

1 **Table S1.** Primers, gDNA concentrations, and primer efficiency for the quantification real-time polymerase chain reaction assay

Microbes	gDNA (ng/10 ul)	Forward (F)/ Reverse (R)	Primer concentration	Primer sequence	Reference	Primer efficiency (%)
Total bacteria	1	F	0.3 uM	CGGCAACGAGCGCAACCC	Denman and McSweeney, 2006, FEMS Microbiol. Ecol. 58:572.	108
		R	0.3 uM	CCATTGTAGCACGTGTGTAGCC		
Total protozoa	20	F	0.3 uM	GCTTTCGWTGGTAGTGTATT	Sylvester et al., 2004, J. Nutr. 134: 3378.	80.3
		R	0.3 uM	CTTGCCCTCYAATCGTWCT		
Total fungi	10	F	0.3 uM	GAGGAAGTAAAAGTCGTAACAAGGTTTC	Denman and McSweeney, 2006, FEMS Microbiol. Ecol. 58:572.	96.1
		R	0.3 uM	CAAATTCACAAAGGGTAGGATGATT		
Methanogenic archaea	5	F	0.5 uM	GAGGAAGGAGTGGACGACGGTA	Ohene-Adjei et al., 2007. Appl. Environ. Microbiol. 73:4609.	95.5
		R	0.5 uM	ACGGGCGGTGTGTGCAAG		
<i>Anaerovibrio lipolytica</i>	10	F	0.5 uM	TGGGTGTTAGAAATGGATTCTAGTG	Khafipour et al., 2009, Appl. Environ. Microbiol. 75:7115	109
		R	0.5 uM	GCACGTCATTCCGTATTAGCAT		
<i>Fibrobacter succinogenes</i>	5	F	0.5 uM	GGAGCGTAGGCGGAGATTCA	Denman and McSweeney, 2006, FEMS Microbiol. Ecol. 58:572.	98.5
		R	0.5 uM	GCCTGCCCCTGAACTATCCA		
<i>Ruminobacter amylophilus</i>	5	F	0.3 uM	CTGGGGAGCTGCCTGAAT	Stevenson and Weimer, 2007, Appl. Microbiol. Biotechnol. 75,165.	91.8
		R	0.3 uM	CATCTGAATGCGACTGGTTG		
<i>Ruminococcus albus</i>	10	F	0.5 uM	CCCTAAAAGCAGTCTTAGTTCG	Koike and Kobayashi, 2001, FEMS Microbiol. Lett. 204:361	98.0
		R	0.5 uM	CCTCCTTGCGGTTAGAACA		
<i>Ruminococcus flavefaciens</i>	5	F	0.5 uM	CGAACGGAGATAATTTGAGTTTACTTAGG	Denman and McSweeney, 2006, FEMS Microbiol. Ecol. 58:572.	92.4
		R	0.5 uM	CGGTCTCTGTATGTTATGAGGTATTACC		
<i>Streptococcus bovis</i>	20	F	0.5 uM	TTCCTAGAGATAGGAAGTTTCTTCGG	Sylvester et al., 2004, J. Nutr. 134:3378	112
		R	0.5 uM	ATGATGGCAACTAACAATAGGGGT		
<i>Succinimonas amylolytica</i>	5	F	0.5 uM	CGTTGGGCGGTCATTGAAAC	Khafipour et al., 2009, Appl. Environ. Microbiol. 75:7115	85.8
		R	0.5 uM	CCTGAGCGTCAGTTACTATCCAGA		
<i>Methanobrevibacter</i> spp.	10	F	0.3 uM	CCTCCGCAATGTGAGAAATCGC	Ramírez-Restrepo et al., 2016, Anim. Feed Sci. Tech. 216: 58.	94.6
		R	0.3 uM	TCWCCAGCAATCCCCACAGTT		

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