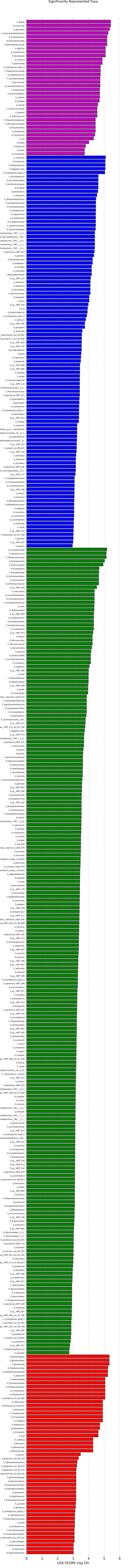
Significantly Represented Taxa
c_Bacilli
p_Firmicutes
D_Bacteria
c_Gammaproteobacteria
p_Proteobacteria
O_Enterobacteriales
f_Enterobacteriaceae
g_Bacillus
s_luteolarus
f_Bacillaceae
s_amylolyticus
g_Lactobacillus
s_Ruminococcus_gnavus
f_Staphylococcaceae
g_Staphylococcus
f_Lactobacillaceae
s_Ruminococcus_obeum
O_Unclassified(Bacteria)
f_Enterococcaceae
s_Holdemania_filiformis
g_Enterococcus
g_Oscillibacter
s_sporogenes
f_Peptostreptococcaceae
s_Clostridium_sporogenes
s_Clostridium_botulinum_B1_353
s_Clostridium_sporogenes_PA_3_679_352
s_Clostridium_botulinum_A2_Kyoto
s_Clostridium_botulinum_A_ATCC_19397
s_Clostridium_botulinum_A_Hall
s_Clostridium_botulinum_A3_Loch_Maree
s_Clostridium_botulinum_B_Eklund_17B
s_Clostridium_botulinum_F_Langeland
s_Clostridium_botulinum_A_ATCC_3502
s_Clostridium_botulinum_B1_Okra
s_Clostridium_botulinum_Ba4_657
s_Clostridium_botulinum_B_ATCC_BAA_1382
s_Clostridium_botulinum_A_ATCC_19397
s_Clostridium_botulinum_A_Hall
s_Clostridium_botulinum_A3_Loch_Maree
s_Clostridium_botulinum_B_Eklund_17B
s_Clostridium_botulinum_F_Langeland
s_Clostridium_botulinum_A_ATCC_3502
s_Clostridium_botulinum_B1_Okra
s_Clostridium_botulinum_Ba4_657
s_Clostridium_botulinum_B_ATCC_BAA_1382
s_Clostridium_botulinum_A2_Kyoto