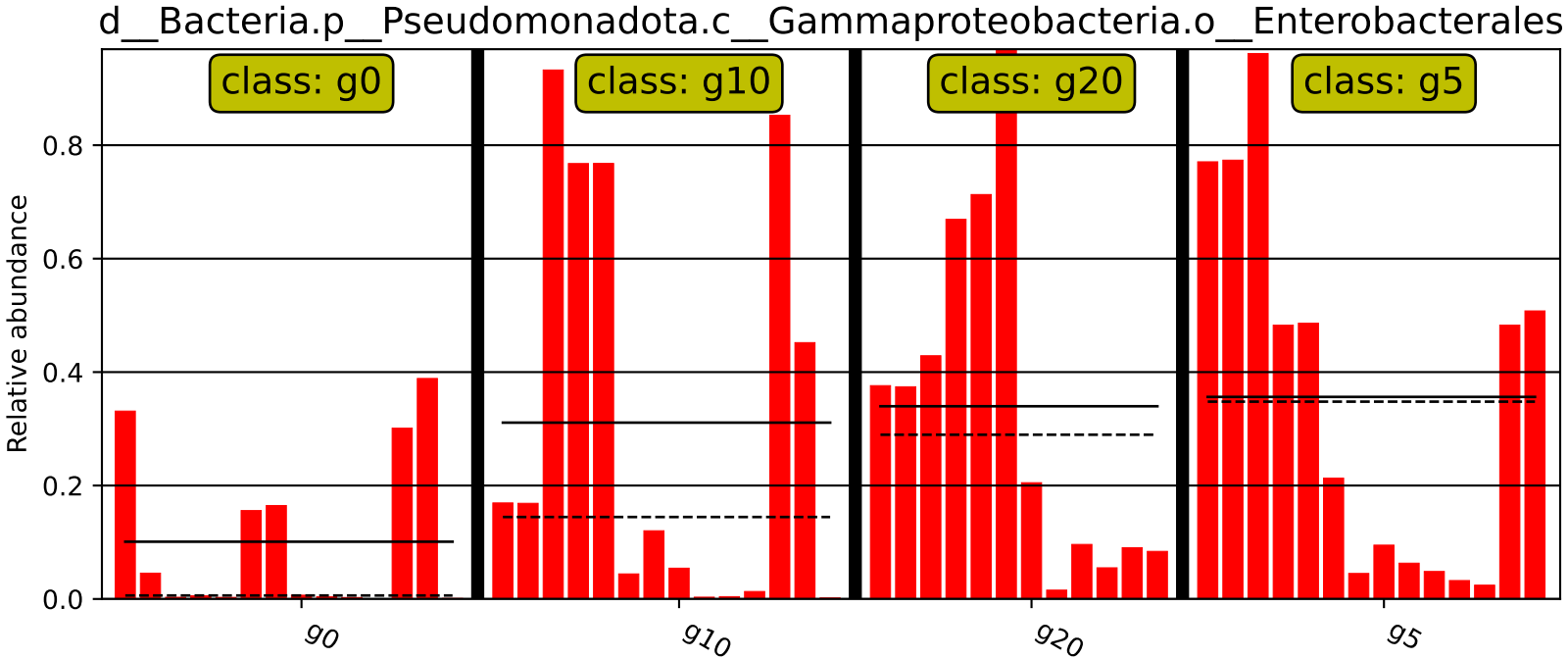
d__Bacteria.p__Pseudomonadota.c__Gammaproteobacteria.o__Enterobacterales
class: g0
class: g10
class: g20
class: g5
Relative abundance
0.0
0.2
0.4
0.6
0.8
g0
g10
g20
g5