

Jean Christophe Lagier

List of Publications by Citations

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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.

383 papers	16,413 citations	55 h-index	122 g-index
420 ext. papers	20,909 ext. citations	6.7 avg, IF	7.03 L-index

#	Paper	IF	Citations
383	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial. <i>International Journal of Antimicrobial Agents</i> , 2020 , 56, 105949	14.3	3043
382	Autoantibodies against type I IFNs in patients with life-threatening COVID-19. <i>Science</i> , 2020 , 370,	33.3	1090
381	Inborn errors of type I IFN immunity in patients with life-threatening COVID-19. <i>Science</i> , 2020 , 370,	33.3	994
380	Microbial culturomics: paradigm shift in the human gut microbiome study. <i>Clinical Microbiology and Infection</i> , 2012 , 18, 1185-93	9.5	722
379	Culture of previously uncultured members of the human gut microbiota by culturomics. <i>Nature Microbiology</i> , 2016 , 1, 16203	26.6	555
378	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
377	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
376	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
375	Culturing the human microbiota and culturomics. <i>Nature Reviews Microbiology</i> , 2018 , 16, 540-550	22.2	303
374	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
373	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
372	Current and past strategies for bacterial culture in clinical microbiology. <i>Clinical Microbiology Reviews</i> , 2015 , 28, 208-36	34	239
371	Human gut microbiota: repertoire and variations. <i>Frontiers in Cellular and Infection Microbiology</i> , 2012 , 2, 136	5.9	207
370	Gut bacterial microbiota and obesity. <i>Clinical Microbiology and Infection</i> , 2013 , 19, 305-13	9.5	191
369	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
368	Systemic <i>Tropheryma whipplei</i> : clinical presentation of 142 patients with infections diagnosed or confirmed in a reference center. <i>Medicine (United States)</i> , 2010 , 89, 337-345	1.8	144
367	High-level colonisation of the human gut by Verrucomicrobia following broad-spectrum antibiotic treatment. <i>International Journal of Antimicrobial Agents</i> , 2013 , 41, 149-55	14.3	131

366	Increased Gut Redox and Depletion of Anaerobic and Methanogenic Prokaryotes in Severe Acute Malnutrition. <i>Scientific Reports</i> , 2016 , 6, 26051	4.9	108
365	Gut microbiota diversity according to dietary habits and geographical provenance. <i>Human Microbiome Journal</i> , 2018 , 7-8, 1-9	5.6	107
364	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
363	The gut microbiota of a patient with resistant tuberculosis is more comprehensively studied by culturomics than by metagenomics. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2013 , 32, 637-45	5.3	104
362	Genome sequence and description of Anaerosalibacter massiliensis sp. nov. <i>New Microbes and New Infections</i> , 2016 , 10, 66-76	4.1	98
361	Non-contiguous finished genome sequence and description of Anaerococcus senegalensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 116-25		93
360	Autoantibodies neutralizing type I IFNs are present in 4% of uninfected individuals over 70 years old and account for 20% of COVID-19 deaths. <i>Science Immunology</i> , 2021 , 6,	2.8	91
359	The contribution of culturomics to the repertoire of isolated human bacterial and archaeal species. <i>Microbiome</i> , 2018 , 6, 94	16.6	88
358	Treatment of classic Whipple's disease: from in vitro results to clinical outcome. <i>Journal of Antimicrobial Chemotherapy</i> , 2014 , 69, 219-27	5.1	86
357	Culturomics identified 11 new bacterial species from a single anorexia nervosa stool sample. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2013 , 32, 1471-81	5.3	85
356	Non-contiguous finished genome sequence and description of Alistipes timonensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 315-24		84
355	Tropheryma whipplei and Whipple's disease. <i>Journal of Infection</i> , 2014 , 69, 103-12	18.9	83
354	The human gut microbiome, a taxonomic conundrum. <i>Systematic and Applied Microbiology</i> , 2015 , 38, 276-86	4.2	82
353	Rapid viral diagnosis and ambulatory management of suspected COVID-19 cases presenting at the infectious diseases referral hospital in Marseille, France, - January 31st to March 1st, 2020: A respiratory virus snapshot. <i>Travel Medicine and Infectious Disease</i> , 2020 , 36, 101632	8.4	82
352	Diet influence on the gut microbiota and dysbiosis related to nutritional disorders. <i>Human Microbiome Journal</i> , 2016 , 1, 3-11	5.6	81
351	Clostridium butyricum Strains and Dysbiosis Linked to Necrotizing Enterocolitis in Preterm Neonates. <i>Clinical Infectious Diseases</i> , 2015 , 61, 1107-15	11.6	76
350	Early Fecal Microbiota Transplantation Improves Survival in Severe Clostridium difficile Infections. <i>Clinical Infectious Diseases</i> , 2018 , 66, 645-650	11.6	74
349	Non contiguous-finished genome sequence and description of Peptoniphilus timonensis sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 1-11		74

348	A Metagenomic Investigation of the Duodenal Microbiota Reveals Links with Obesity. <i>PLoS ONE</i> , 2015 , 10, e0137784	3.7	73
347	Gut Bacteria Missing in Severe Acute Malnutrition, Can We Identify Potential Probiotics by Culturomics?. <i>Frontiers in Microbiology</i> , 2017 , 8, 899	5.7	69
346	Lack of MERS coronavirus but prevalence of influenza virus in French pilgrims after 2013 Hajj. <i>Emerging Infectious Diseases</i> , 2014 , 20, 728-30	10.2	67
345	Evidence of lifetime susceptibility to <i>Tropheryma whipplei</i> in patients with Whipple's disease. <i>Journal of Antimicrobial Chemotherapy</i> , 2011 , 66, 1188-9	5.1	66
344	Abnormal weight gain and gut microbiota modifications are side effects of long-term doxycycline and hydroxychloroquine treatment. <i>Antimicrobial Agents and Chemotherapy</i> , 2014 , 58, 3342-7	5.9	65
343	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
342	Genome sequence and description of <i>Aeromicrobium massiliense</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 246-57		65
341	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
340	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
339	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
338	Failure and relapse after treatment with trimethoprim/sulfamethoxazole in classic Whipple's disease. <i>Journal of Antimicrobial Chemotherapy</i> , 2010 , 65, 2005-12	5.1	59
337	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial		59
336	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
335	Comparison of the gut microbiota of people in France and Saudi Arabia. <i>Nutrition and Diabetes</i> , 2015 , 5, e153	4.7	57
334	Dramatic reduction in <i>Clostridium difficile</i> ribotype 027-associated mortality with early fecal transplantation by the nasogastric route: a preliminary report. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2015 , 34, 1597-601	5.3	57
333	<i>Tropheryma whipplei</i> endocarditis. <i>Emerging Infectious Diseases</i> , 2013 , 19, 1721-30	10.2	56
332	Non-contiguous finished genome sequence and description of <i>Brevibacterium senegalense</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 233-45		56
331	Non contiguous-finished genome sequence and description of <i>Senegalemassilia anaerobia</i> gen. nov., sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 7, 343-56		55

330	Non contiguous-finished genome sequence and description of <i>Cellulomonas massiliensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 258-70		55
329	Culturomics and pyrosequencing evidence of the reduction in gut microbiota diversity in patients with broad-spectrum antibiotics. <i>International Journal of Antimicrobial Agents</i> , 2014 , 44, 117-24	14.3	53
328	Bacterial culture through selective and non-selective conditions: the evolution of culture media in clinical microbiology. <i>New Microbes and New Infections</i> , 2020 , 34, 100622	4.1	52
327	Genome sequence and description of <i>Alistipes senegalensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 6, 1-16		50
326	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
325	Non contiguous-finished genome sequence and description of <i>Alistipes obesi</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 7, 427-39		47
324	Repertoire of human gut microbes. <i>Microbial Pathogenesis</i> , 2017 , 106, 103-112	3.8	46
323	Repertoire of human breast and milk microbiota: a systematic review. <i>Future Microbiology</i> , 2019 , 14, 623-641	2.9	46
322	Non-contiguous finished genome sequence and description of <i>Kurthia massiliensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2012 , 7, 221-32		46
321	Non-contiguous finished genome sequence and description of <i>Paenibacillus senegalensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2012 , 7, 70-81		46
320	Aerobic culture of anaerobic bacteria using antioxidants: a preliminary report. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2014 , 33, 1781-3	5.3	45
319	Faecal microbiota transplantation for stool decolonization of OXA-48 carbapenemase-producing <i>Klebsiella pneumoniae</i> . <i>Journal of Hospital Infection</i> , 2015 , 90, 173-4	6.9	45
318	Molecular studies neglect apparently gram-negative populations in the human gut microbiota. <i>Journal of Clinical Microbiology</i> , 2013 , 51, 3286-93	9.7	45
317	Likely correlation between sources of information and acceptability of A/H1N1 swine-origin influenza virus vaccine in Marseille, France. <i>PLoS ONE</i> , 2010 , 5, e11292	3.7	44
316	Gut microbiota associated with HIV infection is significantly enriched in bacteria tolerant to oxygen. <i>BMJ Open Gastroenterology</i> , 2016 , 3, e000080	3.9	42
315	Gorilla gorilla gorilla gut: a potential reservoir of pathogenic bacteria as revealed using culturomics and molecular tools. <i>Scientific Reports</i> , 2014 , 4, 7174	4.9	42
314	Non-contiguous finished genome sequence and description of <i>Brevibacillus massiliensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 8, 1-14		42
313	Non contiguous-finished genome sequence and description of <i>Dielma fastidiosa</i> gen. nov., sp. nov., a new member of the Family Erysipelotrichaceae. <i>Standards in Genomic Sciences</i> , 2013 , 8, 336-51		41

312	Natural history of COVID-19 and therapeutic options. <i>Expert Review of Clinical Immunology</i> , 2020 , 16, 1159-1184	5.1	41
311	Asymptomatic hypoxia in COVID-19 is associated with poor outcome. <i>International Journal of Infectious Diseases</i> , 2021 , 102, 233-238	10.5	41
310	Successful treatment of immune reconstitution inflammatory syndrome in Whipple's disease using thalidomide. <i>Journal of Infection</i> , 2010 , 60, 79-82	18.9	40
309	Gut microbiome and dietary patterns in different Saudi populations and monkeys. <i>Scientific Reports</i> , 2016 , 6, 32191	4.9	39
308	A quasi-universal medium to break the aerobic/anaerobic bacterial culture dichotomy in clinical microbiology. <i>Clinical Microbiology and Infection</i> , 2016 , 22, 53-58	9.5	39
307	MALDI-TOF identification of the human Gut microbiome in people with and without diarrhea in Senegal. <i>PLoS ONE</i> , 2014 , 9, e87419	3.7	39
306	Probable aircraft transmission of Covid-19 in-flight from the Central African Republic to France. <i>Travel Medicine and Infectious Disease</i> , 2020 , 35, 101643	8.4	38
305	Aerobic culture of methanogenic archaea without an external source of hydrogen. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2016 , 35, 985-91	5.3	37
304	Repertoire of the gut microbiota from stomach to colon using culturomics and next-generation sequencing. <i>BMC Microbiology</i> , 2018 , 18, 157	4.5	37
303	Non contiguous-finished genome sequence and description of <i>Peptoniphilus senegalensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 7, 370-81		34
302	Glycans affect DNA extraction and induce substantial differences in gut metagenomic studies. <i>Scientific Reports</i> , 2016 , 6, 26276	4.9	34
301	Salt in stools is associated with obesity, gut halophilic microbiota and <i>Akkermansia muciniphila</i> depletion in humans. <i>International Journal of Obesity</i> , 2019 , 43, 862-871	5.5	32
300	Testing the repatriated for SARS-Cov2: Should laboratory-based quarantine replace traditional quarantine?. <i>Travel Medicine and Infectious Disease</i> , 2020 , 34, 101624	8.4	31
299	The aerobic activity of metronidazole against anaerobic bacteria. <i>International Journal of Antimicrobial Agents</i> , 2015 , 45, 537-40	14.3	29
298	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
297	Long-term persistence of olfactory and gustatory disorders in COVID-19 patients. <i>Clinical Microbiology and Infection</i> , 2021 , 27, 931-932	9.5	29
296	Non contiguous-finished genome sequence and description of <i>Enorma massiliensis</i> gen. nov., sp. nov., a new member of the Family Coriobacteriaceae. <i>Standards in Genomic Sciences</i> , 2013 , 8, 290-305		28
295	Non contiguous-finished genome sequence and description of <i>Bacillus massilosenegalensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 8, 264-78		28

294	Ultrarapid diagnosis, microscope imaging, genome sequencing, and culture isolation of SARS-CoV-2. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020 , 39, 1601-1603	5.3	27
293	Evidence of SARS-CoV-2 re-infection with a different genotype. <i>Journal of Infection</i> , 2021 , 82, 84-123	18.9	27
292	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
291	Child with liver transplant recovers from COVID-19 infection. A case report. <i>Archives De Pediatrie</i> , 2020 , 27, 275-276	1.8	25
290	First isolation of Tropheryma whipplei from bronchoalveolar fluid and clinical implications. <i>Journal of Infection</i> , 2012 , 65, 275-8	18.9	25
289	Metagenomic and culturomic analysis of gut microbiota dysbiosis during Clostridium difficile infection. <i>Scientific Reports</i> , 2019 , 9, 12807	4.9	24
288	Genome sequence and description of Timonella senegalensis gen. nov., sp. nov., a new member of the suborder Micrococccinae. <i>Standards in Genomic Sciences</i> , 2013 , 8, 318-35		24
287	Human Bacterial Repertoire of the Urinary Tract: a Potential Paradigm Shift. <i>Journal of Clinical Microbiology</i> , 2019 , 57,	9.7	24
286	Non contiguous-finished genome sequence and description of Peptoniphilus obesi sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 7, 357-69		23
285	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
284	New Insights Into the Physiopathology of COVID-19: SARS-CoV-2-Associated Gastrointestinal Illness. <i>Frontiers in Medicine</i> , 2021 , 8, 640073	4.9	23
283	COVID-19 re-infection. <i>European Journal of Clinical Investigation</i> , 2021 , 51, e13537	4.6	22
282	Expression of ACE2, Soluble ACE2, Angiotensin I, Angiotensin II and Angiotensin-(1-7) Is Modulated in COVID-19 Patients. <i>Frontiers in Immunology</i> , 2021 , 12, 625732	8.4	22
281	Non-contiguous finished genome sequence and description of Megasphaera massiliensis sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 8, 525-38		21
280	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
279	Faecal microbiota transplantation shortens the colonisation period and allows re-entry of patients carrying carbapenemase-producing bacteria into medical care facilities. <i>International Journal of Antimicrobial Agents</i> , 2019 , 53, 355-361	14.3	21
278	Whipple's disease and Tropheryma whipplei infections: when to suspect them and how to diagnose and treat them. <i>Current Opinion in Infectious Diseases</i> , 2018 , 31, 463-470	5.4	21
277	Gut microbiota and Clostridium difficile infections. <i>Human Microbiome Journal</i> , 2016 , 2, 10-14	5.6	20

276	Microvirga massiliensis sp. nov., the human commensal with the largest genome. <i>MicrobiologyOpen</i> , 2016 , 5, 307-22	3.4	20
275	Faecal microbiota transplantation: from practice to legislation before considering industrialization. <i>Clinical Microbiology and Infection</i> , 2014 , 20, 1112-8	9.5	20
274	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
273	Microbiota-Orientated Treatments for Major Depression and Schizophrenia. <i>Nutrients</i> , 2020 , 12,	6.7	19
272	Among Live and Dead Bacteria, the Optimization of Sample Collection and Processing Remains Essential in Recovering Gut Microbiota Components. <i>Frontiers in Microbiology</i> , 2019 , 10, 1606	5.7	19
271	Extensive culturomics of 8 healthy samples enhances metagenomics efficiency. <i>PLoS ONE</i> , 2019 , 14, e0233543	3.543	19
270	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
269	Evaluating the serological status of COVID-19 patients using an indirect immunofluorescent assay, France. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021 , 40, 361-371	5.3	18
268	Many More Microbes in Humans: Enlarging the Microbiome Repertoire. <i>Clinical Infectious Diseases</i> , 2017 , 65, S20-S29	11.6	17
267	Melioidosis as a travel-associated infection: Case report and review of the literature. <i>Travel Medicine and Infectious Disease</i> , 2015 , 13, 367-81	8.4	17
266	A severe Whipple disease with an immune reconstitution inflammatory syndrome: an additional case of thalidomide efficiency. <i>Joint Bone Spine</i> , 2014 , 81, 260-2	2.9	17
265	006 RISK FACTORS FOR INFECTIVE ENDOCARDITIS IN PATIENTS WITH STAPHYLOCOCCUS AUREUS BACTEREMIA: A TEACHING HOSPITAL EXPERIENCE. <i>International Journal of Antimicrobial Agents</i> , 2009 , 33, S2-S3	14.3	17
264	Children account for a small proportion of diagnoses of SARS-CoV-2 infection and do not exhibit greater viral loads than adults. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020 , 39, 1983-1987	5.3	17
263	<i>Blautia massiliensis</i> sp. nov., isolated from a fresh human fecal sample and emended description of the genus <i>Blautia</i> . <i>Anaerobe</i> , 2017 , 43, 47-55	2.8	16
262	The Antioxidants Glutathione, Ascorbic Acid and Uric Acid Maintain Butyrate Production by Human Gut Clostridia in The Presence of Oxygen In Vitro. <i>Scientific Reports</i> , 2020 , 10, 7705	4.9	16
261	The proof of concept that culturomics can be superior to metagenomics to study atypical stool samples. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2013 , 32, 1099	5.3	16
260	Non-contiguous finished genome sequence and description of <i>Bacillus massilioanorexius</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 8, 465-79		16
259	Acute infections caused by <i>Tropheryma whipplei</i> . <i>Future Microbiology</i> , 2017 , 12, 247-254	2.9	15

258	From Culturomics to Clinical Microbiology and Forward. <i>Emerging Infectious Diseases</i> , 2018 , 24, 1683-1690.2	10.2	15
257	Case report: Ribavirin and vitamin A in a severe case of measles. <i>Medicine (United States)</i> , 2017 , 96, e9154.8	15.8	15
256	Rise of Microbial Culturomics: Noncontiguous Finished Genome Sequence and Description of Beduini massiliensis gen. nov., sp. nov. <i>OMICS A Journal of Integrative Biology</i> , 2015 , 19, 766-76	3.8	15
255	Optimization and standardization of the culturomics technique for human microbiome exploration. <i>Scientific Reports</i> , 2020 , 10, 9674	4.9	14
254	Isolation and culture of Methanobrevibacter smithii by co-culture with hydrogen-producing bacteria on agar plates. <i>Clinical Microbiology and Infection</i> , 2019 , 25, 1561.e1-1561.e5	9.5	14
253	Non-contiguous finished genome sequence and description of Nosocomiicoccus massiliensis sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 9, 205-19		14
252	Metabarcoding analysis of eukaryotic microbiota in the gut of HIV-infected patients. <i>PLoS ONE</i> , 2018 , 13, e0191913	3.7	14
251	Infections in symptomatic travelers returning from the Arabian peninsula to France: A retrospective cross-sectional study. <i>Travel Medicine and Infectious Disease</i> , 2016 , 14, 414-6	8.4	14
250	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open label non-randomized clinical trial revisited. <i>International Journal of Antimicrobial Agents</i> , 2021 , 57, 106243	14.3	14
249	Escherichia coli spontaneous community-acquired meningitis in adults: A case report and literature review. <i>International Journal of Infectious Diseases</i> , 2018 , 67, 70-74	10.5	14
248	Samples and techniques highlighting the links between obesity and microbiota. <i>Microbial Pathogenesis</i> , 2017 , 106, 119-126	3.8	13
247	Successful Fecal Microbiota Transplantation in a Patient Suffering From Irritable Bowel Syndrome and Recurrent Urinary Tract Infections. <i>Open Forum Infectious Diseases</i> , 2019 , 6, ofz398	1	13
246	Tropheryma whippelii endocarditis relapses after treatment with trimethoprim/sulfamethoxazole. <i>International Journal of Antimicrobial Agents</i> , 2013 , 41, 592-4	14.3	13
245	Clostridium difficile 027 emerging outbreak in Marseille, France. <i>Infection Control and Hospital Epidemiology</i> , 2013 , 34, 1339-41	2	13
244	Clinical outcomes in COVID-19 patients infected with different SARS-CoV-2 variants in Marseille, France. <i>Clinical Microbiology and Infection</i> , 2021 , 27, 1516.e1-1516.e6	9.5	13
243	SARS-CoV-2 Infectivity and Severity of COVID-19 According to SARS-CoV-2 Variants: Current Evidence. <i>Journal of Clinical Medicine</i> , 2021 , 10,	5.1	13
242	Tropheryma whippelii DNA in bronchoalveolar lavage samples: a case control study. <i>Clinical Microbiology and Infection</i> , 2016 , 22, 875-879	9.5	13
241	Tropheryma whippelii associated with diarrhoea in young children. <i>Clinical Microbiology and Infection</i> , 2016 , 22, 869-874	9.5	12

240	Japanese encephalitis in a French traveler to Nepal. <i>Journal of NeuroVirology</i> , 2014 , 20, 99-102	3.9	12
239	Prosthetic joint infection caused by <i>Granulicatella adiacens</i> : a case series and review of literature. <i>BMC Musculoskeletal Disorders</i> , 2017 , 18, 276	2.8	12
238	Genome sequence of <i>Oceanobacillus picturae</i> strain S1, an halophilic bacterium first isolated in human gut. <i>Standards in Genomic Sciences</i> , 2015 , 10, 91		12
237	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
236	Identification of <i>Legionella feeleei</i> cellulitis. <i>Emerging Infectious Diseases</i> , 2011 , 17, 145-6	10.2	12
235	State of the Art in the Culture of the Human Microbiota: New Interests and Strategies. <i>Clinical Microbiology Reviews</i> , 2020 , 34,	34	12
234	<i>Staphylococcus lugdunensis</i> : a neglected pathogen of infections involving fracture-fixation devices. <i>International Orthopaedics</i> , 2017 , 41, 1085-1091	3.8	11
233	A new protectant medium preserving bacterial viability after freeze drying. <i>Microbiological Research</i> , 2020 , 236, 126454	5.3	11
232	<i>Clostridium scindens</i> Is Present in the Gut Microbiota during <i>Clostridium difficile</i> Infection: a Metagenomic and Culturomic Analysis. <i>Journal of Clinical Microbiology</i> , 2018 , 56,	9.7	11
231	Noncontiguous finished genome sequence and description of <i>Paenibacillus ihumii</i> sp. nov. strain AT5. <i>New Microbes and New Infections</i> , 2016 , 10, 142-50	4.1	11
230	Classic Whipple's disease diagnosed by (18)F-fluorodeoxyglucose PET. <i>Lancet Infectious Diseases</i> , 2016 , 16, 130	25.5	11
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227	Meta-analysis on efficacy of amoxicillin in uncomplicated severe acute malnutrition. <i>Microbial Pathogenesis</i> , 2017 , 106, 76-77	3.8	11
226	Fluorescence Hybridization for Diagnosis of Whipple's Disease in Formalin-Fixed Paraffin-Embedded Tissue. <i>Frontiers in Medicine</i> , 2017 , 4, 87	4.9	11
225	Does <i>Tropheryma whipplei</i> contribute to travelers' diarrhea?: A PCR analysis of paired stool samples in French travelers to Senegal. <i>Travel Medicine and Infectious Disease</i> , 2014 , 12, 264-7	8.4	11
224	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
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222	Deciphering the Urinary Microbiota Repertoire by Culturomics Reveals Mostly Anaerobic Bacteria From the Gut. <i>Frontiers in Microbiology</i> , 2020 , 11, 513305	5.7	11
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219	Sewage workers with low antibody responses may be colonized successively by several Tropheryma whipplei strains. <i>International Journal of Infectious Diseases</i> , 2015 , 35, 51-5	10.5	10
218	Noncontiguous finished genome sequence and description of Gabonia massiliensis gen. nov., sp. nov. <i>New Microbes and New Infections</i> , 2016 , 9, 35-44	4.1	10
217	Non-contiguous finished genome sequence and description of Alistipes ihumii sp. nov. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1221-35		10
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195	Non-contiguous finished genome sequence and description of Kurthia senegalensis sp. nov. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1319-30		8
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