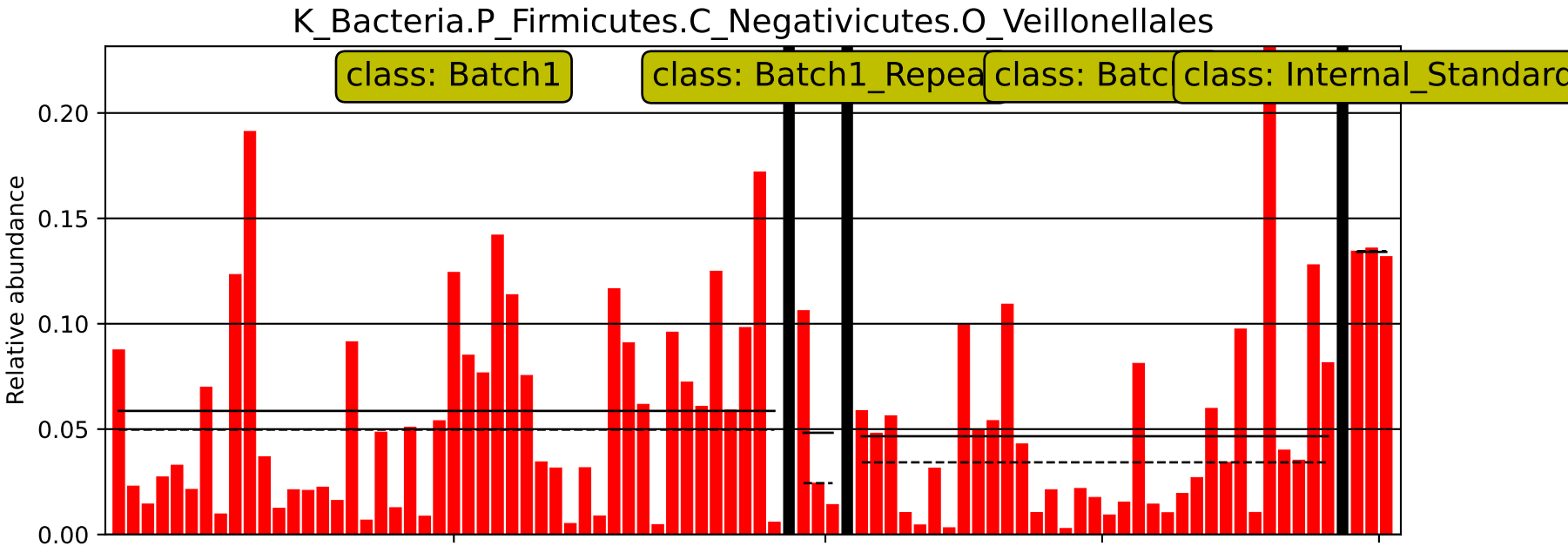
K_Bacteria.P_Firmicutes.C_Negativicutes.O_Veillonellales
class: Batch1
class: Batch1_Repea
class: Batc
class: Internal_Standard
Relative abundance
0.20
0.15
0.10
0.05
0.00