-Specific Monoclonal Antibodies for Rapid Identification of <i>Bartonella quintana </i> Journal, 2000, 7, 21-24.	2.6	18
1444	Detection of Rickettsia prowazekii in Body Lice and Their Feces by Using Monoclonal Antibodies. Journal of Clinical Microbiology, 2002, 40, 3358-3363.	3.9	18
1445	Comparison of intuitive versus systematic strategies for aetiological diagnosis of pericardial effusion. Scandinavian Journal of Infectious Diseases, 2005, 37, 216-220.	1.5	18
1446	Evaluation of Sedimentation Rate, Rheumatoid Factor, C-Reactive Protein, and Tumor Necrosis Factor for the Diagnosis of Infective Endocarditis. Vaccine Journal, 2006, 13, 301-301.	3.1	18
1447	Specific recognition of the major capsid protein of <i>Acanthamoeba polyphaga mimivirus </i> by sera of patients infected by <i>Francisella tularensis </i> . FEMS Microbiology Letters, 2009, 297, 117-123.	1.8	18
1448	How should classic Whipple disease be managed?. Nature Reviews Gastroenterology and Hepatology, 2010, 7, 246-248.	17.8	18
1449	Technology-driven research will dominate hypothesis-driven research: the future of microbiology. Future Microbiology, 2010, 5, 135-137.	2.0	18
1450	Models cannot predict future outbreaks: A/H1N1 virus, the paradigm. European Journal of Epidemiology, 2011, 26, 183-186.	5.7	18
1451	Early Diagnosis of Abscess in Aortic Bioprosthetic Valve by 18F-Fluorodeoxyglucose Positron Emission Tomography-Computed Tomography. Circulation, 2012, 126, e217-20.	1.6	18
1452	Human rickettsioses in the Batna area, eastern Algeria. Ticks and Tick-borne Diseases, 2012, 3, 364-366.	2.7	18
1453	<i>Tropheryma whipplei</i> Pneumonia in a Patient with HIV-2 Infection. American Journal of Respiratory and Critical Care Medicine, 2013, 188, 1036-1037.	5.6	18
1454	Co-Infection with <i>Arsenophonus nasoniae</i> li>and <i>Orientia tsutsugamushi</i> li>in a Traveler. Vector-Borne and Zoonotic Diseases, 2013, 13, 565-571.	1.5	18
1455	Non-contiguous finished genome sequence and description of Nosocomiicoccus massiliensis sp. nov Standards in Genomic Sciences, 2013, 9, 205-219.	1.5	18
1456	Absence of antibodies to Rickettsia spp., Bartonella spp., Ehrlichia spp. and Coxiella burnetii in Tahiti, French Polynesia. BMC Infectious Diseases, 2014, 14, 255.	2.9	18
1457	Genome sequence and description of Nesterenkonia massiliensis sp. nov. strain NP1T. Standards in Genomic Sciences, 2014, 9, 866-882.	1.5	18
1458	Incidental Syphilis Diagnosed by Real-Time PCR Screening of Urine Samples. Journal of Clinical Microbiology, 2015, 53, 3707-3708.	3.9	18

#	Article	IF	CITATIONS
1459	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
1460	High-quality genome sequence and description of Bacillus dielmoensis strain FF4T sp. nov Standards in Genomic Sciences, 2015, 10, 41.	1.5	18
1461	Description of Gabonibacter massiliensis gen. nov., sp. nov., a New Member of the Family Porphyromonadaceae Isolated from the Human Gut Microbiota. Current Microbiology, 2016, 73, 867-877.	2.2	18
1462	Alice's living croquet theory. International Journal of Antimicrobial Agents, 2016, 47, 249.	<b>2.</b> 5	18
1463	Yersinia pestis halotolerance illuminates plague reservoirs. Scientific Reports, 2017, 7, 40022.	3.3	18
1464	<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
1465	Changing Demographics and Prevalence of Body Lice among Homeless Persons, Marseille, France. Emerging Infectious Diseases, 2017, 23, 1894-1897.	4.3	18
1466	MALDI-TOF MS protein profiling for the rapid identification of Chagas disease triatomine vectors and application to the triatomine fauna of French Guiana. Parasitology, 2018, 145, 665-675.	1.5	18
1467	Mutations in GluCl associated with field ivermectin-resistant head lice from Senegal. International Journal of Antimicrobial Agents, 2018, 52, 593-598.	2.5	18
1468	High Concentrations of Serum Soluble E-Cadherin in Patients With Q Fever. Frontiers in Cellular and Infection Microbiology, 2019, 9, 219.	3.9	18
1469	Direct Identification of Pathogens in Urine by Use of a Specific Matrix-Assisted Laser Desorption Ionization–Time of Flight Spectrum Database. Journal of Clinical Microbiology, 2019, 57, .	3.9	18
1470	Atypical Cowpox Virus Infection in Smallpox-Vaccinated Patient, France. Emerging Infectious Diseases, 2019, 25, 212-219.	4.3	18
1471	The role of louse-transmitted diseases in historical plague pandemics. Lancet Infectious Diseases, The, 2021, 21, e17-e25.	9.1	18
1472	Clinical outcomes in COVID-19 patients infected with different SARS-CoV-2 variants in Marseille, France. Clinical Microbiology and Infection, 2021, 27, 1516.e1-1516.e6.	6.0	18
1473	Multifaceted modes of action of the anticancer probiotic Enterococcus hirae. Cell Death and Differentiation, 2021, 28, 2276-2295.	11.2	18
1474	A Possible Role of Remdesivir and Plasma Therapy in the Selective Sweep and Emergence of New SARS-CoV-2 Variants. Journal of Clinical Medicine, 2021, 10, 3276.	2.4	18
1475	A New Method to Extract Dental Pulp DNA: Application to Universal Detection of Bacteria. PLoS ONE, 2007, 2, e1062.	2.5	18
1476	Circulating Matrix Metalloproteinases in Infective Endocarditis: A Possible Marker of the Embolic Risk. PLoS ONE, 2011, 6, e18830.	2.5	18

#	Article	IF	CITATIONS
1477	Molecular Detection of Microorganisms Associated with Small Mammals and Their Ectoparasites in Mali. American Journal of Tropical Medicine and Hygiene, 2020, 103, 2542-2551.	1.4	18
1478	Urinary microbiota and bladder cancer: A systematic review and a focus on uropathogens. Seminars in Cancer Biology, 2022, 86, 875-884.	9.6	18
1479	The severity of the first 207 infections with the SARSâ€CoVâ€2 Omicron BA.2 variant, in Marseille, France, December 2021–FebruaryÂ2022. Journal of Medical Virology, 2022, 94, 3494-3497.	5.0	18
1480	Determination of the genome size of Ehrlichiaspp., using pulsed field gel electrophoresis. FEMS Microbiology Letters, 1999, 176, 73-78.	1.8	17
1481	African Tick-bite Fever in French Travelers. Emerging Infectious Diseases, 2005, 11, 1804-1806.	4.3	17
1482	Western-blot detection of RickA within spotted fever group rickettsiae using a specific monoclonal antibody. FEMS Microbiology Letters, 2008, 286, 257-262.	1.8	17
1483	<i>Rickettsia sibirica</i> subsp. <i>mongolitimonae</i> li>Infection and Retinal Vasculitis. Emerging Infectious Diseases, 2008, 14, 683-684.	4.3	17
1484	Postoperative Panophthalmitis Caused by Whipple Disease. Emerging Infectious Diseases, 2009, 15, 825-827.	4.3	17
1485	Specific and Nonspecific B-Cell Function in the Small Intestines of Patients with Whipple's Disease. Infection and Immunity, 2010, 78, 4589-4592.	2.2	17
1486	<i>Rickettsia felis</i> and <i>Bartonella clarridgeiae</i> in Fleas from New Caledonia. Vector-Borne and Zoonotic Diseases, 2011, 11, 181-183.	1.5	17
1487	Spotted fever group rickettsiae identified in Dermacentor marginatus and Ixodes ricinus ticks in Algeria. Ticks and Tick-borne Diseases, 2012, 3, 380-381.	2.7	17
1488	The proof of concept that culturomics can be superior to metagenomics to study atypical stool samples. European Journal of Clinical Microbiology and Infectious Diseases, 2013, 32, 1099-1099.	2.9	17
1489	Tropheryma whipplei endocarditis relapses after treatment with trimethoprim/sulfamethoxazole. International Journal of Antimicrobial Agents, 2013, 41, 592-594.	2.5	17
1490	Hydrophobicity of imidazole derivatives correlates with improved activity against human methanogenic archaea. International Journal of Antimicrobial Agents, 2013, 41, 544-547.	2.5	17
1491	Codon Usage, Amino Acid Usage, Transfer RNA and Amino-Acyl-tRNA Synthetases in Mimiviruses. Intervirology, 2013, 56, 364-375.	2.8	17
1492	Q Fever Endocarditis and NewCoxiella burnetiiGenotype, Saudi Arabia. Emerging Infectious Diseases, 2014, 20, 726-728.	4.3	17
1493	Draft Genome Sequence of a Human-Associated Isolate of <i>Methanobrevibacter arboriphilicus</i> , the Lowest-G+C-Content Archaeon. Genome Announcements, 2014, 2, .	0.8	17
1494	Non-contiguous finished genome sequence and description of Gorillibacterium massiliense gen. nov, sp. nov., a new member of the family Paenibacillaceae. Standards in Genomic Sciences, 2014, 9, 807-820.	1.5	17

#	Article	IF	Citations
1495	Pathogenic Eukaryotes in Gut Microbiota of Western Lowland Gorillas as Revealed by Molecular Survey. Scientific Reports, 2014, 4, 6417.	3.3	17
1496	<i>Tropheryma whipplei</i> as a Cause of Epidemic Fever, Senegal, 2010–2012. Emerging Infectious Diseases, 2016, 22, 1229-1334.	4.3	17
1497	Compartmentalization in PVC super-phylum: evolution and impact. Biology Direct, 2016, 11, 38.	4.6	17
1498	A Personal View of How Paleomicrobiology Aids Our Understanding of the Role of Lice in Plague Pandemics. Microbiology Spectrum, 2016, 4, .	3.0	17
1499	Low Level of Resistance in Enterococci Isolated in Four Hospitals, Marseille, France. Microbial Drug Resistance, 2016, 22, 218-222.	2.0	17
1500	Clostridium scindens Is Present in the Gut Microbiota during Clostridium difficile Infection: a Metagenomic and Culturomic Analysis. Journal of Clinical Microbiology, 2018, 56, .	3.9	17
1501	Oropharyngeal and nasopharyngeal decontamination with chlorhexidine gluconate in lung cancer surgery: a randomized clinical trial. Intensive Care Medicine, 2018, 44, 578-587.	8.2	17
1502	Insects and the Transmission of Bacterial Agents. Microbiology Spectrum, 2018, 6, .	3.0	17
1503	The hypervirulent Coxiella burnetii Guiana strain compared in silico, inÂvitro and inÂvivo to the Nine Mile and the German strain. Clinical Microbiology and Infection, 2019, 25, 1155.e1-1155.e8.	6.0	17
1504	Bacterial respiratory carriage in French Hajj pilgrims and the effect of pneumococcal vaccine and other individual preventive measures: A prospective cohort survey. Travel Medicine and Infectious Disease, 2019, 31, 101343.	3.0	17
1505	Giant mimiviruses escape many canonical criteria of the virus definition. Clinical Microbiology and Infection, 2019, 25, 147-154.	6.0	17
1506	Metagenomic Analysis of Microdissected Valvular Tissue for Etiological Diagnosis of Blood Culture–Negative Endocarditis. Clinical Infectious Diseases, 2020, 70, 2405-2412.	5.8	17
1507	Parasitic Infections in African Humans and Non-Human Primates. Pathogens, 2020, 9, 561.	2.8	17
1508	Clusters of COVID-19 associated with Purim celebration in the Jewish community in Marseille, France, March 2020. International Journal of Infectious Diseases, 2020, 100, 88-94.	3.3	17
1509	DNA Persistence and Relapses Questions on the Treatment Strategies of Enterococcus Infections of Prosthetic Valves. PLoS ONE, 2012, 7, e53335.	2.5	17
1510	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
1511	Q fever in children in Greece. American Journal of Tropical Medicine and Hygiene, 2004, 70, 540-4.	1.4	17
1512	Tick-Transmitted Infections in Transvaal: Consider <i>Rickettsia africae</i> . Emerging Infectious Diseases, 1999, 5, 178-181.	4.3	16

#	Article	IF	Citations
1513	Intraocular Detection of Bartonella henselae in a Patient with HLA-B27 Uveitis. Journal of Clinical Microbiology, 2004, 42, 1822-1825.	3.9	16
1514	A Century of Rickettsiology: Emerging, Reemerging Rickettsioses, Clinical, Epidemiologic, and Molecular Diagnostic Aspects and Emerging Veterinary Rickettsioses: An Overview. Annals of the New York Academy of Sciences, 2006, 1078, 1-14.	3.8	16
1515	From Cat Scratch Disease to Bartonella henselae Infection. Clinical Infectious Diseases, 2007, 45, 1541-1542.	5.8	16
1516	Rickettsia africaeInfection in Man after Travel to Ethiopia. Emerging Infectious Diseases, 2009, 15, 1867-1869.	4.3	16
1517	African Tick Bite Fever in a Taiwanese Traveler Returning from South Africa: Molecular and Serologic Studies. American Journal of Tropical Medicine and Hygiene, 2009, 81, 735-739.	1.4	16
1518	Molecular Evidence of Bartonella Infection in Domestic Dogs from Algeria, North Africa, by Polymerase Chain Reaction (PCR). American Journal of Tropical Medicine and Hygiene, 2010, 83, 298-300.	1.4	16
1519	Bartonella vinsonii Endocarditis in an Adolescent With Congenital Heart Disease. Pediatric Infectious Disease Journal, 2012, 31, 531-534.	2.0	16
1520	<i>Clostridium difficile</i> O27 Emerging Outbreak in Marseille, France. Infection Control and Hospital Epidemiology, 2013, 34, 1339-1341.	1.8	16
1521	Immune reconstitution inflammatory syndrome associated with bacterial infections. Expert Opinion on Drug Safety, 2014, 13, 341-350.	2.4	16
1522	High Prevalence of Rickettsia typhi and Bartonella Species in Rats and Fleas, Kisangani, Democratic Republic of the Congo. American Journal of Tropical Medicine and Hygiene, 2014, 90, 463-468.	1.4	16
1523	Genome sequence of Oceanobacillus picturae strain S1, an halophilic bacterium first isolated in human gut. Standards in Genomic Sciences, 2015, 10, 91.	1.5	16
1524	Exploring divergent antibiotic resistance genes in ancient metagenomes and discovery of a novel beta″actamase family. Environmental Microbiology Reports, 2016, 8, 886-895.	2.4	16
1525	A Rapid Strategy for the Isolation of New Faustoviruses from Environmental Samples Using <em>Vermamoeba vermiformis</em> . Journal of Visualized Experiments, 2016, , .	0.3	16
1526	Microbial Culturomics Broadens Human Vaginal Flora Diversity: Genome Sequence and Description of <i>Prevotella lascolaii </i> p. nov. Isolated from a Patient with Bacterial Vaginosis. OMICS A Journal of Integrative Biology, 2018, 22, 210-222.	2.0	16
1527	Complexin in ivermectin resistance in body lice. PLoS Genetics, 2018, 14, e1007569.	3.5	16
1528	Molecular evidence of bacteria in Melophagus ovinus sheep keds and Hippobosca equina forest flies collected from sheep and horses in northeastern Algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2019, 65, 103-109.	1.6	16
1529	Miniphocibacter massiliensis gen. nov., sp. nov., a new species isolated from the human gut and its taxonoâ€genomics description. MicrobiologyOpen, 2019, 8, e00735.	3.0	16
1530	Comparison of mortality associated with respiratory viral infections between December 2019 and March 2020 with that of the previous year in Southeastern France. International Journal of Infectious Diseases, 2020, 96, 154-156.	3.3	16

#	Article	IF	CITATIONS
1531	Combination of Hydroxychloroquine Plus Azithromycin As Potential Treatment for COVID-19 Patients: Safety Profile, Drug Interactions, and Management of Toxicity. Microbial Drug Resistance, 2021, 27, 281-290.	2.0	16
1532	Rhizomal Reclassification of Living Organisms. International Journal of Molecular Sciences, 2021, 22, 5643.	4.1	16
1533	Incomplete tricarboxylic acid cycle and proton gradient in <i>Pandoravirus massiliensis</i> : is it still a virus?. ISME Journal, 2022, 16, 695-704.	9.8	16
1534	Respiratory Syncytial Virus Infection: Its Propensity for Bacterial Coinfection and Related Mortality in Elderly Adults. Open Forum Infectious Diseases, 2020, 7, ofaa546.	0.9	16
1535	Chitinophaga vietnamensis sp. nov., a multi-drug resistant bacterium infecting humans. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 1758-1768.	1.7	16
1536	Emerging Rickettsioses., 0,, 17-35.		16
1537	Assessment of MALDI-TOF mass spectrometry for filariae detection in Aedes aegypti mosquitoes. PLoS Neglected Tropical Diseases, 2017, 11, e0006093.	3.0	16
1538	Senegal's Grand Magal of Touba: Syndromic Surveillance during the 2016 Mass Gathering. American Journal of Tropical Medicine and Hygiene, 2020, 102, 476-482.	1.4	16
1539	Biological and Genetic Characterization of Rickettsia sibirica Strains Isolated in the Endemic Area of the North Asian Tick Typhus. American Journal of Tropical Medicine and Hygiene, 1996, 55, 685-692.	1.4	16
1540	Livedo reticularis revealing a latent infective endocarditis due to Coxiella burnetti. Journal of the American Academy of Dermatology, 1999, 41, 842-844.	1.2	15
1541	The Environmental Pathogen Mycobacterium ulcerans Grows in Amphibian Cells at Low Temperatures. Applied and Environmental Microbiology, 2002, 68, 6403-6404.	3.1	15
1542	Interleukinâ€4 Induces <i>Coxiella burnetii</i> Replication in Human Monocytes but not in Macrophages. Annals of the New York Academy of Sciences, 2003, 990, 450-459.	3.8	15
1543	Corynebacterium freneyi Bacteremia. Journal of Clinical Microbiology, 2003, 41, 2777-2778.	3.9	15
1544	Case 5-2007. New England Journal of Medicine, 2007, 356, 715-725.	27.0	15
1545	Comparative genomic analysis of Tropheryma whipplei strains reveals that diversity among clinical isolates is mainly related to the WiSP proteins. BMC Genomics, 2007, 8, 349.	2.8	15
1546	Antibodies against <i>Rickettsia</i> spp. in Hunters, Germany. Emerging Infectious Diseases, 2008, 14, 1961-1963.	4.3	15
1547	Genomotyping of Coxiella burnetii Using Microarrays Reveals a Conserved Genomotype for Hard Tick Isolates. PLoS ONE, 2011, 6, e25781.	2.5	15
1548	Eschar and neck lymphadenopathy caused by Francisella tularensis after a tick bite: a case report. Journal of Medical Case Reports, 2011, 5, 108.	0.8	15

#	Article	IF	CITATIONS
1549	Chronic Q Fever Detection in the Netherlands. Clinical Infectious Diseases, 2011, 53, 1170-1171.	5.8	15
1550	Genome Sequence of Rickettsia conorii subsp. <i>indica</i> , the Agent of Indian Tick Typhus. Journal of Bacteriology, 2012, 194, 3288-3289.	2.2	15
1551	The first reported cases of Q fever endocarditis in Thailand. Gastroenterology Insights, 2012, 4, e7.	1.2	15
1552	Connection of toxin–antitoxin modules to inoculation eschar and arthropod vertical transmission in Rickettsiales. Comparative Immunology, Microbiology and Infectious Diseases, 2013, 36, 199-209.	1.6	15
1553	The Correlation of Q Fever andCoxiella burnetiiDNA in Household Environments in Rural Senegal. Vector-Borne and Zoonotic Diseases, 2013, 13, 70-72.	1.5	15
1554	Tropheryma whippleiGenotypes 1 and 3, Central Europe. Emerging Infectious Diseases, 2013, 19, 341-342.	4.3	15
1555	Bartonella quintana detection in Demodex from erythematotelangiectatic rosacea patients. International Journal of Infectious Diseases, 2014, 29, 176-177.	3.3	15
1556	Non-contiguous finished genome sequence and description of Clostridium saudii sp. nov. Standards in Genomic Sciences, 2014, 9, 8.	1.5	15
1557	Non-contiguous finished genome sequence and description of Anaerococcus provenciensis sp. nov Standards in Genomic Sciences, 2014, 9, 1198-1210.	1.5	15
1558	Rise of Microbial Culturomics: Noncontiguous Finished Genome Sequence and Description of Beduini massiliensisgen. nov., sp. nov OMICS A Journal of Integrative Biology, 2015, 19, 766-776.	2.0	15
1559	Comparison of nasal swabs with throat swabs for the detection of respiratory viruses by real-time reverse transcriptase PCR in adult Hajj pilgrims. Journal of Infection, 2015, 70, 207-210.	3.3	15
1560	Relapse of Tropheryma whipplei endocarditis treated by trimethoprim/sulfamethoxazole, cured by hydroxychloroquine plus doxycycline. International Journal of Infectious Diseases, 2015, 30, 17-19.	3.3	15
1561	How the virophage compels the need to readdress the classification of microbes. Virology, 2015, 477, 119-124.	2.4	15
1562	Coxiella burnetii Induces Inflammatory Interferon-Like Signature in Plasmacytoid Dendritic Cells: A New Feature of Immune Response in Q Fever. Frontiers in Cellular and Infection Microbiology, 2016, 6, 70.	3.9	15
1563	The contribution of genomics to the study of Q fever. Future Microbiology, 2016, 11, 253-272.	2.0	15
1564	MALDI-TOF MS identification of Anopheles gambiae Giles blood meal crushed on Whatman filter papers. PLoS ONE, 2017, 12, e0183238.	2.5	15
1565	Gut microbiota modifications and weight gain in early life. Human Microbiome Journal, 2018, 7-8, 10-14.	3.8	15
1566	Experimental Analysis of Mimivirus Translation Initiation Factor 4a Reveals Its Importance in Viral Protein Translation during Infection of Acanthamoeba polyphaga. Journal of Virology, 2018, 92, .	3.4	15

#	Article	IF	CITATIONS
1567	Corynebacterium fournierii sp. nov., isolated from the female genital tract of a patient with bacterial vaginosis. Antonie Van Leeuwenhoek, 2018, 111, 1165-1174.	1.7	15
1568	Description and genomic characterization of Massiliimalia massiliensis gen. nov., sp. nov., and Massiliimalia timonensis gen. nov., sp. nov., two new members of the family Ruminococcaceae isolated from the human gut. Antonie Van Leeuwenhoek, 2019, 112, 905-918.	1.7	15
1569	Co-infection of bacteria and protozoan parasites in Ixodes ricinus nymphs collected in the Alsace region, France. Ticks and Tick-borne Diseases, 2019, 10, 101241.	2.7	15
1570	The Presence of Acinetobacter baumannii DNA on the Skin of Homeless People and Its Relationship With Body Lice Infestation. Preliminary Results. Frontiers in Cellular and Infection Microbiology, 2019, 9, 86.	3.9	15
1571	Acquisition of respiratory viruses and presence of respiratory symptoms in French pilgrims during the 2016 Hajj: A prospective cohort study. Travel Medicine and Infectious Disease, 2019, 30, 32-38.	3.0	15
1572	Pedobacter schmidteae sp. nov., a new bacterium isolated from the microbiota of the planarian Schmidtea mediterranea. Scientific Reports, 2020, 10, 6113.	3.3	15
1573	Tick-borne relapsing fever Borreliosis, a major public health problem overlooked in Senegal. PLoS Neglected Tropical Diseases, 2021, 15, e0009184.	3.0	15
1574	Implementation of an in-house real-time reverse transcription-PCR assay to detect the emerging SARS-CoV-2 N501Y variants. Journal of Clinical Virology, 2021, 140, 104868.	3.1	15
1575	A Complex Interaction between Rickettsia conorii and Dickkopf-1 – Potential Role in Immune Evasion Mechanisms in Endothelial Cells. PLoS ONE, 2012, 7, e43638.	2.5	15
1576	Murine Typhus, Algeria. Emerging Infectious Diseases, 2008, 14, 676-678.	4.3	15
1577	Molecular identification of a collection of spotted Fever group rickettsiae obtained from patients and ticks from Russia. American Journal of Tropical Medicine and Hygiene, 2006, 74, 440-3.	1.4	15
1578	Prolif erative Glomerulonephritis Revealing Chronic Q Fever. American Journal of Nephrology, 1996, 16, 159-161.	3.1	14
1579	Differentiation of Bartonella Species by a Microimmunofluorescence Assay, Sodium Dodecyl Sulfate-Polyacrylamide Gel Electrophoresis, and Western Immunoblotting. Vaccine Journal, 2000, 7, 617-624.	2.6	14
1580	Cause of Black Death. Lancet Infectious Diseases, The, 2002, 2, 459.	9.1	14
1581	A Pedagogical Farm as a Source of Q Fever in a French City. European Journal of Epidemiology, 2005, 20, 957-961.	5.7	14
1582	Diversity of <i>Borrelia burgdorferi</i> Sensu Lato in Russian Far East. Microbiology and Immunology, 2005, 49, 191-197.	1.4	14
1583	First Identification of Clostridium celerecrescens in Liquid Drained from an Abscess. Journal of Clinical Microbiology, 2005, 43, 3007-3008.	3.9	14
1584	Tick Paralysis by Ixodes holocyclus in a Japanese Traveler Returning from Australia Associated with Rickettsia helvetica Infection. Journal of Travel Medicine, 2006, 10, 61-63.	3.0	14

#	Article	lF	Citations
1585	Impact of the Excision of an Ancient Repeat Insertion on Rickettsia conorii Guanylate Kinase Activity. Molecular Biology and Evolution, 2006, 23, 2112-2122.	8.9	14
1586	Etiological diagnosis of pericardial effusion. Future Microbiology, 2006, 1, 229-239.	2.0	14
1587	Coxiella burnetiiinfection in C57BL/6 mice aged $1\ \rm or\ 14\ months$ . FEMS Immunology and Medical Microbiology, 2007, 50, 396-400.	2.7	14
1588	Therapeutic impact of the correlation of doxycycline serum concentrations and the decline of phase I antibodies in Q fever endocarditis. Journal of Antimicrobial Chemotherapy, 2009, 63, 771-774.	3.0	14
1589	Matrixâ€assisted laser desorption/ionization timeâ€ofâ€flight mass spectrometry for the identification of environmental organisms: the <i>Planctomycetes</i> paradigm. Environmental Microbiology Reports, 2010, 2, 752-760.	2.4	14
1590	Publication biases in probiotics. European Journal of Epidemiology, 2012, 27, 885-886.	5.7	14
1591	Murine typhus in the homeless. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 39-43.	1.6	14
1592	Non-contiguous finished genome sequence and description of Halopiger djelfamassiliensis sp. nov Standards in Genomic Sciences, 2013, 9, 160-174.	1.5	14
1593	Draft Genome Sequencing of Methanobrevibacter oralis Strain JMR01, Isolated from the Human Intestinal Microbiota. Genome Announcements, 2014, 2, .	0.8	14
1594	Molecular Detection of Fastidious and Common Bacteria as Well as Plasmodium spp. in Febrile and Afebrile Children in Franceville, Gabon. American Journal of Tropical Medicine and Hygiene, 2015, 92, 926-932.	1.4	14
1595	Influenza vaccine for Hajj and Umrah pilgrims. Lancet Infectious Diseases, The, 2015, 15, 267.	9.1	14
1596	Paleomicrobiology of Bartonella infections. Microbes and Infection, 2015, 17, 879-883.	1.9	14
1597	Early Endocarditis and Delayed Left Ventricular Pseudoaneurysm Complicating a Transapical Transcatheter Mitral Valve-in-Valve Implantation. Circulation: Cardiovascular Interventions, 2016, 9, .	3.9	14
1598	Anaerococcus rubiinfantis sp. nov., isolated from the gut microbiota of a Senegalese infant with severe acute malnutrition. Anaerobe, 2016, 40, 85-94.	2.1	14
1599	New Species Announcement: a new format to prompt the description of new human microbial species. New Microbes and New Infections, 2017, 15, 136-137.	1.6	14
1600	Olegusella massiliensis gen. nov., sp. nov., strain KHD7 T , a new bacterial genus isolated from the female genital tract of a patient with bacterial vaginosis. Anaerobe, 2017, 44, 87-95.	2.1	14
1601	Evaluation of empirical treatment for blood culture-negative endocarditis. Journal of Antimicrobial Chemotherapy, 2017, 72, 290-298.	3.0	14
1602	Methods for culturing anaerobes from human specimen. Future Microbiology, 2018, 13, 369-381.	2.0	14

#	Article	IF	Citations
1603	Genome sequence and description of Haloferax massiliense sp. nov., a new halophilic archaeon isolated from the human gut. Extremophiles, 2018, 22, 485-498.	2.3	14
1604	Accurate identification of Anopheles gambiae Giles trophic preferences by MALDI-TOF MS. Infection, Genetics and Evolution, 2018, 63, 410-419.	2.3	14
1605	Tick-borne pathogens in removed ticks Veneto, northeastern Italy: A cross-sectional investigation. Travel Medicine and Infectious Disease, 2018, 26, 58-61.	3.0	14
1606	Characterization of a novel Gramâ€stainâ€positive anaerobic coccus isolated from the female genital tract: Genome sequence and description of <i>Murdochiella vaginalis</i> sp. nov MicrobiologyOpen, 2018, 7, e00570.	3.0	14
1607	Passive Filtration, Rapid Scanning Electron Microscopy, and Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry for ⟨i⟩Treponema⟨/i⟩ Culture and Identification from the Oral Cavity. Journal of Clinical Microbiology, 2019, 57, .	3.9	14
1608	Coinfections with SARSâ€CoVâ€2 and other respiratory viruses in Southeastern France: A matter of sampling time. Journal of Medical Virology, 2021, 93, 1878-1881.	5.0	14
1609	Updating the repertoire of cultured bacteria from the human being. Microbial Pathogenesis, 2021, 150, 104698.	2.9	14
1610	Rickettsia fournieri sp. nov., a novel spotted fever group rickettsia from Argas lagenoplastis ticks in Australia. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 3781-3784.	1.7	14
1611	Influence of Long Time Storage in Mineral Water on RNA Stability of Pseudomonas aeruginosa and Escherichia coli after Heat Inactivation. PLoS ONE, 2008, 3, e3443.	2.5	14
1612	Coxiella burnetii (Q Fever). , 2010, , 2511-2519.		14
1613	Q fever endocarditis: over 14 years of surgical experience in a referral center for rickettsioses. Journal of Heart Valve Disease, 2002, 11, 84-90.	0.5	14
1614	Determination of the rpoB gene sequences of Bartonella henselae and Bartonella quintana for phylogenic analysis. Research in Microbiology, 2000, 151, 831-836.	2.1	13
1615	Production of BartonellaGenus-Specific Monoclonal Antibodies. Vaccine Journal, 2001, 8, 847-849.	2.6	13
1616	Experimental Infection of Human Erythrocytes from Alcoholic Patients with <i>Bartonella quintana</i> . Annals of the New York Academy of Sciences, 2003, 990, 605-611.	3.8	13
1617	Lack of bartonella sp. in 167 ixodes ricinus ticks collected in Central Sweden. Scandinavian Journal of Infectious Diseases, 2004, 36, 305-306.	1.5	13
1618	Rickettsia conorii and R. prowazekii Proteome Analysis by 2DE-MS: A Step toward Functional Analysis of Rickettsial Genomes. Annals of the New York Academy of Sciences, 2005, 1063, 90-93.	3.8	13
1619	Molecular basis of Tropheryma whipplei doxycycline susceptibility examined by transcriptional profiling. Journal of Antimicrobial Chemotherapy, 2007, 59, 370-377.	3.0	13
1620	Coxiella burnetii as a possible cause of autoimmune liver disease: a case report. Journal of Medical Case Reports, 2009, 3, 8870.	0.8	13

#	Article	IF	CITATIONS
1621	Detection of Plant DNA in the Bronchoalveolar Lavage of Patients with Ventilator-Associated Pneumonia. PLoS ONE, 2010, 5, e11298.	2.5	13
1622	Emergence of the OXA-23 carbapenemase-encoding gene in multidrug-resistant Acinetobacter baumannii clinical isolates from the Principal Hospital of Dakar, Senegal. International Journal of Infectious Diseases, 2013, 17, e209-e210.	3 <b>.</b> 3	13
1623	Alcohol Disinfection Procedure for Isolating Giant Viruses from Contaminated Samples. Intervirology, 2013, 56, 434-440.	2.8	13
1624	Deciphering Genomic Virulence Traits of a Staphylococcus epidermidis Strain Causing Native-Valve Endocarditis. Journal of Clinical Microbiology, 2013, 51, 1617-1621.	3.9	13
1625	Clinical and Neuropathological Variability in Clinically Isolated Central Nervous System <scp>W</scp> hipple's Disease. Brain Pathology, 2014, 24, 230-238.	4.1	13
1626	Effect of Permethrin–Impregnated Underwear on Body Lice in Sheltered Homeless Persons. JAMA Dermatology, 2014, 150, 273.	4.1	13
1627	Non-contiguous finished genome sequence and description of Collinsella massiliensis sp. nov Standards in Genomic Sciences, 2014, 9, 1144-1158.	1.5	13
1628	Non-contiguous finished genome sequence and description of Kurthia senegalensis sp. nov Standards in Genomic Sciences, 2014, 9, 1321-1332.	1.5	13
1629	A Chlorhexidine- Agar Plate Culture Medium Protocol to Complement Standard Broth Culture of Mycobacterium tuberculosis. Frontiers in Microbiology, 2016, 7, 30.	3.5	13
1630	Genomic Insights into a New Citrobacter koseri Strain Revealed Gene Exchanges with the Virulence-Associated Yersinia pestis pPCP1 Plasmid. Frontiers in Microbiology, 2016, 7, 340.	3 <b>.</b> 5	13
1631	Head lice probably resistant to ivermectin recovered from two rural girls in Dielmo, a village in Sine-Saloum, Senegal. International Journal of Antimicrobial Agents, 2016, 47, 501-502.	2.5	13
1632	Assessment of oral ivermectin versus shampoo in the treatment of pediculosis (head lice infestation) in rural areas of Sine-Saloum, Senegal. International Journal of Antimicrobial Agents, 2016, 48, 627-632.	2.5	13
1633	Co-circulation of Plasmodium and Bacterial DNAs in Blood of Febrile and Afebrile Children from Urban and Rural Areas in Gabon. American Journal of Tropical Medicine and Hygiene, 2016, 95, 123-132.	1.4	13
1634	Host switching of human lice to new world monkeys in South America. Infection, Genetics and Evolution, 2016, 39, 225-231.	2.3	13
1635	First isolation of Akkermansia muciniphila in a blood-culture sample. Clinical Microbiology and Infection, 2017, 23, 682-683.	6.0	13
1636	An Outbreak of Kingella Kingae Infections Complicating a Severe Hand, Foot, And Mouth Disease Outbreak in Nice, France, 2016. Pediatric Infectious Disease Journal, 2017, 36, 530-532.	2.0	13
1637	Low antibodies titer and serological cross-reaction between Coxiella burnetii and Legionella pneumophila challenge the diagnosis of mediastinitis, an emerging Q fever clinical entity. Infection, 2017, 45, 911-915.	4.7	13
1638	Bartonella henselae is usually not viable in lymph nodes of patients with cat scratch disease. European Journal of Clinical Microbiology and Infectious Diseases, 2017, 36, 2207-2213.	2.9	13

#	Article	IF	CITATIONS
1639	Fluorescence In Situ Hybridization for Diagnosis of Whipple's Disease in Formalin-Fixed Paraffin-Embedded Tissue. Frontiers in Medicine, 2017, 4, 87.	2.6	13
1640	Seek and Find! PCR analyses of skin infections in West-European travelers returning from abroad with an eschar. Travel Medicine and Infectious Disease, 2018, 26, 32-36.	3.0	13
1641	Characterization of a New Ezakiella Isolated from the Human Vagina: Genome Sequence and Description of Ezakiella massiliensis sp. nov Current Microbiology, 2018, 75, 456-463.	2.2	13
1642	Measles: is a new vaccine approach needed?. Lancet Infectious Diseases, The, 2018, 18, 1060-1061.	9.1	13
1643	Tropheryma whipplei Increases Expression of Human Leukocyte Antigen-G on Monocytes to Reduce Tumor Necrosis Factor and Promote Bacterial Replication. Gastroenterology, 2018, 155, 1553-1563.	1.3	13
1644	Benefits of antibiotics burden in low-income countries. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E8109-E8110.	7.1	13
1645	Rickettsia massiliae infection after a tick bite on the eyelid. Travel Medicine and Infectious Disease, 2018, 26, 66-68.	3.0	13
1646	Ten-year experience of Q fever endocarditis in a tertiary cardiac center in Saudi Arabia. International Journal of Infectious Diseases, 2019, 88, 21-26.	3.3	13
1647	A transcriptional signature associated with non-Hodgkin lymphoma in the blood of patients with Q fever. PLoS ONE, 2019, 14, e0217542.	2.5	13
1648	Draft genome and description of Merdibacter massiliensis gen.nov., sp. nov., a new bacterium genus isolated from the human ileum. Scientific Reports, 2019, 9, 7931.	3.3	13
1649	Values of diagnostic tests for the various species of spirochetes. Médecine Et Maladies Infectieuses, 2019, 49, 102-111.	5.0	13
1650	Infectious disease symptoms and microbial carriage among French medical students travelling abroad: A prospective study. Travel Medicine and Infectious Disease, 2020, 34, 101548.	3.0	13
1651	Enteroviruses from Humans and Great Apes in the Republic of Congo: Recombination within Enterovirus C Serotypes. Microorganisms, 2020, 8, 1779.	3.6	13
1652	Molecular identification of head lice collected in Franceville (Gabon) and their associated bacteria. Parasites and Vectors, 2020, 13, 410.	2.5	13
1653	Spondylodiscitis complicating infective endocarditis. Heart, 2020, 106, 1914-1918.	2.9	13
1654	Influence of conflicts of interest on public positions in the COVID-19 era, the case of Gilead Sciences. New Microbes and New Infections, 2020, 38, 100710.	1.6	13
1655	Low blood zinc concentrations in patients with poor clinical outcome during SARS-CoV-2 infection: is there a need to supplement with zinc COVID-19 patients?. Journal of Microbiology, Immunology and Infection, 2021, 54, 997-1000.	3.1	13
1656	Multidrug-Resistant Klebsiella pneumoniae Clones from Wild Chimpanzees and Termites in Senegal. Antimicrobial Agents and Chemotherapy, 2021, 65, e0255720.	3.2	13

#	Article	IF	CITATIONS
1657	Cyclosporin A: A Repurposable Drug in the Treatment of COVID-19?. Frontiers in Medicine, 2021, 8, 663708.	2.6	13
1658	Yersinia pestisGenotyping. Emerging Infectious Diseases, 2005, 11, 1318-1319.	4.3	13
1659	Feasibility, Acceptability, and Accuracy of Vaginal Self-Sampling for Screening Human Papillomavirus Types in Women from Rural Areas in Senegal. American Journal of Tropical Medicine and Hygiene, 2019, 100, 1552-1555.	1.4	13
1660	LABORATORY-CONFIRMED MEDITERRANEAN SPOTTED FEVER IN A JAPANESE TRAVELER TO KENYA. American Journal of Tropical Medicine and Hygiene, 2005, 73, 1086-1089.	1.4	13
1661	Experimentally infected human body lice (pediculus humanus humanus) as vectors of Rickettsia rickettsii and Rickettsia conorii in a rabbit model. American Journal of Tropical Medicine and Hygiene, 2006, 74, 521-5.	1.4	13
1662	Sequential Appearance and Isolation of a SARS-CoV-2 Recombinant between Two Major SARS-CoV-2 Variants in a Chronically Infected Immunocompromised Patient. Viruses, 2022, 14, 1266.	3.3	13
1663	Acute meningoencephalitis associated with seroconversion to "Afipia felis". Lancet, The, 1992, 340, 558.	13.7	12
1664	Tick Paralysis by <i>Ixodes holocyclus</i> in a Japanese Traveler Returning from Australia. Annals of the New York Academy of Sciences, 2003, 990, 357-358.	3.8	12
1665	Spotted-fever-group rickettsioses in north Asia. Lancet, The, 2003, 362, 1939.	13.7	12
1666	Eubacterium callanderi Bacteremia: Report of the First Case. Journal of Clinical Microbiology, 2003, 41, 2235-2236.	3.9	12
1667	Culture of C. burnetii from the dental pulp of experimentally infected guinea pigs. Microbial Pathogenesis, 2004, 36, 349-350.	2.9	12
1668	Molecular detection of Bartonella spp. in the dental pulp of stray cats buried for a year. Microbial Pathogenesis, 2005, 38, 47-51.	2.9	12
1669	No link between probiotics and obesity? Author reply. Nature Reviews Microbiology, 2009, 7, 901-901.	28.6	12
1670	Giant Viruses from Amoeba in a Post-Darwinist Viral World. Intervirology, 2010, 53, 251-253.	2.8	12
1671	Comparative Genomics Evidence That Only Protein Toxins are Tagging Bad Bugs. Frontiers in Cellular and Infection Microbiology, $2011,1,7.$	3.9	12
1672	Q Fever in Woolsorters, Belgium. Emerging Infectious Diseases, 2011, 17, 2368-9.	4.3	12
1673	Genomic Comparison of Kingella kingae Strains. Journal of Bacteriology, 2012, 194, 5972-5972.	2.2	12
1674	Query rectal bleeding. Lancet, The, 2012, 380, 446.	13.7	12

#	Article	IF	CITATIONS
1675	The first reported cases of Q fever endocarditis in Thailand. Gastroenterology Insights, 2012, 4, 7.	1.2	12
1676	Foxp3 <sup>+</sup> CD4 <sup>+</sup> CD25 <sup>+</sup> regulatory T cells are increased in patients with <i>Coxiella burnetii</i> endocarditis: Figure 1. FEMS Immunology and Medical Microbiology, 2012, 64, 137-139.	2.7	12
1677	Molecular Detection of Microorganisms in Distal Airways of Patients Undergoing Lung Cancer Surgery. Annals of Thoracic Surgery, 2012, 93, 413-422.	1.3	12
1678	Occam's razor and probiotics activity on Listeria monocytogenes. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E1-E1.	7.1	12
1679	Genome Sequence of Bacillus simplex Strain P558, Isolated from a Human Fecal Sample. Genome Announcements, 2014, 2, .	0.8	12
1680	<i>Clostridium tetani</i> Osteitis without Tetanus. Emerging Infectious Diseases, 2014, 20, 1571-1573.	4.3	12
1681	Identification of Bartonellaein the Soft Tick Species Ornithodoros sonraiin Senegal. Vector-Borne and Zoonotic Diseases, 2014, 14, 26-32.	1.5	12
1682	Hemocytes from Pediculus humanus humanus are hosts for human bacterial pathogens. Frontiers in Cellular and Infection Microbiology, 2014, 4, 183.	3.9	12
1683	Sewage workers with low antibody responses may be colonized successively by several Tropheryma whipplei strains. International Journal of Infectious Diseases, 2015, 35, 51-55.	3.3	12
1684	<i>Rickettsia sibirica mongolitimonae</i> li>Infection, France, 2010–2014. Emerging Infectious Diseases, 2016, 22, 880-882.	4.3	12
1685	Deglycosylation of Tropheryma whipplei biofilm and discrepancies between diagnostic results during Whipple's disease progression. Scientific Reports, 2016, 6, 23883.	3.3	12
1686	Does Bacterial Vaginosis Result From Fecal Transplantation?. Journal of Infectious Diseases, 2016, 214, 1784-1784.	4.0	12
1687	Inediibacterium massiliense gen. nov., sp. nov., a new bacterial species isolated from the gut microbiota of a severely malnourished infant. Antonie Van Leeuwenhoek, 2017, 110, 737-750.	1.7	12
1688	Coxiella burnetii: A Hidden Pathogen in Interstitial Lung Disease?. Clinical Infectious Diseases, 2018, 67, 1120-1124.	5.8	12
1689	Halophilic & amp; halotolerant prokaryotes in humans. Future Microbiology, 2018, 13, 799-812.	2.0	12
1690	Full-length title: NRPPUR database search and in vitro analysis identify an NRPS-PKS biosynthetic gene cluster with a potential antibiotic effect. BMC Bioinformatics, 2018, 19, 463.	2.6	12
1691	Yellow fever: the Pacific should be prepared. Lancet, The, 2018, 392, 2347.	13.7	12
1692	Post-bacterial infection chronic fatigue syndrome is not a latent infection. Médecine Et Maladies Infectieuses, 2019, 49, 140-149.	5.0	12

#	Article	IF	CITATIONS
1693	Low prevalence of resistance genes in sheltered homeless population in Marseille, France, 2014–2018. Infection and Drug Resistance, 2019, Volume 12, 1139-1151.	2.7	12
1694	Seroepidemiological and molecular investigation of spotted fever group rickettsiae and <i>Coxiella burnetii</i> in Sao Tome Island: A One Health approach. Transboundary and Emerging Diseases, 2020, 67, 36-43.	3.0	12
1695	Olfactory and gustative disorders for the diagnosis of COVID-19. Travel Medicine and Infectious Disease, 2020, 37, 101875.	3.0	12
1696	Dual RNase and β-lactamase Activity of a Single Enzyme Encoded in Archaea. Life, 2020, 10, 280.	2.4	12
1697	Proof of Concept of Culturomics Use of Time of Care. Frontiers in Cellular and Infection Microbiology, 2020, 10, 524769.	3.9	12
1698	Interest of bacterial pangenome analyses in clinical microbiology. Microbial Pathogenesis, 2020, 149, 104275.	2.9	12
1699	Capybara and Brush Cutter Involvement in Q Fever Outbreak in Remote Area of Amazon Rain Forest, French Guiana, 2014. Emerging Infectious Diseases, 2020, 26, 993-997.	4.3	12
1700	Different pattern of the second outbreak of COVID-19 in Marseille, France. International Journal of Infectious Diseases, 2021, 102, 17-19.	3.3	12
1701	18F-fluorodeoxyglucose positron emission tomography/computed tomography for the diagnosis of native valve infective endocarditis: A prospective study. Archives of Cardiovascular Diseases, 2021, 114, 211-220.	1.6	12
1702	Role of glyphosate in the emergence of antimicrobial resistance in bacteria?. Journal of Antimicrobial Chemotherapy, 2021, 76, 1655-1657.	3.0	12
1703	Legionella saoudiensis sp. nov., isolated from a sewage water sample. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4367-4371.	1.7	12
1704	Collinsella vaginalis sp. nov. strain Marseille-P2666T, a new member of the Collinsella genus isolated from the genital tract of a patient suffering from bacterial vaginosis. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 949-956.	1.7	12
1705	Lovastatin Protects against Experimental Plague in Mice. PLoS ONE, 2010, 5, e10928.	2.5	12
1706	Case Report: Scalp Eschar and Neck Lymphadenopathy Associated with Bacteremia due to Coxiella-Like Bacteria. American Journal of Tropical Medicine and Hygiene, 2017, 97, 1319-1322.	1.4	12
1707	Sennetsu neorickettsiosis: a probable fish-borne cause of fever rediscovered in Laos. American Journal of Tropical Medicine and Hygiene, 2009, 81, 190-4.	1.4	12
1708	Latent Q fever endocarditis in patients undergoing routine valve surgery. Journal of Heart Valve Disease, 2014, 23, 735-43.	0.5	12
1709	Reduced Transendothelial Migration of Monocytes Infected by Coxiella burnetii. Infection and Immunity, 2000, 68, 3784-3786.	2.2	11
1710	Pyrosequencing identification of Mycobacterium tuberculosis W-Beijing. BMC Research Notes, 2009, 2, 239.	1.4	11

#	Article	IF	CITATIONS
1711	Whipple's endocarditis as a complication of tumour necrosis factor-Â antagonist treatment in a man with ankylosing spondylitis. Rheumatology, 2010, 49, 1600-1602.	1.9	11
1712	Proteomics paves the way for Q fever diagnostics. Genome Medicine, 2011, 3, 50.	8.2	11
1713	Microbe Interactions Undermine Predictions. Science, 2011, 331, 144-145.	12.6	11
1714	Mycoplasma hominis brain abscess following uterus curettage: a case report. Journal of Medical Case Reports, 2011, 5, 278.	0.8	11
1715	<i>Rickettsia felis</i> and <i>Bartonella henselae</i> in Fleas from Lebanon. Vector-Borne and Zoonotic Diseases, 2011, 11, 991-992.	1.5	11
1716	Non-contiguous finished genome sequence and description of Anaerococcus vaginalis. Standards in Genomic Sciences, 2012, 6, 356-365.	1.5	11
1717	How microbiology helps define the rhizome of life. Frontiers in Cellular and Infection Microbiology, 2012, 2, 60.	3.9	11
1718	Isolation of Arsenophonus nasoniae from Ixodes ricinus ticks in Slovakia. Ticks and Tick-borne Diseases, 2012, 3, 367-370.	2.7	11
1719	Infection in homeless people. Lancet Infectious Diseases, The, 2012, 12, 822-823.	9.1	11
1720	Insecticide resistance in mosquitoes and failure of malaria control. Expert Review of Anti-Infective Therapy, 2012, 10, 1379-1381.	4.4	11
1721	Microbial genomics challenge Darwin. Frontiers in Cellular and Infection Microbiology, 2012, 2, 127.	3.9	11
1722	Genetic Recombination Events Between Sympatric Clade A and Clade C Lice in Africa. Journal of Medical Entomology, 2013, 50, 1165-1168.	1.8	11
1723	Use of the plaque assay for testing the antibiotic susceptibility of intracellular bacteria. Future Microbiology, 2013, 8, 1301-1316.	2.0	11
1724	Does Tropheryma whipplei contribute to travelers' diarrhea?: A PCR analysis of paired stool samples in French travelers to Senegal. Travel Medicine and Infectious Disease, 2014, 12, 264-267.	3.0	11
1725	Serum concentration of co-trimoxazole during a high-dosage regimen. Journal of Antimicrobial Chemotherapy, 2014, 69, 757-760.	3.0	11
1726	Non-contiguous finished genome sequence and description of Corynebacterium jeddahense sp. nov Standards in Genomic Sciences, 2014, 9, 987-1002.	1.5	11
1727	Non-contiguous finished genome sequence and description of Clostridium ihumii sp. nov Standards in Genomic Sciences, 2015, 10, 63.	1.5	11
1728	Faecal microbiota transplantation as salvage therapy for fulminant Clostridium difficile infections. International Journal of Antimicrobial Agents, 2015, 46, 227-228.	2.5	11

#	Article	IF	CITATIONS
1729	Rapid Diagnosis of Tuberculosis by Real-Time High-Resolution Imaging of Mycobacterium tuberculosis Colonies. Journal of Clinical Microbiology, 2015, 53, 2693-2696.	3.9	11
1730	Antibiotic susceptibility of Neochlamydia hartmanellae and Parachlamydia acanthamoebae in amoebae. Microbes and Infection, 2015, 17, 761-765.	1.9	11
1731	Murine Typhus, Reunion, France, 2011–2013. Emerging Infectious Diseases, 2015, 21, 316-319.	4.3	11
1732	Reply to Slesak et al.: So much about Rickettsia felis infection to be discovered. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E6595-E6596.	7.1	11
1733	A Regional Outbreak of Clostridium difficile PCR-Ribotype 027 Infections in Southeastern France from a Single Long-Term Care Facility. Infection Control and Hospital Epidemiology, 2016, 37, 1337-1341.	1.8	11
1734	Microbial Culturomics to Map Halophilic Bacterium in Human Gut: Genome Sequence and Description of Oceanobacillus jeddahense sp. nov OMICS A Journal of Integrative Biology, 2016, 20, 248-258.	2.0	11
1735	Olsenella provencensis sp. nov., Olsenella phocaeensis sp. nov., and Olsenella mediterranea sp. nov. isolated from the human colon. Human Microbiome Journal, 2017, 4, 22-23.	3.8	11
1736	Is there a link between urinary microbiota and bladder cancer?. European Journal of Epidemiology, 2017, 32, 255-255.	5.7	11
1737	Multi-omics Analysis Sheds Light on the Evolution and the Intracellular Lifestyle Strategies of Spotted Fever Group Rickettsia spp Frontiers in Microbiology, 2017, 8, 1363.	3.5	11
1738	How mass spectrometric approaches applied to bacterial identification have revolutionized the study of human gut microbiota. Expert Review of Proteomics, 2018, 15, 217-229.	3.0	11
1739	A case of giant cell arteritis associated with culture-proven Coxiella burnetii aortitis. International Journal of Infectious Diseases, 2018, 69, 50-54.	3.3	11
1740	Acute Q Fever Endocarditis: A Paradigm Shift Following the Systematic Use of Transthoracic Echocardiography During Acute Q Fever. Clinical Infectious Diseases, 2019, 69, 1987-1995.	5.8	11
1741	Tularemia: A Case Series of Patients Diagnosed at the National Reference Center for Rickettsioses From 2008 to 2017. Open Forum Infectious Diseases, 2020, 7, ofaa440.	0.9	11
1742	Genomic Characterization of the Novel <i>Bartonella refiksaydamii</i> sp. Isolated from the Blood of a <i>Crocidura suaveolens</i> (Pallas, 1811). Vector-Borne and Zoonotic Diseases, 2021, 21, 432-440.	1.5	11
1743	Acquisition of multidrug-resistant bacteria and colistin resistance genes in French medical students on internships abroad. Travel Medicine and Infectious Disease, 2021, 39, 101940.	3.0	11
1744	Tumor Necrosis Factor Inhibitors Exacerbate Whipple's Disease by Reprogramming Macrophage and Inducing Apoptosis. Frontiers in Immunology, 2021, 12, 667357.	4.8	11
1745	SARS-CoV-2 variant from India to Marseille: The still active role of ports in the introduction of epidemics. Travel Medicine and Infectious Disease, 2021, 42, 102085.	3.0	11
1746	Ambulatory Management of Infected Orthopedic Implants., 0,, 211-230.		11

#	Article	IF	Citations
1747	Prevalence and risk factors for lung involvement on low-dose chest CT (LDCT) in a paucisymptomatic population of 247 patients affected by COVID-19. Insights Into Imaging, 2020, 11, 117.	3.4	11
1748	Implementation of Syndromic Surveillance Systems in Two Rural Villages in Senegal. PLoS Neglected Tropical Diseases, 2016, 10, e0005212.	3.0	11
1749	Vaginimicrobium propionicum gen. nov., sp. nov., a novel propionic acid bacterium derived from human vaginal discharge. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 4091-4097.	1.7	11
1750	Prevalence of human pathogens in cat and dog fleas in New Zealand. New Zealand Medical Journal, 2005, 118, U1754.	0.5	11
1751	Experimental infection of human body lice with Acinetobacter baumannii. American Journal of Tropical Medicine and Hygiene, 2006, 74, 526-31.	1.4	11
1752	Coxiella burnetii stimulates production of RANTES and MCP-1 by mononuclear cells: modulation by adhesion to endothelial cells and its implication in Q fever. European Cytokine Network, 2006, 17, 253-9.	2.0	11
1753	Profile of the Nasopharyngeal Microbiota Affecting the Clinical Course in COVID-19 Patients. Frontiers in Microbiology, 2022, 13, .	3.5	11
1754	Searching for Bacillus anthracis in Suspect Powders: a French Experience. Journal of Clinical Microbiology, 2003, 41, 524-525.	3.9	10
1755	Low seroprevalence of Bartonella species in danish elite orienteers. Scandinavian Journal of Infectious Diseases, 2004, 36, 604-606.	1.5	10
1756	Chlamydialike Organisms and Atherosclerosis. Emerging Infectious Diseases, 2006, 12, 705-706.	4.3	10
1757	Advances in <i>Tropheryma whipplei</i> research: the rush to find biomarkers for Whipple's disease. Future Microbiology, 2007, 2, 631-642.	2.0	10
1758	Microarray for serotyping of Bartonella species. BMC Microbiology, 2007, 7, 59.	3.3	10
1759	Late relapse of Q fever endocarditis. Clinical Research in Cardiology, 2007, 96, 519-521.	3.3	10
1760	Different genes govern Yersinia pestis pathogenicity in Caenorhabditis elegans and human lice. Microbial Pathogenesis, 2008, 44, 435-437.	2.9	10
1761	Intraspecies Diversity of <i>Rickettsia conorii </i>  i>. Journal of Infectious Diseases, 2009, 199, 1097-1098.	4.0	10
1762	Other Tick-Borne Diseases in Europe. Current Problems in Dermatology, 2009, 37, 130-154.	0.7	10
1763	Identification of candidate proteins for the diagnosis of Bartonella henselae infections using an immunoproteomic approach. FEMS Microbiology Letters, 2010, 310, 158-167.	1.8	10
1764	<i>Yersinia pestis</i> DNA Sequences in Late Medieval Skeletal Finds, Bavaria. Emerging Infectious Diseases, 2011, 17, 955-957.	4.3	10

#	Article	IF	Citations
1765	Molecular, Epidemiological, and Clinical Complexities of Predicting Patterns of Infectious Diseases. Frontiers in Microbiology, 2011, 2, 25.	3.5	10
1766	Soluble squalamine tablets for the rapid disinfection of home nebulizers of cystic fibrosis patients. Journal of Cystic Fibrosis, 2012, 11, 555-559.	0.7	10
1767	<i>Rickettsia felis</i> in Fleas, Southern Ethiopia, 2010. Emerging Infectious Diseases, 2012, 18, 1385-6.	4.3	10
1768	Protein candidates for the serodiagnosis of rickettsioses: 1. FEMS Immunology and Medical Microbiology, 2012, 64, 130-133.	2.7	10
1769	Saccharomyces cerevisiae boulardii transient fungemia after intravenous self-inoculation. Medical Mycology Case Reports, 2013, 2, 63-64.	1.3	10
1770	Non contiguous-finished genome sequence and description of Enorma timonensis sp. nov Standards in Genomic Sciences, 2014, 9, 970-986.	1.5	10
1771	Genome sequence and description of Corynebacterium ihumii sp. nov Standards in Genomic Sciences, 2014, 9, 1128-1143.	1.5	10
1772	Genome sequence and description of Bacteroides timonensis sp. nov Standards in Genomic Sciences, 2014, 9, 1181-1197.	1.5	10
1773	Screen-and-treat program by point-of-care of Atopobium vaginae and Gardnerella vaginalis in preventing preterm birth (AuTop trial): study protocol for a randomized controlled trial. Trials, 2015, 16, 470.	1.6	10
1774	Positron emission tomography in the diagnosis of Whipple's endocarditis: a case report. BMC Research Notes, 2015, 8, 56.	1.4	10
1775	Rainfall and Sloth Births in May, Q Fever in July, Cayenne, French Guiana. American Journal of Tropical Medicine and Hygiene, 2015, 92, 979-981.	1.4	10
1776	First Identification of <i>Anaplasma platys</i> in the Blood of Dogs from French Guiana. Vector-Borne and Zoonotic Diseases, 2015, 15, 170-172.	1.5	10
1777	Antibiotic susceptibility determination for six strains of Coxiella burnetii MST 17 from Cayenne, French Guiana. International Journal of Antimicrobial Agents, 2015, 46, 600-602.	2.5	10
1778	The challenges of preexposure prophylaxis for bacterial sexually transmitted infections. Clinical Microbiology and Infection, 2016, 22, 753-756.	6.0	10
1779	A modified multilocus sequence typing protocol to genotype Kingella kingae from oropharyngeal swabs without bacterial isolation. BMC Microbiology, 2017, 17, 200.	3.3	10
1780	Bilateral Ocular Myositis Associated with Whipple's Disease. Ocular Oncology and Pathology, 2017, 3, 17-21.	1.0	10
1781	Genomic analysis of a Raoultella ornithinolytica strain causing prosthetic joint infection in an immunocompetent patient. Scientific Reports, 2018, 8, 9462.	3.3	10
1782	Rickettsia sibirica mongolitimonae human infection: A diagnostic challenge. Travel Medicine and Infectious Disease, 2018, 26, 72-73.	3.0	10

#	Article	IF	CITATIONS
1783	Plague, camels, and lice. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 7620-7621.	7.1	10
1784	Colibacter massiliensis gen. nov. sp. nov., a novel Gram-stain-positive anaerobic diplococcal bacterium, isolated from the human left colon. Scientific Reports, 2019, 9, 17199.	3.3	10
1785	Draft genome and description of Cohnella massiliensis sp. nov., a new bacterial species isolated from the blood culture of a hemodialysis patient. Archives of Microbiology, 2019, 201, 305-312.	2.2	10
1786	From Whipple Disease to Tropheryma whipplei Infection. Clinical Infectious Diseases, 2019, 68, 1098-1099.	5.8	10
1787	Risk factors for symptoms of infection and microbial carriage among French medical students abroad. International Journal of Infectious Diseases, 2020, 100, 104-111.	3.3	10
1788	An outbreak of relapsing fever unmasked by microbial paleoserology, 16th century, France. American Journal of Physical Anthropology, 2020, 173, 784-789.	2.1	10
1789	Insights Into Subspecies Discrimination Potentiality From Bacteria MALDI-TOF Mass Spectra by Using Data Mining and Diversity Studies. Frontiers in Microbiology, 2020, 11, 1931.	3.5	10
1790	Clostridium massiliamazoniense sp. nov., New Bacterial Species Isolated from Stool Sample of a Volunteer Brazilian. Current Microbiology, 2020, 77, 2008-2015.	2.2	10
1791	Gastrointestinal symptoms and the acquisition of enteric pathogens in Hajj pilgrims: a 3-year prospective cohort study. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 315-323.	2.9	10
1792	Potential zoonotic pathogens hosted by endangered bonobos. Scientific Reports, 2021, 11, 6331.	3.3	10
1793	Infective endocarditis with neurological complications: Delaying cardiac surgery is associated with worse outcome. Archives of Cardiovascular Diseases, 2021, 114, 527-536.	1.6	10
1794	Chryseobacterium schmidteae sp. nov. a novel bacterial species isolated from planarian Schmidtea mediterranea. Scientific Reports, 2021, 11, 11002.	3.3	10
1795	Scanning Electron Microscope: A New Potential Tool to Replace Gram Staining for Microbe Identification in Blood Cultures. Microorganisms, 2021, 9, 1170.	3 <b>.</b> 6	10
1796	High-flow oxygen therapy in elderly patients infected with SARS-CoV2 with a contraindication for transfer to an intensive care unit: A preliminary report. International Journal of Infectious Diseases, 2021, 108, 1-3.	3.3	10
1797	Lyophilization to improve the sensitivity of qPCR for bacterial DNA detection in serum: the Q fever paradigm. Journal of Medical Microbiology, 2016, 65, 462-467.	1.8	10
1798	Low-dose chest CT for diagnosing and assessing the extent of lung involvement of SARS-CoV-2 pneumonia using a semi quantitative score. PLoS ONE, 2020, 15, e0241407.	2.5	10
1799	Acute Q Fever Case Detection among Acute Febrile Illness Patients, Thailand, 2002–2005. American Journal of Tropical Medicine and Hygiene, 2018, 98, 252-257.	1.4	10
1800	Laboratory-confirmed Mediterranean spotted fever in a Japanese traveler to Kenya. American Journal of Tropical Medicine and Hygiene, 2005, 73, 1086-9.	1.4	10

#	Article	IF	Citations
1801	Mycoplasma Infections of Aneurysms or Vascular Grafts. Clinical Infectious Diseases, 1999, 28, 694-695.	5.8	9
1802	Use of Macrolides for Q Fever. Antimicrobial Agents and Chemotherapy, 2003, 47, 446-446.	3.2	9
1803	Acute Pericarditis. New England Journal of Medicine, 2005, 352, 1154-1155.	27.0	9
1804	Etiological diagnostic of blood culture negative endocarditis. Enfermedades Infecciosas Y MicrobiologÃa ClÃnica, 2006, 24, 295-296.	0.5	9
1805	Expression of rOmpA and rOmpB Protein in Rickettsia massiliae during the Rhipicephalus turanicus Life Cycle. Annals of the New York Academy of Sciences, 2006, 1078, 352-356.	3.8	9
1806	An immunoproteomic approach for identification of clinical biomarkers of Whipple's disease. Proteomics - Clinical Applications, 2008, 2, 504-516.	1.6	9
1807	First Detection of <i>Wolbachia</i> spp., Including a New Genotype, in Sand Flies Collected in Marseille, France. Journal of Medical Entomology, 2008, 45, 466-469.	1.8	9
1808	Efficacy of antibiotic therapy in polyarthritis: a clue suggesting Whipple's disease. International Journal of Antimicrobial Agents, 2009, 34, 389-390.	2.5	9
1809	The Influence of Rickettsiologists on Post-Modern Microbiology. Frontiers in Cellular and Infection Microbiology, $2011,1,8.$	3.9	9
1810	Genomic Comparison of Rickettsia helvetica and Other Rickettsia Species. Journal of Bacteriology, 2012, 194, 2751-2751.	2.2	9
1811	Overexpression of the Per2 Gene in Male Patients with Acute Q Fever. Journal of Infectious Diseases, 2012, 206, 1768-1770.	4.0	9
1812	Genome Sequence of Rickettsia conorii subsp. israelensis, the Agent of Israeli Spotted Fever. Journal of Bacteriology, 2012, 194, 5130-5131.	2.2	9
1813	Genome Sequence of Diplorickettsia massiliensis, an Emerging Ixodes ricinus-Associated Human Pathogen. Journal of Bacteriology, 2012, 194, 3287-3287.	2.2	9
1814	Imported rickettsioses in Italy. Travel Medicine and Infectious Disease, 2012, 10, 201-204.	3.0	9
1815	A <scp>DNA</scp> microarray for the versatile diagnosis of infectious diarrhea. Apmis, 2013, 121, 634-642.	2.0	9
1816	Non-contiguous finished genome sequence and description of Anaerococcus pacaensis sp. nov., a new species of anaerobic bacterium. Standards in Genomic Sciences, 2013, 8, 548-560.	1.5	9
1817	MALDI-ToF Mass Spectrometry for the Rapid Diagnosis of Cancerous Lung Nodules. PLoS ONE, 2014, 9, e97511.	2.5	9
1818	Rickettsia africae infection complicated with painful sacral syndrome in an Italian traveller returning from Zimbabwe. International Journal of Infectious Diseases, 2014, 29, 194-196.	3.3	9

#	Article	IF	CITATIONS
1819	From the Hajj: it's the flu, idiot. Clinical Microbiology and Infection, 2014, 20, O1.	6.0	9
1820	Are Infectious Disease Doctors Better at Caring for Infectious Diseases Than Other Specialists?. Clinical Infectious Diseases, 2014, 58, 1486-1487.	<b>5.</b> 8	9
1821	Reply. Journal of the American College of Cardiology, 2014, 63, 187-189.	2.8	9
1822	Methods for the discovery of emerging pathogens. Microbial Pathogenesis, 2014, 77, 114-118.	2.9	9
1823	Guidelines need controls. Clinical Microbiology and Infection, 2015, 21, 1043-1044.	6.0	9
1824	Increasing burden of urinary tract infections due to intrinsic colistin-resistant bacteria in hospitals in Marseille, France. International Journal of Antimicrobial Agents, 2015, 45, 144-150.	2.5	9
1825	Doxycycline assay hair samples for testing long-term compliance treatment. Journal of Infection, 2015, 71, 511-517.	3.3	9
1826	Sternoclavicular joint infection caused by Coxiella burnetii: a case report. Journal of Medical Case Reports, 2016, 10, 139.	0.8	9
1827	Implementation and Initial Analysis of a Laboratory-Based Weekly Biosurveillance System, Provence-Alpes-CÃ'te d'Azur, France. Emerging Infectious Diseases, 2017, 23, 582-589.	4.3	9
1828	Case report: Coxiella burnetii vascular infection and lymphoma in the Netherlands. Infection, 2018, 46, 131-134.	4.7	9
1829	Human pegivirus isolates characterized by deep sequencing from hepatitis CÂvirusâ€RNA and human immunodeficiency virusâ€RNA–positive blood donations, France. Journal of Medical Virology, 2019, 91, 38-44.	5.0	9
1830	Introduction to Measurement of Avidity of Anti-Coxiella burnetii IgG in Diagnosis of Q Fever. Journal of Clinical Microbiology, 2019, 57, .	3.9	9
1831	Leptospirosis, one neglected disease in rural Senegal. Veterinary Medicine and Science, 2019, 5, 536-544.	1.6	9
1832	Culturomics provides critical prokaryote strains for anti-Listeria and anti-cancer probiotics. International Journal of Antimicrobial Agents, 2019, 54, 407-409.	2.5	9
1833	<i>Tropheryma whipplei</i> intestinal colonization in Italian and migrant population: a retrospective observational study. Future Microbiology, 2019, 14, 283-292.	2.0	9
1834	No global increase in resistance to antibiotics: a snapshot of resistance from 2001 to 2016 in Marseille, France. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 395-407.	2.9	9
1835	From anaerobes to aerointolerant prokaryotes. Human Microbiome Journal, 2020, 15, 100068.	3.8	9
1836	Rapid Scanning Electron Microscopy Detection and Sequencing of Severe Acute Respiratory Syndrome Coronavirus 2 and Other Respiratory Viruses. Frontiers in Microbiology, 2020, 11, 596180.	3.5	9

#	Article	IF	CITATIONS
1837	A Novel Approach for Detecting Unique Variations among Infectious Bacterial Species in Endocarditic Cardiac Valve Vegetation. Cells, 2020, 9, 1899.	4.1	9
1838	A protein of the metallo-hydrolase/oxidoreductase superfamily with both beta-lactamase and ribonuclease activity is linked with translation in giant viruses. Scientific Reports, 2020, 10, 21685.	3.3	9
1839	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. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 1579-1582.	2.9	9
1840	Bartonella infections diagnosed in the French reference center, 2014–2019, and focus on infections in the immunocompromised. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 2407-2410.	2.9	9
1841	Does SARS-CoV-2 re-infection depend on virus variant?. Clinical Microbiology and Infection, 2021, 27, 1374-1375.	6.0	9
1842	Draft Genome Sequence of Salirhabdus euzebyi Strain Q1438. Microbiology Resource Announcements, 2020, 9, .	0.6	9
1843	Bacteriology, Taxonomy, and Phylogeny of <i>Rickettsia</i> . Infectious Disease and Therapy, 2007, , 1-14.	0.0	9
1844	Other Tick-Borne Rickettsioses. Infectious Disease and Therapy, 2007, , 139-162.	0.0	9
1845	Clinical Aspects, Diagnosis, and Treatment of Q Fever. Infectious Disease and Therapy, 2007, , 291-302.	0.0	9
1846	16S Metagenomic Comparison of Plasmodium falciparum–Infected and Noninfected Anopheles gambiae and Anopheles funestus Microbiota from Senegal. American Journal of Tropical Medicine and Hygiene, 2018, 99, 1489-1498.	1.4	9
1847	Diversity and distribution of ticks from domestic ruminants in Lebanon. Veterinaria Italiana, 2017, 53, 147-155.	0.5	9
1848	Long-Term Persistence of Olfactory and Gustatory Disorders in COVID-19 Patients. Frontiers in Medicine, 2022, 9, 794550.	2.6	9
1849	Pericardial effusion as the initial feature of Q fever. American Heart Journal, 1995, 130, 1308.	2.7	8
1850	Mycoplasma pneumoniae pneumonia following assisted ventilation. American Journal of Medicine, 1996, 101, 165-169.	1.5	8
1851	Monoclonal Antibodies to Immunodominant Epitope of Tropheryma whipplei. Vaccine Journal, 2002, 9, 156-159.	3.1	8
1852	A Homeless Man with Maculopapular Rash Who Died in Marseille, France. Clinical Infectious Diseases, 2004, 38, 1412-1412.	5.8	8
1853	Intervening Sequence Acquired by Lateral Gene Transfer in Tropheryma whipplei Results in 23S rRNA Fragmentation. Applied and Environmental Microbiology, 2005, 71, 6698-6701.	3.1	8
1854	HIV-2 Protease resistance defined in yeast cells. Retrovirology, 2006, 3, 58.	2.0	8

#	Article	IF	CITATIONS
1855	Prospective Evaluation of Rickettsioses in the Trakya (European) Region of Turkey and Atypic Presentations of Rickettsia Conorii. Annals of the New York Academy of Sciences, 2006, 1078, 173-175.	3.8	8
1856	Two Cases of Cellulitis in the Course of African Tick Bite Fever: A Fortuitous Association?. Dermatology, 2008, 217, 140-142.	2.1	8
1857	In vitro activity of pentamidine against Tropheryma whipplei. International Journal of Antimicrobial Agents, 2011, 38, 545-547.	2.5	8
1858	Genotyping Yersinia pestis in historical plague. Lancet Infectious Diseases, The, 2011, 11, 894-895.	9.1	8
1859	Pseudoclavibacter-like subcutaneous infection: a case report. Journal of Medical Case Reports, 2011, 5, 468.	0.8	8
1860	Complete Genome Sequence of Rickettsia slovaca, the Agent of Tick-Borne Lymphadenitis. Journal of Bacteriology, 2012, 194, 1612-1612.	2.2	8
1861	Genome Sequence of Rickettsia australis, the Agent of Queensland Tick Typhus. Journal of Bacteriology, 2012, 194, 5129-5129.	2.2	8
1862	Lamarckian evolution of the giant Mimivirus in allopatric laboratory culture on amoebae. Frontiers in Cellular and Infection Microbiology, 2012, 2, 91.	3.9	8
1863	Tryptose phosphate broth improvesRickettsia felisreplication in mammalian cells: 1. FEMS Immunology and Medical Microbiology, 2012, 64, 111-114.	2.7	8
1864	Point-of-care testing for community-acquired pneumonia. Lancet Infectious Diseases, The, 2013, 13, 647-649.	9.1	8
1865	Partial Disruption of Translational and Posttranslational Machinery Reshapes Growth Rates of Bartonella birtlesii. MBio, 2013, 4, e00115-13.	4.1	8
1866	Staphylococcus aureus subsp. anaerobius strain ST1464 genome sequence. Standards in Genomic Sciences, 2013, 9, 1-11.	1.5	8
1867	Diagnosis ofBartonella henselaeProsthetic Valve Endocarditis in Man, France. Emerging Infectious Diseases, 2014, 20, 1396-1397.	4.3	8
1868	Yersinia pestis and the three plague pandemics. Lancet Infectious Diseases, The, 2014, 14, 918-919.	9.1	8
1869	Non-contiguous finished genome sequence and description of Bacteroides neonati sp. nov., a new species of anaerobic bacterium. Standards in Genomic Sciences, 2014, 9, 794-806.	1.5	8
1870	Studies of Ancient Lice Reveal Unsuspected Past Migrations of Vectors. American Journal of Tropical Medicine and Hygiene, 2015, 93, 623-625.	1.4	8
1871	The detection of vector-borne-disease-related DNA in human stool paves the way to large epidemiological studies. European Journal of Epidemiology, 2015, 30, 1021-1026.	5.7	8
1872	Orientia tsutsugamushiin Lung of Patient with Acute Respiratory Distress Syndrome, France, 2013. Emerging Infectious Diseases, 2015, 21, 373-375.	4.3	8

#	Article	IF	CITATIONS
1873	Rickettsia and Bartonella Species in Fleas from Reunion Island. American Journal of Tropical Medicine and Hygiene, 2015, 92, 617-619.	1.4	8
1874	Saudi Moumouvirus, the First Group B Mimivirus Isolated from Asia. Frontiers in Microbiology, 2016, 07, 2029.	3.5	8
1875	Risungbinella massiliensis sp. nov., a new member of Thermoactinomycetaceae isolated from human gut. Antonie Van Leeuwenhoek, 2016, 109, 773-784.	1.7	8
1876	Genome Sequence of the Tick-Borne Pathogen $\mbox{\ensuremath{$\cdot$}}$ Rickettsia raoultii $\mbox{\ensuremath{$/$}}$ i. Genome Announcements, 2016, 4, .	0.8	8
1877	Human Lice in Paleoentomology and Paleomicrobiology. Microbiology Spectrum, 2016, 4, .	3.0	8
1878	An Alternative Strategy of Preventive Control of Tick-Borne Relapsing Fever in Rural Areas of Sine-Saloum, Senegal. American Journal of Tropical Medicine and Hygiene, 2016, 95, 537-545.	1.4	8
1879	Microbiota, obesity and malnutrition. Microbial Pathogenesis, 2017, 106, 1-2.	2.9	8
1880	Why new antibiotics are not obviously useful now. International Journal of Antimicrobial Agents, 2017, 49, 549-553.	2.5	8
1881	Description of Chryseobacterium timonianum sp. nov., isolated from a patient with pneumonia. Antonie Van Leeuwenhoek, 2017, 110, 1121-1132.	1.7	8
1882	How artificial is the antibiotic resistance definition?. Lancet Infectious Diseases, The, 2017, 17, 690.	9.1	8
1883	Q Fever: Confusion Between Chronic Infection and Chronic Fatigue. Clinical Infectious Diseases, 2017, 65, 1054-1055.	5.8	8
1884	<i>Mycoplasma genitalium,</i> an agent of reemerging sexually transmitted infections. Apmis, 2017, 125, 916-920.	2.0	8
1885	From Expert Protocols to Standardized Management of Infectious Diseases. Clinical Infectious Diseases, 2017, 65, S12-S19.	5.8	8
1886	Hemodialysis vascular graft as a focus of persistent Q fever. Infection, 2018, 46, 881-884.	4.7	8
1887	Unrecognized preâ€transplant disseminated <i>Coxiella burnetti</i> infection diagnosed in a postâ€transplant heartâ€kidney recipient. Transplant Infectious Disease, 2018, 20, e12962.	1.7	8
1888	Dual Genotype <i>Orientia tsutsugamushi</i> Infection in Patient with Rash and Eschar, Vietnam, 2016. Emerging Infectious Diseases, 2018, 24, 1520-1523.	4.3	8
1889	Genome sequence and description of Gracilibacillus timonensissp. nov. strain Marseilleâ€P2481T, a moderate halophilic bacterium isolated from the human gut microflora. MicrobiologyOpen, 2019, 8, e00638.	3.0	8
1890	Alcohol and the global burden of disease. Lancet, The, 2019, 393, 2390-2391.	13.7	8

#	Article	lF	CITATIONS
1891	An Integrative Database of $\hat{l}^2$ -Lactamase Enzymes: Sequences, Structures, Functions, and Phylogenetic Trees. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	8
1892	Whole-Genome Sequence of French Clinical Olivibacter jilunii Strain P8502. Microbiology Resource Announcements, $2019,8,.$	0.6	8
1893	Environmental investigation of respiratory pathogens during the Hajj 2016 and 2018. Travel Medicine and Infectious Disease, 2020, 33, 101500.	3.0	8
1894	Insecticidal Activity of Bacteria from Larvae Breeding Site with Natural Larvae Mortality: Screening of Separated Supernatant and Pellet Fractions. Pathogens, 2020, 9, 486.	2.8	8
1895	Zoonotic Abbreviata caucasica in Wild Chimpanzees (Pan troglodytes verus) from Senegal. Pathogens, 2020, 9, 517.	2.8	8
1896	Epidemiological serosurvey of vector-borne and zoonotic pathogens among homeless people living in shelters in Marseille: cross-sectional one-day surveys (2005–2015). European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1663-1672.	2.9	8
1897	How useful is serology for COVID-19?. International Journal of Infectious Diseases, 2021, 102, 170-171.	3.3	8
1898	FastFung: A novel medium for the culture and isolation of fastidious fungal species from clinical samples. Journal of Microbiological Methods, 2021, 180, 106108.	1.6	8
1899	Effect of hydroxychloroquine and azithromycin on SARS-CoV-2 clearance in COVID-19 patients, a meta-analysis International Journal of Antimicrobial Agents, 2021, 57, 106240.	2.5	8
1900	Human galectin-1 and galectin-3 promote $\langle i \rangle$ Tropheryma whipplei $\langle i \rangle$ infection. Gut Microbes, 2021, 13, 1-15.	9.8	8
1901	Acquisition of multidrug-resistant bacteria and encoding genes among French pilgrims during the 2017 and 2018 Hajj. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 1199-1207.	2.9	8
1902	Negativibacillus massiliensis gen. nov., sp. nov., a New Bacterial Genus Isolated from a Human Left Colon Sample. Microbiology Research, 2021, 12, 29-42.	1.9	8
1903	Planarians (Platyhelminthes)â€"An Emerging Model Organism for Investigating Innate Immune Mechanisms. Frontiers in Cellular and Infection Microbiology, 2021, 11, 619081.	3.9	8
1904	Occurrence of Ten Protozoan Enteric Pathogens in Three Non-Human Primate Populations. Pathogens, 2021, 10, 280.	2.8	8
1905	Rapid Detection of Imipenem Resistance in Gram-Negative Bacteria Using Tabletop Scanning Electron Microscopy: A Preliminary Evaluation. Frontiers in Microbiology, 2021, 12, 658322.	3.5	8
1906	Molecular Characterization and Genetic Diversity of Haplogroup E Human Lice in Guinea, West Africa. Microorganisms, 2021, 9, 257.	3.6	8
1907	Letter to the editor: Plenty of coronaviruses but no SARS-CoV-2. Eurosurveillance, 2020, 25, .	7.0	8
1908	Non-contiguous finished genome sequence and description of Alistipes timonensis sp. nov Standards in Genomic Sciences, 2012, 6, 315-324.	1.5	8

#	Article	IF	CITATIONS
1909	Occurrence of a substitution or deletion of SARS-CoV-2 spike amino acid 677 in various lineages in Marseille, France. Virus Genes, 2022, 58, 53-58.	1.6	8
1910	Adapted Protocol for <i>Saccharibacteria</i> Cocultivation: Two New Members Join the Club of Candidate Phyla Radiation. Microbiology Spectrum, 2021, 9, e0106921.	3.0	8
1911	Serologic response to rickettsial antigens in patients with Astrakhan fever. European Journal of Epidemiology, 1995, 11, 383-387.	5 <b>.</b> 7	7
1912	A "Query―Pancreatitis in a Young Shepherdess: An Uncommon Manifestation of Acute Q Fever. Clinical Infectious Diseases, 1999, 29, 445-446.	5.8	7
1913	Multiplexed Serology in Atypical Bacterial Pneumonia. Annals of the New York Academy of Sciences, 2006, 1078, 530-540.	3.8	7
1914	African Tickbite Fever in Travelers, Swaziland. Emerging Infectious Diseases, 2007, 13, 353-355.	4.3	7
1915	The globalization of intestinal microbiota. European Journal of Clinical Microbiology and Infectious Diseases, 2010, 29, 1049-1050.	2.9	7
1916	Coxiella burnetii Transcriptional Analysis Reveals Serendipity Clusters of Regulation in Intracellular Bacteria. PLoS ONE, 2010, 5, e15321.	2.5	7
1917	Molecular Detection of <i>Bartonella alsatica</i> ii>in Rabbit Fleas, France. Emerging Infectious Diseases, 2010, 16, 2013-2014.	4.3	7
1918	LES ANIMAUX VERTÉBRÉS SONT-ILS RÉSERVOIRS DE RICKETTSI. Bulletin De L'Academie Veterinaire De France, 2010, , 291.	0.0	7
1919	Defining causality in emerging agents of acute bacterial diarrheas: a step beyond the Koch's postulates. Future Microbiology, 2010, 5, 1787-1797.	2.0	7
1920	A viral grandfather: genomics in 2010 contradict Darwin's vision of evolution. European Journal of Clinical Microbiology and Infectious Diseases, 2011, 30, 935-936.	2.9	7
1921	Draft Genome Sequence of Brevibacterium massiliense Strain 541308T. Journal of Bacteriology, 2012, 194, 5151-5152.	2.2	7
1922	Sequence and Annotation of Rickettsia sibirica sibirica Genome. Journal of Bacteriology, 2012, 194, 2377-2377.	2.2	7
1923	Genome Sequence of "Rickettsia sibirica subsp. mongolitimonae― Journal of Bacteriology, 2012, 194, 2389-2390.	2.2	7
1924	Detection of Rickettsia helvetica in Ixodes ricinus ticks collected from Pyrenean chamois in France. Ticks and Tick-borne Diseases, 2012, 3, 387-388.	2.7	7
1925	Tickâ€Borne Relapsing Fever With Cutaneous Eschar andÂRadiculopathy, Ethiopia. Journal of Travel Medicine, 2012, 19, 261-263.	3.0	7
1926	Evaluation of the diagnostic value of fluorescent in situ hybridization in a rat model of bacterial pneumonia. Diagnostic Microbiology and Infectious Disease, 2013, 76, 425-431.	1,8	7

#	Article	IF	CITATIONS
1927	Identification of salivary antigenic markers discriminating host exposition between two European ticks: Rhipicephalus sanguineus and Dermacentor reticulatus. Comparative Immunology, Microbiology and Infectious Diseases, 2013, 36, 39-53.	1.6	7
1928	Incidental diagnosis of colonic tumor by PET/CT in infectious endocarditis. Journal of Infection, 2013, 67, 88-90.	3.3	7
1929	Genome Sequence of Rickettsia tamurae, a Recently Detected Human Pathogen in Japan. Genome Announcements, 2014, 2, .	0.8	7
1930	Uncultured Candidatus Neoehrlichia Mikurensis. Clinical Infectious Diseases, 2014, 59, 1042-1042.	5.8	7
1931	Update on Tick-Borne Rickettsioses around the World: a Geographic Approach. Clinical Microbiology Reviews, 2014, 27, 166-166.	13.6	7
1932	MspA-Mycobacterium tuberculosis-transformant with reduced virulence: The "unbirthday paradigm― Microbial Pathogenesis, 2014, 76, 10-18.	2.9	7
1933	Non-contiguous finished genome sequence and description of Paenibacillus gorillae sp. nov Standards in Genomic Sciences, 2014, 9, 1031-1045.	1.5	7
1934	Doxycycline kills human lice through its activity on their bacterial symbiont. International Journal of Antimicrobial Agents, 2015, 45, 675-676.	2.5	7
1935	Reevaluation of possible outcomes of infections with human immunodeficiency virus. Clinical Microbiology and Infection, 2016, 22, 299-311.	6.0	7
1936	Brevibacterium massiliense bacteremia. IDCases, 2017, 7, 25-26.	0.9	7
1937	Antiphospholipid antibodies proposed in the diagnosis of infective endocarditis. European Journal of Clinical Microbiology and Infectious Diseases, 2017, 36, 1159-1162.	2.9	7
1938	Critical Microbiological View of SER-109. Journal of Infectious Diseases, 2017, 215, 161-162.	4.0	7
1939	Draft Genome Sequence of Kingella negevensis SW7208426, the First European Strain of K. negevensis Isolated from a Healthy Child in Switzerland. Genome Announcements, 2017, 5, .	0.8	7
1940	Molecular Evidence of <i>Rickettsia slovaca</i> in Wild Boar Lice, in Northeastern Algeria. Vector-Borne and Zoonotic Diseases, 2018, 18, 114-116.	1.5	7
1941	Draft genome sequence of Fermentimonas caenicola strain SIT8, isolated from the human gut. Standards in Genomic Sciences, 2018, 13, 8.	1.5	7
1942	Investigation of Pneumonic Plague, Madagascar. Emerging Infectious Diseases, 2018, 24, 183-183.	4.3	7
1943	Western Immunoblotting for the Diagnosis of Enterococcus faecalis and Streptococcus gallolyticus Infective Endocarditis. Frontiers in Cellular and Infection Microbiology, 2019, 9, 314.	3.9	7
1944	Taxonogenomics description of Parabacteroides timonensis sp. nov. isolated from a human stool sample. MicrobiologyOpen, 2019, 8, e00702.	3.0	7

#	Article	IF	CITATIONS
1945	Listeria monocytogenes in human milk in Mali: A potential health emergency. Journal of Infection, 2020, 80, 121-142.	3.3	7
1946	Bouillabaisse or Fish Soup: The Limitations of Meta-analysis Confronted to the Inconsistency of Fecal Microbiota Transplantation Studies. Clinical Infectious Diseases, 2020, 70, 2454-2454.	5.8	7
1947	Early mortality attributable to PICC-lines in 4 public hospitals of Marseille from 2010 to 2016 (Revised) Tj ETQq1	l 0.78431 1:0	4 rgBT /Ove
1948	Listeria monocytogenes detected in vaginal self-samples of 2 women after spontaneous miscarriage, Senegal, West Africa. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 393-394.	2.9	7
1949	Comparison of Three Skin Sampling Methods and Two Media for Culturing Malassezia Yeast. Journal of Fungi (Basel, Switzerland), 2020, 6, 350.	3.5	7
1950	Rickettsia mongolitimonae Encephalitis, Southern France, 2018. Emerging Infectious Diseases, 2020, 26, 362-364.	4.3	7
1951	Alcohol pretreatment of stools effect on culturomics. Scientific Reports, 2020, 10, 5190.	3.3	7
1952	High-speed large-scale automated isolation of SARS-CoV-2 from clinical samples using miniaturized co-culture coupled to high-content screening. Clinical Microbiology and Infection, 2021, 27, 128.e1-128.e7.	6.0	7
1953	Anaerococcus urinimassiliensis sp. nov., a new bacterium isolated from human urine. Scientific Reports, 2021, 11, 2684.	3.3	7
1954	Global Discrepancies between Numbers of Available SARS-CoV-2 Genomes and Human Development Indexes at Country Scales. Viruses, 2021, 13, 775.	3.3	7
1955	SARS-CoV-2 Persistent Viral Shedding in the Context of Hydroxychloroquine-Azithromycin Treatment. Viruses, 2021, 13, 890.	3.3	7
1956	A metallo-β-lactamase enzyme for internal detoxification of the antibiotic thienamycin. Scientific Reports, 2021, 11, 10062.	3.3	7
1957	A Listeria monocytogenes clone in human breast milk associated with severe acute malnutrition in West Africa: A multicentric case-controlled study. PLoS Neglected Tropical Diseases, 2021, 15, e0009555.	3.0	7
1958	Pangenome analysis and virulence profiling of Streptococcus intermedius. BMC Genomics, 2021, 22, 522.	2.8	7
1959	Molecular Detection of Past Pathogens. , 2008, , 55-68.		7
1960	Differences in Levels of Soluble E-selectin and VCAM-1 in Malignant versus non Malignant Mediterranean Spotted Fever. Thrombosis and Haemostasis, 1999, 82, 1610-1613.	3.4	7
1961	PREVALENCE OF HEPATITIS G VIRUS INFECTION IN KIDNEY TRANSPLANT RECIPIENTS. Transplantation, 1997, 64, 537-539.	1.0	7
1962	Increases in the Levels of <i>Coxiella burnetii </i> -Specific Immunoglobulin G1 and G3 Antibodies in Acute Q Fever and Chronic Q Fever. Vaccine Journal, 1998, 5, 814-816.	2.6	7

#	Article	IF	CITATIONS
1963	Whole-Genome Sequence of Haloimpatiens lingqiaonensis Strain P8956. Microbiology Resource Announcements, 2019, 8, .	0.6	7
1964	Molecular characterization of some equine vector-borne diseases and associated arthropods in Egypt. Acta Tropica, 2022, 227, 106274.	2.0	7
1965	Screening and Whole Genome Sequencing of SARS-CoV-2 Circulating During the First Three Waves of the COVID-19 Pandemic in Libreville and the Haut-Ogooué Province in Gabon. Frontiers in Medicine, 2022, 9, .	2.6	7
1966	Letter to the editor: Chronic coxiella burnetii infection mimicking malignant hematologic disease. American Journal of Hematology, 1992, 39, 309-309.	4.1	6
1967	Monoclonal Antibodies to <i>Afipia felis</i> Patative Agent of Cat Scratch Disease. American Journal of Clinical Pathology, 1994, 101, 603-606.	0.7	6
1968	Bartonella quintana, Lice, and Molecular Tools. Journal of Medical Entomology, 2006, 43, 787-787.	1.8	6
1969	Ecology and Molecular Epidemiology of Tick-Borne Rickettsioses and Anaplasmoses with Natural Foci in Russia and Kazakhstan. Annals of the New York Academy of Sciences, 2006, 1078, 299-304.	3.8	6
1970	Validation of a Rickettsia prowazekii-Specific Quantitative Real-Time PCR Cassette and DNA Extraction Protocols Using Experimentally Infected Lice. Annals of the New York Academy of Sciences, 2006, 1078, 617-619.	3.8	6
1971	Murine Model of Infection by Tropheryma whipplei. Infection and Immunity, 2006, 74, 4915-4917.	2.2	6
1972	<i>Bartonella quintana</i> Coinfection in <i>Staphylococcusaureus</i> Endocarditis: Usefulness of Screening in Highâ€Risk Patients?. Clinical Infectious Diseases, 2009, 48, 1332-1333.	5.8	6
1973	Genome Sequence of Rickettsia conorii subsp. caspia, the Agent of Astrakhan Fever. Journal of Bacteriology, 2012, 194, 4763-4764.	2.2	6
1974	Genome Sequence of Reyranella massiliensis, a Bacterium Associated with Amoebae. Journal of Bacteriology, 2012, 194, 5698-5698.	2.2	6
1975	Genomic Comparison of Rickettsia honei Strain RB T and Other Rickettsia Species. Journal of Bacteriology, 2012, 194, 4145-4145.	2.2	6
1976	Protein candidates for Q fever serodiagnosis. FEMS Immunology and Medical Microbiology, 2012, 64, 140-142.	2.7	6
1977	Genome Sequence of Rickettsia gravesii, Isolated from Western Australian Ticks. Genome Announcements, 2013, 1, .	0.8	6
1978	Obesity and stools, the "emperor's new clothing―paradigm. European Journal of Epidemiology, 2015, 30, 1071-1071.	5.7	6
1979	A case of infectious endocarditis due to BCG. International Journal of Infectious Diseases, 2015, 35, 27-28.	3.3	6
1980	Deadly infectious diseases such as Ebola: the parachute paradigm. Clinical Microbiology and Infection, 2015, 21, 389-390.	6.0	6

#	Article	IF	CITATIONS
1981	Coxiella burnetii Endocarditis in a Child Caused by a New Genotype. Pediatric Infectious Disease Journal, 2016, 35, 213-214.	2.0	6
1982	Paleogenetics and Past Infections: the Two Faces of the Coin of Human Immune Evolution. Microbiology Spectrum, 2016, 4, .	3.0	6
1983	Is there a terrible issue with bacterial resistance: pro–con. Clinical Microbiology and Infection, 2016, 22, 403-404.	6.0	6
1984	Contemporary challenges and opportunities in the diagnosis and outbreak detection of multidrug-resistant infectious disease. Expert Review of Molecular Diagnostics, 2016, 16, 1163-1175.	3.1	6
1985	Evaluation of a new extraction protocol for yeast identification by mass spectrometry. Journal of Microbiological Methods, 2016, 129, 61-65.	1.6	6
1986	Monoclonal Antibodies for the Diagnosis of Borrelia crocidurae. American Journal of Tropical Medicine and Hygiene, 2016, 94, 61-67.	1.4	6
1987	Vaginal self-sampling as a diagnosis tool in low-income countries and potential applications for exploring the infectious causes of miscarriage. Future Microbiology, 2017, 12, 609-620.	2.0	6
1988	The study of microbiota needs both microbiologists and medical doctors. Clinical Microbiology and Infection, 2017, 23, 500-501.	6.0	6
1989	Developing new insecticides to prevent chaos: the real future threat. Lancet Infectious Diseases, The, 2017, 17, 804-805.	9.1	6
1990	Transmission of Coxiella burnetii to cage mates using murine animal model. Comparative Immunology, Microbiology and Infectious Diseases, 2017, 50, 29-33.	1.6	6
1991	Coxiella burnetii Multilevel Disk Space Infection, Epidural Abscess, and Vertebral Osteomyelitis Secondary to Contiguous Spread From Infected Abdominal Aortic Aneurysm or Graft: Report of 4 Cases Acquired in the US and Review of the Literature. Open Forum Infectious Diseases, 2017, 4, ofx192.	0.9	6
1992	Koch Postulate: Why Should we Grow Bacteria?. Archives of Medical Research, 2017, 48, 774-779.	3.3	6
1993	PCR Detection of Mimivirus. Emerging Infectious Diseases, 2017, 23, 1044-1045.	4.3	6
1994	Microbial Culturomics Application for Global Health: Noncontiguous Finished Genome Sequence and Description of <i>Pseudomonas massiliensis</i> Strain CB-1 <sup>T</sup> sp. nov. in Brazil. OMICS A Journal of Integrative Biology, 2018, 22, 164-175.	2.0	6
1995	<i>Bacillus kwashiorkori</i> sp. nov., a new bacterial species isolated from a malnourished child using culturomics. MicrobiologyOpen, 2018, 7, e00535.	3.0	6
1996	Asymptomatic carriage of Streptococcus pneumoniae detected by qPCR on the palm of hands of populations in rural Senegal. PLoS Neglected Tropical Diseases, 2018, 12, e0006945.	3.0	6
1997	Tick- and flea-borne rickettsioses in Tizi-Ouzou, Algeria: Implications for travel medicine. Travel Medicine and Infectious Disease, 2018, 26, 51-57.	3.0	6
1998	Isolation of Coxiella burnetii from an acromioclavicular infection with low serological titres. International Journal of Infectious Diseases, 2018, 73, 27-29.	3.3	6

#	Article	IF	CITATIONS
1999	Anaerococcus jeddahensis sp. nov., a New Bacterial Species Isolated From Healthy Nomadic Bedouin Woman From Saudi Arabia. Current Microbiology, 2018, 75, 1419-1428.	2.2	6
2000	Dysgonomonas massiliensis sp. nov., a new species isolated from the human gut and its taxonogenomic description. Antonie Van Leeuwenhoek, 2019, 112, 935-945.	1.7	6
2001	Description of Janibacter massiliensis sp. nov., cultured from the vaginal discharge of a patient with bacterial vaginosis. Antonie Van Leeuwenhoek, 2019, 112, 1147-1159.	1.7	6
2002	Longitudinal monitoring of environmental factors at Culicidae larval habitats in urban areas and their association with various mosquito species using an innovative strategy. Pest Management Science, 2019, 75, 923-934.	3.4	6
2003	Complications of peripheral venous catheters: The need to propose an alternative route of administration. International Journal of Antimicrobial Agents, 2020, 55, 105875.	2.5	6
2004	Temporal and age distributions of SARS-CoV-2 and other coronaviruses, southeastern France. International Journal of Infectious Diseases, 2020, 101, 121-125.	3.3	6
2005	Culturing Ancient Bacteria Carrying Resistance Genes from Permafrost and Comparative Genomics with Modern Isolates. Microorganisms, 2020, 8, 1522.	3.6	6
2006	Hymenopteran Parasitoids of Hard Ticks in Western Africa and the Russian Far East. Microorganisms, 2020, 8, 1992.	3.6	6
2007	Biological Control of Aedes albopictus: Obtained from the New Bacterial Candidates with Insecticidal Activity. Insects, 2020, 11, 403.	2.2	6
2008	Molecular detection of microorganisms in lice collected from farm animals in Northeastern Algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2021, 74, 101569.	1.6	6
2009	Rational for meta-analysis and randomized treatment: the COVID-19 example. Clinical Microbiology and Infection, 2021, 27, 6-8.	6.0	6
2010	Epidemiological and genetic characterization of measles virus circulating strains at Marseille, France during 2017–2019 measles outbreak. Journal of Infection, 2021, 83, 361-370.	3.3	6
2011	Molecular phylogeny of the genus Bartonella: what is the current knowledge?. FEMS Microbiology Letters, 2001, 200, 1-7.	1.8	6
2012	Evaluation of self-collected rectal swabs for the detection of bacteria responsible for sexually transmitted infections in a cohort of HIV-1-infected patients. Journal of Medical Microbiology, 2017, 66, 693-697.	1.8	6
2013	Parallel Decline of Malaria and Rickettsia felis Infections in Senegal. American Journal of Tropical Medicine and Hygiene, 2018, 99, 360-361.	1.4	6
2014	Collinsella ihumii sp. nov., a new anaerobic bacterium isolated from human stool. Archives of Microbiology, 2021, 203, 6315-6322.	2.2	6
2015	Arabiibacter massiliensis gen. nov. sp. nov., New Anaerobic Bacterium Isolated from the Human Gut. Current Microbiology, 2022, 79, 47.	2.2	6
2016	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, 58-9.	1.4	6

#	Article	IF	CITATIONS
2017	Autochthonous epidemic typhus associated with Bartonella quintana bacteremia in a homeless person. American Journal of Tropical Medicine and Hygiene, 2005, 72, 638-9.	1.4	6
2018	Impact of Sex Hormones on Macrophage Responses to Coxiella burnetii. Frontiers in Immunology, 2021, 12, 705088.	4.8	6
2019	Neglectibacter timonensis gen. nov., sp. nov. and Scatolibacter rhodanostii gen. nov., sp. nov., two anaerobic bacteria isolated from human stool samples. Archives of Microbiology, 2022, 204, 45.	2.2	6
2020	Control of common viral epidemics but not of SARS-CoV-2 through the application of hygiene and distancing measures. Journal of Clinical Virology, 2022, 150-151, 105163.	3.1	6
2021	Serologic Diagnosis of Leptospirosis: Comparison of Line Blot and Immunofluorescence Techniques with the Genus-Specific Microscopic Agglutination Test. Journal of Infectious Diseases, 1989, 160, 734-735.	4.0	5
2022	Serotyping Isolates of Anaplasma phagocytophilum by Using Monoclonal Antibodies. Vaccine Journal, 2003, 10, 969-972.	3.1	5
2023	Molecular characterization of resistance to fluoroquinolones in Bartonella henselae and Bartonella quintana. Journal of Antimicrobial Chemotherapy, 2009, 63, 1288-1289.	3.0	5
2024	Recent and future developments in the epidemiology of the infectious diseases. European Journal of Epidemiology, 2009, 24, 393-395.	5.7	5
2025	Global proteomic pattern of <b><i>Tropheryma whipplei</i></b> : A Whipple's disease bacterium. Proteomics, 2009, 9, 1593-1616.	2.2	5
2026	Q fever: a case with a vascular infection complication. BMJ Case Reports, 2010, 2010, bcr0120102690-bcr0120102690.	0.5	5
2027	Viral Endocarditis or Simple Viral Disseminated Infection?. Clinical Infectious Diseases, 2011, 53, 1298-1298.	5.8	5
2028	Genome Sequence of Bartonella birtlesii, a Bacterium Isolated from Small Rodents of the Genus Apodemus. Journal of Bacteriology, 2012, 194, 4779-4779.	2.2	5
2029	Genomic Analysis of Rickettsia japonica Strain YH <sup>T</sup> . Journal of Bacteriology, 2012, 194, 6992-6992.	2.2	5
2030	Megavirales Composing a Fourth Domain of Life: Mimiviridae and Marseilleviridae., 2012,, 217-244.		5
2031	Amazonian Head Lice-Specific Genotypes Are Putatively Pre-Columbian. American Journal of Tropical Medicine and Hygiene, 2013, 88, 1180-1184.	1.4	5
2032	Historical and geographical parallelism between the incidence of dental caries, Streptococcus mutans and sugar intake. European Journal of Epidemiology, 2013, 28, 709-710.	5.7	5
2033	Effect of omeprazole on vacuole size in Coxiella burnetii-infected cells. Journal of Infection, 2013, 66, 288-289.	3.3	5
2034	Infectious endocarditis detected by PET/CT in a patient with a prosthetic knee infection: Case report and review of the literature. Scandinavian Journal of Infectious Diseases, 2013, 45, 570-574.	1.5	5

#	Article	IF	CITATIONS
2035	Non-contiguous genome sequence of Mycobacterium simiae strain DSM 44165T. Standards in Genomic Sciences, 2013, 8, 306-317.	1.5	5
2036	Draft Genome Sequence of Coxiella burnetii Strain Cb196, an Agent of Endocarditis in Saudi Arabia. Genome Announcements, 2014, 2, .	0.8	5
2037	Genome sequence and description of Pantoea septica strain FF5. Standards in Genomic Sciences, 2015, 10, 103.	1.5	5
2038	Ancient Resistome., 2016,, 75-80.		5
2039	Absence of serological evidence of Rickettsia spp., Bartonella spp., Ehrlichia spp. and Coxiella burnetii infections in American Samoa. Ticks and Tick-borne Diseases, 2016, 7, 703-705.	2.7	5
2040	A novel ehrlichial agent detected in tick in French Polynesia. Ticks and Tick-borne Diseases, 2016, 7, 1203-1208.	2.7	5
2041	"Murdochiella vaginalis―sp. nov., a new bacterial species cultivated from the vaginal flora of a woman with bacterial vaginosis. Human Microbiome Journal, 2016, 2, 15-16.	3.8	5
2042	Non-contiguous-Finished Genome Sequence and Description of Paenibacillus camerounensis sp. nov Microbial Ecology, 2016, 71, 990-998.	2.8	5
2043	High Prevalence of Mycoplasma faucium DNA in the Human Oropharynx. Journal of Clinical Microbiology, 2016, 54, 194-196.	3.9	5
2044	Fatal Neisseria macacae infective endocarditis: first report. Infection, 2017, 45, 369-371.	4.7	5
2045	Dental pulp as a source of low-contaminated DNA. Microbial Pathogenesis, 2017, 105, 63-67.	2.9	5
2046	$\hat{A}$ «Veillonella massiliensis $\hat{A}$ », a new anaerobic species isolated from human colostrum. Human Microbiome Journal, 2017, 4, 20-21.	3.8	5
2047	Autochthonous hepatitis E: a common and fatal but neglected emerging disease in France. Clinical Microbiology and Infection, 2017, 23, 898-899.	6.0	5
2048	Treating influenza with antibiotics. International Journal of Antimicrobial Agents, 2017, 50, 505-506.	2.5	5
2049	Blood Culture–Negative Endocarditis, Morocco. Emerging Infectious Diseases, 2017, 23, 1908-1909.	4.3	5
2050	Kocuria massiliensis sp. nov, a new bacterial species isolated from a patient with foot osteomyelitis. Infection, 2018, 46, 259-262.	4.7	5
2051	Comparing two blood culture systems for the detection of bacterial contamination in platelet concentrates. Transfusion, 2018, 58, 2604-2610.	1.6	5
2052	Did Caravaggio die of Staphylococcus aureus sepsis?. Lancet Infectious Diseases, The, 2018, 18, 1178.	9.1	5

#	Article	IF	Citations
2053	<i>Eggerthella timonensis</i> sp. nov, a new species isolated from the stool sample of a pygmy female. MicrobiologyOpen, 2018, 7, e00575.	3.0	5
2054	Experimental Inoculation in Rats and Mice by the Giant Marseillevirus Leads to Long-Term Detection of Virus. Frontiers in Microbiology, 2018, 9, 463.	3.5	5
2055	Mesenteric lymphadenitis as a presenting feature of Whipple's disease: Value of PCR analysis. International Journal of Infectious Diseases, 2018, 75, 15-17.	3.3	5
2056	The Paradigm of the Shadoks and Antibiotic Resistance. Clinical Infectious Diseases, 2019, 69, 1641-1641.	5.8	5
2057	Identification of mixed and successive blood meals of mosquitoes using MALDI-TOF MS protein profiling. Parasitology, 2020, 147, 329-339.	1.5	5
2058	FDG-PET/CT Incidental Detection of Cancer in Patients Investigated for Infective Endocarditis. Frontiers in Medicine, 2020, 7, 535.	2.6	5
2059	T-Bet Controls Susceptibility of Mice to Coxiella burnetii Infection. Frontiers in Microbiology, 2020, 11, 1546.	3.5	5
2060	Major discrepancy between factual antibiotic resistance and consumption in South of France: analysis of 539,037 bacterial strains. Scientific Reports, 2020, 10, 18262.	3.3	5
2061	Promiscuous Enzyme Activity as a Driver of Allo and Iso Convergent Evolution, Lessons from the $\hat{l}^2$ -Lactamases. International Journal of Molecular Sciences, 2020, 21, 6260.	4.1	5
2062	Does spitting in public play a role in transmitting SARS-CoV-2?. Travel Medicine and Infectious Disease, 2020, 36, 101759.	3.0	5
2063	Archeomicrobiology applied to environmental samples. Microbial Pathogenesis, 2020, 143, 104140.	2.9	5
2064	Can hydroxychloroquine be protective against COVID-19-associated thrombotic events?. Journal of Microbiology, Immunology and Infection, 2021, 54, 37-45.	3.1	5
2065	Pedobacter ghigonii sp. nov., Isolated from the Microbiota of the Planarian Schmidtea mediterranea. Microbiology Research, 2021, 12, 268-287.	1.9	5
2066	Assessment of the burden of malaria and bacteraemia by retrospective molecular diagnosis in febrile illnesses and first-line anti-infectives in CÃ te d'Ivoire. Travel Medicine and Infectious Disease, 2021, 43, 102105.	3.0	5
2067	Description of strain FC3T as the neotype strain of Actinobaculum massiliense. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 2702-2703.	1.7	5
2068	Whole-Genome Sequence of French Clinical Peptoniphilus catoniae Strain P8546. Microbiology Resource Announcements, 2019, 8, .	0.6	5
2069	Improving the diagnostic efficiency of H1N1 2009 pandemic flu: analysis of predictive clinical signs through a prospective cohort PLOS Currents, 2009, 1, RRN1120.	1.4	5
2070	Rapid Diagnosis of Lung Tumors, a Feasability Study Using Maldi-Tof Mass Spectrometry. PLoS ONE, 2016, 11, e0155449.	2.5	5

#	Article	lF	CITATIONS
2071	Bartonellaspp. Bacteremia and Rheumatic Symptoms in Patients from Lyme Disease–endemic Region. Emerging Infectious Diseases, 2012, 18, 1919a-1919.	4.3	5
2072	Diagnosis of Louse-Borne Relapsing Fever despite Negative Microscopy in Two Asylum Seekers from Eastern Africa. American Journal of Tropical Medicine and Hygiene, 2017, 97, 1669-1672.	1.4	5
2073	High Circulation of Malaria and Low Prevalence of Bacteremia in Febrile and Afebrile Children in Northeastern Gabon. American Journal of Tropical Medicine and Hygiene, 2020, 102, 121-129.	1.4	5
2074	Isolation of 4000 SARS-CoV-2 shows that contagiousness is associated with viral load, not vaccine or symptomatic status. Emerging Microbes and Infections, 2021, 10, 2276-2278.	6.5	5
2075	Buttiauxella massiliensis sp. nov., Isolated from a Human Bone Infection. Current Microbiology, 2022, 79, 41.	2.2	5
2076	SARS-CoV-2 Vaccination and Protection Against Clinical Disease: A Retrospective Study, Bouches-du-RhĀ'ne District, Southern France, 2021. Frontiers in Microbiology, 2021, 12, 796807.	3.5	5
2077	The emergence, dynamics and significance of SARS-CoV-2 variants. New Microbes and New Infections, 2022, 45, 100962.	1.6	5
2078	Population Diversity of Antibiotic Resistant Enterobacterales in Samples From Wildlife Origin in Senegal: Identification of a Multidrug Resistance Transposon Carrying blaCTX–M–15 in Escherichia coli. Frontiers in Microbiology, 2022, 13, 838392.	3.5	5
2079	Description of <i>Acinetobacter ihumii</i> sp. nov., <i>Microbacterium ihumii</i> sp. nov., and <i>Gulosibacter massiliensis</i> sp. nov., three new bacteria isolated from human blood. FEMS Microbiology Letters, 2022, 369, .	1.8	5
2080	New Beta-lactamases in Candidate Phyla Radiation: Owning Pleiotropic Enzymes Is a Smart Paradigm for Microorganisms with a Reduced Genome. International Journal of Molecular Sciences, 2022, 23, 5446.	4.1	5
2081	Outcomes of 2111 COVID-19 Hospitalized Patients Treated with Hydroxychloroquine/Azithromycin and Other Regimens in Marseille, France, 2020: A Monocentric Retrospective Analysis. Therapeutics and Clinical Risk Management, 0, Volume 18, 603-617.	2.0	5
2082	Are the clinical effects of homoeopathy placebo effects? â€" Authors' reply. Lancet, The, 2005, 366, 2085-2086.	13.7	4
2083	Corpuscular Antigenic Microarray for the Serodiagnosis of Blood Culture-Negative Endocarditis. Annals of the New York Academy of Sciences, 2006, 1078, 595-596.	3.8	4
2084	<i>&gt;Bartonella quintana</i> , Lice, and Molecular Tools. Journal of Medical Entomology, 2006, 43, 787-787.	1.8	4
2085	Creationism—remember the principle of falsifiability. Lancet, The, 2008, 372, 2095-2096.	13.7	4
2086	Emerging Rickettsioses Reach the United States. Clinical Infectious Diseases, 2010, 51, 121-122.	5.8	4
2087	Draft Genome Sequence of Actinomyces massiliensis Strain 4401292T. Journal of Bacteriology, 2012, 194, 5121-5121.	2.2	4
2088	Genome Sequence of Legionella tunisiensis Strain LegM <sup>T</sup> , a New Legionella Species Isolated from Hypersaline Lake Water. Journal of Bacteriology, 2012, 194, 5978-5978.	2.2	4

#	Article	IF	CITATIONS
2089	Genome Sequence of Lactobacillus ingluviei, a Bacterium Associated with Weight Gain in Animals. Journal of Bacteriology, 2012, 194, 5697-5697.	2.2	4
2090	Rapid MALDI-TOF mass spectrometry identification of Leptospira organisms: A reply. Veterinary Microbiology, 2012, 159, 544.	1.9	4
2091	Severe Whipple's disease with acute myocarditis. International Journal of Cardiology, 2012, 159, e41-e42.	1.7	4
2092	Rare occurrence of Whipple Disease in a young female patient with a fatal outcome. Egyptian Journal of Forensic Sciences, 2013, 3, 85-91.	1.0	4
2093	Health-care-associated bloodstream infections in France. Lancet Infectious Diseases, The, 2013, 13, 656.	9.1	4
2094	Vitamin D and Prolonged Treatment with Photosensitivity-Associated Antibiotics. Antimicrobial Agents and Chemotherapy, 2013, 57, 6409-6410.	3.2	4
2095	Draft Genome Sequence of Mycobacterium vulneris DSM 45247 <sup>T</sup> . Genome Announcements, 2014, 2, .	0.8	4
2096	Draft Genome Sequence of Rickettsia aeschlimannii, Associated with Hyalomma marginatum Ticks. Genome Announcements, $2014, 2, \ldots$	0.8	4
2097	Genome Sequence of Borrelia crocidurae Strain 03-02, a Clinical Isolate from Senegal. Genome Announcements, 2014, 2, .	0.8	4
2098	Genome Sequence of Rickettsia hoogstraalii, a Geographically Widely Distributed Tick-Associated Bacterium. Genome Announcements, $2014, 2, \ldots$	0.8	4
2099	Reply to Kampschreur et al. Clinical Infectious Diseases, 2014, 58, 447-448.	5.8	4
2100	Increasing Trend of Invasive Group B Streptococcal Infections, Marseille, France. Clinical Infectious Diseases, 2014, 58, 750-751.	5.8	4
2101	Fetuin is the key for nanon self-propagation. Microbial Pathogenesis, 2014, 73, 25-30.	2.9	4
2102	Genome Sequence of Legionella anisa, Isolated from a Respiratory Sample, Using an Amoebal Coculture Procedure. Genome Announcements, 2014, 2, .	0.8	4
2103	Clostridium polynesiense sp. nov., a new member of the human gut microbiota in French Polynesia. Anaerobe, 2015, 36, 79-87.	2.1	4
2104	Draft Genome Sequence of the Lactobacillus agilis Strain Marseille. Genome Announcements, 2015, 3, .	0.8	4
2105	Draft Genome Sequence of the Lactobacillus mucosae Strain Marseille. Genome Announcements, 2015, 3, .	0.8	4
2106	Noneruptive Fever Revealing Murine Typhus in a Traveler Returning From Tunisia. Journal of Travel Medicine, 2015, 22, 67-69.	3.0	4

#	Article	IF	CITATIONS
2107	Response to a Letter to the Editor by Joachim Denner on HIV infection en route to endogenization: two cases. Clinical Microbiology and Infection, 2015, 21, e35-e37.	6.0	4
2108	Antibiotic prophylaxis of endocarditis. Lancet Infectious Diseases, The, 2016, 16, 773-774.	9.1	4
2109	Hydroxychloroquine susceptibility determination of Coxiella burnetii in human embryonic lung (HEL) fibroblast cells. International Journal of Antimicrobial Agents, 2017, 50, 106-109.	2.5	4
2110	Draft Genome and Description of Eisenbergiella massiliensis Strain AT11T: A New Species Isolated from Human Feces After Bariatric Surgery. Current Microbiology, 2018, 75, 1274-1281.	2.2	4
2111	Limbic encephalitis as a relapse of Whipple's disease with digestive involvement and spondylodiscitis. Infection, 2019, 47, 637-641.	4.7	4
2112	IGF1 levels in children with severe acute malnutrition after nutritional recovery: A good predictor for children's long-term health status. EBioMedicine, 2019, 45, 9-10.	6.1	4
2113	Cellulitis of the face associated with SENLAT caused by Rickettsia slovaca detected by qPCR on scalp eschar swab sample: An unusual case report and review of literature. Ticks and Tick-borne Diseases, 2019, 10, 1142-1145.	2.7	4
2114	Management of patients presenting with generalized musculoskeletal pain and a suspicion of Lyme disease. Médecine Et Maladies Infectieuses, 2019, 49, 157-166.	5.0	4
2115	Oceanobacillus timonensis sp. nov. and Oceanobacillus senegalensis sp. nov., two new moderately halophilic, Gram-stain positive bacteria isolated from stools sample of healthy young Senegalese. Antonie Van Leeuwenhoek, 2019, 112, 785-796.	1.7	4
2116	Fluorescent in situ hybridization can be used as a complementary assay for the diagnosis of Tropheryma whipplei infection. Infection, 2019, 47, 317-321.	4.7	4
2117	Tick-Borne Spotted Fever Rickettsioses. , 2020, , 587-593.		4
2118	Coxiella burnetii endocarditis as a possible cause of ANCA-associated vasculitis. Rheumatology, 2020, 59, e44-e45.	1.9	4
2119	Sunbathing, a possible risk factor of murine typhus infection in Greece. PLoS Neglected Tropical Diseases, 2021, 15, e0009186.	3.0	4
2120	Quantitative Analysis System for Bacterial Cells in SEM Image using Deep Learning., 2021,,.		4
2121	Stool Serology: Development of a Non-Invasive Immunological Method for the Detection of Enterovirus-Specific Antibodies in Congo Gorilla Faeces. Microorganisms, 2021, 9, 810.	3.6	4
2122	Evaluation of Strategies to Fight COVID-19: The French Paradigm. Journal of Clinical Medicine, 2021, 10, 2942.	2.4	4
2123	Vitreoscilla massiliensis sp. nov., Isolated From the Stool of an Amazonian Patient. Current Microbiology, 2021, 78, 3313-3320.	2.2	4
2124	Whole Genome Sequencing of SARS-CoV-2 Strains in COVID-19 Patients From Djibouti Shows Novel Mutations and Clades Replacing Over Time. Frontiers in Medicine, 2021, 8, 737602.	2.6	4

#	Article	IF	Citations
2125	Could β-Lactam Antibiotics Block Humoral Immunity?. Frontiers in Immunology, 2021, 12, 680146.	4.8	4
2126	Orientia tsutsugamushi (Scrub Typhus). , 2010, , 2529-2530.		4
2127	Dynamics and genetic diversity of Haemophilus influenzae carriage among French pilgrims during the 2018 Hajj: A prospective cohort survey. Travel Medicine and Infectious Disease, 2020, 38, 101883.	3.0	4
2128	Predominant Immunoglobulin A Response to Phase II Antigen of Coxiella burnetii in Acute Q Fever. Vaccine Journal, 1999, 6, 173-177.	2.6	4
2129	Flea-Borne Spotted Fever. Infectious Disease and Therapy, 2007, , 87-96.	0.0	4
2130	<i>Rickettsia conorii</i> Infections (Mediterranean Spotted Fever, Israeli Spotted Fever, Indian Tick) Tj ETQq0 0 0	rgBT/Ove	rlock 10 Tf 5
2131	Diagnostic Strategy of Rickettsioses and Ehrlichioses. Infectious Disease and Therapy, 2007, , 315-330.	0.0	4
2132	Point of Care strategy for rapid diagnosis of novel A/H1N1 influenza virus PLOS Currents, 2009, 1, RRN1039.	1.4	4
2133	Using MALDI-TOF spectra in epidemiological surveillance for the detection of bacterial subgroups with a possible epidemic potential. BMC Infectious Diseases, 2021, 21, 1109.	2.9	4
2134	Long-term persistence of symptoms of dyspnoea in COVID-19 patients. International Journal of Infectious Diseases, 2021, , .	3.3	4
2135	Peptostreptococcus faecalis sp. nov., new bacterial species isolated from healthy indigenous congolese volunteer. Heliyon, 2022, 8, e09102.	3.2	4
2136	Serosurvey for Cowdria ruminantium, Coxiella burnetii, and Spotted Fever Group Rickettsiae in Ostriches (Struthio camelus) from Zimbabwe. Avian Diseases, 1996, 40, 448.	1.0	3
2137	Antimicrobial activity against obligate intracellular bacteria. Trends in Microbiology, 2001, 9, 14.	7.7	3
2138	Was the Black Death yersinial plague?. Lancet Infectious Diseases, The, 2003, 3, 328.	9.1	3
2139	Intracellular Life of Coxiella burnetii in Macrophages: Insight into Q Fever Immunopathology. Current Immunology Reviews, 2006, 2, 225-232.	1.2	3
2140	18F-Fluorodeoxyglucose positron emission tomography in Whipple's disease. Scandinavian Journal of Gastroenterology, 2006, 41, 1491-1492.	1.5	3
2141	What makes a virus a virus: reply from Raoult and Forterre. Nature Reviews Microbiology, 2008, 6, 643-643.	28.6	3
2142	Pericardial effusion as the only manifestation of infection with Francisella tularensis: a case report. Journal of Medical Case Reports, 2008, 2, 206.	0.8	3

#	Article	IF	CITATIONS
2143	Q Fever, Free Amoeba, and Air Conditioning. Clinical Infectious Diseases, 2010, 51, 869-870.	5.8	3
2144	Body Lice, Yersinia pestis Orientalis, and Black Death. Emerging Infectious Diseases, 2010, 16, 1650-1651.	4.3	3
2145	Bartonella clarridgeiaein Fleas, Tahiti, French Polynesia. Emerging Infectious Diseases, 2011, 17, 1773-1775.	4.3	3
2146	Genome Sequence of Afipia birgiae, a Rare Bacterium Associated with Amoebae. Journal of Bacteriology, 2012, 194, 7018-7018.	2.2	3
2147	Lack of knowledge can anger patients with chronic diseases. Lancet Infectious Diseases, The, 2012, 12, 654-655.	9.1	3
2148	Birds perching on bushes: Networks to visualize conflicting phylogenetic signals during early avian radiation. Comptes Rendus - Palevol, 2013, 12, 333-337.	0.2	3
2149	Tick-borne Spotted Fever Rickettsioses. , 2013, , 546-552.		3
2150	Noncontiguous Genome Sequence of Mycobacterium septicum Strain DSM 44393 T. Genome Announcements, 2013, 1, .	0.8	3
2151	Draft Genome Sequence of Mycobacterium austroafricanum DSM 44191. Genome Announcements, 2014, 2, .	0.8	3
2152	Editorial: Emerging clones of bacterial epidemics in the genomic area. Clinical Microbiology and Infection, 2014, 20, 371-372.	6.0	3
2153	Throat Swab Samples for Diagnosis of Q Fever. American Journal of Tropical Medicine and Hygiene, 2014, 90, 147-148.	1.4	3
2154	Obtaining informed consent in pediatric clinical trials. Journal of Clinical Epidemiology, 2014, 67, 840-841.	5.0	3
2155	Draft Genome Sequence of Necropsobacter rosorum Strain P709 $\sup T. Genome Announcements, 2014, 2, .$	0.8	3
2156	Kingella Kingae DNA in Langerhans Cell Histiocytosis of Bone. Pediatric Infectious Disease Journal, 2015, 34, 317-318.	2.0	3
2157	Coxiella burnetii (Q Fever). , 2015, , 2208-2216.e2.		3
2158	Infective endocarditis and antibiotic prophylaxis. Lancet, The, 2015, 386, 528.	13.7	3
2159	Influenza-attributable deaths in south-eastern France (1999 to 2010): mortality predictions were undependable. BMC Public Health, 2015, 15, 539.	2.9	3
2160	Positron emission tomography to diagnose chronic Q fever. Médecine Et Maladies Infectieuses, 2015, 45, 420-422.	5.0	3

#	Article	IF	CITATIONS
2161	Emerging Tick-Borne Bacterial Pathogens. , 2016, , 295-310.		3
2162	History of Smallpox and Its Spread in Human Populations. , 0, , 161-172.		3
2163	Prosthetic Valve Endocarditis Caused by Bartonella henselae: A Case Report of Molecular Diagnostics Informing Nonsurgical Management. Open Forum Infectious Diseases, 2016, 3, ofw202.	0.9	3
2164	Draft Genome Sequence of Actinobaculum massiliense Strain FC3. Genome Announcements, 2016, 4, .	0.8	3
2165	New Diagnostic Techniques Highlight the Need for Negative Controls. Clinical Infectious Diseases, 2016, 62, 809.2-809.	5.8	3
2166	High serum CXCL10 in Rickettsia conorii infection is endothelial cell mediated subsequent to whole blood activation. Cytokine, 2016, 83, 269-274.	3.2	3
2167	The mobile phone as a vector for virus RNA: a link with Ebola outbreaks?. Clinical Microbiology and Infection, 2016, 22, 401.	6.0	3
2168	New Laboratory Tools for Emerging Bacterial Challenges. Clinical Infectious Diseases, 2017, 65, S39-S49.	5.8	3
2169	The risk of bioterrorism re-analysed. Clinical Microbiology and Infection, 2017, 23, 351.	6.0	3
2170	Acute Septic Arthritis of the Knee Caused by Kingella kingae in a 5-Year-Old Cameroonian Boy. Frontiers in Pediatrics, 2017, 5, 230.	1.9	3
2171	Marseille scoring system for empiric treatment of infective endocarditis. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 841-849.	2.9	3
2172	Sediminibacillus massiliensis sp. nov., a moderately halophilic, Gram-positive bacterium isolated from a stool sample of a young Senegalese man. Antonie Van Leeuwenhoek, 2018, 111, 1225-1236.	1.7	3
2173	Prevalence of <i>Anaplasmataceae</i> and <i>Filariidae</i> species in unowned and military dogs in New Caledonia. Veterinary Medicine and Science, 2018, 4, 140-149.	1.6	3
2174	First Case of Q Fever Endocarditis Involving the Melody $\hat{A}^{\otimes}$ Transcatheter Pulmonary Valve in an Afebrile Child. Pediatric Cardiology, 2018, 39, 195-197.	1.3	3
2175	Blood Culture–Negative Cardiovascular Infection in a Patient With Multiple Sclerosis. Open Forum Infectious Diseases, 2019, 6, ofz429.	0.9	3
2176	Progenitor mast cells and tryptase in Q fever. Comparative Immunology, Microbiology and Infectious Diseases, 2019, 64, 159-162.	1.6	3
2177	Functional neuroimaging in patients presenting with somatoform disorders: A model for investigating persisting symptoms after tick bites and post-treatment Lyme disease syndrome?. Médecine Et Maladies Infectieuses, 2019, 49, 150-156.	5.0	3
2178	Why do arthropods secrete β-lactams?. International Journal of Antimicrobial Agents, 2019, 53, 370.	2.5	3

#	Article	IF	CITATIONS
2179	Lack of Vibrio cholerae among French pilgrims during the 2017 and 2018 Hajj. Travel Medicine and Infectious Disease, 2020, 36, 101506.	3.0	3
2180	Lancet gate: a matter of fact or a matter of concern. New Microbes and New Infections, 2020, 38, 100758.	1.6	3
2181	Pathogens associated with respiratory, gastrointestinal and febrile illness in patients consulting at Mbacke healthcare centre during the 2018 Grand Magal of Touba: A preliminary study. Travel Medicine and Infectious Disease, 2020, 37, 101820.	3.0	3
2182	Fenollaria timonensis sp. nov., A New Bacterium Isolated from Healthy Human Fresh Stool. Current Microbiology, 2020, 77, 3780-3786.	2.2	3
2183	Investigation of <i>Ctenocephalides felis</i> on domestic dogs and <i>Rickettsia felis</i> infection in the Democratic Republic of Sao Tome and Principe. Zoonoses and Public Health, 2020, 67, 892-902.	2.2	3
2184	A sporadic case of acute Q fever and identification of the animal source of the infection. Folia Microbiologica, 2020, 65, 797-800.	2.3	3
2185	Chloroquine and COVID-19: A western medical and scientific drift?. European Journal of Internal Medicine, 2020, 78, 4-5.	2.2	3
2186	Dramatic HIV DNA degradation associated with spontaneous HIV suppression and disease-free outcome in a young seropositive woman following her infection. Scientific Reports, 2020, 10, 2548.	3.3	3
2187	Persistent Coxiella burnetii cardiovascular infection on Bentall-De Bono prosthesis. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1003-1010.	2.9	3
2188	High-Content Screening, a Reliable System for Coxiella burnetii Isolation from Clinical Samples. Journal of Clinical Microbiology, 2020, 58, .	3.9	3
2189	Full-repertoire comparison of the microscopic objects composing the human gut microbiome with sequenced and cultured communities. Journal of Microbiology, 2020, 58, 377-386.	2.8	3
2190	Clinical efficacy and safety profile of hydroxychloroquine and azithromycin against COVID-19. International Journal of Antimicrobial Agents, 2021, 57, 106242.	2.5	3
2191	Draft Genome Sequence of Herminiimonas contaminans Strain CCM 7991 <sup>T</sup> , a Biopharmaceutical Contaminant. Microbiology Resource Announcements, 2021, 10, .	0.6	3
2192	Sputum proteomic analysis for distinguishing between pulmonary tuberculosis and non-tuberculosis using matrix-assisted laser desorption ionization time-of-flight mass spectrometry (MALDI-TOF MS): preliminary results. Clinical Microbiology and Infection, 2021, 27, 1694.e1-1694.e6.	6.0	3
2193	Is minor surgery safe during the COVID-19 pandemic? A multi-disciplinary study. PLoS ONE, 2021, 16, e0251122.	2.5	3
2194	Clostridium culturomicium sp. nov. and Clostridium jeddahitimonense sp. nov., novel members of the Clostridium genus isolated from the stool of an obese Saudi Arabian. Current Microbiology, 2021, 78, 3586-3595.	2.2	3
2195	Consequences of the COVID-19 Outbreak Lockdown on Non-Viral Infectious Agents as Reported by a Laboratory-Based Surveillance System at the IHU Méditerranée Infection, Marseille, France. Journal of Clinical Medicine, 2021, 10, 3210.	2.4	3
2196	Draft Genome Sequence of Bacillus velezensis Strain Marseille-Q1230, Isolated from a Stool Sample from a Severely Malnourished Child. Microbiology Resource Announcements, 2021, 10, e0051421.	0.6	3

#	Article	IF	CITATIONS
2197	Detection of <i>Coxiella burnetii</i> and <i>Borrelia</i> spp. DNA in Cutaneous Samples and in Household Dust in Rural Areas, Senegal. Vector-Borne and Zoonotic Diseases, 2021, 21, 659-666.	1.5	3
2198	In vitro detection of bacterial contamination in platelet concentrates by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry: a preliminary study. Journal of Medical Microbiology, 2017, 66, 1523-1530.	1.8	3
2199	Pro-apoptotic effect of doxycycline and hydroxychloroquine on B-cell lymphoma induced by C. burnetii. Oncotarget, 2018, 9, 2726-2727.	1.8	3
2200	Louse-Borne Epidemic Typhus. Infectious Disease and Therapy, 2007, , 51-62.	0.0	3
2201	Rickettsia sibirica mongolitimonae infection, Sri Lanka. Journal of Infection in Developing Countries, 2017, 11, 668-671.	1.2	3
2202	First case of imported African tick-bite fever in Poland – Case report. Annals of Agricultural and Environmental Medicine, 2015, 22, 412-413.	1.0	3
2203	Introduction to Rickettsioses, Ehrlichioses, and Anaplasmosis. , 2015, , 2194-2197.		3
2204	Case Report: Vibrio cholerae Biliary Tract Infections in Two North Africans in France. American Journal of Tropical Medicine and Hygiene, 2020, 102, 1306-1308.	1.4	3
2205	Establishing Medical Coverage and Epidemiological Surveillance during the Grand Magal of Touba in Senegal: A Public Health Need. Journal of Epidemiology and Global Health, 2020, 10, 247.	2.9	3
2206	Limited spread of a rare spike E484K-harboring SARS-CoV-2 in Marseille, France. Archives of Virology, 2022, 167, 583.	2.1	3
2207	Introduction of the SARS-CoV-2 Beta variant from Comoros into the Marseille geographical area. Travel Medicine and Infectious Disease, 2022, 46, 102277.	3.0	3
2208	Putative native South Amerindian origin of head lice clade F: evidence from head lice nits infesting human shrunken heads. Scientific Reports, 2022, 12, 4307.	3.3	3
2209	Reverse Genomics: Design of Universal Epitope Sets to Isolate All Saccharibacteria Members from the Human Oral Cavity. Microorganisms, 2022, 10, 602.	3.6	3
2210	High Genetic Diversity and Rickettsia felis in Pediculus humanus Lice Infesting Mbuti (pygmy people), -Democratic Republic of Congo. Frontiers in Cellular and Infection Microbiology, 2022, 12, 834388.	3.9	3
2211	Life-years lost by COVID-19 patients in public hospitals of Marseille (APHM-South-Eastern France): a limited death toll: a retrospective analysis. BMJ Open, 2021, 11, e049475.	1.9	3
2212	Human infections caused by bartonella spp. Part 2. Clinical Microbiology Newsletter, 2000, 22, 9-13.	0.7	2
2213	No Serological Evidence for Rickettsial Diseases among Danish Elite Orienteerers. Annals of the New York Academy of Sciences, 2006, 1078, 150-153.	3.8	2
2214	Doxycycline for Mansonella perstans Infection. New England Journal of Medicine, 2010, 362, 272-273.	27.0	2

#	Article	IF	CITATIONS
2215	Strange World of Emergency Medicine. Journal of Emergency Medicine, 2010, 39, 501.	0.7	2
2216	Being careful with PCR to avoid erroneous discoveries. Scandinavian Journal of Infectious Diseases, 2011, 43, 323-324.	1.5	2
2217	Typhus Group Rickettsioses., 2011,, 329-333.		2
2218	Genome Sequence of Bartonella rattaustraliani, a Bacterium Isolated from an Australian Rat. Journal of Bacteriology, 2012, 194, 7012-7012.	2.2	2
2219	Cross-protection among Rickettsia species and subspecies in a guinea pig model of cutaneous infection. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 551-556.	1.6	2
2220	Genome Sequence of Bartonella rattimassiliensis, a Bacterium Isolated from European Rattus norvegicus. Journal of Bacteriology, 2012, 194, 7013-7013.	2.2	2
2221	Rickettsial Infections., 2012,, 1954-1964.		2
2222	A Need to Discover the World of Giant Viruses. Intervirology, 2013, 56, 347-348.	2.8	2
2223	Non-contiguous finished genome sequence of Phocaeicola abscessus type strain 7401987T. Standards in Genomic Sciences, 2013, 9, 351-358.	1.5	2
2224	Kingella kingae KK247, an Atypical Pulsed-Field Gel Electrophoresis Clone A Strain. Genome Announcements, 2014, 2, .	0.8	2
2225	Reply to Sauvage et al. Journal of Infectious Diseases, 2014, 210, 2018-2019.	4.0	2
2226	Draft Genome Sequence of Mycobacterium farcinogenes NCTC 10955. Genome Announcements, 2014, 2, .	0.8	2
2227	Draft Genome Sequence of Mycobacterium neoaurum Strain DSM 44074 <sup>T</sup> . Genome Announcements, 2014, 2, .	0.8	2
2228	Whipple's disease: surprised by the surprise. Lancet, The, 2014, 384, 1184-1185.	13.7	2
2229	Increase in sexually transmitted infections during Europride 2013 in Marseille, France. Lancet Infectious Diseases, The, 2014, 14, 677-678.	9.1	2
2230	Non-contiguous finished genome sequence of Prevotella timonensis type strain 4401737T. Standards in Genomic Sciences, 2014, 9, 1346-1353.	1.5	2
2231	Reply to Bastien et al: Figure 1 Journal of Infectious Diseases, 2015, 212, 506-508.	4.0	2
2232	Human Lice in Paleoentomology and Paleomicrobiology. , 2016, , 181-190.		2

#	Article	lF	Citations
2233	Draft Genome Sequence of Blautia faecis Strain Marseille-P328, Isolated from the Human Ascending Colon. Genome Announcements, 2016, 4, .	0.8	2
2234	Decreasing level of resistance in invasive Klebsiella pneumoniae strains isolated in Marseille, January 2012–July 2015. SpringerPlus, 2016, 5, 631.	1.2	2
2235	Double paradigm shift for the antibiotics' activity on viruses: Zika's lesson. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E1045-E1045.	7.1	2
2236	Malaria Therapy for Ebola Virus Infection. Clinical Infectious Diseases, 2017, 64, 696-697.	5.8	2
2237	Gonorrhea resistance: don't forget the old chaps. European Journal of Clinical Microbiology and Infectious Diseases, 2017, 36, 2537-2537.	2.9	2
2238	Rickettsia and Rickettsia-Like Organisms. , 2017, , 1666-1675.e1.		2
2239	Rewiring Microbiology and Infection. Clinical Infectious Diseases, 2017, 65, S1-S3.	5.8	2
2240	MimiLook: A Phylogenetic Workflow for Detection of Gene Acquisition in Major Orthologous Groups of Megavirales. Viruses, 2017, 9, 72.	3.3	2
2241	Draft Genome Sequence of Megamonas funiformis Strain Marseille-P3344, Isolated from a Human Fecal Microbiota. Genome Announcements, 2018, 6, .	0.8	2
2242	Rapid identification of microorganisms from platelet concentrates by matrixâ€assisted laser desorption ionization timeâ€ofâ€flight mass spectrometry after shortâ€term incubation on liquid medium. Transfusion, 2018, 58, 766-773.	1.6	2
2243	Genomic and phenotypic description of the newly isolated human species <i>Collinsella bouchesdurhonensis</i> ) sp. nov MicrobiologyOpen, 2018, 7, e00580.	3.0	2
2244	Identification of rickettsial immunoreactive proteins using a proximity ligation assay Western blotting and the traditional immunoproteomic approach. Comparative Immunology, Microbiology and Infectious Diseases, 2018, 58, 17-25.	1.6	2
2245	The functional convergence of antibiotic resistance in $\hat{l}^2\hat{a}\in$ lactamases is not conferred by a simple convergent substitution of amino acid. Evolutionary Applications, 2019, 12, 1812-1822.	3.1	2
2246	Phoenicibacter congonensis gen. nov., sp. nov., a new genus isolated from the human gut and its description using a taxonogenomic approach. Antonie Van Leeuwenhoek, 2019, 112, 775-784.	1.7	2
2247	Resistance to antibiotics of bacteria in tropical countries. Lancet Planetary Health, The, 2019, 3, e238-e239.	11.4	2
2248	Re: chronic Q-fever-related complications and mortality: data from a nationwide cohort. Clinical Microbiology and Infection, 2019, 25, 1433-1435.	6.0	2
2249	Insects and the Transmission of Bacterial Agents. , 0, , 195-202.		2
2250	Detection of Borrelia crocidurae in a vaginal swab after miscarriage, rural Senegal, Western Africa. International Journal of Infectious Diseases, 2020, 91, 261-263.	3.3	2

#	Article	IF	CITATIONS
2251	Current knowledge for the microbiological diagnosis of Tropheryma whipplei infection. Expert Opinion on Orphan Drugs, 2020, 8, 237-244.	0.8	2
2252	Gut dysbiosis in severe acute malnutrition is not an immaturity: The irreversible quantitative-qualitative paradigm shift. Human Microbiome Journal, 2020, 15, 100067.	3.8	2
2253	Response to the use of hydroxychloroquine in combination with azithromycin for patients with COVID-19 is not supported by recent literature International Journal of Antimicrobial Agents, 2021, 57, 106241.	2.5	2
2254	Response to uncertain effect of hydroxychloroquine and azithromycin on SARS-CoV-2 viral load International Journal of Antimicrobial Agents, 2021, 57, 106244.	2.5	2
2255	Lagierella massiliensis gen. nov., sp. nov., Isolated from a Stool Sample. Current Microbiology, 2021, 78, 2481-2487.	2.2	2
2256	COVID-19 Management at IHU Méditerranée Infection: A One-Year Experience. Journal of Clinical Medicine, 2021, 10, 2881.	2.4	2
2257	Gut Microbiota in Military International Travelers with Doxycycline Malaria Prophylaxis: Towards the Risk of a Simpson Paradox in the Human Microbiome Field. Pathogens, 2021, 10, 1063.	2.8	2
2258	Gemella massiliensis sp. nov., a new bacterium isolated from the human sputum. Archives of Microbiology, 2021, 203, 5817-5823.	2.2	2
2259	Determination of the genome size of Ehrlichia spp., using pulsed field gel electrophoresis. FEMS Microbiology Letters, 1999, 176, 73-78.	1.8	2
2260	rpoB Gene Analysis as a Novel Strategy for Identification of Spirochetes from the Genera Borrelia,Treponema, and Leptospira. Journal of Clinical Microbiology, 2000, 38, 3526-3526.	3.9	2
2261	New Tick-Transmitted Rickettsial Diseases. , 2000, , 233-249.		2
2262	Identification of Novel Zoonotic Activity of Bartonella spp., France. Emerging Infectious Diseases, 2016, 22, .	4.3	2
2263	Multispacer sequence typing of Coxiella burnetii from milk and hard tick samples from ruminant farms in Lebanon. Veterinaria Italiana, 2020, 56, 289-296.	0.5	2
2264	Mitral valve repair is better than mitral valve replacement in native mitral valve endocarditis: Results from a prospective matched cohort. Archives of Cardiovascular Diseases, 2022, 115, 160-168.	1.6	2
2265	Value and prognostic impact of a deep learning segmentation model of COVID-19 lung lesions on low-dose chest CT., 2022, 1, 100003.		2
2266	Louse-Associated Bacterial Infections. Infectious Diseases in Clinical Practice, 2000, 9, 281-291.	0.3	1
2267	Comparison of the in vitro efficacy of telithromycin (HMR 3647) and levofloxacin with 22 antibiotic compounds against Bosea and Afipia species. Journal of Antimicrobial Chemotherapy, 2004, 53, 683-685.	3.0	1
2268	Fever and Vesicular Rash in a Traveler Returning from South Africa. Clinical Infectious Diseases, 2004, 39, 700-701.	5.8	1

#	Article	lF	CITATIONS
2269	Rickettsioses in South Korea, Materials and Methods. Emerging Infectious Diseases, 2006, 12, 531-531.	4.3	1
2270	Past Plague. , 2008, , 145-159.		1
2271	Hepatosplenic cat-scratch fever with seropositivity for Bartonella quintana?. Lancet Infectious Diseases, The, 2008, 8, 663.	9.1	1
2272	Bacterial Protein Microarrays for Diagnosis of Infectious Diseases. Current Immunology Reviews, 2008, 4, 28-36.	1.2	1
2273	Quoi de neuf sur les RickettsiosesÂ?. Revue De Medecine Interne, 2009, 30, S19-S21.	1.0	1
2274	Applications of Paleomicrobiology to the Understanding of Emerging and Re-emerging Infectious Diseases. NATO Science for Peace and Security Series A: Chemistry and Biology, 2010, , 91-98.	0.5	1
2275	Long-term outcome of Q fever endocarditis – Authors' reply. Lancet Infectious Diseases, The, 2011, 11, 82.	9.1	1
2276	Draft Genome Sequence of Tsukamurella sp. Strain 1534. Journal of Bacteriology, 2012, 194, 5482-5483.	2.2	1
2277	Draft Genome Sequence of Staphylococcus massiliensis Strain 5402776 <sup>T</sup> . Journal of Bacteriology, 2012, 194, 6984-6985.	2.2	1
2278	Draft Genome Sequences of Actinomyces timonensis Strain 7400942 <sup>T</sup> and Its Prophage. Journal of Bacteriology, 2012, 194, 6613-6614.	2.2	1
2279	Reply. Journal of the American College of Cardiology, 2013, 62, 861-862.	2.8	1
2280	Immunity to measles, diphtheria and tetanus in residents of homeless shelters in Marseilles, France. Journal of Infection, 2013, 66, 189-191.	3.3	1
2281	Bartonellosis, Cat-scratch Disease, Trench Fever, Human Ehrlichiosis. , 2014, , 385-394.e2.		1
2282	Draft Genome Sequence of Mycobacterium cosmeticum DSM 44829. Genome Announcements, 2014, 2, .	0.8	1
2283	Reply to He et al. Clinical Infectious Diseases, 2014, 59, 1347-1348.	5.8	1
2284	La fiÃ"vre Q en 2014 : défidiagnostique et thérapeutique. Revue Francophone Des Laboratoires, 2014, 2014, 51-59.	0.0	1
2285	Common subclinical hypothyroidism during Whipple's disease. BMC Infectious Diseases, 2014, 14, 370.	2.9	1
2286	Non-contiguous finished genome sequence of Corynebacterium timonense type strain 5401744T. Standards in Genomic Sciences, 2014, 9, 948-955.	1,5	1

#	Article	IF	CITATIONS
2287	Draft Genome Sequence of Mycobacterium lentiflavum CSUR P1491. Genome Announcements, 2015, 3, .	0.8	1
2288	Sutton's Law: Keep Going Where The Money Is. Journal of General Internal Medicine, 2015, 30, 1711-1715.	2.6	1
2289	Imbalance of circulating lymphoid cells in Q fever endocarditis. Pathogens and Disease, 2015, 73, 1-3.	2.0	1
2290	The compliance of clinicians and patients cannot be globalized. Clinical Microbiology and Infection, 2015, 21, 391.	6.0	1
2291	Tropheryma whipplei as a causative agent of travelers' diarrhea: Further studies required. Reply to Razavi SM. Travel Medicine and Infectious Disease, 2015, 13, 114.	3.0	1
2292	Extensively Drug-Resistant Tuberculosis: You Can Teach an Old Dog New Tricks. Clinical Infectious Diseases, 2015, 60, 971-971.	5.8	1
2293	Concentrations of doxycycline in cerebrospinal fluid in Whipple's disease. International Journal of Antimicrobial Agents, 2015, 45, 677-678.	2.5	1
2294	Is it the end of the nervous breakdown on avian influenza?. Clinical Microbiology and Infection, 2015, 21, 805.	6.0	1
2295	Past Intestinal Parasites. , 2016, , 143-154.		1
2296	A Personal View of How Paleomicrobiology Aids Our Understanding of the Role of Lice in Plague Pandemics., 0,, 29-37.		1
2297	Parabacteroides timonensis sp. nov., identified in human stool. Human Microbiome Journal, 2016, 2, 1-2.	3.8	1
2298	"Collinsella vaginalis―sp. nov., a new bacterial species cultivated from human female genital tract. Human Microbiome Journal, 2016, 2, 19-20.	3.8	1
2299	Editorial overview: The megaviromes. Current Opinion in Microbiology, 2016, 31, viii-x.	5.1	1
2300	â $\in$ Clostridium massiliodielmoense â $\in$ ™, a new species isolated from the human gutÂmicrobiota. New Microbes and New Infections, 2016, 13, 27-28.	1.6	1
2301	Blood Culture-Negative Endocarditis. , 2016, , 245-258.		1
2302	Motorcycles, Cell Phones, and Electricity Can Dramatically Change the Epidemiology of Infectious Disease in Africa. American Journal of Tropical Medicine and Hygiene, 2017, 96, 16-0290.	1.4	1
2303	Draft Genome Sequence of Agrococcus baldri Strain Marseille-P2731. Genome Announcements, 2017, 5, .	0.8	1
2304	Actinomyces bouchesdurhonensis sp. nov. and Actinomyces mediterranea sp. nov., isolated from human stomach and duodenum. Human Microbiome Journal, 2017, 3, 13-14.	3.8	1

#	Article	IF	CITATIONS
2305	Old Antibiotics for Tuberculosis. Clinical Infectious Diseases, 2017, 64, 983-983.	5.8	1
2306	An uncommon form of a common disease. Journal of Paediatrics and Child Health, 2017, 53, 727-728.	0.8	1
2307	Identification of constraints influencing the bacterial genomes evolution in the PVC super-phylum. BMC Evolutionary Biology, 2017, 17, 75.	3.2	1
2308	Draft Genome Sequence of Ezakiella peruensis Strain M6.X2, a Human Gut Gram-Positive Anaerobic Coccus. Genome Announcements, 2018, 6, .	0.8	1
2309	The â€ <sup>~</sup> Eat me cakeâ€ <sup>™</sup> theory, or genetic cannibalism of the enemy: A cause of vanishing antimicrobial resistance. International Journal of Antimicrobial Agents, 2018, 52, 441-442.	2.5	1
2310	Bacterial Cocktail to Treat Clostridium difficile Infection: Primum Non Nocere. Clinical Infectious Diseases, 2018, 67, 1799.	5.8	1
2311	Whipple's Disease: Diagnostic Value of rpoB Gene PCR from Peripheral Blood Mononuclear Cells. Molecular Diagnosis and Therapy, 2018, 22, 459-469.	3.8	1
2312	Of ignorance and blindness: The Lyme disease paradigm. Médecine Et Maladies Infectieuses, 2019, 49, 85-86.	5.0	1
2313	Value of mathematical models for epidemics: the plague paradigm. Clinical Microbiology and Infection, 2019, 25, 120.	6.0	1
2314	Relevance of Medical Big Data Analysis Depends on Clinical Accuracy: The Q Fever Paradigm. Clinical Infectious Diseases, 2019, 68, 169-170.	5.8	1
2315	The living croquet theory: The Staphylococcus aureus paradigm. International Journal of Antimicrobial Agents, 2019, 53, 724-725.	2.5	1
2316	Acquisition of respiratory and gastrointestinal pathogens among health care workers during the 2015 Hajj season. American Journal of Infection Control, 2019, 47, 1071-1076.	2.3	1
2317	Antibiotic resistance, stewardship, and consumption. Lancet Planetary Health, The, 2019, 3, e67.	11.4	1
2318	Epidemic Louse-Borne Typhus. , 2020, , 577-579.		1
2319	Klenkia terrae resistant to DNA extraction in germ-free mice stools illustrates the extraction pitfall faced by metagenomics. Scientific Reports, 2020, 10, 10228.	3.3	1
2320	The impact of daily soap use in rural areas of Senegal on respiratory infectious diseases, fevers and skin microbiota. International Journal of Infectious Diseases, 2020, 96, 408-415.	3.3	1
2321	Gorillibacterium timonense sp. nov., isolated from an obese patient. Archives of Microbiology, 2020, 202, 1223-1229.	2.2	1
2322	Faecal microbiota transplantations and urinary tract infections. Lancet, The, 2020, 395, 270-271.	13.7	1

#	Article	IF	CITATIONS
2323	Hydroxychloroquine Failure: The End Does Not Justify the Means. Clinical Infectious Diseases, 2021, 72, e439-e439.	5.8	1
2324	Response to advances statistical methods and designs for clinical trials for COVID-19. International Journal of Antimicrobial Agents, 2021, 57, 106235.	2.5	1
2325	Safety profile of hydroxychloroquine and azithromycin combined treatment in COVID-19 patients. International Journal of Antimicrobial Agents, 2021, 57, 106236.	2.5	1
2326	Re: †Effect of hydroxychloroquine with or without azithromycin on the mortality of COVID-19 patients' by Fiolet etÂal Clinical Microbiology and Infection, 2021, 27, 132-133.	6.0	1
2327	Reply to Lebeaux D, Revest M. No evidence of clinical benefits of early treatment of COVID-19 patients with hydroxychloroquine and azithromycin. Travel Medicine and Infectious Disease, 2021, 39, 101954.	3.0	1
2328	Draft Genome Sequence of Comamonas aquatilis Strain LK (= CSUR P6418 = CECT 9772), Isolated Planarian Schmidtea mediterranea. Microbiology Resource Announcements, 2021, 10, .	from the	1
2329	Effect of hydroxychloroquine and azithromycin on the viral clearance of SARS-CoV-2: response to Hervé Seligmann. International Journal of Antimicrobial Agents, 2021, 57, 106306.	2.5	1
2330	Enteric pathogenic bacteria and resistance gene carriage in the homeless population in Marseille, France. Acta Microbiologica Et Immunologica Hungarica, 2021, 68, 7-13.	0.8	1
2331	Contagion Management at the Méditerranée Infection University Hospital Institute. Journal of Clinical Medicine, 2021, 10, 2627.	2.4	1
2332	Limosilactobacillus caccae sp. nov., a new bacterial species isolated from the human gut microbiota. FEMS Microbiology Letters, 2021, 368, .	1.8	1
2333	Is Acanthamoeba polyphaga Mimivirus an Emerging Causative Agent of Pneumonia?., 0,, 33-51.		1
2334	Dental Pulp as a Tool for the Retrospective Diagnosis of Infectious Diseases. , 2008, , 175-196.		1
2335	Scrub typhus and other tropical rickettsioses. , 2010, , 1231-1237.		1
2336	Bacterial genomes., 2010,, 86-91.		1
2337	Orientia tsutsugamushi (Scrub Typhus). , 2015, , 2225-2226.		1
2338	Pediatric scrub typhus in Indian Himalayas. Indian Journal of Pediatrics, 0, , .	0.8	1
2339	Bacterial Infections in Humans and Nonhuman Primates from Africa: Expanding the Knowledge. Yale Journal of Biology and Medicine, 2021, 94, 227-248.	0.2	1
2340	Genome analysis as a new and powerful tool to design suitable culture media. Application to tropheryma whipplei. Discovery Medicine, 2003, 3, 32-3.	0.5	1

#	Article	IF	CITATIONS
2341	Culturomics revealed the bacterial constituents of the microbiota of a 10-year-old laboratory culture of planarian species S. mediterranea. Scientific Reports, 2021, 11, 24311.	3.3	1
2342	Human ehrlichiosis with features of toxic shock syndrome. American Journal of Medicine, 1995, 99, 107.	1.5	0
2343	Life with Rickettsiae. Infectious Diseases in Clinical Practice, 2000, 9, 364-367.	0.3	0
2344	Bartonella Infections Resurgence in the New Century. , 2004, , 35-68.		0
2345	Genomics ofRickettsia., 2006,,.		0
2346	Bartonelloses., 2008,, 25-31.		0
2347	Rickettsioses., 2008,, 38-48.		0
2348	Transmissible cancer in Africa. Lancet, The, 2009, 374, 2052.	13.7	0
2349	Sound and Fury, Clarified. Science, 2012, 336, 152-155.	12.6	0
2350	Bartonella Infections., 2012,, 1906-1911.		0
2351	Non transmissible diseases are often transmissible. European Journal of Epidemiology, 2012, 27, 243-245.	5.7	0
2352	Antibiotic susceptibility and intracellular localization of <i>Diplorickettsia massiliensis </i> Immunology and Medical Microbiology, 2012, 64, 48-56.	2.7	0
2353	Impact factors of scientific journals: the new season. Clinical Microbiology and Infection, 2013, 19, 989-990.	6.0	0
2354	Digestive microbiota and its influence on health: Facts and myths. Microbial Pathogenesis, 2013, 61-62, A1.	2.9	0
2355	Rickettsial infections., 0,, 322-329.		0
2356	Treatment of Intracellular Infections. , 2014, , 323-335.		0
2357	Genome Sequence of Legionella massiliensis, Isolated from a Cooling Tower Water Sample. Genome Announcements, 2014, 2, .	0.8	0
2358	Draft Genome Sequence of Mycobacterium asiaticum Strain DSM 44297. Genome Announcements, 2014, 2, .	0.8	0

#	Article	IF	CITATIONS
2359	Draft Genome Sequence of Mycobacterium mageritense DSM 44476 <sup>T</sup> . Genome Announcements, 2014, 2, .	0.8	0
2360	Draft Genome Sequence of Mycobacterium triplex DSM 44626. Genome Announcements, 2014, 2, .	0.8	0
2361	Editorial. Microbial Pathogenesis, 2014, 77, 113.	2.9	0
2362	688: Atopobium vaginae reduces interval to delivery in high risk pregnancies. American Journal of Obstetrics and Gynecology, 2014, 210, S338-S339.	1.3	0
2363	Draft Genome Sequence of Mycobacterium europaeum Strain CSUR P1344. Genome Announcements, 2015, 3, .	0.8	O
2364	What is hot in infectious diseases?. Clinical Microbiology and Infection, 2015, 21, 803-804.	6.0	0
2365	The ignored pandemic of typhoid. Clinical Microbiology and Infection, 2015, 21, 877.	6.0	O
2366	Paleopathology of Human Infections: Old Bones, Antique Books, Ancient and Modern Molecules. , 2016, , 93-106.		0
2367	Sources of Materials for Paleomicrobiology. , 2016, , 39-50.		O
2368	Past Bartonelloses. , 2016, , 107-111.		0
2369	Paleomicrobiology Data: Authentification and Interpretation. , 2016, , 51-58.		0
2370	Human Coprolites as a Source for Paleomicrobiology. , 0, , 59-74.		0
2371	The History of Epidemic Typhus. , 0, , 81-92.		O
2372	"Nigerium massiliense―gen. nov., sp. nov., a new bacterium isolated from the gut from a patient with acute malnutrition. Human Microbiome Journal, 2016, 1, 12-13.	3.8	0
2373	Rickettsia conorii is a potent complement activator in vivo and combined inhibition of complement and CD14 is required for attenuation of the cytokine response ex vivo. Immunobiology, 2016, 221, 1204-1205.	1.9	O
2374	Antimicrobial Therapy in Infective Endocarditis., 2016,, 275-279.		0
2375	Microbiological Diagnosis in Infective Endocarditis. , 2016, , 31-36.		O
2376	New prevention strategies for sexually transmitted diseases. Clinical Microbiology and Infection, 2016, 22, 752.	6.0	0

#	Article	IF	Citations
2377	Questioning effectiveness of vaccines against malaria. Lancet Infectious Diseases, The, 2017, 17, 22.	9.1	O
2378	Draft Genome Sequence of Streptomyces specialis Type Strain GW41-1564 (DSM 41924). Genome Announcements, 2017, 5, .	0.8	0
2379	Peptoniphilus duodeni sp. nov., a new bacterial species identified in human duodenum. Human Microbiome Journal, 2017, 3, 9-10.	3.8	0
2380	Bacterial Genomes. , 2017, , 62-67.e1.		0
2381	Human papillomavirus vaccine: Urgent need to promote gender parity. European Journal of Epidemiology, 2018, 33, 259-261.	5.7	0
2382	Does antibiotic prophylaxis really prevent streptococci infective endocarditis?. Heart, 2018, 104, 360-360.	2.9	0
2383	Should evidence-based medicine be considered as a truth in clinical microbiology?. International Journal of Antimicrobial Agents, 2018, 51, 278-279.	2.5	0
2384	Endocarditis Prophylaxis. Clinical Infectious Diseases, 2018, 66, 982-982.	5.8	0
2385	Draft genome and description of Chryseobacterium phocaeense sp. nov.: a new bacterial species isolated from the sputum of a cystic fibrosis patient. Archives of Microbiology, 2019, 201, 1361-1368.	2.2	0
2386	Vaccination against the big three killers: an illusion?. Clinical Microbiology and Infection, 2019, 25, 654-655.	6.0	0
2387	Failure of metagenomics in detecting emerging pathogens, the Clostridium difficile paradigm. Journal of Infection, 2019, 78, 409-421.	3.3	0
2388	Should we fear gonorrhoea?. Lancet Infectious Diseases, The, 2019, 19, 1286-1287.	9.1	0
2389	A case of sepsis in a 17th century man from Porto Ercole – Author's reply. Lancet Infectious Diseases, The, 2019, 19, 25-26.	9.1	0
2390	Epidemiology of human common coronavirus acquisition in pilgrims. Travel Medicine and Infectious Disease, 2020, 37, 101845.	3.0	0
2391	Vibrio vulnificus casualties during the American Civil War. Lancet Infectious Diseases, The, 2020, 20, 170-171.	9.1	O
2392	Draft genome and description of Negativicoccus massiliensis strain Marseille-P2082, a new species isolated from the gut microbiota of an obese patient. Antonie Van Leeuwenhoek, 2020, 113, 997-1008.	1.7	0
2393	Rickettsioses: "A Treasure Is Hidden in This Garden― Clinical Infectious Diseases, 2021, 72, 1179-1180.	5.8	0
2394	Be Careful With Big Data: Reanalysis of Patient Characteristics and Outcomes of 11 721 Patients With Coronavirus Disease 2019 Hospitalized Across the United States. Clinical Infectious Diseases, 2021, 72, e928-e928.	5.8	0

#	Article	IF	CITATIONS
2395	Response to effect estimation of hydroxychloroquine for COVID-19: a secondary analysis of an open label non-randomized clinical trial International Journal of Antimicrobial Agents, 2021, 57, 106237.	2.5	0
2396	Interpretation of SARS-CoV-2 PCR results for the diagnosis of COVID-19. International Journal of Antimicrobial Agents, 2021, 57, 106238.	2.5	0
2397	Draft Genome Sequence of Comamonas jiangduensis Strain YW1 <sup>T</sup> . Microbiology Resource Announcements, 2021, 10, .	0.6	0
2398	Draft Genome Sequence of Vogesella oryzae L3B39 <sup>T</sup> , Isolated from the Rhizosphere of Saline-Tolerant Pokkali Rice. Microbiology Resource Announcements, 2021, 10, .	0.6	0
2399	HydroxychloroquineÂ+Âazithromycin treatment in elderly patients. International Journal of Antimicrobial Agents, 2021, 57, 106313.	2.5	0
2400	Genomic description and characterization of Nigeribacterium massiliense gen. nov., sp. nov., isolated from the human gut. Microbes and Infection, 2021, 23, 104842.	1.9	0
2401	Detection of zoonotic pathogens in animals performed at the University Hospital Institute Méditerranée Infection (Marseille – France). One Health, 2021, 12, 100210.	3.4	0
2402	Rapid Isothermal Amplification for the Buccal Detection SARS-CoV-2 in the Context of Out-Patient COVID-19 Screening. Journal of Clinical Medicine, 2021, 10, 2643.	2.4	0
2403	A review of in vitro attempts to develop the axenic culture of Treponema pallidum and genomics-based suggestions to achieve this elusive goal. Journal of Medical Microbiology, 2021, 70, .	1.8	0
2404	Evaluation of pain susceptibility by taking blood pressure in patients with infections. Medicine (United) Tj ETQq0	0 0 rgBT /	Overlock 10
2405	Real world and hyper reality. Travel Medicine and Infectious Disease, 2021, 43, 102122.	3.0	0
2406	Clinical and Epidemiological Changes in French Soldiers After Deployment: Impact of Doxycycline Malaria Prophylaxis on Body Weight. Military Medicine, 2023, 188, e1084-e1093.	0.8	0
2407	Les rickettsioses en réanimation. , 2010, , 83-95.		0
2408	Rickettsia and rickettsia-like organisms. , 2010, , 1807-1816.		0
2409	Mimivirus., 2011,, 877-891.		0
2410	Epidemic Louse-borne Typhus. , 2013, , 535-538.		0
2411	Staphylococcus aureus subsp. anaerobius strain ST1464 genome sequence. Standards in Genomic Sciences, 2013, 9, 1-11.	1.5	0
2412	Mediterranean Spotted Fever and Other Tick-Borne Rickettsioses., 0,, 302-327.		0

#	Article	IF	Citations
2413	Rickettsia akari (Rickettsialpox)., 2015,, 2206-2207.		0
2414	Genomes of Rickettsia Species. , 2016, , 447-462.		0
2415	Global Warming and Global Decrease in Vector-Borne Disease Prevalence and Mortality. Journal of Infectious Diseases, 2017, 215, 660-661.	4.0	0
2416	A concise overview on tick-borne human infections in Europe: a focus on Lyme borreliosis and tick-borne Rickettsia spp Microbiology Australia, 2018, 39, 207.	0.4	0
2417	Konateibacter massiliensis gen. nov. sp. nov. and Paenibacillus faecalis sp. nov., Two New Species Isolated from the Stool Samples of Infants Suffering from Marasmus. Current Microbiology, 2022, 79, 68.	2.2	0
2418	Differential word expression analyses highlight plague dynamics during the second pandemic. Royal Society Open Science, 2022, 9, 210039.	2.4	0
2419	Unusual gram-positive spiral-shaped bacilli detected in a positive blood culture. Clinical Microbiology and Infection, 2022, , .	6.0	0
2420	Re: 'treatment of Coxiella burnetii endocarditis with hydroxychloroquine' by Stahl et al. Clinical Microbiology and Infection, 2022, , .	6.0	0
2421	Antilogic, a new supervised machine learning software for the automatic interpretation of antibiotic susceptibility testing in clinical microbiology: proof-of-concept on three frequently isolated bacterial species. Clinical Microbiology and Infection, 2022, , .	6.0	0
2422	Bhargavaea massiliensis sp. nov. and Dietzia massiliensis sp. nov., Novel Bacteria Species Isolated from Human Urine Samples in Nigeria. Current Microbiology, 2022, 79, 18.	2.2	0
2423	Towards 21st century microbiology in Africa. Medecine Et Sante Tropicales, 2019, 29, 340-342.	0.3	0
2424	La transformation d'une bactérie en cellule virale à noyau relance l'hypothèse de l'eucaryogenèse virale. Virologie, 2017, 21, 157-159.	0.1	0
2425	Who Were Hospitalized Deceased Patients from COVID-19 During the First Year of Pandemic? Retrospective Analysis of $1104$ Deceased Patients in South of France. Journal of Epidemiology and Global Health, $2022$ , , .	2.9	0
2426	Draft Genome Sequence of Corynebacterium sanguinis Strain Marseille-P8776. Microbiology Resource Announcements, 2022, 11, e0000822.	0.6	0
2427	The absolute number of leukocytes per vial as a major cause of early false positive blood cultures: proof-of-concept and application. European Journal of Clinical Microbiology and Infectious Diseases, 2022, 41, 951-959.	2.9	O
2428	Antimicrobial susceptibility testing for Gram positive cocci towards vancomycin using scanning electron microscopy. Current Research in Microbial Sciences, 2022, 3, 100154.	2.3	0