

**Table S1** Primer sequence, annealing temperature, and size of PCR products

Target	Primer sequence	Tm (°C)	Size (bp)	Reference
Total bacteria	ACTCCTACGGGAGGCAGCAGT GTATTACCGCGGCTGCTGGCAC	56	200	Tannock, Munro, Harmsen, Welling, Smart and Gopal [1]
<i>Bifidobacterium</i> spp.	TCGCGTC)C/T(GGTGTGAAAG CCACATCCAGC)A/G(TCCAC	58	243	Rinttilä, Kassinen, Malinen, Krogius and Palva [2]
<i>Bacteroides fragilis</i> group	ATAGCCTTCGAAAGRAAGAT CCAGTATCAACTGCAATTAA	50	501	Matsuki, Watanabe, Fujimoto, Miyamoto, Takada, Matsumoto, Oyaizu and Tanaka [3]
<i>Enterobacteriaceae</i>	CATTGACGTTACCCGCAGAAGAA GC CTCTACGAGACTCAAGCTTGC	57	195	Bartosch, Fite, Macfarlane and McMurdo [4]
<i>Clostridium coccoides</i> – <i>Eubacteria. rectale</i> group	CGGTACCTGACTAAGAACG AGTTT(C/T)ATTCTTGCAGACG	51	429	Rinttilä, Kassinen, Malinen, Krogius and Palva [2]
<i>Clostridium leptum</i> group	GCACAAGCAGTGGAGT CTTCCTCCGTTTGCAA	55	239	Matsuki, Watanabe,

					Fujimoto, Takada and Tanaka [5]
<i>F. prausnitzii</i> -	ACCATGAGAGCCGGGGGG	57	100	Lund, Bjerrum and	
<i>S. variabile</i> group	GGTTACCTTGTACGACTT			Pedersen [6]	
<i>Lactobacillus</i> group	AGCAGTAGGGAATCTTCCA CACCGCTACACATGGAG	53	341	Walter, Hertel, Tannock, Lis, Munro and Hammes [7]	
<i>Prevotella</i>	CACRGTAACGATGGATGCC GGTCGGGTTGCAGACC	62	513	Matsuki, Watanabe, Fujimoto, Miyamoto, Takada, Matsumoto, Oyaizu and Tanaka [3]	

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## References for Table S1

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