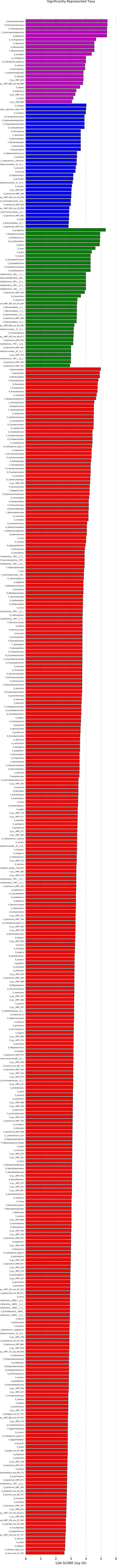
Significantly Represented Taxa
0_Enterobacteriales
F_Enterobacteriaceae
P_Proteobacteria
C_Gammaproteobacteria
O_Aeromonadales
S_Aeromonas
S_Aeromonas_sp_NR
S_Rahnella
S_Serratia
S_Pantoea
S_Enterobacter
S_Klebsiella
S_Citrobacter
S_Escherichia
S_Shigella
S_Salmonella
S_Yersinia
S_Photorhabdus
S_Pectobacterium
S_Erwinia
S_Dickeya
S_Pantoea_sp_At_9_b
S_Enterobacter_sp_638
S_Cronobacter
S_Enterobacter_cloacae_complex
S_Escherichia_coli
S_Shigella_flexneri_2a_str_301
S_Shigella_sonnei_Ss_046
S_Yersinia_pestis_CO_92_biovar_Orientalis
S_Yersinia_pestis_KIM10+
S_Yersinia_pestis_CO_92
S_Yersinia_pestis_KIM5_P12A
S_Yersinia_pestis_Antiqua
S_Yersinia_pestis_Nepal516
S_Yersinia_pestis_Pestoides_F
S_Yersinia_pestis_Angola
S_Yersinia_pestis_Medievalis_Harbin_35
S_Yersinia_pestis_Pestoides_A
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S_Yersinia_pestis_CO_92_plasmid_pCD1
S_Yersinia_pestis_KIM5_P12A_plasmid_pCD1
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LDA SCORE (log 10)