ns (-log(g-val)*sign(coeff)) Methanobacteria|D_3_Methanobacteriales|D_4_Methanobacteriaceae
Methanobacteria|D_3_Methanobacteriales|D_4_Methanobacteriaceae|D_5_Methanobrevibacter
Thermoplasmata|D_3_Methanomassiliicoccae|SD_4_Methanomethylophilaceae
Thermoplasmata|D_3_Methanomassiliicoccae|SD_4_Methanomethylophilaceae|D_5_Candidatus Methanogranum
Corriobacteria|D_3_Corriobacteriales|D_4_Eggerthellaceae|D_5_Senegalimassilia
Corriobacteria|D_3_Corriobacteriales|D_4_Eggerthellaceae|D_5_Incultured Euryarchaeota D Euryarchaeota D 0 Archaea D Describilitates Describilitate Archaea Eurýarchaeota Archaea Bacteria Bacteria Bacteria Bacteria 1 Bacteria D Bacteria D Bacteria 0_Bacteria D 0_Bacteria D Bacteria Bacteria D Bacteria D 0 Bacteria D 0 Bacteria D 0 Bacteria D BacterialD Bacteria D 1 Firmicutes D Christensenellaceae
Christensenellaceae|D 5_christensenellaceae R-7 group
Christensenellaceae|D 5_christensenellaceae|D 5_christensene|D 5_christensene|D 5_christensene|D 5_christensene|D 5_christen Bacteria 0 Bacteria D 1 Firmicutes D
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D-5 [Eubacterium] ruminantium group
D-5 Agathobacter
D-5 Anaerostipes
D-5 CAG-56
D-5 CAG-56
D-5 Coprococcus 1
D-5 Fusicatenibacter Firmicutes D Clostridiales D Clostridiales D 0_Bacteria 0_Bacteria Clostridia D Lachnospiraceae D Clostridia D Clostridia D Lachnosbiraceae Bacteria Firmicutes D Clostridiales Lachnospiraceae Clostridia D Clostridia D Clostridia D D_0_Bacteria D_0_Bacteria Firmicutes D Firmicutes D Clostridiales D Clostridiales D Clostridiales D Lachnospiraceae Lachnospiraceae Bacteria Firmicutes D Lachnospiraceae Clostridiales D Clostridiales D Clostridiales D 0 Bacteria 0 Bacteria Firmicutes D Firmicutes D Clostridia D Lachnospiraceae D_5_Lachnospiraceae UCG-003 Lachnospiraceae D_5_Oribacterium D Bacteria D - Firmicutes D 2 Clostridial D 3 Clostridiales D 4 Lachnospiraceae|D 5 Orbiacterium
D 0 Bacteria D 1 Firmicutes D 2 Clostridial D 3 Clostridiales D 4 Peptostreptococcaceae|D 5 Paeniclostridium
D 0 Bacteria D 1 Firmicutes D 2 Clostridial D 3 Clostridiales D 4 Peptostreptococcaceae|D 5 Paeniclostridium
D 0 Bacteria D 1 Firmicutes D 2 Clostridial D 3 Clostridiales D 4 Ruminococcaceae|D 5 Paeniclostridium
D 0 Bacteria D 1 Firmicutes D 2 Clostridial D 3 Clostridiales D 4 Ruminococcaceae|D 5 Paeniclostridium
D 0 Bacteria D 1 Firmicutes D 2 Clostridial D 3 Clostridiales D 4 Ruminococcaceae|D 5 Paeniclostridium
D 0 Bacteria D 1 Firmicutes D 2 Clostridiales D 4 Ruminococcaceae|D 5 Paeniclostridium
D 0 Bacteria D 1 Firmicutes D 2 Clostridiales D 4 Ruminococcaceae|D 5 Ruminoclostridium
D 0 Bacteria D 1 Firmicutes D 2 Clostridiales D 3 Clostridiales D 4 Ruminococcaceae|D 5 Ruminococcaceae|UCG-010
D 0 Bacteria D 1 Firmicutes D 2 Clostridiales D 3 Clostridiales D 4 Ruminococcaceae|D 5 Ruminococcaceae|UCG-013
D 0 Bacteria D 1 Firmicutes D 2 Erysipelotrichial D 3 Erysipelotrichales D 4 Erysipelotrichaee|D 5 Asteroleplasma
D 0 Bacteria D 1 Firmicutes D 2 Erysipelotrichial D 3 Erysipelotrichales D 4 Erysipelotrichaeeae|D 5 Erysipelotrichae Clostridia D Clostridia D Lacnnospiraceae|D_5_Vribacterium Peptostreptococcaceae|D_5_Paeniclostridium Peptostreptococcaceae|D_5_Romboutsia Ruminococcaceae Bacteria Firmicutes D