

Fadi Bittar

List of Publications by Year in Descending Order

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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.

89
papers

2,661
citations

20
h-index

50
g-index

98
ext. papers

3,347
ext. citations

5.1
avg. IF

4.71
L-index

#	Paper	IF	Citations
89	Reverse Genomics: Design of Universal Epitope Sets to Isolate All Members from the Human Oral Cavity.. <i>Microorganisms</i> , 2022 , 10,	4.9	2
88	Emergence of <i>Candida auris</i> in intensive care units in Algeria.. <i>Mycoses</i> , 2022 ,	5.2	3
87	New Beta-lactamases in Candidate Phyla Radiation: Owing Pleiotropic Enzymes Is a Smart Paradigm for Microorganisms with a Reduced Genome. <i>International Journal of Molecular Sciences</i> , 2022 , 23, 5446	6.3	0
86	Occurrence of Ten Protozoan Enteric Pathogens in Three Non-Human Primate Populations. <i>Pathogens</i> , 2021 , 10,	4.5	3
85	Drug Repurposing in Medical Mycology: Identification of Compounds as Potential Antifungals to Overcome the Emergence of Multidrug-Resistant Fungi. <i>Pharmaceuticals</i> , 2021 , 14,	5.2	6
84	Rhizomal Reclassification of Living Organisms. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	6
83	SCA Medium: A New Culture Medium for the Isolation of All Clades. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021 , 7,	5.6	4
82	FastFung: A novel medium for the culture and isolation of fastidious fungal species from clinical samples. <i>Journal of Microbiological Methods</i> , 2021 , 180, 106108	2.8	2
81	Development and standardization of a specific real-time PCR assay for the rapid detection of <i>Candida auris</i> . <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021 , 40, 1547-1551	5.3	7
80	Adapted Protocol for Cocultivation: Two New Members Join the Club of Candidate Phyla Radiation.. <i>Microbiology Spectrum</i> , 2021 , 9, e0106921	8.9	2
79	Small and Equipped: the Rich Repertoire of Antibiotic Resistance Genes in Candidate Phyla Radiation Genomes. <i>MSystems</i> , 2021 , e0089821	7.6	4
78	<i>Gorillibacterium timonense</i> sp. nov., isolated from an obese patient. <i>Archives of Microbiology</i> , 2020 , 202, 1223-1229	3	0
77	Identification of repositionable drugs with novel antimycotic activity by screening the Prestwick Chemical Library against emerging invasive moulds. <i>Journal of Global Antimicrobial Resistance</i> , 2020 , 21, 314-317	3.4	12
76	Comparison of Three Skin Sampling Methods and Two Media for Culturing Yeast. <i>Journal of Fungi (Basel, Switzerland)</i> , 2020 , 6,	5.6	2
75	<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
74	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
73	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

72	In vitro polymyxin activity against clinical multidrug-resistant fungi. <i>Antimicrobial Resistance and Infection Control</i> , 2019 , 8, 66	6.2	25
71	A hospital qPCR-based survey of 10 gastrointestinal parasites in routine diagnostic screening, Marseille, France. <i>Epidemiology and Infection</i> , 2019 , 147, e100	4.3	9
70	Repurposing of Ribavirin as an Adjunct Therapy against Invasive Strains in an Study. <i>Antimicrobial Agents and Chemotherapy</i> , 2019 , 63,	5.9	12
69	Draft genome and description of <i>Chryseobacterium phocaeense</i> 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	
68	Colonization Is Associated with Increased Diversity and Altered Gut Bacterial Communities in Healthy Malian Children. <i>Microorganisms</i> , 2019 , 7,	4.9	13
67	Draft genome and description of <i>Cohnella massiliensis</i> sp. nov., a new bacterial species isolated from the blood culture of a hemodialysis patient. <i>Archives of Microbiology</i> , 2019 , 201, 305-312	3	8
66	Taxonogenomics description of <i>Parabacteroides timonensis</i> sp. nov. isolated from a human stool sample. <i>MicrobiologyOpen</i> , 2019 , 8, e00702	3.4	5
65	sp. nov., taxonogenomics description of a new bacterial species isolated from human gut. <i>New Microbes and New Infections</i> , 2018 , 23, 7-16	4.1	8
64	Draft Genome Sequence of "Nocardia suismassiliense" Strain S-137 (CSUR P4007). <i>Genome Announcements</i> , 2018 , 6,		3
63	First Draft Genome Sequences of Two Strains from Laos and Cambodia. <i>Genome Announcements</i> , 2018 , 6,		1
62	<i>Bacillus kwashiorkori</i> sp. nov., a new bacterial species isolated from a malnourished child using culturomics. <i>MicrobiologyOpen</i> , 2018 , 7, e00535	3.4	5
61	Evaluation of two DNA extraction methods for the PCR-based detection of eukaryotic enteric pathogens in fecal samples. <i>BMC Research Notes</i> , 2018 , 11, 206	2.3	25
60	Draft Genome and Description of <i>Eisenbergiella massiliensis</i> Strain AT11: A New Species Isolated from Human Feces After Bariatric Surgery. <i>Current Microbiology</i> , 2018 , 75, 1274-1281	2.4	3
59	Metabarcoding analysis of eukaryotic microbiota in the gut of HIV-infected patients. <i>PLoS ONE</i> , 2018 , 13, e0191913	3.7	14
58	Characterization of a New <i>Ezakiella</i> Isolated from the Human Vagina: Genome Sequence and Description of <i>Ezakiella massiliensis</i> sp. nov. <i>Current Microbiology</i> , 2018 , 75, 456-463	2.4	7
57	Description of <i>Mediterraneibacter massiliensis</i> , gen. nov., sp. nov., a new genus isolated from the gut microbiota of an obese patient and reclassification of <i>Ruminococcus faecis</i> , <i>Ruminococcus lactaris</i> , <i>Ruminococcus torques</i> , <i>Ruminococcus gnavus</i> and <i>Clostridium glycyrrhizinilyticum</i> as <i>Mediterraneibacter faecis</i> comb. nov., <i>Mediterraneibacter lactaris</i> comb. nov., <i>Mediterraneibacter torques</i> comb. nov., <i>Mediterraneibacter gnavus</i> comb. nov. and <i>Mediterraneibacter glycyrrhizinilyticum</i> comb. nov. <i>MicrobiologyOpen</i> , 2018 , 7, e00575	2.1	18
56	<i>Eggerthella timonensis</i> sp. nov, a new species isolated from the stool sample of a pygmy female. <i>MicrobiologyOpen</i> , 2018 , 7, e00575 <i>Journal of Microbiology and Biotechnology</i> , 2018 , 111, 2107-2128	3.4	4
55	Repertoire of human gut microbes. <i>Microbial Pathogenesis</i> , 2017 , 106, 103-112	3.8	46

54	Molecular survey of <i>Dirofilaria immitis</i> and <i>Dirofilaria repens</i> by new real-time TaqMan PCR assay in dogs and mosquitoes (Diptera: Culicidae) in Corsica (France). <i>Veterinary Parasitology</i> , 2017 , 235, 1-7	2.8	19
53	Noncontiguous finished genome sequence and description of sp. nov. isolated from the human stomach. <i>New Microbes and New Infections</i> , 2017 , 15, 77-88	4.1	5
52	Description of <i>Chryseobacterium timonianum</i> sp. nov., isolated from a patient with pneumonia. <i>Antonie Van Leeuwenhoek</i> , 2017 , 110, 1121-1132	2.1	7
51	<i>Olsenella provencensis</i> sp. nov., <i>Olsenella phocaeensis</i> sp. nov., and <i>Olsenella mediterranea</i> sp. nov. isolated from the human colon. <i>Human Microbiome Journal</i> , 2017 , 4, 22-23	5.6	7
50	gen. nov., sp. nov., a new member of the family isolated from the left colon of a 27-year-old woman. <i>New Microbes and New Infections</i> , 2017 , 16, 63-72	4.1	1
49	<i>Hugonella massiliensis</i> gen. nov., sp. nov., genome sequence, and description of a new strictly anaerobic bacterium isolated from the human gut. <i>MicrobiologyOpen</i> , 2017 , 6, e00458	3.4	16
48	gen. nov., sp. nov., a new bacterial genus isolated from the human gut. <i>New Microbes and New Infections</i> , 2017 , 16, 30-31	4.1	
47	Noncontiguous finished genome sequence and description of sp. nov. isolated from human duodenum. <i>New Microbes and New Infections</i> , 2017 , 16, 3-12	4.1	2
46	Many More Microbes in Humans: Enlarging the Microbiome Repertoire. <i>Clinical Infectious Diseases</i> , 2017 , 65, S20-S29	11.6	17
45	Culturomics and Amplicon-based Metagenomic Approaches for the Study of Fungal Population in Human Gut Microbiota. <i>Scientific Reports</i> , 2017 , 7, 16788	4.9	47
44	Discovery of a new species within genus in a stool sample from Gabon: sp. nov. <i>New Microbes and New Infections</i> , 2017 , 15, 3-5	4.1	2
43	Performance of Real-Time Polymerase Chain Reaction Assays for the Detection of 20 Gastrointestinal Parasites in Clinical Samples from Senegal. <i>American Journal of Tropical Medicine and Hygiene</i> , 2017 , 97, 173-182	3.2	17
42	a new species identified from human stool from an obese patient before bariatric surgery. <i>New Microbes and New Infections</i> , 2016 , 14, 67-68	4.1	1
41	Culture of previously uncultured members of the human gut microbiota by culturomics. <i>Nature Microbiology</i> , 2016 , 1, 16203	26.6	555
40	<i>Anaerococcus rubiinfantis</i> sp. nov., isolated from the gut microbiota of a Senegalese infant with severe acute malnutrition. <i>Anaerobe</i> , 2016 , 40, 85-94	2.8	6
39	Non-contiguous-Finished Genome Sequence and Description of <i>Paenibacillus camerounensis</i> sp. nov. <i>Microbial Ecology</i> , 2016 , 71, 990-8	4.4	3
38	" <i>Actinomyces ihumii</i> ," a new bacterial species isolated from the digestive microbiota of a HIV-infected patient. <i>New Microbes and New Infections</i> , 2016 , 12, 71-2	4.1	2
37	Non contiguous-finished genome sequence and description of <i>Microbacterium gorillae</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2016 , 11, 32		12

36	"Raoultibacter massiliensis" gen. nov., sp. nov., a new bacterium isolated from the human gut of a Saudi Bedouin. <i>New Microbes and New Infections</i> , 2016 , 14, 1-3	4.1	5
35	<i>Clostridium massiliodiemoense</i> a new species isolated from the human gut microbiota. <i>New Microbes and New Infections</i> , 2016 , 13, 27-8	4.1	1
34	<i>Eisenbergiella massiliensis</i> a new species isolated from human stool collected after bariatric surgery. <i>New Microbes and New Infections</i> , 2016 , 13, 15-6	4.1	13
33	Evaluation of a new extraction protocol for yeast identification by mass spectrometry. <i>Journal of Microbiological Methods</i> , 2016 , 129, 61-65	2.8	5
32	Repertory of eukaryotes (eukaryome) in the human gastrointestinal tract: taxonomy and detection methods. <i>Parasite Immunology</i> , 2016 , 38, 12-36	2.2	34
31	Reply to Bastien et al. <i>Journal of Infectious Diseases</i> , 2015 , 212, 506-8	7	2
30	Wild gorillas as a potential reservoir of <i>Leishmania major</i> . <i>Journal of Infectious Diseases</i> , 2015 , 211, 267-73		19
29	Non contiguous-finished genome sequence and description of <i>Bacillus jeddahensis</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2015 , 10, 47		8
28	Cutaneous and pulmonary dirofilariasis due to <i>Dirofilaria repens</i> . <i>British Journal of Dermatology</i> , 2015 , 173, 788-91	4	15
27	Mansonella, including a Potential New Species, as Common Parasites in Children in Gabon. <i>PLoS Neglected Tropical Diseases</i> , 2015 , 9, e0004155	4.8	22
26	Looking in apes as a source of human pathogens. <i>Microbial Pathogenesis</i> , 2014 , 77, 149-54	3.8	13
25	Non-contiguous finished genome sequence and description of <i>Gorillibacterium massiliense</i> gen. nov, sp. nov., a new member of the family Paenibacillaceae. <i>Standards in Genomic Sciences</i> , 2014 , 9, 807-20		8
24	Non-contiguous finished genome sequence and description of <i>Paenibacillus gorillae</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1031-45		3
23	Detection of termites and other insects consumed by African great apes using molecular fecal analysis. <i>Scientific Reports</i> , 2014 , 4, 4478	4.9	31
22	Pathogenic eukaryotes in gut microbiota of western lowland gorillas as revealed by molecular survey. <i>Scientific Reports</i> , 2014 , 4, 6417	4.9	14
21	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
20	Phenotypic and genotypic properties of <i>Microbacterium yannicii</i> , a recently described multidrug resistant bacterium isolated from a lung transplanted patient with cystic fibrosis in France. <i>BMC Microbiology</i> , 2013 , 13, 97	4.5	16
19	<i>Pandoraea pulmonicola</i> chronic colonization in a cystic fibrosis patient, France. <i>New Microbes and New Infections</i> , 2013 , 1, 27-9	4.1	11

18	Non-contiguous finished genome sequence and description of <i>Bacillus massiliogorillae</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2013 , 9, 93-105		23
17	Molecular detection of microorganisms in distal airways of patients undergoing lung cancer surgery. <i>Annals of Thoracic Surgery</i> , 2012 , 93, 413-22	2.7	10
16	Microbial culturomics: paradigm shift in the human gut microbiome study. <i>Clinical Microbiology and Infection</i> , 2012 , 18, 1185-93	9.5	722
15	Molecular detection of eukaryotes in a single human stool sample from Senegal. <i>PLoS ONE</i> , 2012 , 7, e40888	9.8	62
14	Detection and accurate identification of new or emerging bacteria in cystic fibrosis patients. <i>Clinical Microbiology and Infection</i> , 2010 , 16, 809-20	9.5	52
13	Outbreak of <i>Corynebacterium pseudodiphtheriticum</i> infection in cystic fibrosis patients, France. <i>Emerging Infectious Diseases</i> , 2010 , 16, 1231-6	10.2	39
12	<i>Nocardia farcinica</i> lung infection in a patient with cystic fibrosis: a case report. <i>Journal of Medical Case Reports</i> , 2010 , 4, 84	1.2	12
11	Hygiene of nasal masks used at home for non-invasive ventilation in children. <i>Journal of Hospital Infection</i> , 2010 , 76, 187-8	6.9	2
10	Microbial diversity in the sputum of a cystic fibrosis patient studied with 16S rDNA pyrosequencing. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2009 , 28, 1151-4	5.3	79
9	MALDI-TOF-MS for rapid detection of staphylococcal Panton-Valentine leukocidin. <i>International Journal of Antimicrobial Agents</i> , 2009 , 34, 467-70	14.3	73
8	Genomic analysis of an emerging multiresistant <i>Staphylococcus aureus</i> strain rapidly spreading in cystic fibrosis patients revealed the presence of an antibiotic inducible bacteriophage. <i>Biology Direct</i> , 2009 , 4, 1	7.2	71
7	First isolation of two colistin-resistant emerging pathogens, <i>Brevundimonas diminuta</i> and <i>Ochrobactrum anthropi</i> , in a woman with cystic fibrosis: a case report. <i>Journal of Medical Case Reports</i> , 2008 , 2, 373	1.2	36
6	<i>Acetobacter indonesiensis</i> pneumonia after lung transplant. <i>Emerging Infectious Diseases</i> , 2008 , 14, 997-8	8.2	17
5	Molecular detection of multiple emerging pathogens in sputa from cystic fibrosis patients. <i>PLoS ONE</i> , 2008 , 3, e2908	3.7	170
4	<i>Inquilinus limosus</i> and cystic fibrosis. <i>Emerging Infectious Diseases</i> , 2008 , 14, 993-5	10.2	22
3	Unexpected enzyme TEM-126: role of mutation Asp179Glu. <i>Antimicrobial Agents and Chemotherapy</i> , 2005 , 49, 4280-7	5.9	13
2	Adapted protocol for Saccharibacteria co-cultivation: two new members join the club of Candidate Phyla radiation		1
1	Small and equipped: the rich repertoire of antibiotic resistance genes in Candidate Phyla Radiation genomes		2

