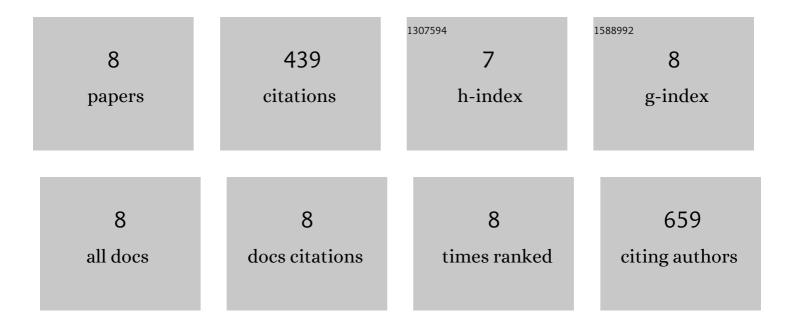


List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2094147/publications.pdf Version: 2024-02-01



YANGLI

#	Article	IF	CITATIONS
1	The rumen microbiome inhibits methane formation through dietary choline supplementation. Scientific Reports, 2021, 11, 21761.	3.3	3
2	Use of Lactic Acid Bacteria to Reduce Methane Production in Ruminants, a Critical Review. Frontiers in Microbiology, 2019, 10, 2207.	3.5	53
3	Gene and transcript abundances of bacterial type III secretion systems from the rumen microbiome are correlated with methane yield in sheep. BMC Research Notes, 2017, 10, 367.	1.4	8
4	Complete Genome Sequence of Methanogenic Archaeon ISO4-G1, a Member of the <i>Methanomassiliicoccales</i> , Isolated from a Sheep Rumen. Genome Announcements, 2016, 4, .	0.8	9
5	Rumen metagenome and metatranscriptome analyses of low methane yield sheep reveals a Sharpea-enriched microbiome characterised by lactic acid formation and utilisation. Microbiome, 2016, 4, 56.	11.1	268
6	The complete genome sequence of the methanogenic archaeon ISO4-H5 provides insights into the methylotrophic lifestyle of a ruminal representative of the Methanomassiliicoccales. Standards in Genomic Sciences, 2016, 11, 59.	1.5	41
7	Investigation of the Essentiality of Clutamate Racemase in Mycobacterium smegmatis. Journal of Bacteriology, 2014, 196, 4239-4244.	2.2	15
8	The Complete Genome Sequence of Methanobrevibacter sp. AbM4. Standards in Genomic Sciences, 2013, 8, 215-227.	1.5	42