Sean P Kennedy

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4278698/publications.pdf

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1163117 1125743 6,211 13 8 13 citations h-index g-index papers 14 14 14 10422 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Richness of human gut microbiome correlates with metabolic markers. Nature, 2013, 500, 541-546.	27.8	3,641
2	Dietary intervention impact on gut microbial gene richness. Nature, 2013, 500, 585-588.	27.8	1,485
3	Identification and assembly of genomes and genetic elements in complex metagenomic samples without using reference genomes. Nature Biotechnology, 2014, 32, 822-828.	17.5	909
4	Whole-genome mapping of 5′ RNA ends in bacteria by tagged sequencing: a comprehensive view in <i>Enterococcus faecalis</i> . Rna, 2015, 21, 1018-1030.	3.5	59
5	Evidence for contamination as the origin for bacteria found in human placenta rather than a microbiota. PLoS ONE, 2020, 15, e0237232.	2.5	42
6	RadD Contributes to R-Loop Avoidance in Sub-MIC Tobramycin. MBio, 2019, 10, .	4.1	17
7	Alteration of the gut fecal microbiome in children living with HIV on antiretroviral therapy in Yaounde, Cameroon. Scientific Reports, 2021, 11, 7666.	3.3	15
8	Sequana Coverage: Detection and Characterization of Genomic Variations using Running Median and Mixture Models. GigaScience, 2018, 7, .	6.4	11
9	Dynamics of extended-spectrum beta-lactamase-producing Enterobacterales colonization in long-term carriers following travel abroad. Microbial Genomics, 2021, 7, .	2.0	11
10	Protein supplementation during an energy-restricted diet induces visceral fat loss and gut microbiota amino acid metabolism activation: a randomized trial. Scientific Reports, 2021, 11, 15620.	3.3	9
11	Fitness Restoration of a Genetically Tractable Enterococcus faecalis V583 Derivative To Study Decoration-Related Phenotypes of the Enterococcal Polysaccharide Antigen. MSphere, 2019, 4, .	2.9	6
12	Dynamic insights on transcription initiation and RNA processing during bacterial adaptation. Rna, 2020, 26, 382-395.	3.5	4
13	Complete Genome and Methylome Analysis of the Box-Shaped Halophilic Archaeon Haloarcula sinaiiensis ATCC 33800. Microbiology Resource Announcements, 2021, 10, e0061921.	0.6	2