

Pierre E Fournier

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801
papers

28,827
citations

86
h-index

149
g-index

865
ext. papers

33,983
ext. citations

6.3
avg, IF

7.1
L-index

#	Paper	IF	Citations
801	Ongoing revolution in bacteriology: routine identification of bacteria by matrix-assisted laser desorption ionization time-of-flight mass spectrometry. <i>Clinical Infectious Diseases</i> , 2009 , 49, 543-51	11.6	1396
800	Update on tick-borne rickettsioses around the world: a geographic approach. <i>Clinical Microbiology Reviews</i> , 2013 , 26, 657-702	34	783
799	The epidemiology and control of <i>Acinetobacter baumannii</i> in health care facilities. <i>Clinical Infectious Diseases</i> , 2006 , 42, 692-9	11.6	625
798	Culture of previously uncultured members of the human gut microbiota by culturomics. <i>Nature Microbiology</i> , 2016 , 1, 16203	26.6	555
797	Comparative genomics of multidrug resistance in <i>Acinetobacter baumannii</i> . <i>PLoS Genetics</i> , 2006 , 2, e7	6	553
796	The rebirth of culture in microbiology through the example of culturomics to study human gut microbiota. <i>Clinical Microbiology Reviews</i> , 2015 , 28, 237-64	34	490
795	Clinical and microbiological effect of a combination of hydroxychloroquine and azithromycin in 80 COVID-19 patients with at least a six-day follow up: A pilot observational study. <i>Travel Medicine and Infectious Disease</i> , 2020 , 34, 101663	8.4	479
794	Diagnosis of Q fever. <i>Journal of Clinical Microbiology</i> , 1998 , 36, 1823-34	9.7	404
793	Cultivation of the bacillus of Whipple's disease. <i>New England Journal of Medicine</i> , 2000 , 342, 620-5	59.2	382
792	Mechanisms of evolution in <i>Rickettsia conorii</i> and <i>R. prowazekii</i> . <i>Science</i> , 2001 , 293, 2093-8	33.3	365
791	Q fever 1985-1998. Clinical and epidemiologic features of 1,383 infections. <i>Medicine (United States)</i> , 2000 , 79, 109-23	1.8	353
790	Comprehensive diagnostic strategy for blood culture-negative endocarditis: a prospective study of 819 new cases. <i>Clinical Infectious Diseases</i> , 2010 , 51, 131-40	11.6	322
789	Differentiation of spotted fever group rickettsiae by sequencing and analysis of restriction fragment length polymorphism of PCR-amplified DNA of the gene encoding the protein rOmpA. <i>Journal of Clinical Microbiology</i> , 1996 , 34, 2058-65	9.7	315
788	Identification of rare pathogenic bacteria in a clinical microbiology laboratory: impact of matrix-assisted laser desorption ionization-time of flight mass spectrometry. <i>Journal of Clinical Microbiology</i> , 2013 , 51, 2182-94	9.7	312
787	Culturing the human microbiota and culturomics. <i>Nature Reviews Microbiology</i> , 2018 , 16, 540-550	22.2	303
786	Early treatment of COVID-19 patients with hydroxychloroquine and azithromycin: A retrospective analysis of 1061 cases in Marseille, France. <i>Travel Medicine and Infectious Disease</i> , 2020 , 35, 101738	8.4	300
785	A polyphasic strategy incorporating genomic data for the taxonomic description of novel bacterial species. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014 , 64, 384-391	2.2	298

784	Gene sequence-based criteria for identification of new rickettsia isolates and description of <i>Rickettsia heilongjiangensis</i> sp. nov. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 5456-65	9.7	289
783	MALDI-TOF-mass spectrometry applications in clinical microbiology. <i>Future Microbiology</i> , 2010 , 5, 1733-54	9	266
782	Guidelines for the diagnosis of tick-borne bacterial diseases in Europe. <i>Clinical Microbiology and Infection</i> , 2004 , 10, 1108-32	9.5	260
781	Diagnosis of 22 new cases of Bartonella endocarditis. <i>Annals of Internal Medicine</i> , 1996 , 125, 646-52	8	255
780	Genome sequence of <i>Rickettsia bellii</i> illuminates the role of amoebae in gene exchanges between intracellular pathogens. <i>PLoS Genetics</i> , 2006 , 2, e76	6	239
779	Characterization of the naturally occurring oxacillinase of <i>Acinetobacter baumannii</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2005 , 49, 4174-9	5.9	229
778	Clinical features and prognostic factors of listeriosis: the MONALISA national prospective cohort study. <i>Lancet Infectious Diseases</i> , 2017 , 17, 510-519	25.5	227
777	<i>Rickettsia africae</i> , a tick-borne pathogen in travelers to sub-Saharan Africa. <i>New England Journal of Medicine</i> , 2001 , 344, 1504-10	59.2	225
776	Phylogenetic analysis of spotted fever group rickettsiae by study of the outer surface protein rOmpA. <i>International Journal of Systematic Bacteriology</i> , 1998 , 48 Pt 3, 839-49		225
775	The genome sequence of <i>Rickettsia felis</i> identifies the first putative conjugative plasmid in an obligate intracellular parasite. <i>PLoS Biology</i> , 2005 , 3, e248	9.7	224
774	Risks factors and prevention of Q fever endocarditis. <i>Clinical Infectious Diseases</i> , 2001 , 33, 312-6	11.6	224
773	Modern clinical microbiology: new challenges and solutions. <i>Nature Reviews Microbiology</i> , 2013 , 11, 574-85	25.2	220
772	Bacterial strain typing in the genomic era. <i>FEMS Microbiology Reviews</i> , 2009 , 33, 892-916	15.1	213
771	Value and limitations of the Duke criteria for the diagnosis of infective endocarditis. <i>Journal of the American College of Cardiology</i> , 1999 , 33, 2023-9	15.1	207
770	Rickettsioses and the international traveler. <i>Clinical Infectious Diseases</i> , 2004 , 39, 1493-9	11.6	184
769	Warmer weather linked to tick attack and emergence of severe rickettsioses. <i>PLoS Neglected Tropical Diseases</i> , 2008 , 2, e338	4.8	183
768	A comprehensive repertoire of prokaryotic species identified in human beings. <i>Lancet Infectious Diseases</i> , 2015 , 15, 1211-1219	25.5	170
767	Modification of the diagnostic criteria proposed by the Duke Endocarditis Service to permit improved diagnosis of Q fever endocarditis. <i>American Journal of Medicine</i> , 1996 , 100, 629-33	2.4	170

766	A flea-associated Rickettsia pathogenic for humans. <i>Emerging Infectious Diseases</i> , 2001 , 7, 73-81	10.2	169
765	Epidemiologic and clinical characteristics of Bartonella quintana and Bartonella henselae endocarditis: a study of 48 patients. <i>Medicine (United States)</i> , 2001 , 80, 245-51	1.8	166
764	Outcomes of 3,737 COVID-19 patients treated with hydroxychloroquine/azithromycin and other regimens in Marseille, France: A retrospective analysis. <i>Travel Medicine and Infectious Disease</i> , 2020 , 36, 101791	8.4	162
763	African tick bite fever. <i>Lancet Infectious Diseases</i> , 2003 , 3, 557-64	25.5	159
762	Evidence of Rickettsia helvetica infection in humans, eastern France. <i>Emerging Infectious Diseases</i> , 2000 , 6, 389-92	10.2	157
761	Significant molecular and systemic adaptations after repeated sprint training in hypoxia. <i>PLoS ONE</i> , 2013 , 8, e56522	3.7	155
760	Description of Tropheryma whipplei gen. nov., sp. nov., the Whipple's disease bacillus. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2001 , 51, 1471-1479	2.2	153
759	Outcome and treatment of Bartonella endocarditis. <i>Archives of Internal Medicine</i> , 2003 , 163, 226-30		151
758	Spotless rickettsiosis caused by Rickettsia slovaca and associated with Dermacentor ticks. <i>Clinical Infectious Diseases</i> , 2002 , 34, 1331-6	11.6	151
757	Comparison of PCR and serology assays for early diagnosis of acute Q fever. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 5094-8	9.7	148
756	rpoB gene sequence-based identification of aerobic Gram-positive cocci of the genera Streptococcus, Enterococcus, Gemella, Abiotrophia, and Granulicatella. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 497-504	9.7	146
755	Complementarity between targeted real-time specific PCR and conventional broad-range 16S rDNA PCR in the syndrome-driven diagnosis of infectious diseases. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2015 , 34, 561-70	5.3	142
754	Mimivirus and the emerging concept of "giant" virus. <i>Virus Research</i> , 2006 , 117, 133-44	6.4	141
753	Genotyping, Orientalis-like Yersinia pestis, and plague pandemics. <i>Emerging Infectious Diseases</i> , 2004 , 10, 1585-92	10.2	141
752	African tick bite fever in travelers to rural sub-Equatorial Africa. <i>Clinical Infectious Diseases</i> , 2003 , 36, 1411-7	11.6	141
751	Diagnosis and management of Q fever--United States, 2013: recommendations from CDC and the Q Fever Working Group. <i>MMWR Recommendations and Reports</i> , 2013 , 62, 1-30	37.3	141
750	Molecular detection of Coxiella burnetii in the sera of patients with Q fever endocarditis or vascular infection. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 4919-24	9.7	133
749	The bacterial pangenome as a new tool for analysing pathogenic bacteria. <i>New Microbes and New Infections</i> , 2015 , 7, 72-85	4.1	130

748	Evidence for louse-transmitted diseases in soldiers of Napoleon's Grand Army in Vilnius. <i>Journal of Infectious Diseases</i> , 2006 , 193, 112-20	7	129
747	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. <i>FEMS Microbiology Reviews</i> , 2010 , 34, 281-294	15.1	122
746	Asynchrony between the rates of standing height gain and bone mass accumulation during puberty. <i>Osteoporosis International</i> , 1997 , 7, 525-32	5.3	120
745	First documented human <i>Rickettsia aeschlimannii</i> infection. <i>Emerging Infectious Diseases</i> , 2002 , 8, 748-9	10.2	120
744	<i>Rickettsia raoultii</i> sp. nov., a spotted fever group rickettsia associated with Dermacentor ticks in Europe and Russia. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008 , 58, 1635-9	2.2	116
743	Spotted fever rickettsioses in southern and eastern Europe. <i>FEMS Immunology and Medical Microbiology</i> , 2007 , 49, 2-12		113
742	Influence of the timing of cardiac surgery on the outcome of patients with infective endocarditis and stroke. <i>Clinical Infectious Diseases</i> , 2013 , 56, 209-17	11.6	111
741	<i>Coxiella burnetii</i> infection of aortic aneurysms or vascular grafts: report of 30 new cases and evaluation of outcome. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2007 , 26, 635-40	5.3	111
740	Aneruptive fever associated with antibodies to <i>Rickettsia helvetica</i> in Europe and Thailand. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 816-8	9.7	110
739	HACEK infective endocarditis: characteristics and outcomes from a large, multi-national cohort. <i>PLoS ONE</i> , 2013 , 8, e63181	3.7	108
738	Lymphangitis-associated rickettsiosis, a new rickettsiosis caused by <i>Rickettsia sibirica mongolotimonae</i> : seven new cases and review of the literature. <i>Clinical Infectious Diseases</i> , 2005 , 40, 1435-44	11.6	108
737	From culturomics to taxonomogenomics: A need to change the taxonomy of prokaryotes in clinical microbiology. <i>Anaerobe</i> , 2015 , 36, 73-8	2.8	106
736	Diagnosis of infectious endocarditis in patients undergoing valve surgery. <i>American Journal of Medicine</i> , 2005 , 118, 230-8	2.4	106
735	Genetic identification of rickettsiae isolated from ticks in Japan. <i>Journal of Clinical Microbiology</i> , 2002 , 40, 2176-81	9.7	106
734	Burden of emerging anaerobes in the MALDI-TOF and 16S rRNA gene sequencing era. <i>Anaerobe</i> , 2011 , 17, 106-12	2.8	105
733	<i>Rickettsia felis</i> infection acquired in Europe and documented by polymerase chain reaction. <i>Emerging Infectious Diseases</i> , 2002 , 8, 207-8	10.2	105
732	Human pathogens in body and head lice. <i>Emerging Infectious Diseases</i> , 2002 , 8, 1515-8	10.2	104
731	Selfish DNA in protein-coding genes of <i>Rickettsia</i> . <i>Science</i> , 2000 , 290, 347-50	33.3	104

730	HIV-2 integrase gene polymorphism and phenotypic susceptibility of HIV-2 clinical isolates to the integrase inhibitors raltegravir and elvitegravir in vitro. <i>Journal of Antimicrobial Chemotherapy</i> , 2008 , 62, 914-20	5.1	101
729	Genome sequence and description of <i>Anaerosolibacter massiliensis</i> sp. nov. <i>New Microbes and New Infections</i> , 2016 , 10, 66-76	4.1	98
728	Prognosis and treatment of paraquat poisoning: a review of 28 cases. <i>Journal of Toxicology: Clinical Toxicology</i> , 1982 , 19, 461-74		97
727	Non-steroidal anti-inflammatory drugs for athletes: an update. <i>Annals of Physical and Rehabilitation Medicine</i> , 2010 , 53, 278-82, 282-8	3.8	95
726	Analysis of the <i>Rickettsia africae</i> genome reveals that virulence acquisition in <i>Rickettsia</i> species may be explained by genome reduction. <i>BMC Genomics</i> , 2009 , 10, 166	4.5	95
725	Acute tick-borne rickettsiosis caused by <i>Rickettsia heilongjiangensis</i> in Russian Far East. <i>Emerging Infectious Diseases</i> , 2004 , 10, 810-7	10.2	95
724	Molecular evolution of rickettsia surface antigens: evidence of positive selection. <i>Molecular Biology and Evolution</i> , 2005 , 22, 2073-83	8.3	95
723	Cautionary tale of using 16S rRNA gene sequence similarity values in identification of human-associated bacterial species. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015 , 65, 1929-1934	2.2	94
722	Proposal to create subspecies of <i>Rickettsia conorii</i> based on multi-locus sequence typing and an emended description of <i>Rickettsia conorii</i> . <i>BMC Microbiology</i> , 2005 , 5, 11	4.5	94
721	Genome analysis of microorganisms living in amoebae reveals a melting pot of evolution. <i>FEMS Microbiology Reviews</i> , 2010 , 34, 281-94	15.1	94
720	Non-contiguous finished genome sequence and description of <i>Anaerococcus senegalensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 116-25		93
719	Isolation of <i>Rickettsia parkeri</i> and identification of a novel spotted fever group <i>Rickettsia</i> sp. from Gulf Coast ticks (<i>Amblyomma maculatum</i>) in the United States. <i>Applied and Environmental Microbiology</i> , 2010 , 76, 2689-96	4.8	91
718	Outbreak of <i>Rickettsia africae</i> infections in participants of an adventure race in South Africa. <i>Clinical Infectious Diseases</i> , 1998 , 27, 316-23	11.6	90
717	Respiratory viruses and bacteria among pilgrims during the 2013 Hajj. <i>Emerging Infectious Diseases</i> , 2014 , 20, 1821-7	10.2	87
716	Genomic variation of <i>Bartonella henselae</i> strains detected in lymph nodes of patients with cat scratch disease. <i>Journal of Clinical Microbiology</i> , 2002 , 40, 1023-30	9.7	87
715	Current knowledge on phylogeny and taxonomy of <i>Rickettsia</i> spp. <i>Annals of the New York Academy of Sciences</i> , 2009 , 1166, 1-11	6.5	85
714	Non-contiguous finished genome sequence and description of <i>Alistipes timonensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 315-24		84
713	The human gut microbiome, a taxonomic conundrum. <i>Systematic and Applied Microbiology</i> , 2015 , 38, 276-86	4.2	82

712	What does the future hold for clinical microbiology?. <i>Nature Reviews Microbiology</i> , 2004 , 2, 151-9	22.2	82
711	Use of genome selected repeated sequences increases the sensitivity of PCR detection of <i>Tropheryma whipplei</i> . <i>Journal of Clinical Microbiology</i> , 2004 , 42, 401-3	9.7	81
710	<i>Coxiella burnetii</i> infection of aneurysms or vascular grafts: report of seven cases and review. <i>Clinical Infectious Diseases</i> , 1998 , 26, 116-21	11.6	81
709	Clinical detection and characterization of bacterial pathogens in the genomics era. <i>Genome Medicine</i> , 2014 , 6, 114	14.4	80
708	Evaluation of the Panbio COVID-19 Rapid Antigen Detection Test Device for the Screening of Patients with COVID-19. <i>Journal of Clinical Microbiology</i> , 2021 , 59,	9.7	79
707	<i>Bartonella</i> , a common cause of endocarditis: a report on 106 cases and review. <i>Journal of Clinical Microbiology</i> , 2015 , 53, 824-9	9.7	77
706	Use of highly variable intergenic spacer sequences for multispacer typing of <i>Rickettsia conorii</i> strains. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 5757-66	9.7	75
705	Non contiguous-finished genome sequence and description of <i>Peptoniphilus timonensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 1-11		74
704	Real-time PCR for universal antibiotic susceptibility testing. <i>Journal of Antimicrobial Chemotherapy</i> , 2004 , 54, 538-41	5.1	74
703	Phylogenetic classification of <i>Bartonella</i> species by comparing groEL sequences. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2002 , 52, 165-71	2.2	74
702	Diagnosis of <i>Bartonella</i> endocarditis by a real-time nested PCR assay using serum. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 919-25	9.7	73
701	Detection of members of the genera <i>Rickettsia</i> , <i>Anaplasma</i> , and <i>Ehrlichia</i> in ticks collected in the Asiatic part of Russia. <i>Annals of the New York Academy of Sciences</i> , 2006 , 1078, 378-83	6.5	72
700	Naming of <i>Rickettsiae</i> and rickettsial diseases. <i>Annals of the New York Academy of Sciences</i> , 2005 , 1063, 1-12	6.5	72
699	Metagenomic analysis of brain abscesses identifies specific bacterial associations. <i>Clinical Infectious Diseases</i> , 2012 , 54, 202-10	11.6	71
698	Bacterial genome sequencing and its use in infectious diseases. <i>Lancet Infectious Diseases</i> , 2007 , 7, 711-23	25.5	71
697	Suicide PCR on skin biopsy specimens for diagnosis of rickettsioses. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 3428-34	9.7	71
696	Genotyping of human lice suggests multiple emergencies of body lice from local head louse populations. <i>PLoS Neglected Tropical Diseases</i> , 2010 , 4, e641	4.8	69
695	Advantages and limitations of genomics in prokaryotic taxonomy. <i>Clinical Microbiology and Infection</i> , 2013 , 19, 790-5	9.5	68

694	Genotyping reveals a wide heterogeneity of <i>Tropheryma whippelii</i> . <i>Microbiology (United Kingdom)</i> , 2008 , 154, 521-527	2.9	68
693	Quantitative analysis of valvular lesions during <i>Bartonella</i> endocarditis. <i>American Journal of Clinical Pathology</i> , 2000 , 114, 880-9	1.9	67
692	Update on blood culture-negative endocarditis. <i>Médecine Et Maladies Infectieuses</i> , 2015 , 45, 1-8	4	65
691	Non-contiguous finished genome sequence and description of <i>Clostridium senegalense</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 386-95		65
690	Genome sequence and description of <i>Aeromicrobium massiliense</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 246-57		65
689	An experimental model of human body louse infection with <i>Rickettsia prowazekii</i> . <i>Journal of Infectious Diseases</i> , 2002 , 186, 1639-46	7	65
688	Blood culture-negative endocarditis: Improving the diagnostic yield using new diagnostic tools. <i>Medicine (United States)</i> , 2017 , 96, e8392	1.8	63
687	Culture and immunological detection of <i>Tropheryma whippelii</i> from the duodenum of a patient with Whipple disease. <i>JAMA - Journal of the American Medical Association</i> , 2001 , 285, 1039-43	27.4	63
686	Systematic PCR detection in culture-negative osteoarticular infections. <i>American Journal of Medicine</i> , 2013 , 126, 1143.e25-33	2.4	62
685	Non contiguous-finished genome sequence and description of <i>Bacillus timonensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 346-55		62
684	Non-contiguous finished genome sequence and description of <i>Herbaspirillum massiliense</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 200-9		62
683	Quantitative detection of <i>Tropheryma whippelii</i> DNA by real-time PCR. <i>Journal of Clinical Microbiology</i> , 2002 , 40, 1119-20	9.7	62
682	Identification of Novel Zoonotic Activity of <i>Bartonella</i> spp., France. <i>Emerging Infectious Diseases</i> , 2016 , 22, 457-62	10.2	62
681	Impact of early valve surgery on outcome of <i>Staphylococcus aureus</i> prosthetic valve infective endocarditis: analysis in the International Collaboration of Endocarditis-Pro prospective Cohort Study. <i>Clinical Infectious Diseases</i> , 2015 , 60, 741-9	11.6	61
680	<i>Lactobacillus rhamnosus</i> bacteremia: an emerging clinical entity. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2012 , 31, 2469-80	5.3	61
679	Non contiguous-finished genome sequence and description of <i>Enterobacter massiliensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 7, 399-412		59
678	<i>Rickettsia tamurae</i> sp. nov., isolated from <i>Amblyomma testudinarium</i> ticks. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2006 , 56, 1673-1675	2.2	59
677	Non-contiguous finished genome sequence and description of <i>Paenibacillus senegalensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 70-81		58

676	Do we need new antibiotics?. <i>Clinical Microbiology and Infection</i> , 2016 , 22, 408-15	9.5	57
675	Tropheryma whipplei endocarditis. <i>Emerging Infectious Diseases</i> , 2013 , 19, 1721-30	10.2	56
674	Non-contiguous finished genome sequence and description of Brevibacterium senegalense sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 233-45		56
673	Non contiguous-finished genome sequence and description of Senegalemassilia anaerobia gen. nov., sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 7, 343-56		55
672	Non contiguous-finished genome sequence and description of Cellulomonas massiliensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 258-70		55
671	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. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 2221-3	9.7	55
670	A Concise Review of the Epidemiology and Diagnostics of Rickettsioses: Rickettsia and Orientia spp. <i>Journal of Clinical Microbiology</i> , 2018 , 56,	9.7	55
669	Serological and molecular evidence of Rickettsia helvetica in Denmark. <i>Scandinavian Journal of Infectious Diseases</i> , 2004 , 36, 559-63		53
668	"Candidatus Rickettsia tarasevichiae" in Ixodes persulcatus ticks collected in Russia. <i>Annals of the New York Academy of Sciences</i> , 2003 , 990, 162-72	6.5	53
667	MOLECULAR DETECTION OF BARTONELLA QUINTANA, B. ELIZABETHAE, B. KOEHLERAE, B. DOSHIAE, B. TAYLORII, AND RICKETTSIA FELIS IN RODENT FLEAS COLLECTED IN KABUL, AFGHANISTAN. <i>American Journal of Tropical Medicine and Hygiene</i> , 2006 , 74, 436-439	3.2	53
666	The role of Streptococcus intermedius in brain abscess. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2013 , 32, 477-83	5.3	52
665	Molecular analysis of pericardial fluid: a 7-year experience. <i>European Heart Journal</i> , 2006 , 27, 1942-6	9.5	52
664	Rapid identification of Legionella species by mass spectrometry. <i>Journal of Medical Microbiology</i> , 2010 , 59, 273-284	3.2	51
663	Genome sequence and description of Alistipes senegalensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 1-16		50
662	Tick-borne rickettsioses in international travellers. <i>International Journal of Infectious Diseases</i> , 2004 , 8, 139-46	10.5	50
661	Genetic classification of "Rickettsia heilongjiangii" and "Rickettsia hulunii," two Chinese spotted fever group rickettsiae. <i>Journal of Clinical Microbiology</i> , 2000 , 38, 3498-501	9.7	50
660	Detection and culture of Bartonella quintana, Serratia marcescens, and Acinetobacter spp. from decontaminated human body lice. <i>Journal of Clinical Microbiology</i> , 2001 , 39, 1707-9	9.7	49
659	Clinical efficacy of chloroquine derivatives in COVID-19 infection: comparative meta-analysis between the big data and the real world. <i>New Microbes and New Infections</i> , 2020 , 38, 100709	4.1	48

658	Rickettsia africae, Western Africa. <i>Emerging Infectious Diseases</i> , 2010 , 16, 571-3	10.2	48
657	Myocarditis, a rare but severe manifestation of Q fever: report of 8 cases and review of the literature. <i>Clinical Infectious Diseases</i> , 2001 , 32, 1440-7	11.6	48
656	Value of microimmunofluorescence for diagnosis and follow-up of Bartonella endocarditis. <i>Vaccine Journal</i> , 2002 , 9, 795-801		48
655	Non contiguous-finished genome sequence and description of Alistipes obesi sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 7, 427-39		47
654	Rickettsia felis, Bartonella henselae, and B. clarridgeiae, New Zealand. <i>Emerging Infectious Diseases</i> , 2004 , 10, 967-8	10.2	47
653	Mink, SARS-CoV-2, and the Human-Animal Interface. <i>Frontiers in Microbiology</i> , 2021 , 12, 663815	5.7	47
652	Rapid and cost-effective identification of Bartonella species using mass spectrometry. <i>Journal of Medical Microbiology</i> , 2009 , 58, 1154-1159	3.2	46
651	Wohlfahrtiimonas chitiniclastica bacteremia in homeless woman. <i>Emerging Infectious Diseases</i> , 2009 , 15, 985-7	10.2	46
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649	Protein coding palindromes are a unique but recurrent feature in Rickettsia. <i>Genome Research</i> , 2002 , 12, 808-16	9.7	46
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