

Species

- SP1 *Staphylococcus warneri*
- SP10 *Listeria monocytogenes*
- SP11 *Fusobacterium nucleatum*
- SP12 *Prevotella oris*
- SP13 *Alloprevotella* sp._HMT_473
- SP14 *Leptotrichia* sp._HMT_212
- SP15 *Streptococcus mitis*
- SP16 *Streptococcus oralis*
- SP18 *Veillonella parvula*
- SP19 *Escherichia fergusonii*
- SP2 *Streptococcus sanguinis*
- SP20 *Shigella sonnei*
- SP21 *Schaalia odontolytica*
- SP22 *Lautropia mirabilis*
- SP23 *Streptococcus* sp._HMT_064
- SP24 *Streptococcus gordonii*
- SP25 *Bacillus halotolerans*
- SP26 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_957
- SP27 *Capnocytophaga gingivalis*
- SP28 *Truepera radiovictrix*
- SP29 *Limosilactobacillus fermentum*
- SP3 *Salmonella enterica*
- SP30 *Leptotrichia* sp._HMT_221
- SP31 *Pseudomonas aeruginosa*
- SP32 *Neisseria sicca*
- SP33 *Actinomyces* sp._HMT_448
- SP34 *Capnocytophaga leadbetteri*
- SP35 *Schaalia* sp._HMT_180
- SP36 *Enterococcus faecalis*
- SP37 *Corynebacterium matruchotii*
- SP38 *Campylobacter gracilis*
- SP39 *Actinomyces* sp._HMT_170
- SP40 *Porphyromonas gingivalis*
- SP41 *Actinomyces oris*
- SP42 *Streptococcus sobrinus*
- SP43 *Leptotrichia hongkongensis*
- SP44 *Streptococcus salivarius*
- SP45 *Prevotella melaninogenica*
- SP46 *Kingella oralis*
- SP47 *Neisseria cinerea*
- SP48 *Rothia dentocariosa*
- SP49 *Escherichia coli*
- SP50 *Actinomyces* sp._HMT_169
- SP51 *Leptotrichia wadei*
- SP52 *Streptococcus chosunense*
- SP53 *Leptotrichia* sp._HMT_392
- SP54 *Streptococcus intermedius*
- SP55 *Granulicatella adiacens*
- SP56 *Streptococcus oralis*_subsp._*tigurinus*_clade_070
- SP57 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_346
- SP58 *Streptococcus cristatus*
- SP59 *Ottowia* sp._HMT_894
- SP60 *Abiotrophia defectiva*
- SP61 *Streptococcus mutans*
- SP62 *Prevotella intermedia*
- SP63 *Schaalia meyeri*
- SP64 *Erythrobacter aureus*
- SP65 *Actinomyces viscosus*
- SP66 *Saccharibacteria*_(TM7)_[G-1] bacterium_HMT_348
- SP67 *Parvimonas micra*
- SP7 *Prevotella nigrescens*
- SP8 *Allobacillus halotolerans*
- SP9 *Neisseria subflava*
- SPN1 *Frondebacter mangrovi*_nov_92.771%
- SPN2 *Actinomyces* sp._HMT_175_nov_97.633%
- SPN3 *Saccharibacteria*_(TM7)_[G-6] bacterium_HMT_870_nov_97.093%
- SPN4 *Aeromicrobium panaciterrae*_nov_96.122%
- SPP1 *Veillonella dispar*_parvula
- SPP2 *Staphylococcus argenteus_aureus_roterodami*