

Sean P Kennedy

List of Publications by Year in descending order

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13
papers

6,211
citations

1163117

8
h-index

1125743

13
g-index

14
all docs

14
docs citations

14
times ranked

10422
citing authors

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