

pecies

SP134	Treponema sp._HMT_237	SP279	Prevotella lusca	SP6	Prevotella denticola
SP135	Porphyromonas gingivalis	SP28	Leptotrichia sp._HMT_212	SP80	Prevotella buccae
SP136	Capnocytophaga sp._HMT_863	SP280	Mogibacterium timidum	SP81	Porphyromonas endodontalis
SP137	Bacteroidetes_[G-3] bacterium_HMT_503	SP281	Slackia exigua	SP82	Schaalia odontolytica
SP138	Corynebacterium durum	SP282	Treponema maltophilum	SP83	Alloprevotella rava
SP139	Lachnospiraceae_[G-3] bacterium_HMT_100	SP283	Veillonellaceae_[G-1] bacterium_HMT_145	SP84	Scardovia wiggisiae
SP14	Leptotrichia buccalis	SP284	Capnocytophaga sp._HMT_412	SP85	Alloprevotella tanneriae
SP140	Arachnia rubra	SP285	Peptococcus sp._HMT_168	SP86	Porphyromonas pasteri
SP141	Porphyromonas sp._HMT_275	SP286	Prevotella sp._HMT_304	SP87	Selenomonas sp._HMT_138
SP142	Leptotrichia sp._HMT_417	SP287	Bacteroidetes_[G-5] bacterium_HMT_511	SP88	Parvimonas sp._HMT_110
SP143	Schaalia sp._HMT_877	SP288	Gracilibacteria_(GN02)_[G-1] bacterium_HMT_871	SP89	Actinomyces oris
SP144	Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349	SP29	Selenomonas infelix	SP9	Bacteroidales_[G-2] bacterium_HMT_274
SP145	Anaeroglobus geminatus	SP290	Capnocytophaga sp._HMT_338	SP90	Abiotrophia defectiva
SP147	Peptostreptococcaceae_[XII][G-6] [Eubacterium]_nodatum	SP291	Lachnoanaerobaculum sp._HMT_083	SP91	Actinomyces sp._HMT_171
SP148	Actinomyces dentalis	SP292	Pseudoleptotrichia goodfellowii	SP92	Shuttleworthia satelles
SP149	Leptotrichia trevisanii	SP293	Neisseria sp._HMT_018	SP93	Prevotella sp._HMT_300
SP15	Actinomyces johnsonii	SP294	Actinomyces sp._HMT_448	SP94	Campylobacter concisus
SP150	Aggregatibacter aphrophilus	SP295	Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870	SP95	Prevotella sp._HMT_292
SP151	Selenomonas sp._HMT_134	SP296	Bacteroidetes_[G-3] bacterium_HMT_365	SP96	Eikenella corrodens
SP152	Schaalia georgiae	SP297	Selenomonas sp._HMT_936	SP97	Lachnoanaerobaculum umeaense
SP153	Aggregatibacter sp._HMT_513	SP298	Capnocytophaga haemolytica	SP98	Stomatobaculum longum
SP154	Haemophilus haemolyticus	SP3	Streptococcus mutans	SP99	Selenomonas artemidis
SP155	Selenomonas sp._HMT_479	SP30	Gemella sanguinis	SPN100	Selenomonas noxia_nov_95.971%
SP156	Capnocytophaga sp._HMT_336	SP300	Atopobium sp._HMT_199	SPN101	Actinomyces dentalis_nov_97.683%
SP157	Fretibacterium sp._HMT_359	SP301	Streptococcus sp._HMT_056	SPN102	Abiotrophia defectiva_nov_97.026%
SP158	Anaerolineae_[G-1] bacterium_HMT_439	SP302	Mogibacterium neglectum	SPN103	Prevotella veroralis_nov_97.727%
SP159	Peptostreptococcaceae_[XII][G-5] [Eubacterium]_saphenum	SP303	Lautropia dentalis	SPN104	Actinomyces sp._HMT_448_nov_97.121%
SP16	Streptococcus anginosus	SP305	Neisseria cinerea	SPN105	Lacrimispora xylanolytica_nov_89.105%
SP160	Peptostreptococcaceae_[XII][G-9] [Eubacterium]_brachy	SP306	Prevotella pallens	SPN106	Lactonifactor longoviformis_nov_88.803%
SP161	Granulicatella adiacens	SP307	Mycoplasma salivarium	SPN108	Actinomyces israelii_nov_97.217%
SP162	Tannerella sp._HMT_286	SP31	Leptotrichia sp._HMT_498	SPN109	Wolinella succinogenes_nov_97.320%
SP163	Treponema sp._HMT_238	SP310	Treponema sp._HMT_256	SPN110	Bacteroidetes_[G-5] bacterium_HMT_505_nov_97.787%
SP164	Leptotrichia hofstadii	SP311	Selenomonas sp._HMT_481	SPN111	Flavonifractor plautii_nov_93.898%
SP165	Streptococcus intermedius	SP313	Aggregatibacter sp._HMT_898	SPN112	Treponema maltophilum_nov_97.656%
SP166	Schaalia sp._HMT_178	SP314	Selenomonas sp._HMT_133	SPN113	Porphyromonas catoniae_nov_97.148%
SP168	Olsenella sp._HMT_807	SP315	Prevotella sp._HMT_315	SPN114	Actinomyces dentalis_nov_97.876%
SP17	Haemophilus parainfluenzae	SP316	Tannerella sp._HMT_808	SPN115	Selenomonas sp._HMT_919_nov_97.901%
SP170	Peptostreptococcaceae_[XII][G-3] bacterium_HMT_495	SP317	Bifidobacterium dentium	SPN116	Prevotella oris_nov_96.395%
SP171	Capnocytophaga sp._HMT_332	SP318	Peptostreptococcaceae_[XII][G-4] bacterium_HMT_103	SPN117	Cardiobacterium hominis_nov_96.545%
SP172	Alloprevotella sp._HMT_912	SP319	Colibacter massiliensis	SPN119	Selenomonas sp._HMT_936_nov_96.756%
SP173	Mogibacterium vesicum	SP32	Fusobacterium nucleatum	SPN120	Cardiobacterium hominis_nov_96.750%
SP174	Actinomyces sp._HMT_897	SP320	Aggregatibacter sp._HMT_512	SPN13	Actinomyces israelii_nov_96.507%
SP175	Prevotella pleuritidis	SP321	Streptococcus australis	SPN130	Actinomyces sp._HMT_175_nov_97.633%
SP176	Treponema sp._HMT_257	SP322	Erysipelotrichaceae_[G-1] bacterium_HMT_905	SPN14	Actinomyces viscosus_nov_96.226%
SP177	Actinomyces sp._HMT_525	SP323	Streptococcus cristatus_clade_578	SPN141	Porphyromonas catoniae_nov_96.604%
SP178	Veillonellaceae_[G-1] bacterium_HMT_132	SP324	Treponema sp._HMT_231	SPN152	Capnocytophaga gingivalis_nov_97.665%
SP179	Gemella haemolysans	SP327	Streptococcus vestibularis	SPN158	Corynebacterium durum_nov_97.379%
SP18	Solobacterium moorei	SP328	Olsenella uli	SPN164	Prevotella oris_nov_96.893%
SP180	Clostridiales_[F-1][G-1] bacterium_HMT_093	SP329	Haemophilus parahaemolyticus	SPN176	Streptococcus sanguinis_nov_97.727%
SP181	Aggregatibacter segnis	SP33	Selenomonas noxia	SPN188	Sneathia sanguinegens_nov_91.287%
SP182	Ligilactobacillus salivarius	SP330	Veillonellaceae_[G-1] bacterium_HMT_129	SPN198	Treponema sp._HMT_927_nov_91.732%
SP183	Capnocytophaga ochracea	SP331	Leptotrichia sp._HMT_223	SPN2	Leptotrichia sp._HMT_215_nov_97.500%
SP184	Peptococcus sp._HMT_167	SP333	Tannerella sp._HMT_916	SPN210	Capnocytophaga gingivalis_nov_90.856%
SP185	Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347	SP334	Cryptobacterium curtum	SPN223	Prevotella sp._HMT_300_nov_97.835%
SP186	Actinomyces timonensis	SP335	Centipeda periodontii	SPN26	Leptotrichia hofstadii_nov_96.800%
SP187	Alloprevotella sp._HMT_308	SP336	Leptotrichia sp._HMT_219	SPN38	Actinomyces dentalis_nov_97.881%
SP188	Fretibacterium sp._HMT_362	SP338	Treponema vincentii	SPN43	Fusobacterium nucleatum_nov_97.813%
SP189	Peptoniphilaceae_[G-1] bacterium_HMT_113	SP339	Bulleidia extructa	SPN50	Capnocytophaga gingivalis_nov_95.914%
SP19	Capnocytophaga sputigena	SP34	Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957	SPN62	Prevotella veroralis_nov_96.388%
SP190	Mitsuokella sp._HMT_521	SP340	Prevotella baroniae	SPN66	Actinomyces weissii_nov_94.220%
SP191	Fusobacterium canifelinum	SP342	Capnocytophaga sp._HMT_901	SPN67	Corynebacterium matruchotii_nov_97.638%
SP192	Lachnoanaerobaculum saburreum	SP343	Desulfobulbus sp._HMT_041	SPN68	Selenomonas sp._HMT_137_nov_95.992%
SP193	Prevotella sp._HMT_526	SP344	Prevotella sp._HMT_472	SPN69	Actinomyces israelii_nov_94.881%
SP194	Bacteroidetes_[G-5] bacterium_HMT_505	SP347	Catonella sp._HMT_451	SPN70	Actinomyces naeslundii_nov_97.606%
SP195	Rothia dentocariosa	SP348	Treponema sp._HMT_239	SPN71	Actinomyces viscosus_nov_96.226%
SP196	Neisseria bacilliformis	SP349	Leptotrichia shahii	SPN72	Corynebacterium matruchotii_nov_97.642%
SP197	Dialister sp._HMT_119	SP35	Leptotrichia wadei	SPN73	Selenomonas sp._HMT_137_nov_97.137%
SP198	Stomatobaculum sp._HMT_097	SP351	Peptostreptococcaceae_[XIII][G-6] [Eubacterium]_minutum	SPN74	Peptidobacterium cupriacola_nov_96.571%