Didier Raoult

List of Publications by Year in descending order

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2,427 papers

147,088 citations

159 h-index 280 g-index

2515 all docs

2515 docs citations

2515 times ranked

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.	1.1	3,955
2	Gut microbiome influences efficacy of PD-1–based immunotherapy against epithelial tumors. Science, 2018, 359, 91-97.	6.0	3,689
3	Anticancer immunotherapy by CTLA-4 blockade relies on the gut microbiota. Science, 2015, 350, 1079-1084.	6.0	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.	2.9	1,638
5	The abundance and variety of carbohydrate-active enzymes in the human gut microbiota. Nature Reviews Microbiology, 2013, 11, 497-504.	13.6	1,240
6	Update on Tick-Borne Rickettsioses around the World: a Geographic Approach. Clinical Microbiology Reviews, 2013, 26, 657-702.	5.7	1,033
7	The 1.2-Megabase Genome Sequence of Mimivirus. Science, 2004, 306, 1344-1350.	6.0	959
8	Tick-Borne Rickettsioses around the World: Emerging Diseases Challenging Old Concepts. Clinical Microbiology Reviews, 2005, 18, 719-756.	5.7	920
9	Microbial culturomics: paradigm shift in the human gut microbiome study. Clinical Microbiology and Infection, 2012, 18, 1185-1193.	2.8	905
10	Microorganisms Resistant to Free-Living Amoebae. Clinical Microbiology Reviews, 2004, 17, 413-433.	5.7	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.	1.8	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.	1.1	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.	1.3	767
14	A Giant Virus in Amoebae. Science, 2003, 299, 2033-2033.	6.0	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.	1.1	735
16	Culture of previously uncultured members of the human gut microbiota by culturomics. Nature Microbiology, 2016, 1, 16203.	5.9	735
17	Chloroquine and hydroxychloroquine as available weapons to fight COVID-19. International Journal of Antimicrobial Agents, 2020, 55, 105932.	1.1	724
18	Comparative Genomics of Multidrug Resistance in Acinetobacter baumannii. PLoS Genetics, 2006, 2, e7.	1.5	677

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19	Endocarditis Due to Rare and Fastidious Bacteria. Clinical Microbiology Reviews, 2001, 14, 177-207.	5.7	668
20	Enterococcus hirae and Barnesiella intestinihominis Facilitate Cyclophosphamide-Induced Therapeutic Immunomodulatory Effects. Immunity, 2016, 45, 931-943.	6.6	645
21	From Q Fever to Coxiella burnetii Infection: a Paradigm Change. Clinical Microbiology Reviews, 2017, 30, 115-190.	5 . 7	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.	5.7	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.	1.5	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.	13.9	574
26	Pig Liver Sausage as a Source of Hepatitis E Virus Transmission to Humans. Journal of Infectious Diseases, 2010, 202, 825-834.	1.9	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.	1.6	567
28	Diagnosis of Q Fever. Journal of Clinical Microbiology, 1998, 36, 1823-1834.	1.8	529
29	Q fever. Veterinary Microbiology, 2010, 140, 297-309.	0.8	525
30	The microbiome in cancer immunotherapy: Diagnostic tools and therapeutic strategies. Science, 2018, 359, 1366-1370.	6.0	525
31	Culturing the human microbiota and culturomics. Nature Reviews Microbiology, 2018, 16, 540-550.	13.6	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.	0.8	510
33	The virophage as a unique parasite of the giant mimivirus. Nature, 2008, 455, 100-104.	13.7	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.	3.3	482
35	Q Fever 1985-1998: Clinical and Epidemiologic Features of 1,383 Infections. Medicine (United States), 2000, 79, 109-123.	0.4	459
36	Cultivation of the Bacillus of Whipple's Disease. New England Journal of Medicine, 2000, 342, 620-625.	13.9	458

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37	Recommendations for Treatment of Human Infections Caused by Bartonella Species. Antimicrobial Agents and Chemotherapy, 2004, 48, 1921-1933.	1.4	456
38	rpoB sequence analysis as a novel basis for bacterial identification. Molecular Microbiology, 1997, 26, 1005-1011.	1.2	450
39	Positron Emission Tomography/Computed Tomography forÂDiagnosis of Prosthetic Valve Endocarditis. Journal of the American College of Cardiology, 2013, 61, 2374-2382.	1.2	440
40	Comprehensive Diagnostic Strategy for Blood Culture–Negative Endocarditis: A Prospective Study of 819 New Cases. Clinical Infectious Diseases, 2010, 51, 131-140.	2.9	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.	1.5	410
42	Mechanisms of Evolution in Rickettsia conorii and R. prowazekii. Science, 2001, 293, 2093-2098.	6.0	408
43	Laboratory diagnosis of rickettsioses: current approaches to diagnosis of old and new rickettsial diseases. Journal of Clinical Microbiology, 1997, 35, 2715-2727.	1.8	397
44	Chikungunya Outbreaks $\hat{a} \in \mathbb{C}^n$ The Globalization of Vectorborne Diseases. New England Journal of Medicine, 2007, 356, 769-771.	13.9	394
45	Echocardiography predicts embolic events in infective endocarditis. Journal of the American College of Cardiology, 2001, 37, 1069-1076.	1.2	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.	1.8	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.	3.3	385
48	Blood Culture-Negative Endocarditis in a Reference Center. Medicine (United States), 2005, 84, 162-173.	0.4	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.	1.5	372
50	Comparative meta-analysis of the effect of Lactobacillus species on weight gain in humans and animals. Microbial Pathogenesis, 2012, 53, 100-108.	1.3	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.	1.8	362
52	Management of infective endocarditis: challenges and perspectives. Lancet, The, 2012, 379, 965-975.	6.3	359
53	Redefining viruses: lessons from Mimivirus. Nature Reviews Microbiology, 2008, 6, 315-319.	13.6	358
54	Current and Past Strategies for Bacterial Culture in Clinical Microbiology. Clinical Microbiology Reviews, 2015, 28, 208-236.	5.7	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, .	5.6	357
56	Bartonella (Rochalimaea) quintanaEndocarditis in Three Homeless Men. New England Journal of Medicine, 1995, 332, 419-423.	13.9	355
57	Chloroquine for the 2019 novel coronavirus SARS-CoV-2. International Journal of Antimicrobial Agents, 2020, 55, 105923.	1.1	354
58	Gene Sequence-Based Criteria for Identification of New Rickettsia Isolates and Description of Rickettsia heilongjiangensis sp. nov. Journal of Clinical Microbiology, 2003, 41, 5456-5465.	1.8	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.	1.1	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.	1.1	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.	0.8	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.	0.8	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.	0.6	328
64	Guidelines for the diagnosis of tick-borne bacterial diseases in Europe. Clinical Microbiology and Infection, 2004, 10, 1108-1132.	2.8	328
65	Whipple's disease: new aspects of pathogenesis and treatment. Lancet Infectious Diseases, The, 2008, 8, 179-190.	4.6	328
66	Q Fever. Clinical Infectious Diseases, 1995, 20, 489-496.	2.9	322
67	Diagnosis of 22 New Cases of Bartonella Endocarditis. Annals of Internal Medicine, 1996, 125, 646.	2.0	321
68	Sequencing of the rpoB Gene and Flanking Spacers for Molecular Identification of Acinetobacter Species. Journal of Clinical Microbiology, 2006, 44, 827-832.	1.8	321
69	Eukaryotic large nucleo-cytoplasmic DNA viruses: Clusters of orthologous genes and reconstruction of viral genome evolution. Virology Journal, 2009, 6, 223.	1.4	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.	1.0	320
71	The Body Louse as a Vector of Reemerging Human Diseases. Clinical Infectious Diseases, 1999, 29, 888-911.	2.9	316
72	Comparative Analysis of Acinetobacters: Three Genomes for Three Lifestyles. PLoS ONE, 2008, 3, e1805.	1.1	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.	1.1	312
74	Fleas and flea-borne diseases. International Journal of Infectious Diseases, 2010, 14, e667-e676.	1.5	312
75	MALDI-TOF-mass spectrometry applications in clinical microbiology. Future Microbiology, 2010, 5, 1733-1754.	1.0	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.	4.3	306
78	The relationship between gut microbiota and weight gain in humans. Future Microbiology, 2012, 7, 91-109.	1.0	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.	3.3	301
80	The Q fever epidemic in The Netherlands: history, onset, response and reflection. Epidemiology and Infection, 2011, 139, 1-12.	1.0	298
81	Genome of Acanthamoeba castellanii highlights extensive lateral gene transfer and early evolution of tyrosine kinase signaling. Genome Biology, 2013, 14, R11.	13.9	296
82	Novel Chikungunya Virus Variant in Travelers Returning from Indian Ocean Islands. Emerging Infectious Diseases, 2006, 12, 1493-1499.	2.0	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.	2.9	289
84	Molecular identification by "suicide PCR" of Yersinia pestis as the agent of Medieval Black Death. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 12800-12803.	3.3	288
85	Genome Sequence of Rickettsia bellii Illuminates the Role of Amoebae in Gene Exchanges between Intracellular Pathogens. PLoS Genetics, 2006, 2, e76.	1.5	286
86	ChronicBartonella quintanaBacteremia in Homeless Patients. New England Journal of Medicine, 1999, 340, 184-189.	13.9	285
87	Rickettsia africae,a Tick-Borne Pathogen in Travelers to Sub-Saharan Africa. New England Journal of Medicine, 2001, 344, 1504-1510.	13.9	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.	4.6	282
89	Bacterial strain typing in the genomic era. FEMS Microbiology Reviews, 2009, 33, 892-916.	3.9	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.	2.0	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.	1.2	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.	1.8	271
93	Coronavirus infections: Epidemiological, clinical and immunological features and hypotheses. Cell Stress, 2020, 4, 66-75.	1.4	271
94	Molecular Quantification of <i> Gardnerella vaginalis </i> and <i> Atopobium vaginae </i> Loads to Predict Bacterial Vaginosis. Clinical Infectious Diseases, 2008, 47, 33-43.	2.9	267
95	Risks Factors and Prevention of Q Fever Endocarditis. Clinical Infectious Diseases, 2001, 33, 312-316.	2.9	264
96	Modern clinical microbiology: new challenges and solutions. Nature Reviews Microbiology, 2013, 11, 574-585.	13.6	264
97	Whipple's disease. Lancet, The, 2003, 361, 239-246.	6.3	263
98	Gene-sequence-based criteria for species definition in bacteriology: the Bartonella paradigm. Trends in Microbiology, 2003, 11, 318-321.	3.5	259
99	The rpoB gene as a tool for clinical microbiologists. Trends in Microbiology, 2009, 17, 37-45.	3.5	257
100	"Megavirales― a proposed new order for eukaryotic nucleocytoplasmic large DNA viruses. Archives of Virology, 2013, 158, 2517-2521.	0.9	256
101	rpoB Gene Sequence-Based Identification of Staphylococcus Species. Journal of Clinical Microbiology, 2002, 40, 1333-1338.	1.8	255
102	Dramatic Reduction in Infective Endocarditis–Related Mortality With a Management-Based Approach. Archives of Internal Medicine, 2009, 169, 1290.	4.3	255
103	Characterization of the Naturally Occurring Oxacillinase of Acinetobacter baumannii. Antimicrobial Agents and Chemotherapy, 2005, 49, 4174-4179.	1.4	254
104	Understanding the Cholera Epidemic, Haiti. Emerging Infectious Diseases, 2011, 17, 1161-1168.	2.0	252
105	Human Gut Microbiota: Repertoire and Variations. Frontiers in Cellular and Infection Microbiology, 2012, 2, 136.	1.8	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.	1.8	251
107	A comprehensive repertoire of prokaryotic species identified in human beings. Lancet Infectious Diseases, The, 2015, 15, 1211-1219.	4.6	250
108	Tick- and flea-borne rickettsial emerging zoonoses. Veterinary Research, 2005, 36, 469-492.	1.1	248

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109	Tailed giant Tupanvirus possesses the most complete translational apparatus of the known virosphere. Nature Communications, 2018, 9, 749.	5.8	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.	1.1	246
111	In vitro testing of combined hydroxychloroquine and azithromycin on SARS-CoV-2 shows synergistic effect. Microbial Pathogenesis, 2020, 145, 104228.	1.3	246
112	Wind in November, Q Fever in December. Emerging Infectious Diseases, 2004, 10, 1264-1269.	2.0	244
113	rpoB Gene Sequencing for Identification of Corynebacterium Species. Journal of Clinical Microbiology, 2004, 42, 3925-3931.	1.8	243
114	The Genome Sequence of Rickettsia felis Identifies the First Putative Conjugative Plasmid in an Obligate Intracellular Parasite. PLoS Biology, 2005, 3, e248.	2.6	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.	0.7	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.	1.8	232
117	Warmer Weather Linked to Tick Attack and Emergence of Severe Rickettsioses. PLoS Neglected Tropical Diseases, 2008, 2, e338.	1.3	228
118	Antibiotic discovery: history, methods and perspectives. International Journal of Antimicrobial Agents, 2019, 53, 371-382.	1.1	223
119	Rickettsioses and the International Traveler. Clinical Infectious Diseases, 2004, 39, 1493-1499.	2.9	222
120	Rickettsial evolution in the light of comparative genomics. Biological Reviews, 2011, 86, 379-405.	4.7	219
121	The two faces of interleukin 10 in human infectious diseases. Lancet Infectious Diseases, The, 2006, 6, 557-569.	4.6	218
122	Cross-reactivity between tumor MHC class l–restricted antigens and an enterococcal bacteriophage. Science, 2020, 369, 936-942.	6.0	217
123	Gut microbiota signatures are associated with toxicity to combined CTLA-4 and PD-1 blockade. Nature Medicine, 2021, 27, 1432-1441.	15.2	216
124	Use of rpoB Gene Analysis for Detection and Identification of Bartonella Species. Journal of Clinical Microbiology, 2001, 39, 430-437.	1.8	211
125	<i>Coxiella burnetii</i> Genotyping. Emerging Infectious Diseases, 2005, 11, 1211-1217.	2.0	210
126	Structural Studies of the Giant Mimivirus. PLoS Biology, 2009, 7, e1000092.	2.6	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.	1.5	209
128	Epidemiologic and Clinical Characteristics of Bartonella quintana and Bartonella henselae Endocarditis. Medicine (United States), 2001, 80, 245-251.	0.4	207
129	A Flea-Associated Rickettsia Pathogenic for Humans. Emerging Infectious Diseases, 2001, 7, 73-81.	2.0	207
130	Teicoplanin: an alternative drug for the treatment of COVID-19?. International Journal of Antimicrobial Agents, 2020, 55, 105944.	1.1	205
131	Outcome and Treatment of Bartonella Endocarditis. Archives of Internal Medicine, 2003, 163, 226.	4.3	202
132	<i>Propionibacterium acnes</i> Postoperative Shoulder Arthritis: An Emerging Clinical Entity. Clinical Infectious Diseases, 2008, 46, 1884-1886.	2.9	202
133	Molecular Detection of Multiple Emerging Pathogens in Sputa from Cystic Fibrosis Patients. PLoS ONE, 2008, 3, e2908.	1.1	201
134	African tick bite fever. Lancet Infectious Diseases, The, 2003, 3, 557-564.	4.6	199
135	Rickettsial Infections and Fever, Vientiane, Laos. Emerging Infectious Diseases, 2006, 12, 256-262.	2.0	197
136	Long-term outcome of Q fever endocarditis: a 26-year personal survey. Lancet Infectious Diseases, The, 2010, 10, 527-535.	4.6	197
137	Systemic Tropheryma whipplei. Medicine (United States), 2010, 89, 337-345.	0.4	195
138	Infections in the homeless. Lancet Infectious Diseases, The, 2001, 1, 77-84.	4.6	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.	3.3	194
140	Gut Bacteria Composition Drives Primary Resistance to Cancer Immunotherapy in Renal Cell Carcinoma Patients. European Urology, 2020, 78, 195-206.	0.9	192
141	<i>Rickettsia slovaca</i> and <i>R. raoultii</i> ii> Tick <i>-</i> borne Rickettsioses. Emerging Infectious Diseases, 2009, 15, 1105-1108.	2.0	191
142	Faustovirus, an Asfarvirus-Related New Lineage of Giant Viruses Infecting Amoebae. Journal of Virology, 2015, 89, 6585-6594.	1.5	191
143	Epidemic typhus. Lancet Infectious Diseases, The, 2008, 8, 417-426.	4.6	189
144	Massive comparative genomic analysis reveals convergent evolution of specialized bacteria. Biology Direct, 2009, 4, 13.	1.9	187

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145	<i>Rickettsia massiliae</i> Human Isolation. Emerging Infectious Diseases, 2006, 12, 174-175.	2.0	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.	1.1	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.	0.6	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.	1.0	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.	1.8	183
150	Comparison of PCR and Serology Assays for Early Diagnosis of Acute Q Fever. Journal of Clinical Microbiology, 2003, 41, 5094-5098.	1.8	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.	0.7	182
152	Coxiella burnetii in Humans and Ticks in Rural Senegal. PLoS Neglected Tropical Diseases, 2010, 4, e654.	1.3	181
153	Genome-based design of a cell-free culture medium for Tropheryma whipplei. Lancet, The, 2003, 362, 447-449.	6.3	180
154	Mediterranean Spotted Fever: Clinical, Laboratory and Epidemiological Features of 199 Cases. American Journal of Tropical Medicine and Hygiene, 1986, 35, 845-850.	0.6	179
155	Spotless Rickettsiosis Caused byRickettsia slovacaand Associated withDermacentorTicks. Clinical Infectious Diseases, 2002, 34, 1331-1336.	2.9	177
156	African Tick Bite Fever in Travelers to Rural Sub-Equatorial Africa. Clinical Infectious Diseases, 2003, 36, 1411-1417.	2.9	177
157	Molecular diagnosis of bloodstream infections caused by non-cultivable bacteria. International Journal of Antimicrobial Agents, 2007, 30, 7-15.	1.1	176
158	Gut microbiota and malnutrition. Microbial Pathogenesis, 2017, 106, 127-138.	1.3	173
159	Etiologic Diagnosis of 204 Pericardial Effusions. Medicine (United States), 2003, 82, 385-391.	0.4	170
160	Value of <i>Tropheryma whipplei </i> Value of <i>Tropheryma whipplei Value of <i>Tropheryma whipplei Value of <i>Tropheryma whipple Diagnosis of Whipple Disease: Usefulness of Saliva and Stool Specimens for Firstâ€Line Screening. Clinical Infectious Diseases, 2008, 47, 659-667.</i></i></i>	2.9	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.	1.8	167
162	Reductive Genome Evolution from the Mother of Rickettsia. PLoS Genetics, 2007, 3, e14.	1.5	167

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163	Pathogenicity and treatment of Bartonella infections. International Journal of Antimicrobial Agents, 2014, 44, 16-25.	1.1	167
164	Genotyping, Orientalis-like <i>Yersinia pestis</i> , and Plague Pandemics. Emerging Infectious Diseases, 2004, 10, 1585-1592.	2.0	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.	1.3	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.	0.8	161
167	<i>Tropheryma whipplei</i> Twist: A Human Pathogenic Actinobacteria With a Reduced Genome. Genome Research, 2003, 13, 1800-1809.	2.4	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.	2.9	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.	1.8	157
170	Prevalence of Asymptomatic <i>Tropheryma whipplei</i> Primates. Journal of Infectious Diseases, 2008, 197, 880-887.	1.9	157
171	Q fever in children. Lancet Infectious Diseases, The, 2002, 2, 686-691.	4.6	155
172	Evidence for Louseâ€Transmitted Diseases in Soldiers of Napoleon's Grand Army in Vilnius. Journal of Infectious Diseases, 2006, 193, 112-120.	1.9	155
173	Related actions of probiotics and antibiotics on gut microbiota and weight modification. Lancet Infectious Diseases, The, 2013, 13, 889-899.	4.6	154
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