Didier Raoult

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2,384 papers	111,188 citations	144 h-index	248 g-index
2,514 ext. papers	132,365 ext. citations	7.8 avg, IF	8.75 L-index

#	Paper	IF	Citations
2384	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	14.3	3043
2383	Gut microbiome influences efficacy of PD-1-based immunotherapy against epithelial tumors. *Science*, **2018**, 359, 91-97	33.3	2203
2382	Anticancer immunotherapy by CTLA-4 blockade relies on the gut microbiota. *Science*, **2015**, 350, 1079-84	33.3	1689
2381	Ongoing revolution in bacteriology: routine identification of bacteria by matrix-assisted laser desorption ionization time-of-flight mass spectrometry. *Clinical Infectious Diseases*, **2009**, 49, 543-51	11.6	1396
2380	The abundance and variety of carbohydrate-active enzymes in the human gut microbiota. *Nature Reviews Microbiology*, **2013**, 11, 497-504	22.2	811
2379	Update on tick-borne rickettsioses around the world: a geographic approach. *Clinical Microbiology Reviews*, **2013**, 26, 657-702	34	783
2378	The 1.2-megabase genome sequence of Mimivirus. *Science*, **2004**, 306, 1344-50	33.3	774
2377	Tick-borne rickettsioses around the world: emerging diseases challenging old concepts. *Clinical Microbiology Reviews*, **2005**, 18, 719-56	34	767
2376	Microorganisms resistant to free-living amoebae. *Clinical Microbiology Reviews*, **2004**, 17, 413-33	34	767
2375	16S ribosomal DNA sequence analysis of a large collection of environmental and clinical unidentifiable bacterial isolates. *Journal of Clinical Microbiology*, **2000**, 38, 3623-30	9.7	723
2374	Microbial culturomics: paradigm shift in the human gut microbiome study. *Clinical Microbiology and Infection*, **2012**, 18, 1185-93	9.5	722
2373	New insights on the antiviral effects of chloroquine against coronavirus: what to expect for COVID-19?. *International Journal of Antimicrobial Agents*, **2020**, 55, 105938	14.3	663
2372	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	3.7	588
2371	A giant virus in amoebae. *Science*, **2003**, 299, 2033	33.3	558
2370	Culture of previously uncultured members of the human gut microbiota by culturomics. *Nature Microbiology*, **2016**, 1, 16203	26.6	555
2369	Comparative genomics of multidrug resistance in Acinetobacter baumannii. *PLoS Genetics*, **2006**, 2, e7	6	553
2368	Endocarditis due to rare and fastidious bacteria. *Clinical Microbiology Reviews*, **2001**, 14, 177-207	34	544

2367	Pig liver sausage as a source of hepatitis E virus transmission to humans. *Journal of Infectious Diseases*, **2010**, 202, 825-34	7	502
2366	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	5.3	494
2365	The rebirth of culture in microbiology through the example of culturomics to study human gut microbiota. *Clinical Microbiology Reviews*, **2015**, 28, 237-64	34	490
2364	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	8.4	479
2363	Whipple's disease. *New England Journal of Medicine*, **2007**, 356, 55-66	59.2	452
2362	Risk of embolism and death in infective endocarditis: prognostic value of echocardiography: a prospective multicenter study. *Circulation*, **2005**, 112, 69-75	16.7	446
2361	Obesity-associated gut microbiota is enriched in Lactobacillus reuteri and depleted in Bifidobacterium animalis and Methanobrevibacter smithii. *International Journal of Obesity*, **2012**, 36, 817-25	5.5	412
2360	Methanomassiliicoccus luminyensis gen. nov., sp. nov., a methanogenic archaeon isolated from human faeces. *International Journal of Systematic and Evolutionary Microbiology*, **2012**, 62, 1902-1907	2.2	410
2359	Q Fever. *Veterinary Microbiology*, **2010**, 140, 297-309	3.3	409
2358	Diagnosis of Q fever. *Journal of Clinical Microbiology*, **1998**, 36, 1823-34	9.7	404
2357	rpoB sequence analysis as a novel basis for bacterial identification. *Molecular Microbiology*, **1997**, 26, 1005-11	4.1	402
2356	The virophage as a unique parasite of the giant mimivirus. *Nature*, **2008**, 455, 100-4	50.4	389
2355	From Q Fever to Coxiella burnetii Infection: a Paradigm Change. *Clinical Microbiology Reviews*, **2017**, 30, 115-190	34	388
2354	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, 12168-73	11.5	383
2353	Cultivation of the bacillus of Whipple's disease. *New England Journal of Medicine*, **2000**, 342, 620-5	59.2	382
2352	Enterococcus hirae and Barnesiella intestinihominis Facilitate Cyclophosphamide-Induced Therapeutic Immunomodulatory Effects. *Immunity*, **2016**, 45, 931-943	32.3	376
2351	Mechanisms of evolution in Rickettsia conorii and R. prowazekii. *Science*, **2001**, 293, 2093-8	33.3	365
2350	Recommendations for treatment of human infections caused by Bartonella species. *Antimicrobial Agents and Chemotherapy*, **2004**, 48, 1921-33	5.9	357

2349	Q fever 1985-1998. Clinical and epidemiologic features of 1,383 infections. *Medicine (United States)*, **2000**, 79, 109-23	1.8	353
2348	Chikungunya outbreaks--the globalization of vectorborne diseases. *New England Journal of Medicine*, **2007**, 356, 769-71	59.2	346
2347	The microbiome in cancer immunotherapy: Diagnostic tools and therapeutic strategies. *Science*, **2018**, 359, 1366-1370	33.3	341
2346	Laboratory diagnosis of rickettsioses: current approaches to diagnosis of old and new rickettsial diseases. *Journal of Clinical Microbiology*, **1997**, 35, 2715-27	9.7	323
2345	Comprehensive diagnostic strategy for blood culture-negative endocarditis: a prospective study of 819 new cases. *Clinical Infectious Diseases*, **2010**, 51, 131-40	11.6	322
2344	Positron emission tomography/computed tomography for diagnosis of prosthetic valve endocarditis: increased valvular 18F-fluorodeoxyglucose uptake as a novel major criterion. *Journal of the American College of Cardiology*, **2013**, 61, 2374-82	15.1	318
2343	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, 2058-65	9.7	315
2342	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, 2182-94	9.7	312
2341	Blood culture-negative endocarditis in a reference center: etiologic diagnosis of 348 cases. *Medicine (United States)*, **2005**, 84, 162-173	1.8	305
2340	Culturing the human microbiota and culturomics. *Nature Reviews Microbiology*, **2018**, 16, 540-550	22.2	303
2339	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	8.4	300
2338	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	2.2	298
2337	Echocardiography predicts embolic events in infective endocarditis. *Journal of the American College of Cardiology*, **2001**, 37, 1069-76	15.1	293
2336	Bartonella (Rochalimaea) quintana endocarditis in three homeless men. *New England Journal of Medicine*, **1995**, 332, 419-23	59.2	292
2335	Comparative meta-analysis of the effect of Lactobacillus species on weight gain in humans and animals. *Microbial Pathogenesis*, **2012**, 53, 100-8	3.8	290
2334	Gene sequence-based criteria for identification of new rickettsia isolates and description of Rickettsia heilongjiangensis sp. nov. *Journal of Clinical Microbiology*, **2003**, 41, 5456-65	9.7	289
2333	Redefining viruses: lessons from Mimivirus. *Nature Reviews Microbiology*, **2008**, 6, 315-9	22.2	285
2332	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	3.7	277

2331	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, 133-43	2.2	276
2330	Sequencing of the rpoB gene and flanking spacers for molecular identification of Acinetobacter species. *Journal of Clinical Microbiology*, **2006**, 44, 827-32	9.7	275
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2328	Q fever. *Clinical Infectious Diseases*, **1995**, 20, 489-95; quiz 496	11.6	272
2327	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, 21848-53	11.5	270
2326	MALDI-TOF-mass spectrometry applications in clinical microbiology. *Future Microbiology*, **2010**, 5, 1733-54	2.9	266
2325	Whipple's disease: new aspects of pathogenesis and treatment. *Lancet Infectious Diseases, The*, **2008**, 8, 179-90	25.5	265
2324	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	8.5	264
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2321	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	14.3	259
2320	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, 925-32	25.5	258
2319	Diagnosis of 22 new cases of Bartonella endocarditis. *Annals of Internal Medicine*, **1996**, 125, 646-52	8	255
2318	Epidemiologic features and clinical presentation of acute Q fever in hospitalized patients: 323 French cases. *American Journal of Medicine*, **1992**, 93, 427-34	2.4	250
2317	Treatment of Q fever endocarditis: comparison of 2 regimens containing doxycycline and ofloxacin or hydroxychloroquine. *Archives of Internal Medicine*, **1999**, 159, 167-73		249
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2315	Impact of cerebrovascular complications on mortality and neurologic outcome during infective endocarditis: a prospective multicentre study. *European Heart Journal*, **2007**, 28, 1155-61	9.5	247
2314	Comparative analysis of Acinetobacters: three genomes for three lifestyles. *PLoS ONE*, **2008**, 3, e1805	3.7	246

2313	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, 12637-40	11.5	243
2312	Staphylococcus aureus native valve infective endocarditis: report of 566 episodes from the International Collaboration on Endocarditis Merged Database. *Clinical Infectious Diseases*, **2005**, 41, 507-14	11.6	241
2311	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, 12800-3	11.5	241
2310	The Q fever epidemic in The Netherlands: history, onset, response and reflection. *Epidemiology and Infection*, **2011**, 139, 1-12	4.3	240
2309	Novel chikungunya virus variant in travelers returning from Indian Ocean islands. *Emerging Infectious Diseases*, **2006**, 12, 1493-9	10.2	240
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2307	Current and past strategies for bacterial culture in clinical microbiology. *Clinical Microbiology Reviews*, **2015**, 28, 208-36	34	239
2306	Genome sequence of Rickettsia bellii illuminates the role of amoebae in gene exchanges between intracellular pathogens. *PLoS Genetics*, **2006**, 2, e76	6	239
2305	Amoebal coculture of "Mycobacterium massiliense" sp. nov. from the sputum of a patient with hemoptoic pneumonia. *Journal of Clinical Microbiology*, **2004**, 42, 5493-501	9.7	239
2304	Molecular detection of Bartonella quintana, B. koehlerae, B. henselae, B. clarridgeiae, Rickettsia felis, and Wolbachia pipientis in cat fleas, France. *Emerging Infectious Diseases*, **2003**, 9, 338-42	10.2	237
2303	The relationship between gut microbiota and weight gain in humans. *Future Microbiology*, **2012**, 7, 91-109	2.9	235
2302	Fleas and flea-borne diseases. *International Journal of Infectious Diseases*, **2010**, 14, e667-76	10.5	233
2301	Eukaryotic large nucleo-cytoplasmic DNA viruses: clusters of orthologous genes and reconstruction of viral genome evolution. *Virology Journal*, **2009**, 6, 223	6.1	229
2300	Characterization of the naturally occurring oxacillinase of Acinetobacter baumannii. *Antimicrobial Agents and Chemotherapy*, **2005**, 49, 4174-9	5.9	229
2299	Rickettsia africae, a tick-borne pathogen in travelers to sub-Saharan Africa. *New England Journal of Medicine*, **2001**, 344, 1504-10	59.2	225
2298	The genome sequence of Rickettsia felis identifies the first putative conjugative plasmid in an obligate intracellular parasite. *PLoS Biology*, **2005**, 3, e248	9.7	224
2297	Risks factors and prevention of Q fever endocarditis. *Clinical Infectious Diseases*, **2001**, 33, 312-6	11.6	224
2296	rpoB gene sequence-based identification of Staphylococcus species. *Journal of Clinical Microbiology* , **2002**, 40, 1333-8	9.7	221

2295	Modern clinical microbiology: new challenges and solutions. *Nature Reviews Microbiology*, **2013**, 11, 574-85	22.2	220
2294	Molecular quantification of Gardnerella vaginalis and Atopobium vaginae loads to predict bacterial vaginosis. *Clinical Infectious Diseases*, **2008**, 47, 33-43	11.6	219
2293	Tick- and flea-borne rickettsial emerging zoonoses. *Veterinary Research*, **2005**, 36, 469-92	3.8	215
2292	Bacterial strain typing in the genomic era. *FEMS Microbiology Reviews*, **2009**, 33, 892-916	15.1	213
2291	rpoB gene sequencing for identification of Corynebacterium species. *Journal of Clinical Microbiology*, **2004**, 42, 3925-31	9.7	213
2290	Gene-sequence-based criteria for species definition in bacteriology: the Bartonella paradigm. *Trends in Microbiology*, **2003**, 11, 318-21	12.4	213
2289	Chronic Bartonella quintana bacteremia in homeless patients. *New England Journal of Medicine*, **1999**, 340, 184-9	59.2	210
2288	The rpoB gene as a tool for clinical microbiologists. *Trends in Microbiology*, **2009**, 17, 37-45	12.4	208
2287	Whipple's disease. *Lancet, The*, **2003**, 361, 239-46	40	208
2286	Human gut microbiota: repertoire and variations. *Frontiers in Cellular and Infection Microbiology*, **2012**, 2, 136	5.9	207
2285	Value and limitations of the Duke criteria for the diagnosis of infective endocarditis. *Journal of the American College of Cardiology*, **1999**, 33, 2023-9	15.1	207
2284	Genome of Acanthamoeba castellanii highlights extensive lateral gene transfer and early evolution of tyrosine kinase signaling. *Genome Biology*, **2013**, 14, R11	18.3	205
2283	Understanding the cholera epidemic, Haiti. *Emerging Infectious Diseases*, **2011**, 17, 1161-8	10.2	201
2282	The two faces of interleukin 10 in human infectious diseases. *Lancet Infectious Diseases, The*, **2006**, 6, 557-69	25.5	200
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2280	Wind in November, Q fever in December. *Emerging Infectious Diseases*, **2004**, 10, 1264-9	10.2	188
2279	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, 1018-28	9.7	187
2278	Rickettsioses and the international traveler. *Clinical Infectious Diseases*, **2004**, 39, 1493-9	11.6	184

2277	Detection of ehrlichiae in African ticks by polymerase chain reaction. *Transactions of the Royal Society of Tropical Medicine and Hygiene*, **2000**, 94, 707-8	2	184
2276	Culture of Bartonella quintana and Bartonella henselae from human samples: a 5-year experience (1993 to 1998). *Journal of Clinical Microbiology*, **1999**, 37, 1899-905	9.7	184
2275	Warmer weather linked to tick attack and emergence of severe rickettsioses. *PLoS Neglected Tropical Diseases*, **2008**, 2, e338	4.8	183
2274	Use of rpoB gene analysis for detection and identification of Bartonella species. *Journal of Clinical Microbiology*, **2001**, 39, 430-7	9.7	182
2273	Rickettsial infections and fever, Vientiane, Laos. *Emerging Infectious Diseases*, **2006**, 12, 256-62	10.2	179
2272	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, 500-7	14.3	177
2271	Structural studies of the giant mimivirus. *PLoS Biology*, **2009**, 7, e92	9.7	176
2270	Dramatic reduction in infective endocarditis-related mortality with a management-based approach. *Archives of Internal Medicine*, **2009**, 169, 1290-8		176
2269	Coxiella burnetii genotyping. *Emerging Infectious Diseases*, **2005**, 11, 1211-7	10.2	175
2268	Propionibacterium acnes postoperative shoulder arthritis: an emerging clinical entity. *Clinical Infectious Diseases*, **2008**, 46, 1884-6	11.6	174
2267	In vitro testing of combined hydroxychloroquine and azithromycin on SARS-CoV-2 shows synergistic effect. *Microbial Pathogenesis*, **2020**, 145, 104228	3.8	171
2266	A comprehensive repertoire of prokaryotic species identified in human beings. *Lancet Infectious Diseases, The*, **2015**, 15, 1211-1219	25.5	170
2265	Rickettsial evolution in the light of comparative genomics. *Biological Reviews*, **2011**, 86, 379-405	13.5	170
2264	Molecular detection of multiple emerging pathogens in sputa from cystic fibrosis patients. *PLoS ONE*, **2008**, 3, e2908	3.7	170
2263	A flea-associated Rickettsia pathogenic for humans. *Emerging Infectious Diseases*, **2001**, 7, 73-81	10.2	169
2262	Epidemiologic and clinical characteristics of Bartonella quintana and Bartonella henselae endocarditis: a study of 48 patients. *Medicine (United States)*, **2001**, 80, 245-51	1.8	166
2261	Long-term outcome of Q fever endocarditis: a 26-year personal survey. *Lancet Infectious Diseases, The*, **2010**, 10, 527-35	25.5	165
2260	Rickettsia slovaca and R. raoultii in tick-borne Rickettsioses. *Emerging Infectious Diseases*, **2009**, 15, 1105-8	10.2	164

2259	Genome-based design of a cell-free culture medium for Tropheryma whipplei. *Lancet, The*, **2003**, 362, 447-9	40	163
2258	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	8.4	162
2257	Massive comparative genomic analysis reveals convergent evolution of specialized bacteria. *Biology Direct*, **2009**, 4, 13	7.2	162
2256	African tick bite fever. *Lancet Infectious Diseases, The*, **2003**, 3, 557-64	25.5	159
2255	Rickettsia massiliae human isolation. *Emerging Infectious Diseases*, **2006**, 12, 174-5	10.2	155
2254	Outcome and treatment of Bartonella endocarditis. *Archives of Internal Medicine*, **2003**, 163, 226-30		151
2253	Spotless rickettsiosis caused by Rickettsia slovaca and associated with Dermacentor ticks. *Clinical Infectious Diseases*, **2002**, 34, 1331-6	11.6	151
2252	Infections in the homeless. *Lancet Infectious Diseases, The*, **2001**, 1, 77-84	25.5	150
2251	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, 18078-83	11.5	148
2250	Comparison of PCR and serology assays for early diagnosis of acute Q fever. *Journal of Clinical Microbiology*, **2003**, 41, 5094-8	9.7	148
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2247	Molecular diagnosis of bloodstream infections caused by non-cultivable bacteria. *International Journal of Antimicrobial Agents*, **2007**, 30 Suppl 1, S7-15	14.3	145
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2245	Validation of partial rpoB gene sequence analysis for the identification of clinically important and emerging Acinetobacter species. *Microbiology (United Kingdom)*, **2009**, 155, 2333-2341	2.9	144
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2234	Tailed giant Tupanvirus possesses the most complete translational apparatus of the known virosphere. *Nature Communications*, **2018**, 9, 749	17.4	136
2233	Reclassification of giant viruses composing a fourth domain of life in the new order Megavirales. *Intervirology*, **2012**, 55, 321-32	2.5	134
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2231	Tropheryma whipplei Twist: a human pathogenic Actinobacteria with a reduced genome. *Genome Research*, **2003**, 13, 1800-9	9.7	132
2230	High-level colonisation of the human gut by Verrucomicrobia following broad-spectrum antibiotic treatment. *International Journal of Antimicrobial Agents*, **2013**, 41, 149-55	14.3	131
2229	Coxiella burnetii in humans and ticks in rural Senegal. *PLoS Neglected Tropical Diseases*, **2010**, 4, e654	4.8	131
2228	Emerging rickettsioses of the Thai-Myanmar border. *Emerging Infectious Diseases*, **2003**, 9, 592-5	10.2	131
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2226	Mediterranean spotted fever: clinical, laboratory and epidemiological features of 199 cases. *American Journal of Tropical Medicine and Hygiene*, **1986**, 35, 845-50	3.2	130
2225	Evidence for louse-transmitted diseases in soldiers of Napoleon's Grand Army in Vilnius. *Journal of Infectious Diseases*, **2006**, 193, 112-20	7	129
2224	Related actions of probiotics and antibiotics on gut microbiota and weight modification. *Lancet Infectious Diseases, The*, **2013**, 13, 889-99	25.5	128

2223	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	6	128
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2221	PCR detection of bacteria on cardiac valves of patients with treated bacterial endocarditis. *Journal of Clinical Microbiology*, **2005**, 43, 163-7	9.7	127
2220	Palaeomicrobiology: current issues and perspectives. *Nature Reviews Microbiology*, **2005**, 3, 23-35	22.2	124
2219	Mimivirus in pneumonia patients. *Emerging Infectious Diseases*, **2005**, 11, 449-52	10.2	123
2218	Comparison between rpoB and 16S rRNA gene sequencing for molecular identification of 168 clinical isolates of Corynebacterium. *Journal of Clinical Microbiology*, **2005**, 43, 1934-6	9.7	123
2217	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. *FEMS Microbiology Reviews*, **2010**, 34, 281-294	15.1	122
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2215	Effect of sex on Coxiella burnetii infection: protective role of 17beta-estradiol. *Journal of Infectious Diseases*, **2004**, 189, 339-45	7	121
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2213	Survival of environmental mycobacteria in Acanthamoeba polyphaga. *Applied and Environmental Microbiology*, **2006**, 72, 5974-81	4.8	119
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