

Table S1

a. Primers used for qPCR - based quantification of *Aggregatibacter actinomycetemcomitans* and *Filifactor alocis* in accordance to Kirakodu et al. [18] and Siquiera & Rocas [19].

b. Cycle settings for quantification of *A. actinomycetemcomitans* and *F. alocis* in accordance to Kirakodu et al. [18] and Siquiera & Rocas [19].

a	Forward	Reverse
Kirakodu <i>Aa</i> (<i>ItxA</i>)	CTAGGTATTGCGAAACAATTTG	CCTGAAATTAAGCTGGTAATC
Siquiera and Rocas <i>Fa</i>	AGGCAGCTTGCCATACTGCG	ACTGTTAGCAACTACCGATGT

b	Kirakodu (<i>Aa</i>)	Siquiera and Rocas (<i>Fa</i>)
Hold/time	95°/10 min	95°/2 min
Cycling/time	95°/10 s	95°/30 s
Cycling/time	55°/5 s	55°/1 min (72°/2 min)
Cycling/time	72°/10 s	72°/2 min
Cycles	45	40

Table S2

a. Primers used for detection of the *ltx* promoter gene of *A. actinomycescomitans* and the *ftxA* gene of *F. alocis* in accordance to Poulsen et. al. [20] and Oscarsson et al. [15].

b. Cycle settings for detection of the *ltx promoter gene* of *A. actinomycescomitans* and the *ftxA* gene of *F. alocis* in accordance to Poulsen et. al. [20] and Oscarsson et al. [15].

a	Forward	Reverse
Poulsen <i>ltx- promoter</i>	GCCGACACCAAAGACAAAGTCT	GCCCATAACCAAGCCACATAC
Oscarsson <i>ftxA</i>	GGCTCAGATACCTACTTCTTC	GAAGGCTATGATTTGATTGTTTCC

b	Poulsen (<i>ltx promoter</i>)	Oscarsson (<i>ftxA</i>)
Hold/time	94°/5 min	95°/1 min
Cycling/time	94°/1 min	95°/30 s
Cycling/time	60°/1 min	54°/30 s
Cycling/time	72°/2 min	72°/1 min
Cycling/time	72°/8 min	72°/7 min
Cycles	30	35