

Fehmida Bibi

List of Publications by Year in descending order

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471061

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docs citations

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citing authors

#	ARTICLE	IF	CITATIONS
1	Arabiibacter massiliensis gen. nov. sp. nov., New Anaerobic Bacterium Isolated from the Human Gut. Current Microbiology, 2022, 79, 47.	1.0	6
2	Bacterial Community and Genomic Analysis of Carbapenem-Resistant Acinetobacter baumannii Isolates from the Environment of a Health Care Facility in the Western Region of Saudi Arabia. Pharmaceuticals, 2022, 15, 611.	1.7	1
3	Analysis of bacterial communities in sponges and coral inhabiting Red Sea, using barcoded 454 pyrosequencing. Saudi Journal of Biological Sciences, 2021, 28, 847-854.	1.8	5
4	Assessing the diversity of bacterial communities from marine sponges and their bioactive compounds. Saudi Journal of Biological Sciences, 2021, 28, 2747-2754.	1.8	6
5	Draft genome sequence of colistin-resistant and extended-spectrum β -lactamase (ESBL)-producing multidrug-resistant Escherichia coli isolated from poultry meat. Journal of Global Antimicrobial Resistance, 2021, 27, 112-114.	0.9	6
6	Bacterial diversity and the antimicrobial resistome in the southwestern highlands of Saudi Arabia. Saudi Journal of Biological Sciences, 2021, 29, 2138-2147.	1.8	2
7	Pyrosequencing reveals sponge specific bacterial communities in marine sponges of Red Sea, Saudi Arabia. Saudi Journal of Biological Sciences, 2020, 27, 67-73.	1.8	5
8	Exome Analysis Identified Novel Homozygous Splice Site Donor Alteration in NT5C2 Gene in a Saudi Family Associated With Spastic Diplegia Cerebral Palsy, Developmental Delay, and Intellectual Disability. Frontiers in Genetics, 2020, 11, 14.	1.1	14
9	Antimicrobial activity of bacteria from marine sponge Suberea mollis and bioactive metabolites of Vibrio sp. EA348. Saudi Journal of Biological Sciences, 2020, 27, 1139-1147.	1.8	32
10	Comparative metagenomics and characterization of antimicrobial resistance genes in pasteurized and homemade fermented Arabian laban. Food Research International, 2020, 137, 109639.	2.9	11
11	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. MicrobiologyOpen, 2019, 8, e00758.	1.2	19
12	Taxonomic diversity of antimicrobial-resistant bacteria and genes in the Red Sea coast. Science of the Total Environment, 2019, 677, 474-483.	3.9	23
13	Culturomics-Based Taxonomic Diversity of Bacterial Communities in the Hot Springs of Saudi Arabia. OMICS A Journal of Integrative Biology, 2019, 23, 17-27.	1.0	13
14	Halophytes-associated endophytic and rhizospheric bacteria: diversity, antagonism and metabolite production. Biocontrol Science and Technology, 2018, 28, 192-213.	0.5	18
15	Diversity and antagonistic potential of bacteria isolated from marine grass Halodule uninervis. 3 Biotech, 2018, 8, 48.	1.1	18
16	Detection of genetic alterations in gastric cancer patients from Saudi Arabia using comparative genomic hybridization (CGH). PLoS ONE, 2018, 13, e0202576.	1.1	12
17	Microbial Flora Associated with the Halophyte <i>Salsola imbricate</i> and Its Biotechnical Potential. Frontiers in Microbiology, 2018, 9, 65.	1.5	26
18	Anaerococcus jeddahensis sp. nov., a New Bacterial Species Isolated From Healthy Nomadic Bedouin Woman From Saudi Arabia. Current Microbiology, 2018, 75, 1419-1428.	1.0	6

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19	Chromosomal Micro-aberration in a Saudi Family with Juvenile Myoclonic Epilepsy. <i>CNS and Neurological Disorders - Drug Targets</i> , 2018, 16, 1010-1017.	0.8	7
20	Diversity of antagonistic bacteria isolated from medicinal plant <i>Peganum harmala</i> L.. <i>Saudi Journal of Biological Sciences</i> , 2017, 24, 1288-1293.	1.8	17
21	Comparative bacterial community analysis in relatively pristine and anthropogenically influenced mangrove ecosystems on the Red Sea. <i>Canadian Journal of Microbiology</i> , 2017, 63, 649-660.	0.8	23
22	Metformin attenuate PTZ-induced apoptotic neurodegeneration in human cortical neuronal cells. <i>Pakistan Journal of Medical Sciences</i> , 2017, 33, 581-585.	0.3	18
23	Detection and Genotyping of <i>Helicobacter pylori</i> among Gastric ulcer and Cancer Patients from Saudi Arabia. <i>Pakistan Journal of Medical Sciences</i> , 2017, 33, 320-324.	0.3	15
24	Gut microbiome and dietary patterns in different Saudi populations and monkeys. <i>Scientific Reports</i> , 2016, 6, 32191.	1.6	55
25	Culture of previously uncultured members of the human gut microbiota by culturomics. <i>Nature Microbiology</i> , 2016, 1, 16203.	5.9	735
26	microRNA analysis of gastric cancer patients from Saudi Arabian population. <i>BMC Genomics</i> , 2016, 17, 751.	1.2	36
27	Microbial Culturomics to Map Halophilic Bacterium in Human Gut: Genome Sequence and Description of <i>Oceanobacillus jeddahense</i> sp. nov.. <i>OMICS A Journal of Integrative Biology</i> , 2016, 20, 248-258.	1.0	11
28	Non contiguous-finished genome sequence and description of <i>Bacillus jeddahensis</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2015, 10, 47.	1.5	22
29	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.	1.5	16
30	Rise of Microbial Culturomics: Noncontiguous Finished Genome Sequence and Description of <i>Beduini massiliensis</i> gen. nov., sp. nov.. <i>OMICS A Journal of Integrative Biology</i> , 2015, 19, 766-776.	1.0	15
31	Composition of soil microbiome along elevation gradients in southwestern highlands of Saudi Arabia. <i>BMC Microbiology</i> , 2015, 15, 65.	1.3	37
32	Genome Sequence of <i>Bacillus simplex</i> Strain P558, Isolated from a Human Fecal Sample. <i>Genome Announcements</i> , 2014, 2, .	0.8	12
33	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, 117-124.	1.1	84
34	Non-contiguous finished genome sequence and description of <i>Clostridium saudii</i> sp. nov. <i>Standards in Genomic Sciences</i> , 2014, 9, 8.	1.5	15
35	Non-contiguous finished genome sequence and description of <i>Corynebacterium jeddahense</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2014, 9, 987-1002.	1.5	11
36	Non contiguous-finished genome sequence and description of <i>Clostridium jeddahense</i> sp. nov.. <i>Standards in Genomic Sciences</i> , 2014, 9, 1003-1019.	1.5	26

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37	Martellella endophytica sp. nov., an antifungal bacterium associated with a halophyte. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 2914-2919.	0.8	21
38	Diversity and Characterization of Endophytic Bacteria Associated with Tidal Flat Plants and their Antagonistic Effects on Oomycetous Plant Pathogens. Plant Pathology Journal, 2012, 28, 20-31.	0.7	69