

## Species

- SP1 *Staphylococcus saprophyticus*
- SP11 *Ileibacterium valens*
- SP17 *Bacteroides thetaiotaomicron*
- SP18 *Akkermansia muciniphila*
- SP19 *Staphylococcus ureilyticus*
- SP2 *Ligilactobacillus murinus*
- SP20 *Bifidobacterium pseudolongum*
- SP26 *Sutterella* sp.\_str.\_cont1.66
- SP28 *Propionibacterium acnes*
- SP3 *Streptococcus species1\_MOT12*
- SP32 *Acutalibacter muris*
- SP33 *Acinetobacter johnsonii*
- SP4 *Escherichia coli*
- SP5 *Corynebacterium mastitidis*
- SP7 *Streptococcus thoraltensis*
- SP8 *Streptococcus danieliae*
- SPN106 *Duncaniella freteri*\_nov\_90.512%
- SPN107 *Dubosiella newyorkensis*\_nov\_87.451%
- SPN113 *Duncaniella freteri*\_nov\_91.255%
- SPN122 *Duncaniella freteri*\_nov\_83.146%
- SPN131 *Erysipelatoclostridium* sp.\_str.\_HGF2\_nov\_88.787%
- SPN137 *Duncaniella freteri*\_nov\_86.567%
- SPN153 *Duncaniella freteri*\_nov\_84.815%
- SPN154 *Duncaniella freteri*\_nov\_87.500%
- SPN155 *Duncaniella freteri*\_nov\_87.617%
- SPN156 *Duncaniella freteri*\_nov\_87.194%
- SPN157 *Lacnocyclus hithewayi*\_nov\_94.212%
- SPN159 *Alistipes finegoldii*\_nov\_92.789%
- SPN160 *Acetivibrio cellulolyticus*\_nov\_83.851%
- SPN161 *Eubacterium siraeum*\_nov\_91.252%
- SPN181 *Duncaniella freteri*\_nov\_91.667%
- SPN20 *Duncaniella freteri*\_nov\_88.476%
- SPN212 *Duncaniella freteri*\_nov\_86.989%
- SPN223 *Gabonia massiliensis*\_nov\_86.891%
- SPN244 *Alistipes senegalensis*\_nov\_93.846%
- SPN252 *Duncaniella freteri*\_nov\_93.974%
- SPN261 *Duncaniella freteri*\_nov\_94.151%
- SPN271 *Duncaniella freteri*\_nov\_93.585%
- SPN31 *Turicibacter sanguinis*\_nov\_95.635%
- SPN41 *Duncaniella freteri*\_nov\_88.743%
- SPN48 *Alistipes finegoldii*\_nov\_93.690%
- SPN56 *Duncaniella freteri*\_nov\_87.079%
- SPN70 *Duncaniella freteri*\_nov\_87.896%
- SPN77 *Duncaniella freteri*\_nov\_85.661%
- SPN84 *Alistipes senegalensis*\_nov\_94.073%
- SPN99 *Hathewayia proteolytica*\_nov\_83.768%
- SPP2 *Escherichia Shigella coli flexneri sonnei*
- SPP4 *Cutibacterium Propionibacterium acnes*
- SPPN9 *multigenus multispecies\_sppn9\_2\_nov\_92.263%*