

Probe ID/Sample	Periodontitis subject #1	Periodontitis subject #2	Periodontitis subject #3	Periodontitis subject #4	Periodontitis subject #5, site 1	Periodontitis subject #5, site 2	Periodontitis subject #5, site 3	Periodontitis subject #5, site 4	severe periodontitis subject #6	severe periodontitis subject #7	severe periodontitis subject #8	severe periodontitis subject #9	severe periodontitis subject #10
16S Universal I_E29	5	5	5	5	5	5	5	5	5	5	5	5	5
Bacteroidetes sp AU126 I_K76	0	0	2	0	0	2	0	0	2	4	1	3	0
Bacteroidetes sp X083 II_X17	0	0	0	0	0	2	1	0	0	0	0	0	0
Capnocytophaga Cluster II_AA89	0	0	0	0	0	0	0	0	5	5	4	3	5
Sphaerocytophaga sp strain S3 II_X23	0	0	0	0	0	0	0	0	0	1	0	0	0
Capnocytophaga sputigena I_W46	0	0	0	0	0	0	0	0	5	3	4	1	2
Porphyromonas endodontalis IV_W78	0	1	0	0	0	2	3	2	4	5	5	0	0
Porphyromonas gingivalis I_X21	1	0	0	2	2	3	4	3	0	0	0	0	0
Porphyromonas catoniae_CW034 I_N	0	0	0	0	0	0	0	0	3	0	0	3	0
Prevotella Cluster VI_AA44	0	0	0	0	0	0	0	3	2	0	0	2	2
Prevotella Cluster I_Y64	0	0	0	0	0	0	0	0	3	1	3	3	4
Prevotella sp strain B31FD II_AB21	0	0	0	0	0	0	0	0	0	0	0	0	2
Prevotella denticola I_K73	0	0	0	2	0	0	0	0	0	0	0	1	0
Prevotella intermedia II_AB92	0	2	0	0	0	0	0	0	3	0	0	4	0
Prevotella Cluster II_Y65	0	0	0	0	0	0	0	0	2	2	2	2	2
Prevotella melaninogenica_BE073 I_T	0	0	0	0	0	0	0	0	0	0	0	2	3
Prevotella nigrescens I_W40	0	0	0	0	0	0	0	0	0	0	3	2	2
Prevotella oralis I_X76	0	0	0	0	0	0	0	0	2	0	0	0	0
Prevotella oris F045_Y66	0	0	0	0	0	0	0	4	1	2	0	2	3
Prevotella oulora I_AA52	0	0	0	2	0	0	0	4	0	0	0	0	0
Prevotella pallens_DR022 I_R67	0	0	0	2	0	0	0	0	0	0	0	0	0
Prevotella sp BE073_FO012_AA53	0	0	0	0	0	0	0	0	2	2	2	0	0
Prevotella sp BI027 I_X15	0	0	0	0	0	0	0	1	0	0	0	0	0
Prevotella sp CY006_FL019_L10	0	0	0	0	0	0	0	0	0	0	0	1	0
Prevotella sp DR022_Y67	0	0	0	0	0	0	0	0	2	2	3	2	2
Prevotella sp DO027 I_Q99	0	0	0	0	0	0	0	0	2	2	0	1	1
Prevotella sp DO045 II_X59	0	0	0	0	0	0	0	1	0	1	0	1	1
Prevotella tanneriae II_AB03	0	0	0	0	0	0	0	0	0	2	0	0	0
Tannerella forsythensis I_X56	0	2	2	2	4	4	3	3	4	5	4	5	0
Tannerella sp BU063 I_T83	0	0	0	0	0	0	0	0	3	0	0	3	2
Aggregatibacter actinomycetemcomitans	0	0	0	0	0	0	0	0	2	2	5	2	0
Haemophilus parainfluenzae_aphroph	0	0	0	0	0	0	0	0	4	4	3	2	0
Aggregatibacter segnis I_P05	0	0	0	0	0	0	0	0	0	1	2	2	2
Haemophilus sp BJ095 I_AA97	0	0	0	0	0	0	0	0	0	2	1	2	0

Campylobacter concisus I_O46	0	0	0	0	0	0	0	0	0	2	4	1	
Campylobacter gracilis II_X34	0	0	0	2	0	0	1	3	3	4	5	5	3
Campylobacter rectus_concisus II_X3	4	4	4	3	4	5	5	3	5	5	5	5	5
Campylobacter Cluster I_T87	0	0	0	0	1	2	3	0	3	4	4	3	1
Campylobacter Cluster II_X37	0	0	0	0	2	3	3	2	4	4	5	3	2
Campylobacter showae II_X35	0	0	0	1	0	0	0	0	2	4	5	4	0
Negative Control I_AC32	0	0	0	0	0	0	0	0	0	0	0	0	0
Lautropia mirabilis II_X44	0	0	0	0	0	0	0	0	0	2	0	0	0
Neisseria Cluster III_O45	0	0	0	0	0	0	0	0	0	0	2	0	0
Neisseria flavescens I_AA76	2	0	0	0	0	0	0	0	0	0	0	0	0
Desulfobulbus sp R004 I_K70	2	2	0	0	2	0	0	0	2	0	0	0	0
Kingella denitrificans I_W30	0	0	0	0	0	0	0	0	0	0	0	0	2
Dialister invisus I_P73	0	2	0	0	1	4	2	0	0	0	0	2	0
Dialister pneumosintes I_X78	0	0	0	2	0	0	0	2	0	2	2	2	0
Selenomonas Cluster I_AC13	0	0	0	0	0	0	0	2	0	0	2	2	0
Selenomonas infelix I_O54	1	2	0	2	0	0	0	3	2	2	3	4	2
Selenomonas noxia III_AC04	0	0	0	0	0	0	0	2	0	0	1	2	1
Selenomonas sp CS002 I_P86	0	0	0	0	0	0	0	0	1	0	0	1	0
Selenomonas sp DD020 I_Y73	0	0	0	0	0	0	0	0	0	0	0	2	0
Selenomonas sp EW084_DS071 I_Q5	0	0	0	0	0	0	0	0	0	0	0	2	0
Selenomonas sp EZ011_DM071_W86	0	0	0	0	0	2	2	1	0	0	0	0	0
Selenomonas sputigena I_K65	0	0	0	2	0	0	0	2	2	1	2	2	2
Selenomonas strain GAA14 I_AB58	0	0	0	2	0	3	2	3	0	3	2	2	0
Veillonella Cluster II_Q67	0	0	0	2	0	0	0	0	4	2	3	5	5
Veillonella Cluster IV_AC37	0	0	0	0	0	0	0	0	2	2	2	3	4
Veillonella atypica II_W88	0	0	0	3	0	0	2	0	0	0	0	4	0
Veillonella parvula I_M04	0	0	0	0	0	0	0	0	0	0	0	2	2
Catonella morbi BR036 I_O56	0	0	0	2	1	1	1	2	2	3	2	3	0
Eubacterium brachy II_AC03	0	2	0	0	0	0	3	0	4	0	0	4	0
Eubacterium clone 7 69 I_AB31	0	0	0	0	0	0	0	0	0	0	0	2	0
Eubacterium nodatum I_Y46	0	0	0	1	4	4	4	5	2	0	0	2	0
Eubacterium PUS9170_MCE10174 II	0	0	0	0	2	2	2	0	0	0	0	2	0
Eubacterium saburreum_BE088 I_AB	0	0	0	0	0	0	0	0	0	0	0	2	2
Eubacterium saphenum I_Y48	2	1	0	1	3	3	2	2	0	0	0	4	0
Eubacterium sp BB124 I_AA65	0	0	0	0	0	0	0	0	0	2	2	0	0
Eubacterium sp BB142 I_K98	0	0	0	0	0	0	0	0	0	1	0	0	0
Eubacterium sp strain A3MT I_L01	0	0	0	0	0	0	0	0	0	3	0	0	0
Eubacterium yurii_A3MT I_W84	0	0	0	0	0	0	0	2	2	4	4	2	0
Filifactor alocis III_AB94	4	4	2	2	5	5	5	5	2	4	2	4	0
Gemella haemolysans_sanguis I_K63	0	0	0	0	0	0	0	0	5	4	2	2	3
Gemella morbillorum I_K64	1	2	0	2	0	0	0	1	5	4	2	5	4
Granulicatella adiacens_elegans II_W	0	0	0	0	0	0	0	0	0	1	0	2	1
Granulicatella elegans II_AB29	0	0	0	0	0	0	0	0	0	2	0	4	2
Mogibacterium tidmidum II_AB26	0	0	0	0	4	3	1	3	0	0	0	0	0
Peptostreptococcus micros I_L97	3	5	5	4	5	5	5	5	5	3	4	5	3
Peptostreptococcus stomatis_AA55	0	0	0	0	1	0	2	2	2	0	0	2	0
Peptostreptococcus sp DA014_P72	1	2	0	0	0	0	0	0	0	0	0	0	0
Pseudoramibacter alactolyticus II_AB7	0	0	0	0	3	4	2	3	0	0	0	0	0
Streptococcus Cluster III_Q65	1	1	2	2	1	2	3	2	5	5	5	5	5

Strep gordonii anginosus_X11	0	0	0	0	0	0	0	0	1	0	0	1	0
Streptococcus oralis_F46	0	0	0	0	0	0	0	0	5	5	4	4	4
Streptococcus Cluster II_Q59	0	0	0	2	0	0	2	0	3	4	3	4	4
Streptococcus sp C6_7A_N20	0	0	0	0	0	0	0	0	3	3	2	4	0
Streptococcus intermedius_constellatu	0	2	2	2	2	2	1	4	5	5	5	2	4
Streptococcus mitis biovar 2 I_Q64	0	0	0	0	0	0	0	0	3	1	1	4	3
Streptococcus Cluster I_AB98	0	0	0	0	0	0	0	0	3	3	0	2	0
Streptococcus anginosus_gordonii I_F	0	0	0	1	0	0	0	0	4	3	3	4	3
Streptococcus Cluster IV_Q62	0	1	3	2	0	0	0	1	5	5	5	5	5
Streptococcus anginosus I_AB84	0	0	0	0	0	0	0	0	1	3	1	4	2
Streptococcus anginosus_intermedius	0	0	1	0	0	0	0	0	4	3	4	3	3
Streptococcus australis T1 E5 II_AB83	0	0	0	0	0	0	0	0	1	2	0	1	2
Streptococcus constellatus I_AB81	0	0	0	0	0	0	0	0	2	1	0	0	0
Streptococcus intermedius_constellatu	0	2	2	2	2	2	1	4	5	5	5	2	4
Streptococcus cristatus_BM053 II_AA	0	0	0	0	0	0	0	0	2	0	0	4	4
Streptococcus mutans I_X04	0	0	0	0	0	0	0	0	0	1	0	0	2
Streptococcus parasanguinis II_AB05	0	0	0	0	0	0	1	0	2	2	2	2	2
Streptococcus salivarius_FO042 I_E3	0	0	0	0	0	0	3	0	0	2	0	0	5
Streptococcus sanguinis II_AB78	0	0	0	0	0	0	1	0	3	1	4	1	2
Streptococcus infantis FN042_Y74	0	0	0	0	0	0	0	0	2	0	0	1	0
Synergistes Cluster I_D70	0	0	0	0	2	2	2	1	0	0	0	2	0
Synergistes sp BA121 genomosp C1	0	0	0	0	0	0	1	1	0	0	0	1	0
Synergistes sp BH007 I_AC53	0	0	0	0	1	1	1	0	0	0	0	0	0
Synergistes Cluster II_AC56	0	0	0	0	1	0	0	0	0	0	0	0	0
Fusobacterium I035 I_K61	0	0	0	0	0	0	0	0	0	0	2	0	0
Fusobacterium nucleatum ss polymor	0	0	0	1	0	0	0	2	2	0	5	3	3
Fusobacterium periodonticum II_R20	0	0	0	0	0	0	0	0	0	0	1	4	0
Leptotrichia Cluster I_AA45	0	0	0	2	0	0	0	0	2	1	2	3	2
Leptotrichia hofstadii_FAC5 II_AA58	0	0	0	0	0	0	0	0	0	0	1	1	0
Treponema Cluster I_AA63	0	0	0	0	0	0	0	0	0	1	0	1	0
Treponema maltophilum II_Q01	0	0	0	0	0	0	0	0	0	0	3	2	0
Treponema 5 All_AC41	0	0	0	0	0	0	0	0	0	0	0	1	0
Treponema parvum 5 Smibert 2 I_AC3	0	0	0	0	1	1	0	0	0	0	0	0	0
Treponema socranskii all ss I_Y95	0	0	0	0	0	0	0	0	1	1	1	1	0
Treponema sp AT039 AU076 F89	0	0	0	0	2	2	2	0	1	0	0	0	0
Treponema 5 19 D36ER1_AC40	0	0	0	0	0	0	0	0	0	0	0	2	0
TM7 sp AH040_BS003 I_W85	0	0	0	1	0	0	0	0	0	0	0	0	0
Actinomyces gerensceriae I_E46	0	0	0	0	0	0	0	0	0	0	2	1	2
Propionibacterium propionicum II_AB	0	0	0	0	0	0	0	0	2	2	2	1	2
Rothia dentocariosa_mucilaginoso I_E	0	0	0	0	0	0	0	2	0	1	2	0	0
Rothia mucilaginoso III_AB62	0	0	0	0	0	0	0	1	0	0	0	0	0