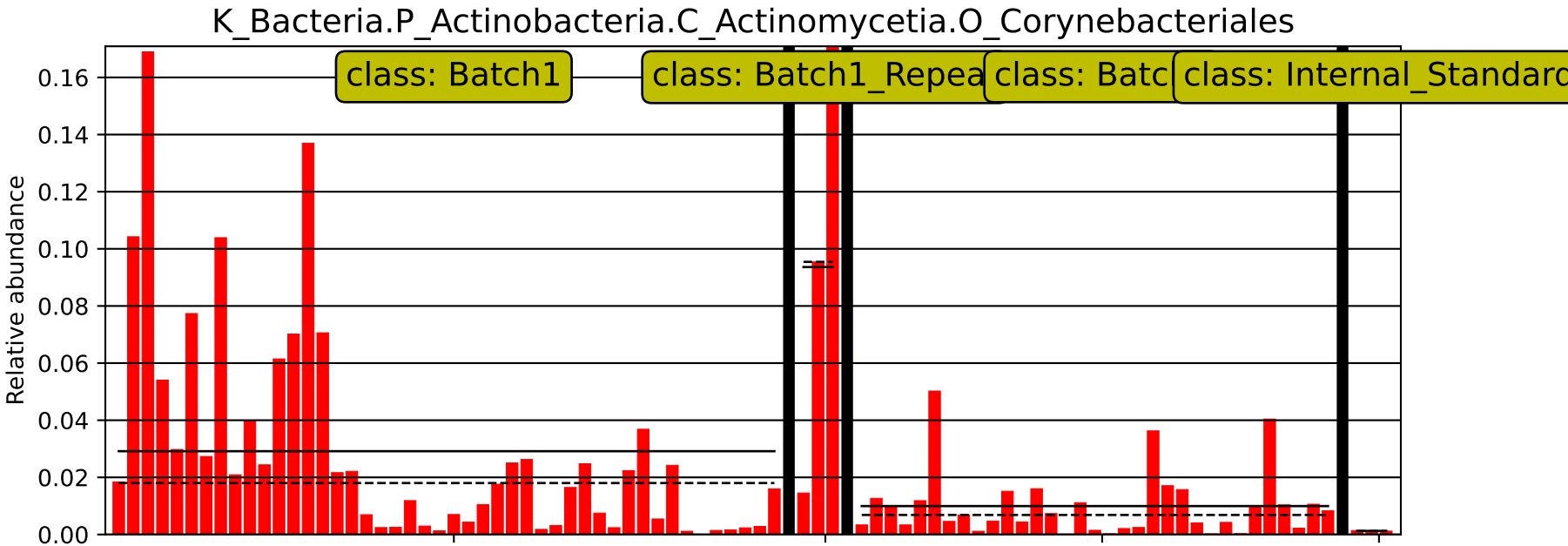
K_Bacteria.P_Actinobacteria.C_Actinomycetia.O_Corynebacteriales
class: Batch1
class: Batch1_Repea
class: Batc
class: Internal_Standard
Relative abundance
0.16
0.14
0.12
0.10
0.08
0.06
0.04
0.02
0.00