Supplementary Table S1. Taxonomic composition of the subgingival microbiome.

Genus	Prevalence (# samples)	Adjusted p-value	Average relative abundance at the diseased state	Average relative abundance at the resolved state
Actinomyces	47	0.018	4.50%	8.82%
Fusobacterium	45	0.519	12.47%	10.94%
Capnocytophaga	44	0.177	4.53%	7.09%
Streptococcus	41	0.024	3.18%	7.86%
Prevotella	39	0.365	13.94%	10.81%
Porphyromonas	38	0.004	11.65%	2.17%
Campylobacter	38	0.546	1.92%	1.72%
Corynebacterium	37	0.095	3.48%	7.75%
Leptotrichia	37	0.929	4.11%	4.20%
Treponema	34	0.007	9.56%	2.92%
Selenomonas	33	0.519	2.77%	3.40%
Neisseria	29	0.365	2.61%	4.32%
Veillonella	24	0.193	1.84%	5.55%
Tannerella	22	0.002	2.16%	0.48%
Aggregatibacter	21	0.457	1.13%	1.55%
Dialister	16	0.046	1.37%	0.57%
Eubacterium	16	0.183	0.99%	0.74%
Rothia	15	0.010	1.28%	3.50%
Synergistes	14	0.001	1.89%	0.09%
Parvimonas	14	0.102	0.95%	0.60%
Gemella	14	0.374	0.70%	1.00%
Kingella	14	0.519	0.53%	0.68%
Haemophilus	12	0.098	0.47%	5.74%
Cardiobacterium	11	0.652	0.82%	0.67%
Filifactor	9	0.004	0.78%	0.05%
Catonella	8	0.365	0.61%	0.49%
Granulicatella	5	0.065	0.25%	0.61%
Actinobaculum	5	0.374	0.73%	0.24%
Propionibacterium	5	0.399	0.59%	0.34%
Bergeyella	4	0.023	0.18%	0.51%
Olsenella	4	0.033	0.34%	0.08%
Lautropia	4	0.116	0.10%	0.35%
Peptostreptococcus	3	0.024	0.47%	0.12%
Oribacterium	3	0.502	0.24%	0.39%
Mycoplasma	2	0.018	0.23%	0.00%
Peptococcus	2	0.497	0.35%	0.24%
Eikenella	2	0.624	0.25%	0.21%
Shuttleworthia	2	0.885	0.16%	0.18%