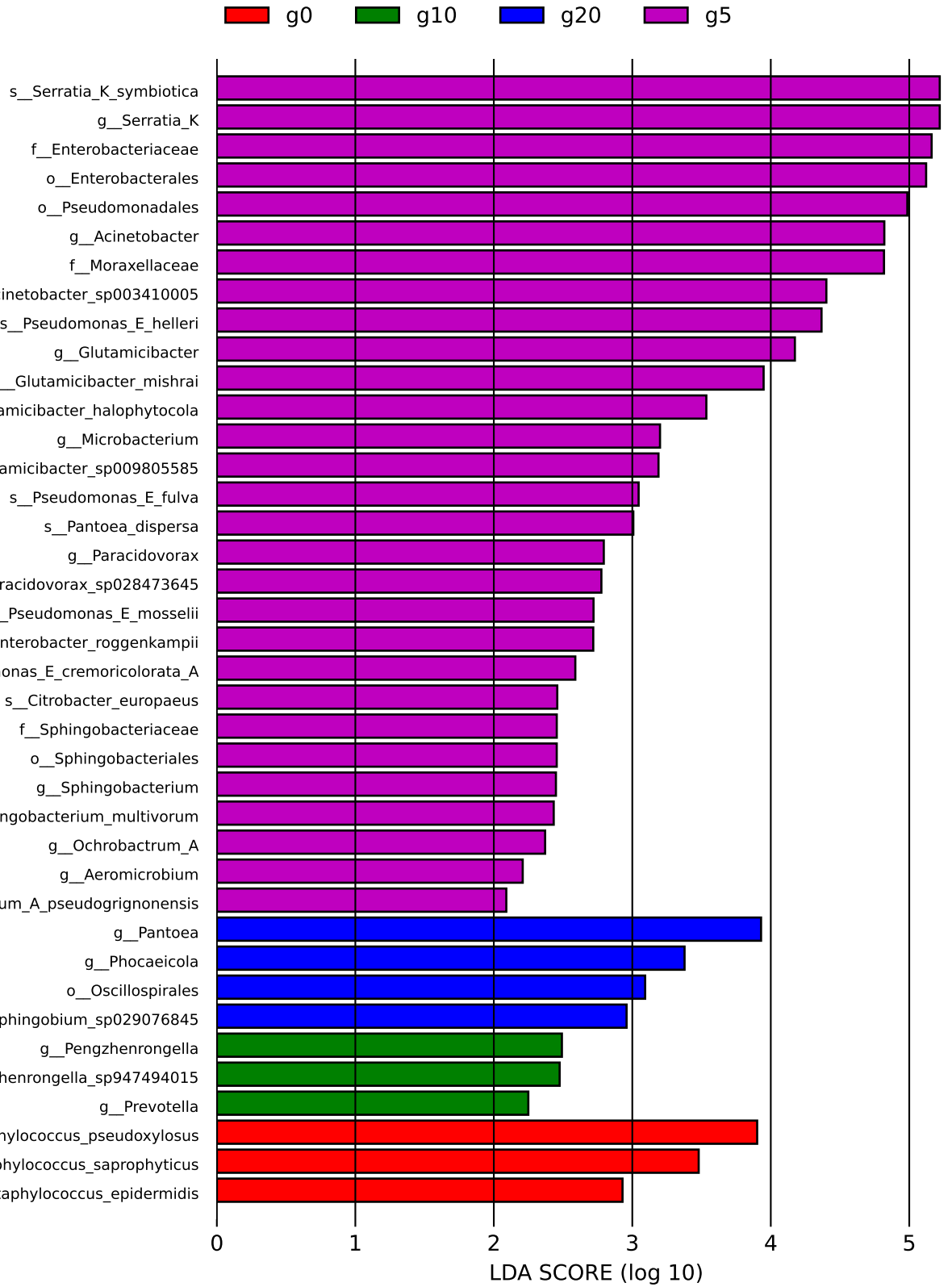
g0
g10
g20
g5
s__Serratia_K_symbiotica
g__Serratia_K
f__Enterobacteriaceae
o__Enterobacterales
o__Pseudomonadales
g__Acinetobacter
f__Moraxellaceae
inetobacter_sp003410005
s__Pseudomonas_E_helleri
g__Glutamicibacter
__Glutamicibacter_mishrai
amicibacter_halophytocola
g__Microbacterium
amicibacter_sp009805585
s__Pseudomonas_E_fulva
s__Pantoea_dispersa
g__Paracidovorax
racidovorax_sp028473645
_Pseudomonas_E_mosselii
nterobacter_roggenkampii
onas_E_cremoricolorata_A
s__Citrobacter_europaeus
f__Sphingobacteriaceae
o__Sphingobacteriales
g__Sphingobacterium
ngobacterium_multivorum
g__Ochrobactrum_A
g__Aeromicrobium
um_A_pseudogrignonensis
g__Pantoea
g__Phocaeicola
o__Oscillospirales
phingobium_sp029076845
g__Pengzhenrongella
henrongella_sp947494015
g__Prevotella
hylococcus_pseudoxylosus
phylococcus_saprophyticus
taphylococcus_epidermidis
0
1
2
3
4
5
LDA SCORE (log 10)