



Relative abundance of a bacterial taxon of a bacterial taxon Min. Max.

OTU ID Taxonomic amotation Max.
326702 Faecalibacterium praussitzii 194745 Ruminococcus sp. 5 1 39BFAA 189827 Ruminococcus sp. 5 1 39BFAA 186234 Ruminococcus sp. 5 1 39BFAA 185951 Clostridiales sp. 212619 Ruminococcus sp. 5 1 39BFAA 185951 Clostridiales sp. 2126191 Ruminococcus torques 267610 Catenibacterium mitisuokai 361809 Ruminococcus torques 261912 Dorea formicigenerans 191687 Dorea formicigenerans 191687 Dorea formicigenerans 191687 Dorea formicigenerans 191687 Haemophilus parainfiluenzae 3126141 Clostridium sp. 148099 Weissella cibaria 470139 Clostridium ramosum 53785 Bifidobacterium sp. 15141 Lactobacillus mucosae 108747 Streptococcus thermophilus 561443 Streptococcus thermophilus 5786143 Streptococcus thermophilus 17899 Staphylococcus sp.