

Carly P Rosewarne

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4906211/publications.pdf>

Version: 2024-02-01

16
papers

1,433
citations

623734

14
h-index

940533

16
g-index

16
all docs

16
docs citations

16
times ranked

2345
citing authors

#	ARTICLE	IF	CITATIONS
1	Effect of Fecal Microbiota Transplantation on 8-Week Remission in Patients With Ulcerative Colitis. JAMA - Journal of the American Medical Association, 2019, 321, 156.	7.4	548
2	Cultivation and sequencing of rumen microbiome members from the Hungate1000 Collection. Nature Biotechnology, 2018, 36, 359-367.	17.5	414
3	In Vitro Response of Rumen Microbiota to the Antimethanogenic Red Macroalga <i>Asparagopsis taxiformis</i> . Microbial Ecology, 2018, 75, 811-818.	2.8	42
4	Culture- and metagenomics-enabled analyses of the <i>Methanosphaera</i> genus reveals their monophyletic origin and differentiation according to genome size. ISME Journal, 2018, 12, 2942-2953.	9.8	24
5	Fiber Supplements Derived From Sugarcane Stem, Wheat Dextrin and Psyllium Husk Have Different In Vitro Effects on the Human Gut Microbiota. Frontiers in Microbiology, 2018, 9, 1618.	3.5	25
6	Genomic insights into the carbohydrate catabolism of <i>Cairneyella variabilis</i> gen. nov. sp. nov., the first reports from a genome of an ericoid mycorrhizal fungus from the southern hemisphere. Mycorrhiza, 2016, 26, 345-352.	2.8	18
7	Enhancing biogenic methane generation from a brown coal by combining different microbial communities. International Journal of Coal Geology, 2016, 154-155, 107-110.	5.0	23
8	Analysis of the bovine rumen microbiome reveals a diversity of Sus-like polysaccharide utilization loci from the bacterial phylum <i>Bacteroidetes</i> . Journal of Industrial Microbiology and Biotechnology, 2014, 41, 601-606.	3.0	41
9	Draft Genome Sequence of <i>Methanobacterium</i> sp. Maddingley, Reconstructed from Metagenomic Sequencing of a Methanogenic Microbial Consortium Enriched from Coal-Seam Gas Formation Water. Genome Announcements, 2013, 1, .	0.8	10
10	Draft Genome Sequence of <i>Clostridium</i> sp. Maddingley, Isolated from Coal-Seam Gas Formation Water. Genome Announcements, 2013, 1, .	0.8	9
11	Draft Genome Sequence of <i>Treponema</i> sp. Strain JC4, a Novel Spirochete Isolated from the Bovine Rumen. Journal of Bacteriology, 2012, 194, 4130-4130.	2.2	33
12	High-Yield and Phylogenetically Robust Methods of DNA Recovery for Analysis of Microbial Biofilms Adherent to Plant Biomass in the Herbivore Gut. Microbial Ecology, 2011, 61, 448-454.	2.8	33
13	Class 1 integrons in benthic bacterial communities: abundance, association with Tn402-like transposition modules and evidence for coselection with heavy-metal resistance. FEMS Microbiology Ecology, 2010, 72, 35-46.	2.7	130
14	Cytotoxic G-rich oligodeoxynucleotides: putative protein targets and required sequence motif. Nucleic Acids Research, 2007, 35, 4562-4572.	14.5	21
15	ST64B is a defective bacteriophage in <i>Salmonella enterica</i> serovar Typhimurium DT64 that encodes a functional immunity region capable of mediating phage-type conversion. International Journal of Medical Microbiology, 2004, 294, 59-63.	3.6	15
16	Genomic Structure of the <i>Salmonella enterica</i> Serovar Typhimurium DT 64 Bacteriophage ST64T: Evidence for Modular Genetic Architecture. Journal of Bacteriology, 2003, 185, 3473-3475.	2.2	47