



Group3

- Saliva Preterm Baby (+)
- Saliva Preterm Baby (-)
- Saliva On-term Baby (-)

- Enterococcus faecalis
- Staphylococcus hominis
- Corynebacterium coyleae
- Glutamicibacter soli
- Tepidiphilus succinatimandens
- Cutibacterium avidum
- Corynebacterium matruchotii
- Neisseria mucosa
- Escherichia coli
- Enterobacter hormaechei
- Bradyrhizobium valentinum
- Ligilactobacillus murinus
- Janibacter cremeus
- Staphylococcus caprae
- Klebsiella pneumoniae
- Streptococcus danieliae
- Varibaculum cambriense
- Staphylococcus capitis
- Corynebacterium tuberculoostearicum
- Fusobacterium nucleatum
- Facklamia languida
- Citrobacter murlinae
- Rothia mucilaginosa
- Actinomyces sp._HMT_448
- Pantoea alii
- Streptococcus parasanguinis_clade_411
- Streptococcus sp._HMT_066
- Methylophilus leisingeri
- Neisseriaceae_[G-1]_bacterium_HMT_174
- Cutibacterium acnes
- Bradyrhizobium lupini
- Enterococcus gallinarum
- Acinetobacter lwoffii
- Enterobacter cloacae
- Streptococcus mitis
- Prevotella oralis
- Corynebacterium durum
- Bifidobacterium pseudolongum
- Cutibacterium namnetense
- Microbacterium maritipicum
- Corynebacterium ureicelerivorans
- Corynebacterium aurimucosum
- Jeotgalicoccus halotolerans
- Corynebacterium jeikeium
- Triticum aestivum
- Bifidobacterium longum
- Faecalibaculum rodentium
- Bacillus subtilis
- Corynebacterium afermentans
- Salmonella enterica
- Peribacillus frigitolerans
- Rhodococcus qingshengii
- Streptococcus thoraltensis
- Streptococcus lactarius
- Acinetobacter pittii
- Actinidia eriantha
- Enterococcus casseliflavus
- Staphylococcus warneri
- Bradyrhizobium japonicum
- Staphylococcus pasteurii
- Enterobacter mori
- Tannerella serpentiformis
- Mammaliococcus sciuri
- Shigella flexneri
- Corynebacterium kroppenstedtii
- Corynebacterium pilbarensis
- Mammaliococcus lentus
- Mediterraneibacter faecis
- Escherichia fergusonii
- Bifidobacterium bifidum
- Streptococcus chosunense
- Enterobacter kobei
- Massilia arenae
- Lawsonella clevelandensis
- Sphingomonas leidyi
- Streptococcus oralis
- Streptococcus mutans
- Bacillus halotolerans
- Abiotrophia defectiva
- Staphylococcus saprophyticus
- Pseudoramibacter alactolyticus
- Corynebacterium pseudodiphtheriticum
- Lactiplantibacillus plantarum
- Prevotella sp._HMT_314
- Kocuria indica_nov_94.872%
- Corynebacterium argentoratense_nov_95.850%
- Daejeonella oryzae_nov_85.551%
- Thermodesulfobium acidiphilum_nov_81.729%
- Arthrosira platensis_nov_88.987%
- Sporosarcina sp._MOT-205_nov_96.360%
- Brevibacterium paucivorans_nov_97.368%
- Streptococcus danieliae_nov_95.594%
- Clostridium saccharoperbutylacetonicum_nov_95.800%
- Microbacterium saccharophilum_nov_76.981%
- Staphylococcus argenteus_aureus_roterodami
- Enterobacter_Leclercia adecarboxylata_cloacae
- Streptococcus infantis_infantis_clade_638
- Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro...(6 sp)
- Terrimonas multispecies_sppn3_2_nov_90.822%
- Gloeobacter multispecies_sppn4_2_nov_84.254%

Samples