



Description2  
 RC\_VP\_PP  
 PT\_VP\_PP



- Veillonellaceae\_[G-1] bacterium\_HMT\_145
- Actinomyces israelii
- Neisseria subflava
- Rothia mucilaginosa
- Campylobacter concisus
- Fusobacterium nucleatum\_subsp.\_animalis
- Micrococcus cohnii
- Neisseria oralis
- Prevotella loescheii
- Peptostreptococcaceae\_[XI][G-5] [Eubacterium]\_saphenum
- Mycoplasma faucium
- Pelomonas aquatica
- Prevotella multisaccharivorax
- Campylobacter gracilis
- Selenomonas noxia
- Peptostreptococcaceae\_[XI][G-9] [Eubacterium]\_brachy
- Prevotella pleuritidis
- Staphylococcus saprophyticus
- Streptococcus sanguinis
- Abiotrophia defectiva
- Porphyromonas pasteri
- Fusobacterium periodonticum
- Aggregatibacter aphrophilus
- Olsenella uli
- Neisseria elongata
- Treponema socranskii
- Bacteroidetes\_[G-3] bacterium\_HMT\_280
- Bacteroidaceae\_[G-1] bacterium\_HMT\_272
- Corynebacterium durum
- Tannerella sp.\_HMT\_808
- Kingella denitrificans
- Stutzerimonas stutzeri
- Actinomyces massiliensis
- Klebsiella pneumoniae
- Pyramidobacter piscolens
- Corynebacterium matruchotii
- Lachnospiraceae\_[G-7] bacterium\_HMT\_086
- Veillonellaceae\_[G-1] bacterium\_HMT\_155
- Parascardovia denticolens
- Stomatobaculum sp.\_HMT\_373
- Peptococcus sp.\_HMT\_167
- Oribacterium sp.\_HMT\_078
- Micrococcus aloeverae
- Peptostreptococcaceae\_[XI][G-3] bacterium\_HMT\_495
- Desulfovibrio desulfuricans
- Lachnospiraceae\_[G-8] bacterium\_HMT\_500
- Enterococcus faecalis
- Olsenella profusa
- Veillonella parvula
- Pseudomonas putida
- Olsenella sp.\_HMT\_807
- Stenotrophobacter terrae
- Treponema sp.\_HMT\_254
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_869
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_346
- Peptostreptococcaceae\_[XI][G-6] [Eubacterium]\_nodatum
- Veillonellaceae\_[G-1] bacterium\_HMT\_135
- Alloprevotella tannerae
- Parvimonas sp.\_HMT\_110
- Shuttleworthia satelles
- Peptostreptococcaceae\_[XI][G-6] [Eubacterium]\_minutum
- Capnocytophaga sp.\_HMT\_338
- Porphyromonas gingivalis
- Streptococcus mutans
- Bacteroidetes\_[G-5] bacterium\_HMT\_511
- Atopobium sp.\_HMT\_199
- Sphingomonas hankookensis
- Peptoniphilaceae\_[G-1] bacterium\_HMT\_113
- Haemophilus parainfluenzae
- Treponema sp.\_HMT\_951
- Johnsonella ignava
- Atopobium sp.\_HMT\_416
- Fretibacterium sp.\_HMT\_360
- Actinomyces sp.\_HMT\_169
- Aggregatibacter sp.\_HMT\_458
- Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_349
- Bacteroidetes\_[G-3] bacterium\_HMT\_281
- Peptostreptococcaceae\_[XI][G-7] bacterium\_HMT\_081
- Capnocytophaga granulosa
- Pseudomonas aeruginosa
- Oribacterium asaccharolyticum
- Pseudomonas koreensis
- Treponema sp.\_HMT\_257
- Treponema sp.\_HMT\_270
- Erysipelotrichaceae\_[G-1] bacterium\_HMT\_904
- Lachnoanaerobaculum saburreum
- Veillonella dispar
- Mogibacterium timidum
- Lautropia mirabilis
- Peptostreptococcaceae\_[XI][G-4] bacterium\_HMT\_369
- Pseudoramibacter alactolyticus
- Paracoccus speluncae
- Parvimonas micra
- Peptostreptococcaceae\_[XI][G-4] bacterium\_HMT\_103
- Desulfohalobium japonicum\_nov\_86.848%
- Parvimonas micra\_nov\_95.426%
- Peptostreptococcaceae\_[XI][G-9] [Eubacterium]\_brachy\_nov\_96.05%
- Peptostreptococcaceae\_[XI][G-6] [Eubacterium]\_nodatum\_nov\_95.52%
- Parvimonas micra\_nov\_93.971%
- Aridibacter famidurans\_nov\_96.552%

Species

Samples