

Species

SP116	Leptotrichia wadei	SP217	Streptococcus anginosus	SP74	Peptoanaerobacter [Eubacterium] yurii
SP117	Actinomyces sp._HMT_896	SP22	Stomatobaculum sp._HMT_097	SP75	Porphyromonas sp._HMT_930
SP118	Prevotella loeschii	SP220	Porphyromonas endodontalis	SP76	Peptostreptococcaceae_XI[G-9] [Eubacterium]_brachy
SP119	Mogibacterium vesicum	SP222	Streptococcus infantis_clade_431	SP77	Tannerella sp._HMT_286
SP12	Selenomonas noxia	SP223	Lactobacillus vaginalis	SP78	Campylobacter showae
SP120	Capnocytophaga sp._HMT_901	SP224	Alloprevotella sp._HMT_912	SP79	Prevotella saccharolytica
SP121	Schaalia georgiae	SP225	Saccharibacteria_(TM7)_ [G-6] bacterium_HMT_870	SP8	Porphyromonas pasteri
SP122	Streptococcus sobrinus	SP226	Prevotella vespertina	SP80	Actinomyces sp._HMT_175
SP123	Fusobacterium sp._HMT_203	SP227	Selenomonas sp._HMT_920	SP81	Treponema socranskii
SP124	Streptococcus australis	SP228	Peptostreptococcus stomatis	SP82	Treponema sp._HMT_231
SP125	Selenomonas flueggei	SP229	Streptococcus downii	SP83	Actinomyces massiliensis
SP126	Leptotrichia sp._HMT_219	SP23	Bergeyella sp._HMT_322	SP84	Gemella haemolysans
SP127	Absconditabacteria_(SR1)_ [G-1] bacterium_HMT_345	SP230	Johnsonella sp._HMT_166	SP85	Actinomyces sp._HMT_897
SP128	Capnocytophaga sputigena	SP231	Haemophilus sp._HMT_908	SP86	Aggregatibacter sp._HMT_458
SP129	Selenomonas sputigena	SP234	Actinomyces sp._HMT_448	SP87	Actinomyces dentalis
SP13	Prevotella sp._HMT_317	SP236	Gemella sanguinis	SP88	Propionibacterium acidifaciens
SP130	Porphyromonas sp._HMT_278	SP237	Johnsonella ignava	SP89	Leptotrichia sp._HMT_225
SP131	Aggregatibacter sp._HMT_512	SP238	Lachnospiraceae_ [G-2] bacterium_HMT_088	SP9	Veillonella parvula
SP132	Rothia dentocariosa	SP239	Olsenella sp._HMT_807	SP90	Lachnoanaerobaculum saburreum
SP133	Campylobacter gracilis	SP24	Aggregatibacter sp._HMT_513	SP91	Selenomonas artemidis
SP134	Schaalia odontolytica	SP241	Saccharibacteria_(TM7)_ [G-8] bacterium_HMT_955	SP92	Streptococcus salivarius
SP135	Peptidiphaga sp._HMT_183	SP242	Porphyromonas sp._HMT_284	SP93	Bergeyella sp._HMT_900
SP136	Fusobacterium nucleatum_subsp._vincentii	SP243	Absconditabacteria_(SR1)_ [G-1] bacterium_HMT_875	SP97	Selenomonas sp._HMT_892
SP137	Saccharibacteria_(TM7)_ [G-3] bacterium_HMT_351	SP244	Capnocytophaga sp._HMT_335	SP98	Granulicatella elegans
SP138	Selenomonas sp._HMT_138	SP246	Ruminococcaceae_ [G-2] bacterium_HMT_085	SP99	Streptococcus sp._HMT_066
SP139	Tannerella serpentiformis	SP248	Haemophilus sputorum	SPN1	Streptococcus sanguinis_nov_97.727%
SP14	Actinomyces sp._HMT_169	SP249	Haemophilus haemolyticus	SPN100	Selenomonas noxia_nov_97.623%
SP140	Saccharibacteria_(TM7)_ [G-1] bacterium_HMT_957	SP25	Streptococcus cristatus	SPN101	Selenomonas artemidis_nov_97.137%
SP141	Saccharibacteria_(TM7)_ [G-1] bacterium_HMT_347	SP250	Alloprevotella sp._HMT_913	SPN102	Streptococcus gordonii_nov_96.846%
SP142	Scardovia wiggisiae	SP252	Leptotrichia sp._HMT_221	SPN103	Corynebacterium matruchotii_nov_97.835%
SP143	Neisseria perflava	SP253	Megasphaera micronuciformis	SPN104	Actinomyces naeslundii_nov_97.238%
SP144	Parvimonas sp._HMT_110	SP254	Selenomonas dianae	SPN106	Leptotrichia wadei_nov_97.595%
SP145	Actinomyces sp._HMT_414	SP255	Bergeyella sp._HMT_206	SPN115	Cardiobacterium hominis_nov_96.161%
SP146	Limosilactobacillus fermentum	SP257	Alloprevotella sp._HMT_914	SPN13	Corynebacterium matruchotii_nov_97.830%
SP147	Selenomonas sp._HMT_137	SP258	Capnocytophaga sp._HMT_863	SPN34	Actinomyces sp._HMT_175_nov_97.633%
SP148	Capnocytophaga sp._HMT_332	SP259	Alloprevotella sp._HMT_308	SPN41	Corynebacterium durum_nov_97.984%
SP149	Lautropia mirabilis	SP26	Streptococcus mitis	SPN46	Actinomyces viscosus_nov_97.159%
SP15	Leptotrichia hongkongensis	SP262	Lancefieldella rimae	SPN52	Kingella oralis_nov_97.826%
SP150	Neisseria elongata	SP263	Rothia mucilaginosa	SPN57	Lacrimispora xylanolytica_nov_88.911%
SP151	Prevotella histicola	SP264	Gracilibacteria_(GN02)_ [G-1] bacterium_HMT_871	SPN61	Selenomonas noxia_nov_96.691%
SP152	Capnocytophaga granulosa	SP266	Neisseria cinerea	SPN63	Corynebacterium matruchotii_nov_97.847%
SP153	Actinomyces gerencseriae	SP267	Bifidobacterium dentium	SPN64	Selenomonas noxia_nov_97.258%
SP154	Capnocytophaga sp._HMT_864	SP27	Actinomyces sp._HMT_525	SPN65	Capnocytophaga gingivalis_nov_97.276%
SP155	Prevotella oulorum	SP277	Selenomonas sp._HMT_136	SPN66	Selenomonas noxia_nov_95.971%
SP156	Schaalia sp._HMT_180	SP278	Veillonella atypica	SPN67	Capnocytophaga gingivalis_nov_95.914%
SP157	Prevotella sp._HMT_314	SP28	Lachnospiraceae_ [G-3] bacterium_HMT_100	SPN68	Fusobacterium nucleatum_nov_97.018%
SP158	Prevotella sp._HMT_300	SP281	Lactiplantibacillus plantarum	SPN69	Porphyromonas sp._HMT_284_nov_97.633%
SP159	Rothia aeria	SP282	Selenomonas sp._HMT_481	SPN70	Moraxella oblonga_nov_93.812%
SP16	Fusobacterium nucleatum	SP283	Selenomonas sp._HMT_146	SPN71	Streptococcus oralis_nov_96.226%
SP160	Schaalia meyeri	SP284	Streptococcus gwangjuense	SPN72	Capnocytophaga gingivalis_nov_91.051%
SP161	Pseudoleptotrichia goodfellowii	SP289	Lancefieldella parvula	SPN74	Actinomyces israelii_nov_97.584%
SP162	Parvimonas micra	SP29	Arachnia rubra	SPN75	Actinomyces oris_nov_97.778%
SP163	Leptotrichia sp._HMT_215	SP3	Lachnoanaerobaculum gingivalis	SPN76	Porphyromonas catoniae_nov_97.529%
SP164	Prevotella sp._HMT_475	SP30	Kingella denitrificans	SPN77	Selenomonas sp._HMT_137_nov_97.710%
SP165	Lactobacillus gasserii	SP304	Cutibacterium acnes	SPN78	Porphyromonas catoniae_nov_97.529%
SP166	Solobacterium moorei	SP31	Saccharibacteria_(TM7)_ [G-1] bacterium_HMT_346	SPN79	Prevotella veroralis_nov_96.780%
SP167	Actinomyces sp._HMT_170	SP32	Actinomyces johnsonii	SPN80	Porphyromonas catoniae_nov_97.719%
SP168	Ruminococcaceae_ [G-1] bacterium_HMT_075	SP33	Cardiobacterium valvarum	SPN81	Aggregatibacter sp._HMT_949_nov_96.457%
SP169	Streptococcus intermedius	SP34	Capnocytophaga sp._HMT_336	SPN82	Selenomonas sp._HMT_920_nov_97.909%
SP17	Leptotrichia sp._HMT_417	SP35	Capnocytophaga sp._HMT_324	SPN83	Actinomyces viscosus_nov_96.212%
SP170	Prevotella sp._HMT_472	SP36	Neisseria mucosa	SPN85	Actinomyces sp._HMT_175_nov_97.847%
SP171	Streptococcus parasanguinis	SP37	Actinomyces oris	SPN86	Porphyromonas catoniae_nov_97.533%
SP172	Actinomyces sp._HMT_171	SP38	Leptotrichia sp._HMT_392	SPN87	Schaalia georgiae_nov_97.471%
SP173	Streptococcus oralis_subsp._tigurinus_clade_070	SP39	Neisseria sicca	SPN88	Leptotrichia sp._HMT_215_nov_97.280%
SP174	Actinomyces viscosus	SP4	Lachnoanaerobaculum umeaense	SPN89	Leptotrichia sp._HMT_392_nov_97.510%
SP176	Schaalia sp._HMT_178	SP40	Porphyromonas sp._HMT_277	SPN90	Selenomonas artemidis_nov_97.519%
SP177	Oribacterium sp._HMT_078	SP41	Leptotrichia buccalis	SPN91	Mollicutes_ [G-2] bacterium_MOT-188_nov_82.759%