Fadi Bittar

List of Publications by Year
in descending order

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91 papers	3,715 citations	279487 23 h-index	138251 58 g-index
99 all docs	99 docs citations	99 times ranked	4672 citing authors

#	Article	IF	Citations
1	Reverse Genomics: Design of Universal Epitope Sets to Isolate All Saccharibacteria Members from the Human Oral Cavity. Microorganisms, 2022, 10, 602.	1.6	3
2	Emergence of Candida auris in intensive care units in Algeria. Mycoses, 2022, 65, 753-759.	1.8	10
3	New Beta-lactamases in Candidate Phyla Radiation: Owning Pleiotropic Enzymes Is a Smart Paradigm for Microorganisms with a Reduced Genome. International Journal of Molecular Sciences, 2022, 23, 5446.	1.8	5
4	Candidate Phyla Radiation, an Underappreciated Division of the Human Microbiome, and Its Impact on Health and Disease. Clinical Microbiology Reviews, 2022, 35, .	5.7	19
5	FastFung: A novel medium for the culture and isolation of fastidious fungal species from clinical samples. Journal of Microbiological Methods, 2021, 180, 106108.	0.7	8
6	Development and standardization of a specific real-time PCR assay for the rapid detection of Candida auris. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 1547-1551.	1.3	8
7	Occurrence of Ten Protozoan Enteric Pathogens in Three Non-Human Primate Populations. Pathogens, 2021, 10, 280.	1.2	8
8	Drug Repurposing in Medical Mycology: Identification of Compounds as Potential Antifungals to Overcome the Emergence of Multidrug-Resistant Fungi. Pharmaceuticals, 2021, 14, 488.	1.7	24
9	Rhizomal Reclassification of Living Organisms. International Journal of Molecular Sciences, 2021, 22, 5643.	1.8	16
10	SCA Medium: A New Culture Medium for the Isolation of All Candida auris Clades. Journal of Fungi (Basel, Switzerland), 2021, 7, 433.	1.5	7
11	Adapted Protocol for Saccharibacteria Cocultivation: Two New Members Join the Club of Candidate Phyla Radiation. Microbiology Spectrum, 2021, 9, e0106921.	1.2	8
12	Small and Equipped: the Rich Repertoire of Antibiotic Resistance Genes in Candidate Phyla Radiation Genomes. MSystems, 2021, 6, e0089821.	1.7	13
13	Identification of repositionable drugs with novel antimycotic activity by screening the Prestwick Chemical Library against emerging invasive moulds. Journal of Global Antimicrobial Resistance, 2020, 21, 314-317.	0.9	21
14	Comparison of Three Skin Sampling Methods and Two Media for Culturing Malassezia Yeast. Journal of Fungi (Basel, Switzerland), 2020, 6, 350.	1.5	7
15	Gorillibacterium timonense sp. nov., isolated from an obese patient. Archives of Microbiology, 2020, 202, 1223-1229.	1.0	1
16	Repurposing of Ribavirin as an Adjunct Therapy against Invasive Candida Strains in an In Vitro Study. Antimicrobial Agents and Chemotherapy, 2019, 63, .	1.4	19
17	Draft genome and description of Chryseobacterium phocaeense sp. nov.: a new bacterial species isolated from the sputum of a cystic fibrosis patient. Archives of Microbiology, 2019, 201, 1361-1368.	1.0	0
18	Phoenicibacter congonensis gen. nov., sp. nov., a new genus isolated from the human gut and its description using a taxonogenomic approach. Antonie Van Leeuwenhoek, 2019, 112, 775-784.	0.7	2

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19	Noncontiguous finished genome sequence and description of Raoultibacter massiliensis gen. nov., sp. nov. and Raoultibacter timonensis sp. nov, two new bacterial species isolated from the human gut. MicrobiologyOpen, 2019, 8, e00758.	1.2	19
20	Non-contiguous finished genome sequencing and description of Enterococcus timonensis sp. nov. isolated from human sputum. New Microbes and New Infections, 2019, 29, 100532.	0.8	2
21	In vitro polymyxin activity against clinical multidrug-resistant fungi. Antimicrobial Resistance and Infection Control, 2019, 8, 66.	1.5	41
22	A hospital qPCR-based survey of 10 gastrointestinal parasites in routine diagnostic screening, Marseille, France. Epidemiology and Infection, 2019, 147, e100.	1.0	13
23	Blastocystis Colonization Is Associated with Increased Diversity and Altered Gut Bacterial Communities in Healthy Malian Children. Microorganisms, 2019, 7, 649.	1.6	35
24	Draft genome and description of Cohnella massiliensis sp. nov., a new bacterial species isolated from the blood culture of a hemodialysis patient. Archives of Microbiology, 2019, 201, 305-312.	1.0	10
25	Taxonogenomics description of Parabacteroides timonensis sp. nov. isolated from a human stool sample. MicrobiologyOpen, 2019, 8, e00702.	1.2	7
26	â€˜Cellulomonas timonensisâ€™ sp. nov., taxonogenomics description of a new bacterial species isolated from human gut. New Microbes and New Infections, 2018, 23, 7-16.	0.8	10
27	Draft Genome Sequence of â€œNocardia suismassilienseâ€•Strain S-137 (CSUR P4007). Genome Announcements, 2018, 6, .	0.8	3
28	First Draft Genome Sequences of Two Bartonella tribocorum Strains from Laos and Cambodia. Genome Announcements, 2018, 6, .	0.8	1
29	Bacillus kwashiorkori sp. nov., a new bacterial species isolated from a malnourished child using culturomics. MicrobiologyOpen, 2018, 7, e00535.	1.2	6
30	Characterization of a New Ezakiella Isolated from the Human Vagina: Genome Sequence and Description of Ezakiella massiliensis sp. nov.. Current Microbiology, 2018, 75, 456-463.	1.0	13
31	Description of Mediterraneibacter massiliensis gen. nov., sp. nov., a new genus isolated from the gut microbiota of an obese patient and reclassification of Ruminococcus faecis, Ruminococcus lactaris, Ruminococcus torques, Ruminococcus gnavus and Clostridium glycyrrhizinilyticum as Mediterraneibacter faecis comb. nov., Mediterraneibacter lactaris comb. nov., Mediterraneibacter torques comb. nov., Mediterraneibacter gnavus comb. nov. and Mediterraneibacter glycyrrhizinilyticus comb. nov.. Antonie Van Leeuwenhoek, 2018, 111, 2107-2128.	0.7	87
32	Eggerthella timonensis sp. nov, a new species isolated from the stool sample of a pygmy female. MicrobiologyOpen, 2018, 7, e00575.	1.2	5
33	Evaluation of two DNA extraction methods for the PCR-based detection of eukaryotic enteric pathogens in fecal samples. BMC Research Notes, 2018, 11, 206.	0.6	42
34	Draft Genome and Description of Eisenbergiella massiliensis Strain AT11T: A New Species Isolated from Human Feces After Bariatric Surgery. Current Microbiology, 2018, 75, 1274-1281.	1.0	4
35	Metabarcoding analysis of eukaryotic microbiota in the gut of HIV-infected patients. PLoS ONE, 2018, 13, e0191913.	1.1	23
36	Repertoire of human gut microbes. Microbial Pathogenesis, 2017, 106, 103-112.	1.3	70

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37	Molecular survey of Dirofilaria immitis and Dirofilaria repens by new real-time TaqManÂ® PCR assay in dogs and mosquitoes (Diptera: Culicidae) in Corsica (France). Veterinary Parasitology, 2017, 235, 1-7.	0.7	24
38	Noncontiguous finished genome sequence and description of Streptococcus timonensis sp. nov. isolated from the human stomach. New Microbes and New Infections, 2017, 15, 77-88.	0.8	7
39	Description of Chryseobacterium timonianum sp. nov., isolated from a patient with pneumonia. Antonie Van Leeuwenhoek, 2017, 110, 1121-1132.	0.7	8
40	Olsenella provencensis sp. nov., Olsenella phocaeensis sp. nov., and Olsenella mediterranea sp. nov. isolated from the human colon. Human Microbiome Journal, 2017, 4, 22-23.	3.8	11
41	Massilioclostridium coli gen. nov., sp. nov., a new member of the Clostridiaceae family isolated from the left colon of a 27-year-old woman. New Microbes and New Infections, 2017, 16, 63-72.	0.8	1
42	Hugonella massiliensis gen. nov., sp. nov., genome sequence, and description of a new strictly anaerobic bacterium isolated from the human gut. MicrobiologyOpen, 2017, 6, e00458.	1.2	18
43	Marseillibacter massiliensis gen. nov., sp. nov., a new bacterial genus isolated from the human gut. New Microbes and New Infections, 2017, 16, 30-31.	0.8	0
44	Noncontiguous finished genome sequence and description of Fusobacterium massiliense sp. nov. isolated from human duodenum. New Microbes and New Infections, 2017, 16, 3-12.	0.8	2
45	Many More Microbes in Humans: Enlarging the Microbiome Repertoire. Clinical Infectious Diseases, 2017, 65, S20-S29.	2.9	20
46	Culturomics and Amplicon-based Metagenomic Approaches for the Study of Fungal Population in Human Gut Microbiota. Scientific Reports, 2017, 7, 16788.	1.6	78
47	Discovery of a new species within Bacillus genus in a stool sample from Gabon: â€œBacillus massiliogabonensisâ€•sp. nov.. New Microbes and New Infections, 2017, 15, 3-5.	0.8	2
48	Performance of Real-Time Polymerase Chain Reaction Assays for the Detection of 20 Gastrointestinal Parasites in Clinical Samples from Senegal. American Journal of Tropical Medicine and Hygiene, 2017, 97, 173-182.	0.6	23
49	â€œActinomyces ihumii,â€•a new bacterial species isolated from the digestive microbiota of a HIV-infected patient. New Microbes and New Infections, 2016, 12, 71-72.	0.8	2
50	Non contiguous-finished genome sequence and description of Microbacterium gorillae sp. nov.. Standards in Genomic Sciences, 2016, 11, 32.	1.5	24
51	â€œRaoultibacter massiliensisâ€•gen. nov., sp. nov., a new bacterium isolated from the human gut of a Saudi Bedouin. New Microbes and New Infections, 2016, 14, 1-3.	0.8	8
52	â€˜ Clostridium massiliodielmoense â€™, a new species isolated from the human gutÂmicrobiota. New Microbes and New Infections, 2016, 13, 27-28.	0.8	1
53	â€˜ Eisenbergiella massiliensis â€™, a new species isolated from human stool collected after bariatric surgery. New Microbes and New Infections, 2016, 13, 15-16.	0.8	25
54	Evaluation of a new extraction protocol for yeast identification by mass spectrometry. Journal of Microbiological Methods, 2016, 129, 61-65.	0.7	6

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55	Repertory of eukaryotes (eukaryome) in the human gastrointestinal tract: taxonomy and detection methods. Parasite Immunology, 2016, 38, 12-36.	0.7	47
56	â€œRuminococcus phoceensis,â€•a new species identified from human stool from an obese patient before bariatric surgery. New Microbes and New Infections, 2016, 14, 67-68.	0.8	6
57	Culture of previously uncultured members of the human gut microbiota by culturomics. Nature Microbiology, 2016, 1, 16203.	5.9	735
58	Anaerococcus rubiinfantis sp. nov., isolated from the gut microbiota of a Senegalese infant with severe acute malnutrition. Anaerobe, 2016, 40, 85-94.	1.0	14
59	Non-contiguous-Finished Genome Sequence and Description of Paenibacillus camerounensis sp. nov.. Microbial Ecology, 2016, 71, 990-998.	1.4	5
60	Non contiguous-finished genome sequence and description of Bacillus jeddahensis sp. nov.. Standards in Genomic Sciences, 2015, 10, 47.	1.5	22
61	Cutaneous and pulmonary dirofilariasis due to Dirofilaria repens. British Journal of Dermatology, 2015, 173, 788-791.	1.4	18
62	Reply to Bastien et al: Figure 1.. Journal of Infectious Diseases, 2015, 212, 506-508.	1.9	2
63	Wild Gorillas as a Potential Reservoir ofLeishmania major. Journal of Infectious Diseases, 2015, 211, 267-273.	1.9	25
64	Mansonella, including a Potential New Species, as Common Parasites in Children in Gabon. PLoS Neglected Tropical Diseases, 2015, 9, e0004155.	1.3	35
65	Looking in apes as a source of human pathogens. Microbial Pathogenesis, 2014, 77, 149-154.	1.3	21
66	Non-contiguous finished genome sequence and description of Gorillibacterium massiliense gen. nov, sp. nov., a new member of the family Paenibacillaceae. Standards in Genomic Sciences, 2014, 9, 807-820.	1.5	17
67	Non-contiguous finished genome sequence and description of Paenibacillus gorillae sp. nov.. Standards in Genomic Sciences, 2014, 9, 1031-1045.	1.5	7
68	Detection of Termites and Other Insects Consumed by African Great Apes using Molecular Fecal Analysis. Scientific Reports, 2014, 4, 4478.	1.6	35
69	Pathogenic Eukaryotes in Gut Microbiota of Western Lowland Gorillas as Revealed by Molecular Survey. Scientific Reports, 2014, 4, 6417.	1.6	17
70	Gorilla gorilla gorilla gut: a potential reservoir of pathogenic bacteria as revealed using culturomics and molecular tools. Scientific Reports, 2014, 4, 7174.	1.6	53
71	Phenotypic and genotypic properties of Microbacterium yannicii, a recently described multidrug resistant bacterium isolated from a lung transplanted patient with cystic fibrosis in France. BMC Microbiology, 2013, 13, 97.	1.3	22
72	Pandoraea pulmonicola chronic colonization in a cystic fibrosis patient, France. New Microbes and New Infections, 2013, 1, 27-29.	0.8	20

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73	Non-contiguous finished genome sequence and description of Bacillus massiliogorillae sp. nov.. Standards in Genomic Sciences, 2013, 9, 93-105.	1.5	32
74	Microbial culturomics: paradigm shift in the human gut microbiome study. Clinical Microbiology and Infection, 2012, 18, 1185-1193.	2.8	905
75	Molecular Detection of Microorganisms in Distal Airways of Patients Undergoing Lung Cancer Surgery. Annals of Thoracic Surgery, 2012, 93, 413-422.	0.7	12
76	Molecular Detection of Eukaryotes in a Single Human Stool Sample from Senegal. PLoS ONE, 2012, 7, e40888.	1.1	86
77	Hygiene of nasal masks used at home for non-invasive ventilation in children. Journal of Hospital Infection, 2010, 76, 187-188.	1.4	3
78	Detection and accurate identification of new or emerging bacteria in cystic fibrosis patients. Clinical Microbiology and Infection, 2010, 16, 809-820.	2.8	64
79	Outbreak ofCorynebacterium pseudodiphtheriticumInfection in Cystic Fibrosis Patients, France. Emerging Infectious Diseases, 2010, 16, 1231-1236.	2.0	46
80	Nocardia farcinica lung infection in a patient with cystic fibrosis: a case report. Journal of Medical Case Reports, 2010, 4, 84.	0.4	15
81	Microbial diversity in the sputum of a cystic fibrosis patient studied with 16S rDNA pyrosequencing. European Journal of Clinical Microbiology and Infectious Diseases, 2009, 28, 1151-1154.	1.3	81
82	MALDI-TOF-MS for rapid detection of staphylococcal Pantonâ€"Valentine leukocidin. International Journal of Antimicrobial Agents, 2009, 34, 467-470.	1.1	93
83	Genomic analysis of an emerging multiresistant Staphylococcus aureus strain rapidly spreading in cystic fibrosis patients revealed the presence of an antibiotic inducible bacteriophage. Biology Direct, 2009, 4, 1.	1.9	113
84	First isolation of two colistin-resistant emerging pathogens, Brevundimonas diminuta and Ochrobactrum anthropi, in a woman with cystic fibrosis: a case report. Journal of Medical Case Reports, 2008, 2, 373.	0.4	45
85	Molecular detection of multiple emerging pathogens in sputa from cystic fibrosis patients. Journal of Cystic Fibrosis, 2008, 7, S41.	0.3	0
86	Emergence and spread of a phylogenetic cluster of Corynebacteria in cystic fibrosis patients. Journal of Cystic Fibrosis, 2008, 7, S49.	0.3	0
87	Inquilinus limosus: an easily missed emerging resistant respiratory pathogen in cystic fibrosis patients. Journal of Cystic Fibrosis, 2008, 7, S50.	0.3	1
88	Acetobacter indonesiensisPneumonia after Lung Transplant. Emerging Infectious Diseases, 2008, 14, 997-998.	2.0	19
89	Molecular Detection of Multiple Emerging Pathogens in Sputa from Cystic Fibrosis Patients. PLoS ONE, 2008, 3, e2908.	1.1	201
90	Inquilinus limosusand Cystic Fibrosis. Emerging Infectious Diseases, 2008, 14, 993-995.	2.0	28

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91	Unexpected Enzyme TEM-126: Role of Mutation Asp179Glu. Antimicrobial Agents and Chemotherapy, 2005, 49, 4280-4287.	1.4	17