

Pierre E Fournier

List of Publications by Year in Descending Order

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Version: 2024-04-26

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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	Buttiauxella massiliensis sp. nov., Isolated from a Human Bone Infection.. <i>Current Microbiology</i> , 2022 , 79, 41	2.4	0
800	Arabiibacter massiliensis gen. nov. sp. nov., New Anaerobic Bacterium Isolated from the Human Gut.. <i>Current Microbiology</i> , 2022 , 79, 47	2.4	2
799	Limited spread of a rare spike E484K-harboring SARS-CoV-2 in Marseille, France.. <i>Archives of Virology</i> , 2022 , 167, 583	2.6	1
798	Characteristics of the first 1,119 SARS-CoV-2 Omicron variant cases, in Marseille, France, November-December 2021.. <i>Journal of Medical Virology</i> , 2022 ,	19.7	12
797	Introduction of the SARS-CoV-2 Beta variant from Comoros into the Marseille geographical area.. <i>Travel Medicine and Infectious Disease</i> , 2022 , 102277	8.4	0
796	Q fever endocarditis in India: A report of two cases.. <i>Indian Journal of Medical Microbiology</i> , 2022 ,	1.3	
795	Direct diagnosis of Pasteurella multocida meningitis using next-generation sequencing.. <i>Lancet Microbe</i> , 2022 , 3, e6	22.2	2
794	SARS-CoV-2 reinfection and COVID-19 severity.. <i>Emerging Microbes and Infections</i> , 2022 , 1-41	18.9	5
793	sp. nov., new bacterial species isolated from healthy indigenous congolese volunteer.. <i>Heliyon</i> , 2022 , 8, e09102	3.6	1
792	Genomic evolution and adaptation of arthropod-associated Rickettsia.. <i>Scientific Reports</i> , 2022 , 12, 38074.9	4.9	3
791	First cases of infection with the 21L/BA.2 Omicron variant in Marseille, France.. <i>Journal of Medical Virology</i> , 2022 ,	19.7	6
790	Severity of the first 207 infections with the SARS-CoV-2 Omicron BA.2 variant, in Marseille, France, December, 2021-February, 2022.. <i>Journal of Medical Virology</i> , 2022 ,	19.7	1
789	Culture and identification of a "Deltamicron" SARS-CoV-2 in a three cases cluster in southern France.. <i>Journal of Medical Virology</i> , 2022 ,	19.7	3
788	High rate of reinfection with the SARS-CoV-2 Omicron variant.. <i>Journal of Infection</i> , 2022 ,	18.9	1
787	Draft Genome Sequence of Corynebacterium sanguinis Strain Marseille-P8776.. <i>Microbiology Resource Announcements</i> , 2022 , 11, e0000822	1.3	
786	SARS-CoV-2 Vaccination and Protection Against Clinical Disease: A Retrospective Study, Bouches-du-Rhône District, Southern France, 2021.. <i>Frontiers in Microbiology</i> , 2021 , 12, 796807	5.7	0
785	Analysis of SARS-CoV-2 Variants From 24,181 Patients Exemplifies the Role of Globalization and Zoonosis in Pandemics.. <i>Frontiers in Microbiology</i> , 2021 , 12, 786233	5.7	11

784	Occurrence of a substitution or deletion of SARS-CoV-2 spike amino acid 677 in various lineages in Marseille, France. <i>Virus Genes</i> , 2021 , 1	2.3	2
783	<i>Collinsella ihumii</i> sp. nov., a new anaerobic bacterium isolated from human stool. <i>Archives of Microbiology</i> , 2021 , 203, 6315-6322	3	0
782	Real-time whole genome sequencing direct diagnosis of <i>Streptococcus pneumoniae</i> meningitis: A case report. <i>Journal of Infection</i> , 2021 ,	18.9	1
781	Pneumococcal pneumonia among shipyard workers: Inside the features of disease onset. <i>Travel Medicine and Infectious Disease</i> , 2021 , 44, 102183	8.4	1
780	Using MALDI-TOF spectra in epidemiological surveillance for the detection of bacterial subgroups with a possible epidemic potential. <i>BMC Infectious Diseases</i> , 2021 , 21, 1109	4	3
779	Flying Fox Hemolytic Fever, Description of a New Zoonosis Caused by Candidatus <i>Mycoplasma haemohominis</i> . <i>Clinical Infectious Diseases</i> , 2021 , 73, e1445-e1453	11.6	3
778	Automated Western immunoblotting detection of anti-SARS-CoV-2 serum antibodies. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021 , 40, 1309-1317	5.3	8
777	Introduction into the Marseille geographical area of a mild SARS-CoV-2 variant originating from sub-Saharan Africa: An investigational study. <i>Travel Medicine and Infectious Disease</i> , 2021 , 40, 101980	8.4	21
776	Planarians (Platyhelminthes)-An Emerging Model Organism for Investigating Innate Immune Mechanisms. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021 , 11, 619081	5.9	4
775	Screening of SARS-CoV-2 among homeless people, asylum-seekers and other people living in precarious conditions in Marseille, France, March-April 2020. <i>International Journal of Infectious Diseases</i> , 2021 , 105, 1-6	10.5	7
774	Mink, SARS-CoV-2, and the Human-Animal Interface. <i>Frontiers in Microbiology</i> , 2021 , 12, 663815	5.7	47
773	<i>Pedobacter ghigonii</i> sp. nov., Isolated from the Microbiota of the Planarian <i>Schmidtea mediterranea</i> . <i>Microbiology Research</i> , 2021 , 12, 268-287	1	2
772	<i>Metabacillus schmidteae</i> sp. nov., Cultivated from Planarian <i>Schmidtea mediterranea</i> Microbiota. <i>Microbiology Research</i> , 2021 , 12, 299-316	1	3
771	<i>Bartonella</i> infections diagnosed in the French reference center, 2014-2019, and focus on infections in the immunocompromised. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021 , 40, 2407-2410	5.3	4
770	Genomic description and characterization of <i>Nigeribacterium massiliense</i> gen. nov., sp. nov., isolated from the human gut. <i>Microbes and Infection</i> , 2021 , 23, 104842	9.3	
769	Evaluation of factors influencing tick bites and tick-borne infections: a longitudinal study. <i>Parasites and Vectors</i> , 2021 , 14, 289	4	0
768	<i>Chryseobacterium schmidteae</i> sp. nov. a novel bacterial species isolated from planarian <i>Schmidtea mediterranea</i> . <i>Scientific Reports</i> , 2021 , 11, 11002	4.9	1
767	Is minor surgery safe during the COVID-19 pandemic? A multi-disciplinary study. <i>PLoS ONE</i> , 2021 , 16, e0251122	3.7	2

766	Lagierella massiliensis gen. nov., sp. nov., Isolated from a Stool Sample. <i>Current Microbiology</i> , 2021 , 78, 2481-2487	2.4	0
765	LamPORE SARS-CoV-2 diagnosis and genotyping: A preliminary report. <i>Journal of Clinical Virology</i> , 2021 , 138, 104815	14.5	2
764	Viral RNA in City Wastewater as a Key Indicator of COVID-19 Recrudescence and Containment Measures Effectiveness. <i>Frontiers in Microbiology</i> , 2021 , 12, 664477	5.7	10
763	New human-associated species of the family and proposal to reclassify members of the genus. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021 , 71,	2.2	3
762	Emergence and outcomes of the SARS-CoV-2 'Marseille-4' variant. <i>International Journal of Infectious Diseases</i> , 2021 , 106, 228-236	10.5	23
761	Implementation of an in-house real-time reverse transcription-PCR assay for the rapid detection of the SARS-CoV-2 Marseille-4 variant. <i>Journal of Clinical Virology</i> , 2021 , 139, 104814	14.5	13
760	Contagion Management at the Mditerranè Infection University Hospital Institute. <i>Journal of Clinical Medicine</i> , 2021 , 10,	5.1	1
759	Vitreoscilla massiliensis sp. nov., Isolated From the Stool of an Amazonian Patient. <i>Current Microbiology</i> , 2021 , 78, 3313-3320	2.4	0
758	Pangenome analysis and virulence profiling of Streptococcus intermedius. <i>BMC Genomics</i> , 2021 , 22, 522	4.5	0
757	Implementation of an in-house real-time reverse transcription-PCR assay to detect the emerging SARS-CoV-2 N501Y variants. <i>Journal of Clinical Virology</i> , 2021 , 140, 104868	14.5	10
756	Draft genome of strain 4T19 isolated from cotton-waste composts for mushroom cultivation. <i>New Microbes and New Infections</i> , 2021 , 42, 100894	4.1	
755	Genomic Characterization of the Novel sp. Isolated from the Blood of a (Pallas, 1811). <i>Vector-Borne and Zoonotic Diseases</i> , 2021 , 21, 432-440	2.4	2
754	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
753	Statins potentiate the antibacterial effect of platelets on. <i>Platelets</i> , 2021 , 32, 671-676	3.6	1
752	Human galectin-1 and galectin-3 promote infection. <i>Gut Microbes</i> , 2021 , 13, 1-15	8.8	3
751	Anaerococcus urinimassiliensis sp. nov., a new bacterium isolated from human urine. <i>Scientific Reports</i> , 2021 , 11, 2684	4.9	4
750	Early combination therapy with hydroxychloroquine and azithromycin reduces mortality in 10,429 COVID-19 outpatients. <i>Reviews in Cardiovascular Medicine</i> , 2021 , 22, 1063-1072	3.9	3
749	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. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021 , 40, 1579-1582	5.3	3

748	Draft Genome Sequence of <i>Comamonas aquatilis</i> Strain LK (= CSUR P6418 = CECT 9772), Isolated from the Planarian <i>Schmidtea mediterranea</i> . <i>Microbiology Resource Announcements</i> , 2021 , 10,	1.3	1
747	Draft Genome Sequence of <i>Herminiimonas contaminans</i> Strain CCM 7991, a Biopharmaceutical Contaminant. <i>Microbiology Resource Announcements</i> , 2021 , 10,	1.3	2
746	SARS-CoV-2 variant from India to Marseille: The still active role of ports in the introduction of epidemics. <i>Travel Medicine and Infectious Disease</i> , 2021 , 42, 102085	8.4	8
745	Draft Genome Sequence of <i>Bacillus velezensis</i> Strain Marseille-Q1230, Isolated from a Stool Sample from a Severely Malnourished Child. <i>Microbiology Resource Announcements</i> , 2021 , 10, e0051421	1.3	0
744	<i>Gemella massiliensis</i> sp. nov., a new bacterium isolated from the human sputum. <i>Archives of Microbiology</i> , 2021 , 203, 5817-5823	3	1
743	Whole Genome Sequencing of SARS-CoV-2 Strains in COVID-19 Patients From Djibouti Shows Novel Mutations and Clades Replacing Over Time. <i>Frontiers in Medicine</i> , 2021 , 8, 737602	4.9	1
742	Spreading of a new SARS-CoV-2 N501Y spike variant in a new lineage. <i>Clinical Microbiology and Infection</i> , 2021 , 27, 1352.e1-1352.e5	9.5	20
741	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. <i>Infection, Genetics and Evolution</i> , 2021 , 95, 105092	4.5	7
740	Culturomics revealed the bacterial constituents of the microbiota of a 10-year-old laboratory culture of planarian species <i>S. mediterranea</i> .. <i>Scientific Reports</i> , 2021 , 11, 24311	4.9	
739	Incidence and Outcome of Coinfections with SARS-CoV-2 and Rhinovirus.. <i>Viruses</i> , 2021 , 13,	6.2	4
738	<i>Neglectibacter timonensis</i> gen. nov., sp. nov. and <i>Scatolibacter rhodanostii</i> gen. nov., sp. nov., two anaerobic bacteria isolated from human stool samples.. <i>Archives of Microbiology</i> , 2021 , 204, 45	3	0
737	Description of sp. nov., a new halophilic bacterium isolated from Senegalian human stool. <i>New Microbes and New Infections</i> , 2020 , 38, 100799	4.1	0
736	Investigation of <i>Ctenocephalides felis</i> on domestic dogs and <i>Rickettsia felis</i> infection in the Democratic Republic of Sao Tome and Principe. <i>Zoonoses and Public Health</i> , 2020 , 67, 892-902	2.9	1
735	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
734	Interest of bacterial pangenome analyses in clinical microbiology. <i>Microbial Pathogenesis</i> , 2020 , 149, 104275	3.8	2
733	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
732	sp. nov., a new bacterium isolated from the human gut. <i>New Microbes and New Infections</i> , 2020 , 34, 100657	4.1	2
731	sp. nov., a new bacterium isolated from human gut microbiota. <i>New Microbes and New Infections</i> , 2020 , 35, 100655	4.1	0

730	Systematic Nucleotide Exchange Analysis of ESTs From the Human Cancer Genome Project Report: Origins of 347 Unknown ESTs Indicate Putative Transcription of Non-Coding Genomic Regions. <i>Frontiers in Genetics</i> , 2020 , 11, 42	4.5	2
729	<i>Clostridium massiliamazoniense</i> sp. nov., New Bacterial Species Isolated from Stool Sample of a Volunteer Brazilian. <i>Current Microbiology</i> , 2020 , 77, 2008-2015	2.4	4
728	Detailed description of strain SIT17 a bacterium isolated from the human gut. <i>New Microbes and New Infections</i> , 2020 , 37, 100700	4.1	
727	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
726	Description of sp. nov., a new bacterium isolated from human rectum. <i>New Microbes and New Infections</i> , 2020 , 36, 100703	4.1	0
725	<i>Gorillibacterium timonense</i> sp. nov., isolated from an obese patient. <i>Archives of Microbiology</i> , 2020 , 202, 1223-1229	3	0
724	gen. nov., sp. nov., a new bacterium isolated from the human gut microbiota. <i>New Microbes and New Infections</i> , 2020 , 34, 100645	4.1	1
723	Actinomycetoma Caused by <i>Actinomadura mexicana</i> , A Neglected Entity in the Caribbean. <i>Emerging Infectious Diseases</i> , 2020 , 26, 379-380	10.2	4
722	Epidemiological serosurvey of vector-borne and zoonotic pathogens among homeless people living in shelters in Marseille: cross-sectional one-day surveys (2005-2015). <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020 , 39, 1663-1672	5.3	1
721	gen. nov., sp. nov., a new bacterium isolated from the human microbiota. <i>New Microbes and New Infections</i> , 2020 , 35, 100664	4.1	0
720	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
719	<i>Pedobacter schmidteae</i> sp. nov., a new bacterium isolated from the microbiota of the planarian <i>Schmidtea mediterranea</i> . <i>Scientific Reports</i> , 2020 , 10, 6113	4.9	6
718	High-Content Screening, a Reliable System for <i>Coxiella burnetii</i> Isolation from Clinical Samples. <i>Journal of Clinical Microbiology</i> , 2020 , 58,	9.7	2
717	Survival of <i>Rickettsia conorii</i> in artificially contaminated whole and leukoreduced canine blood units during the storage period. <i>Parasites and Vectors</i> , 2020 , 13, 118	4	3
716	Multispacer sequence typing of <i>Coxiella burnetii</i> from milk and hard tick samples from ruminant farms in Lebanon. <i>Veterinaria Italiana</i> , 2020 , 56, 289-296	1	1
715	Draft Genome Sequence of <i>Salirhabdus euzebyi</i> Strain Q1438. <i>Microbiology Resource Announcements</i> , 2020 , 9,	1.3	5
714	gen. nov., sp. nov., a novel propionic acid bacterium derived from human vaginal discharge. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020 , 70, 4091-4097	2.2	1
713	Case Report: Biliary Tract Infections in Two North Africans in France. <i>American Journal of Tropical Medicine and Hygiene</i> , 2020 , 102, 1306-1308	3.2	0

712	Dynamics and genetic diversity of Haemophilus influenzae carriage among French pilgrims during the 2018 Hajj: A prospective cohort survey. <i>Travel Medicine and Infectious Disease</i> , 2020 , 38, 101883	8.4	4
711	Genome sequence-based criteria for demarcation and definition of species in the genus. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020 , 70, 1738-1750	2.2	12
710	sp. nov. a multi-drug resistant bacterium infecting humans. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020 , 70, 1758-1768	2.2	2
709	Phenotypic and molecular characterization of macrolide resistance mechanisms among isolated in Tunisia. <i>Journal of Medical Microbiology</i> , 2020 , 69, 505-520	3.2	2
708	Early mortality attributable to PICC-lines in 4 public hospitals of Marseille from 2010 to 2016 (Revised V3). <i>Medicine (United States)</i> , 2020 , 99, e18494	1.8	3
707	Effect of antiplatelet agents on platelet antistaphylococcal capacity: An in vitro study. <i>International Journal of Antimicrobial Agents</i> , 2020 , 55, 105890	14.3	4
706	Taxono-genomics and description of sp. nov., a new bacterium isolated from the gut of a healthy infant. <i>New Microbes and New Infections</i> , 2020 , 33, 100631	4.1	1
705	Identification of Noncanonical Transcripts Produced by Systematic Nucleotide Exchanges in HIV-Associated Centroblastic Lymphoma. <i>DNA and Cell Biology</i> , 2020 , 39, 1444-1448	3.6	2
704	Culturing Ancient Bacteria Carrying Resistance Genes from Permafrost and Comparative Genomics with Modern Isolates. <i>Microorganisms</i> , 2020 , 8,	4.9	1
703	Description of sp. nov., sp. nov. and sp. nov., new anaerobic bacteria isolated from human stool samples. <i>New Microbes and New Infections</i> , 2020 , 37, 100719	4.1	0
702	Genome sequence and description of gen. nov., sp. nov., a new bacterium isolated from a human stool sample. <i>New Microbes and New Infections</i> , 2020 , 37, 100720	4.1	1
701	Tularemia: A Case Series of Patients Diagnosed at the National Reference Center for Rickettsioses From 2008 to 2017. <i>Open Forum Infectious Diseases</i> , 2020 , 7, ofaa440	1	5
700	Contribution of VitaPCR SARS-CoV-2 to the emergency diagnosis of COVID-19. <i>Journal of Clinical Virology</i> , 2020 , 133, 104682	14.5	16
699	Utilization of Galectins by Pathogens for Infection. <i>Frontiers in Immunology</i> , 2020 , 11, 1877	8.4	7
698	Type 1 Diabetes in People Hospitalized for COVID-19: New Insights From the CORONADO Study. <i>Diabetes Care</i> , 2020 , 43, e174-e177	14.6	26
697	Fenollaria timonensis sp. nov., A New Bacterium Isolated from Healthy Human Fresh Stool. <i>Current Microbiology</i> , 2020 , 77, 3780-3786	2.4	1
696	Seroepidemiological and molecular investigation of spotted fever group rickettsiae and Coxiella burnetii in Sao Tome Island: A One Health approach. <i>Transboundary and Emerging Diseases</i> , 2020 , 67 Suppl 2, 36-43	4.2	6
695	Q Fever 2020 , 599-601		0

694	Mediastinitis in the intensive care unit patient: a narrative review. <i>Clinical Microbiology and Infection</i> , 2020 , 26, 26-34	9.5	15
693	Tick-Borne Spotted Fever Rickettsioses 2020 , 587-593		1
692	Draft genome and description of <i>Negativicoccus massiliensis</i> strain Marseille-P2082, a new species isolated from the gut microbiota of an obese patient. <i>Antonie Van Leeuwenhoek</i> , 2020 , 113, 997-1008	2.1	
691	sp. nov., and sp. nov., two new species isolated from obese patients. <i>New Microbes and New Infections</i> , 2019 , 31, 100581	4.1	0
690	Rickettsial infections of the central nervous system. <i>PLoS Neglected Tropical Diseases</i> , 2019 , 13, e0007462	6.8	22
689	Western Immunoblotting for the Diagnosis of and Infective Endocarditis. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019 , 9, 314	5.9	4
688	gen. nov., sp. nov., a new bacterium isolated from the vagina flora and its taxono-genomic description. <i>New Microbes and New Infections</i> , 2019 , 32, 100601	4.1	1
687	sp. nov., a new bacterium isolated from stool from a healthy Senegalese child. <i>New Microbes and New Infections</i> , 2019 , 31, 100588	4.1	2
686	sp. nov., a new bacterium isolated from the human microbiota. <i>New Microbes and New Infections</i> , 2019 , 32, 100592	4.1	
685	sp. nov., a new bacterium isolated from a fresh human stool specimen. <i>New Microbes and New Infections</i> , 2019 , 32, 100602	4.1	1
684	<i>Phoenicibacter congonensis</i> gen. nov., sp. nov., a new genus isolated from the human gut and its description using a taxonogenomic approach. <i>Antonie Van Leeuwenhoek</i> , 2019 , 112, 775-784	2.1	2
683	Description and genomic characterization of <i>Massiliimalia massiliensis</i> gen. nov., sp. nov., and <i>Massiliimalia timonensis</i> gen. nov., sp. nov., two new members of the family Ruminococcaceae isolated from the human gut. <i>Antonie Van Leeuwenhoek</i> , 2019 , 112, 905-918	2.1	3
682	Noncontiguous finished genome sequence and description of <i>Raoultibacter massiliensis</i> gen. nov., sp. nov. and <i>Raoultibacter timonensis</i> sp. nov, two new bacterial species isolated from the human gut. <i>MicrobiologyOpen</i> , 2019 , 8, e00758	3.4	5
681	Draft genome and description of <i>Merdibacter massiliensis</i> gen.nov., sp. nov., a new bacterium genus isolated from the human ileum. <i>Scientific Reports</i> , 2019 , 9, 7931	4.9	13
680	Genome sequence and description of sp. nov., a new anaerobic bacterium isolated from the human gut. <i>New Microbes and New Infections</i> , 2019 , 31, 100571	4.1	1
679	sp. nov., a new halophilic bacterium isolated from the stool of a healthy 11-year-old boy. <i>New Microbes and New Infections</i> , 2019 , 30, 100541	4.1	
678	Taxono-genomics description of sp. nov., strain Marseille-P4006, a new species isolated from the female genital tract of a patient with bacterial vaginosis. <i>New Microbes and New Infections</i> , 2019 , 29, 100534	4.1	2
677	gen. nov., sp. nov.; gen. nov., sp. nov.; gen. nov., sp. nov. and sp. nov. isolated from breast milk from healthy breastfeeding African women. <i>New Microbes and New Infections</i> , 2019 , 29, 100537	4.1	2

676	Non-contiguous finished genome sequencing and description of sp. nov. isolated from human sputum. <i>New Microbes and New Infections</i> , 2019 , 29, 100532	4.1	1
675	Genome sequence and description of gen. nov., sp. nov., a new bacterial genus isolated from human left colon. <i>New Microbes and New Infections</i> , 2019 , 29, 100520	4.1	2
674	sp. nov., a succinate-producing bacterium isolated from human stool from an obese patient after bariatric surgery. <i>New Microbes and New Infections</i> , 2019 , 29, 100508	4.1	10
673	The association between vegetation size and surgical treatment on 6-month mortality in left-sided infective endocarditis. <i>European Heart Journal</i> , 2019 , 40, 2243-2251	9.5	13
672	Description of Janibacter massiliensis sp. nov., cultured from the vaginal discharge of a patient with bacterial vaginosis. <i>Antonie Van Leeuwenhoek</i> , 2019 , 112, 1147-1159	2.1	3
671	Genome sequence and description of Gracilibacillus timonensis sp. nov. strain Marseille-P2481, a moderate halophilic bacterium isolated from the human gut microflora. <i>MicrobiologyOpen</i> , 2019 , 8, e00638	3.4	5
670	Description of three new Peptoniphilus species cultured in the vaginal fluid of a woman diagnosed with bacterial vaginosis: Peptoniphilus pacaensis sp. nov., Peptoniphilus raoultii sp. nov., and Peptoniphilus vaginalis sp. nov. <i>MicrobiologyOpen</i> , 2019 , 8, e00661	3.4	12
669	Human pegivirus isolates characterized by deep sequencing from hepatitis C virus-RNA and human immunodeficiency virus-RNA-positive blood donations, France. <i>Journal of Medical Virology</i> , 2019 , 91, 38-44	19.7	5
668	sp. nov., sp. nov. and sp. nov., isolated from vaginal swabs from healthy Senegalese women. <i>New Microbes and New Infections</i> , 2019 , 31, 100585	4.1	4
667	Draft genome and description of Chryseobacterium phocaeense sp. nov.: a new bacterial species isolated from the sputum of a cystic fibrosis patient. <i>Archives of Microbiology</i> , 2019 , 201, 1361-1368	3	
666	sp. nov., a new bacterial species isolated from human stool sample. <i>New Microbes and New Infections</i> , 2019 , 31, 100575	4.1	3
665	sp. nov., a new bacterial species isolated from human urine sample. <i>New Microbes and New Infections</i> , 2019 , 31, 100576	4.1	4
664	gen. nov., sp. nov., a new bacterium isolated from the human gut microbiota. <i>New Microbes and New Infections</i> , 2019 , 31, 100574	4.1	
663	Optimized fixation of actin filaments for improved indirect immunofluorescence staining of rickettsiae. <i>BMC Research Notes</i> , 2019 , 12, 657	2.3	2
662	sp. nov., a new bacterium isolated from human urine with culturomics. <i>New Microbes and New Infections</i> , 2019 , 32, 100591	4.1	2
661	sp. nov., a new bacterium isolated from a stool specimen of a healthy woman. <i>New Microbes and New Infections</i> , 2019 , 32, 100607	4.1	3
660	sp. nov., a new bacteria species isolated from the human gut microbiota. <i>New Microbes and New Infections</i> , 2019 , 32, 100610	4.1	1
659	gen. nov., a new genesis and sp. nov., a new bacterium isolated from the human oral microbiota. <i>New Microbes and New Infections</i> , 2019 , 32, 100611	4.1	

658	Rickettsia 2019 , 1-43		0
657	Whole-Genome Sequence of Haloimpatiens lingqiaonensis Strain P8956. <i>Microbiology Resource Announcements</i> , 2019 , 8,	1.3	7
656	Whole-Genome Sequence of French Clinical Peptoniphilus catoniae Strain P8546. <i>Microbiology Resource Announcements</i> , 2019 , 8,	1.3	4
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450	Patterns of <i>Kingella kingae</i> Disease Outbreaks. <i>Pediatric Infectious Disease Journal</i> , 2016 , 35, 340-6	3.4	31
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346	Non contiguous-finished genome sequence and description of <i>Enorma timonensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2014 , 9, 970-86		4
345	Genome sequence and description of <i>Corynebacterium ihumii</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1128-43		6
344	Genome sequence and description of <i>Bacteroides timonensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1181-97		6
343	Non contiguous-finished genome sequence and description of <i>Clostridium jeddahense</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1003-19		19
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300	Non-contiguous finished genome sequence and description of <i>Holdemania massiliensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 9, 395-409		12
299	Non contiguous-finished genome sequence and description of <i>Peptoniphilus obesi</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2013 , 7, 357-369		29

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255	Non-contiguous finished genome sequence and description of Alistipes timonensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 315-24		84
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253	Non-contiguous finished genome sequence and description of Clostridium senegalense sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 386-95		65
252	Non contiguous-finished genome sequence and description of Bacillus timonensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 346-55		62
251	Non contiguous-finished genome sequence and description of Peptoniphilus timonensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 1-11		74
250	Non-contiguous finished genome sequence and description of Herbaspirillum massiliense sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 200-9		62
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