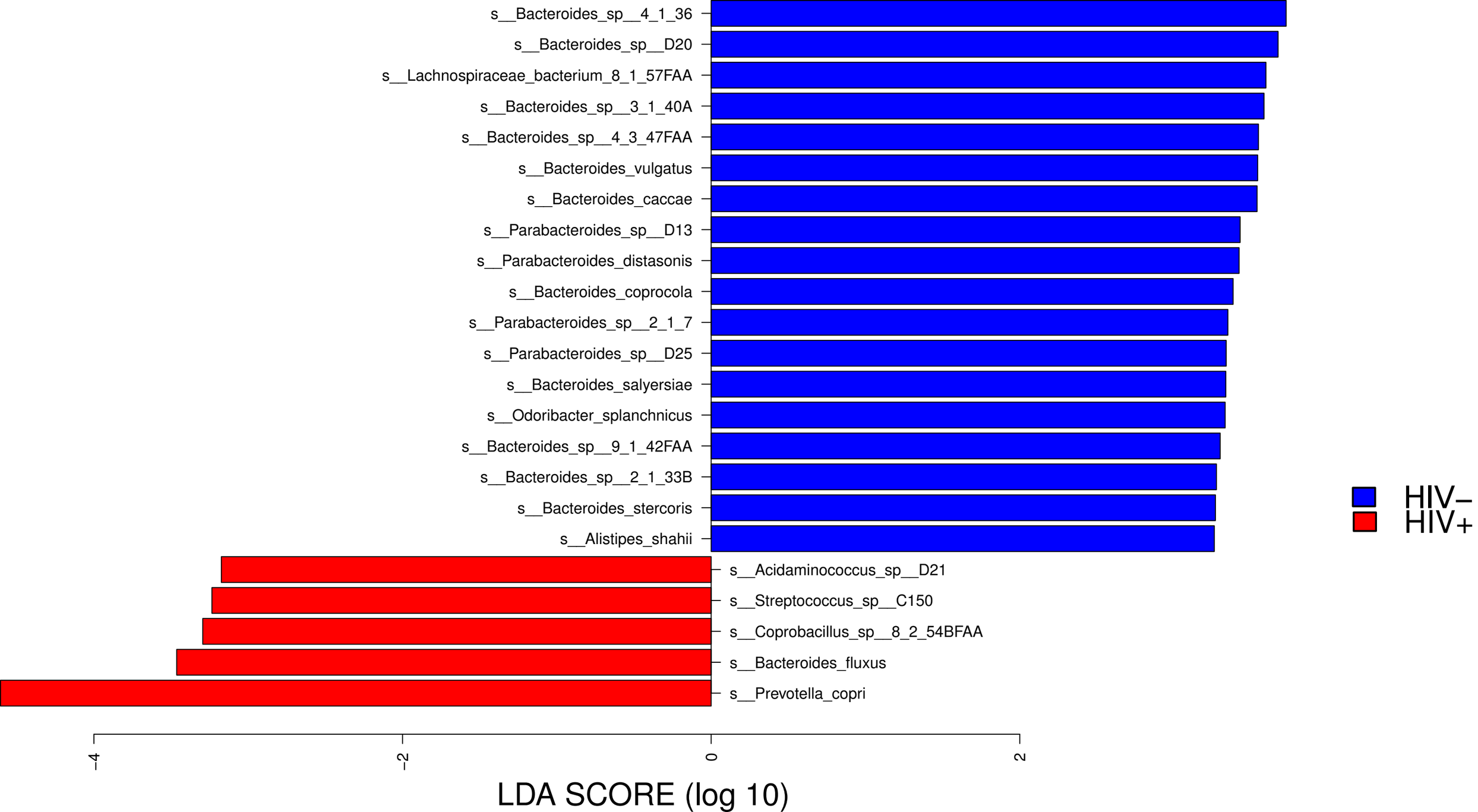
s__Bacteroides_sp__4_1_36
s__Bacteroides_sp__D20
s__Lachnospiraceae_bacterium_8_1_57FAA
s__Bacteroides_sp__3_1_40A
s__Bacteroides_sp__4_3_47FAA
s__Bacteroides_vulgatus
s__Bacteroides_caccae
s__Parabacteroides_sp__D13
s__Parabacteroides_distasonis
s__Bacteroides_coprocola
s__Parabacteroides_sp__2_1_7
s__Parabacteroides_sp__D25
s__Bacteroides_salyersiae
s__Odoribacter_splanchnicus
s__Bacteroides_sp__9_1_42FAA
s__Bacteroides_sp__2_1_33B
s__Bacteroides_stercoris
s__Alistipes_shahii
s__Acidaminococcus_sp__D21
s__Streptococcus_sp__C150
s__Coprobacillus_sp__8_2_54BFAA
s__Bacteroides_fluxus
s__Prevotella_copri
HIV−
HIV+
−4
−2
0
2
LDA SCORE (log 10)