

Radwa A Hanafy

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

460
citations

840585

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887953

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24
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24
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24
times ranked

376
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Pecoramyces ruminantium</i> , gen. nov., sp. nov., an anaerobic gut fungus from the feces of cattle and sheep. <i>Mycologia</i> , 2017, 109, 231-243.	0.8	89
2	Horizontal Gene Transfer as an Indispensable Driver for Evolution of Neocallimastigomycota into a Distinct Gut-Dwelling Fungal Lineage. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	1.4	61
3	<i>Feramyces austinii</i> , gen. nov., sp. nov., an anaerobic gut fungus from rumen and fecal samples of wild Barbary sheep and fallow deer. <i>Mycologia</i> , 2018, 110, 513-525.	0.8	55
4	Seven new Neocallimastigomycota genera from wild, zoo-housed, and domesticated herbivores greatly expand the taxonomic diversity of the phylum. <i>Mycologia</i> , 2020, 112, 1212-1239.	0.8	50
5	A fast and reliable procedure for spore collection from anaerobic fungi: Application for RNA uptake and long-term storage of isolates. <i>Journal of Microbiological Methods</i> , 2016, 127, 206-213.	0.7	33
6	Assessing anaerobic gut fungal diversity in herbivores using <i>D1</i> / <i>D2</i> large ribosomal subunit sequencing and multi-year isolation. <i>Environmental Microbiology</i> , 2020, 22, 3883-3908.	1.8	32
7	Molecular Dating of the Emergence of Anaerobic Rumen Fungi and the Impact of Laterally Acquired Genes. <i>MSystems</i> , 2019, 4, .	1.7	28
8	<i>Aestipascuomyces dupliciliberans</i> gen. nov, sp. nov., the First Cultured Representative of the Uncultured SK4 Clade from Aoudad Sheep and Alpaca. <i>Microorganisms</i> , 2020, 8, 1734.	1.6	21
9	<i>Anaeromyces contortus</i> , sp. nov., a new anaerobic gut fungal species (Neocallimastigomycota) isolated from the feces of cow and goat. <i>Mycologia</i> , 2018, 110, 502-512.	0.8	18
10	<i>Paucimyces polynucleatus</i> gen. nov, sp. nov., a novel polycentric genus of anaerobic gut fungi from the faeces of a wild blackbuck antelope. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	14
11	Draft genome sequence of <i>Pseudomonas moraviensis</i> strain Devor implicates metabolic versatility and bioremediation potential. <i>Genomics Data</i> , 2016, 9, 154-159.	1.3	11
12	Taxonomy of the anaerobic gut fungi (Neocallimastigomycota): a review of classification criteria and description of current taxa. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	0.8	11
13	Draft genome sequence of <i>Microbacterium oleivorans</i> strain Wellendorf implicates heterotrophic versatility and bioremediation potential. <i>Genomics Data</i> , 2016, 10, 54-60.	1.3	9
14	Draft genome sequence and detailed analysis of <i>Pantoea eucrina</i> strain Russ and implication for opportunistic pathogenesis. <i>Genomics Data</i> , 2016, 10, 63-68.	1.3	6
15	Draft genome sequence of <i>Micrococcus luteus</i> strain O'Kane implicates metabolic versatility and the potential to degrade polyhydroxybutyrates. <i>Genomics Data</i> , 2016, 9, 148-153.	1.3	5
16	Draft genome sequence of <i>Staphylococcus hominis</i> strain Hudgins isolated from human skin implicates metabolic versatility and several virulence determinants. <i>Genomics Data</i> , 2016, 10, 91-96.	1.3	2
17	Draft Genome of the <i>Arthrobacter</i> sp. Strain Edens01. <i>Genome Announcements</i> , 2015, 3, .	0.8	1
18	The Draft Genome Sequence of <i>Xanthomonas</i> sp. Strain Mitacek01 Expands the Pangenome of a Genus of Plant Pathogens. <i>Genome Announcements</i> , 2015, 3, .	0.8	0