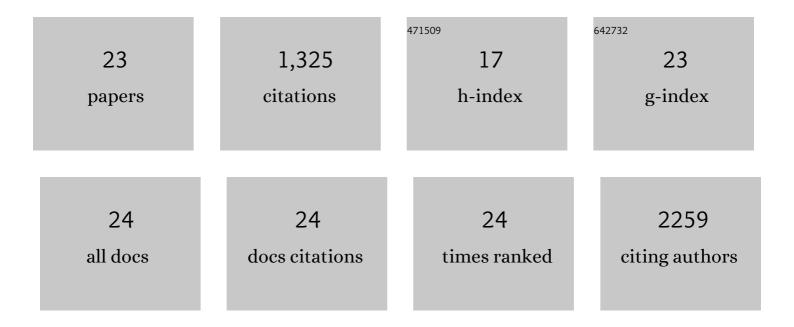
## Krithivasan Sankaranarayanan

List of Publications by Year in descending order

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Krithivasan

#	Article	IF	CITATIONS
1	Subsistence strategies in traditional societies distinguish gut microbiomes. Nature Communications, 2015, 6, 6505.	12.8	449
2	Intrinsic challenges in ancient microbiome reconstruction using 16S rRNA gene amplification. Scientific Reports, 2015, 5, 16498.	3.3	153
3	Differential preservation of endogenous human and microbial DNA in dental calculus and dentin. Scientific Reports, 2018, 8, 9822.	3.3	88
4	The evolution and changing ecology of the African hominid oral microbiome. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	74
5	Comparison of extraction methods for recovering ancient microbial DNA from paleofeces. American Journal of Physical Anthropology, 2020, 171, 275-284.	2.1	71
6	Gut Microbiome Diversity among Cheyenne and Arapaho Individuals from Western Oklahoma. Current Biology, 2015, 25, 3161-3169.	3.9	69
7	Successful enrichment and recovery of whole mitochondrial genomes from ancient human dental calculus. American Journal of Physical Anthropology, 2016, 160, 220-228.	2.1	68
8	Insights into human evolution from ancient and contemporary microbiome studies. Current Opinion in Genetics and Development, 2016, 41, 14-26.	3.3	49
9	The dental calculus metabolome in modern and historic samples. Metabolomics, 2017, 13, 134.	3.0	44
10	Mapping of host-parasite-microbiome interactions reveals metabolic determinants of tropism and tolerance in Chagas disease. Science Advances, 2020, 6, eaaz2015.	10.3	39
11	Biogeographic study of human gut-associated crAssphage suggests impacts from industrialization and recent expansion. PLoS ONE, 2020, 15, e0226930.	2.5	38
12	Reclassification of Facklamia ignava, Facklamia sourekii and Facklamia tabacinasalis as Falseniella ignava gen. nov., comb. nov., Hutsoniella sourekii gen. nov., comb. nov., and Ruoffia tabacinasalis gen. nov., comb. nov., and description of Ruoffia halotolerans sp. nov., isolated from hypersaline Inland Sea of Qatar. Antonie Van Leeuwenhoek, 2021, 114, 1181-1193.	1.7	28
13	Oral microbiome diversity among Cheyenne and Arapaho individuals from Oklahoma. American Journal of Physical Anthropology, 2016, 161, 321-327.	2.1	25
14	The efficacy of whole human genome capture on ancient dental calculus and dentin. American Journal of Physical Anthropology, 2019, 168, 496-509.	2.1	24
15	Characterization of Ancient DNA Supports Long-Term Survival of Haloarchaea. Astrobiology, 2014, 14, 553-560.	3.0	22
16	Ningiella ruwaisensis gen. nov., sp. nov., a member of the family Alteromonadaceae isolated from marine water of the Arabian Gulf. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 4130-4138.	1.7	22
17	The earliest farmers of northwest China exploited grain-fed pheasants not chickens. Scientific Reports, 2020, 10, 2556.	3.3	18
18	Shifts in gut and vaginal microbiomes are associated with cancer recurrence time in women with ovarian cancer. PeerJ, 2021, 9, e11574.	2.0	16

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#	Article	IF	CITATIONS
19	Inherited and somatic mitochondrial DNA mutations in Guam amyotrophic lateral sclerosis and parkinsonism-dementia. Neurological Sciences, 2011, 32, 883-892.	1.9	13
20	Analysis of global human gut metagenomes shows that metabolic resilience potential for short-chain fatty acid production is strongly influenced by lifestyle. Scientific Reports, 2021, 11, 1724.	3.3	11
21	"The dead shall be raised": Multidisciplinary analysis of human skeletons reveals complexity in 19th century immigrant socioeconomic history and identity in New Haven, Connecticut. PLoS ONE, 2019, 14, e0219279.	2.5	2
22	What ancient DNA reveals about the ubiquitous rockfish of the Pacific Coast of North America. Archaeological and Anthropological Sciences, 2022, 14, 1.	1.8	1
23	Associations of urinary microbiome profiles with gestational diabetes mellitus in an Oklahoma case-control study. ISEE Conference Abstracts, 2021, 2021, .	0.0	0