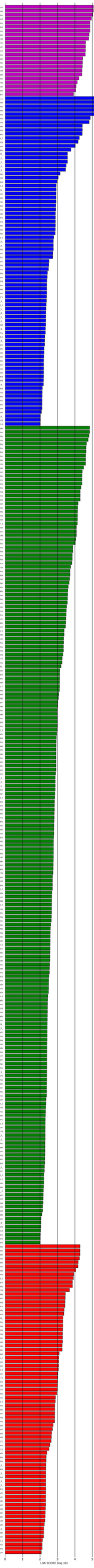
Significantly Represented Taxa


G_Streptococcus
F_Streptococcaceae
O_Lactobacillales
Ga_Tenui_855_sno_01_101
O_Enterobacterales
G_Serratia
F_Enterobacteriaceae
G_Citrobacter
Ga_Staphylococcus_sno_07_140
G_Brevibacterium
G_Brevibacterium
Ga_Corynebacterium_sno_01_153
F_Brevibacteriaceae
Pseudoalteromonadaceae
G_Alteromonadales
G_Pseudoalteromonas
Ga_Haemophilus_sno_05_549
Ga_Haemophilus_sno_01_180
G_Bacteroides
Ga_Bacteroides_sno_01_180
G_Actinomyces
F_Actinomycetaceae
Ga_Haemophilus_sno_04_565
sp_Haemophilus_sspp_01
G_Pseudomonas
F_Pseudomonadaceae
G_Pseudomonas
Pseudomonas_aeruginosa
F_Caenimonadaceae
F_Pasteurellaceae
F_Carnobacteriaceae
G_Aerococcus
Ga_Serratia_haemolyticum
Ga_Haemophilus_sspp_02
G_Bacteria
G_Rothia_mucilaginosa
G_Granulicatella_elegans
Ga_Eubacterium_TM7_
G_TM7_G_1
F_TM7_F_1
O_TM7_O_1
G_Selenomonas
O_Clostridiales
G_Lachnospiraceae_G_
F_Peptostreptococcaceae
G_Peptostreptococcus
G_Peptostreptococcus
F_Peptostreptococcaceae
G_Parvimonas
F_Synergistaceae
G_Fenestrimonas_micra
F_Aerococcaceae
G_Abiotrophia
G_Weissella_diffecticus
F_Synergistaceae
F_Cryptobacteriaceae
G_Sporobacter
Ga_Haemophilus_sspp_03
G_TM7_G_3
F_TM7_F_3
O_Lachnospiraceae
Ga_Lachnospiraceae_sno_01_101
Ga_Lachnospiraceae_sno_02_101
Ga_Lachnospiraceae_sno_03_101
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Ga_Lachnospiraceae_sno_87_101
F_DMR2_F_1
Ga_Ruminococcus_sno_01_101
F_DMR2_F_3
O_DMR2_O_1
G_Lactococcus_G_6
Ga_Lactococcus_sno_01_101
G_DMR2_G_1
G_Ruminococcus_G_6
Ga_Ruminococcus_sno_02_101
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