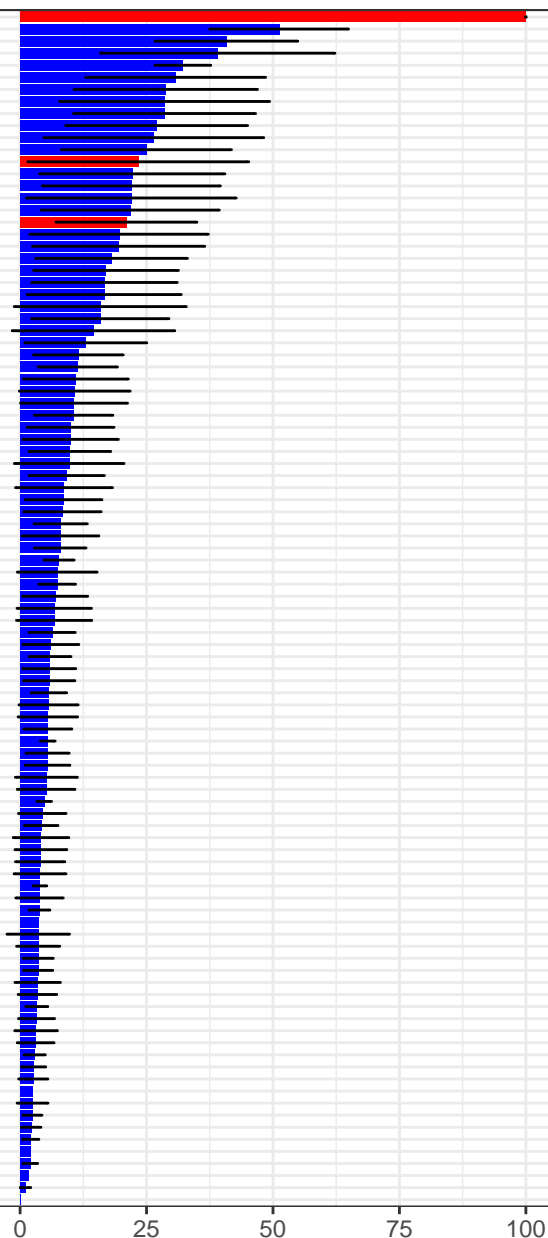


Feature Importance

Feature

* s_Veillonella parvula A
 * s_Dysosporobacterium sp000716993
 * s_Lachnospira foodsaie
 * s_Fusicatenibacter saccharovorans
 * s_Bacteroides uniformis
 * s_Streptococcus anginosus
 * s_Faecalibacterium prausnitzii
 * s_Ruminococcus Boudavv
 * s_Eisenbergiia sp000000000
 * s_Phocaeibacter plebeius
 * s_Erysipelatoclostridium faecis
 * s_Streptococcus faecalis
 * s_Veillonella nakazawae
 * s_Acetatifactor fressinai
 * s_Gemmiger sp000540777
 * s_Streptococcaceae
 * s_Bacteroides metaluminicoccus
 * s_Faecalibacterium prausnitzii
 * s_Faecalibacterium prausnitzii
 * s_Faecalibacterium prausnitzii
 * s_Bifidobacterium longum
 * s_Phacellobacterium faecis
 * s_Dorea formicigenerans
 * s_LMG5141 sp000000000
 * s_Christidium gressin
 * s_Phocaeibacter plebeius
 * s_Lachnospira lachnospira
 * s_Faecalibacterium prausnitzii
 * s_Roseburia inresinai
 * s_Copromonas sp000000000
 * s_Agathobacter faecis
 * s_Bacteroides xylanolyticus
 * s_Gemmiger fressinai
 * s_Bifidobacterium longum
 * s_Veillonellaceae
 * s_Mediterraneibacter faecis
 * s_Lachnospira eldens
 * s_Eubacterium g. veillonis
 * s_Bacteroides finegoldii
 * s_Alistipes finegoldii
 * s_Roseburia inresinai
 * s_Parabacteroides distasonis
 * s_UBA5201 sp000000000
 * s_Granulicatella sp000705833
 * s_KLE161 sp000000000
 * s_Faecalibacterium prausnitzii
 * s_Veillonella parvula
 * s_Copromonas sp000000000
 * s_Roseburia inresinai
 * s_Dysosporobacterium saccharovorans
 * s_Acetatifactor sp000000000
 * s_Agathobacter faecis
 * s_Odoribacter solanchoicus
 * s_Dorea A. longicatenata
 * s_Faecalibacterium prausnitzii
 * s_Phocaeibacter plebeius
 * s_Bacteroides cellulosilyticus
 * s_Bifidobacterium longum
 * s_Ruminococcus B. bromilii
 * s_Haemophilus B. bifidobacterium
 * s_Enterococcus sp000000000
 * s_Faecalibacterium prausnitzii



Increase

condition.control
 condition.disease

Mean Relative Importance