Tulika Prakash

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2045900/publications.pdf

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	1478505	1125743
175	6	13
citations	h-index	g-index
19	19	345
docs citations	times ranked	citing authors
	citations 19	175 6 citations h-index 19 19

#	Article	IF	CITATIONS
1	Impact of dietary fructooligosaccharides (FOS) on murine gut microbiota and intestinal IgA secretion. 3 Biotech, 2022, 12, 56.	2.2	2
2	Shotgun metagenomic sequencing revealed the prebiotic potential of a grain-based diet in mice. Scientific Reports, 2022, 12, 6748.	3.3	2
3	Gut microbiota alternation under the intestinal epithelium-specific knockout of mouse Piga gene. Scientific Reports, 2022, 12, .	3.3	3
4	Decoding the roles of extremophilic microbes in the anaerobic environments: Past, Present, and Future. Current Research in Microbial Sciences, 2022, , 100146.	2.3	1
5	Comparative gut microbiome analysis of the Prakriti and Sasang systems reveals functional level similarities in constitutionally similar classes. 3 Biotech, 2020, 10, 379.	2.2	4
6	In silico functional and evolutionary analyses of rubber oxygenases (RoxA and RoxB). 3 Biotech, 2020, 10, 376.	2.2	2
7	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. PLoS ONE, 2020, 15, e0228358.	2.5	17
8	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice., 2020, 15, e0228358.		0
9	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		O
10	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice., 2020, 15, e0228358.		0
11	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		O
12	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		0
13	Association of colitis with gut-microbiota dysbiosis in clathrin adapter AP-1B knockout mice. , 2020, 15, e0228358.		O
14	Functional signature analysis of extreme Prakriti endo-phenotypes in gut microbiome of western Indian rural population. Bioinformation, 2019, 15, 490-505.	0.5	6
15	Exploration of Survival Traits, Probiotic Determinants, Host Interactions, and Functional Evolution of Bifidobacterial Genomes Using Comparative Genomics. Genes, 2018, 9, 477.	2.4	16
16	Metabolic and taxonomic insights into the Gram-negative natural rubber degrading bacterium Steroidobacter cummioxidans sp. nov., strain 35Y. PLoS ONE, 2018, 13, e0197448.	2.5	26
17	Enterotype Variations of the Healthy Human Gut Microbiome in Different Geographical Regions. Bioinformation, 2018, 14, 560-573.	0.5	65
18	Comparative Genomics of Herpesviridae Family to Look for Potential Signatures of Human Infecting Strains. International Journal of Genomics, 2016, 2016, 1-10.	1.6	26