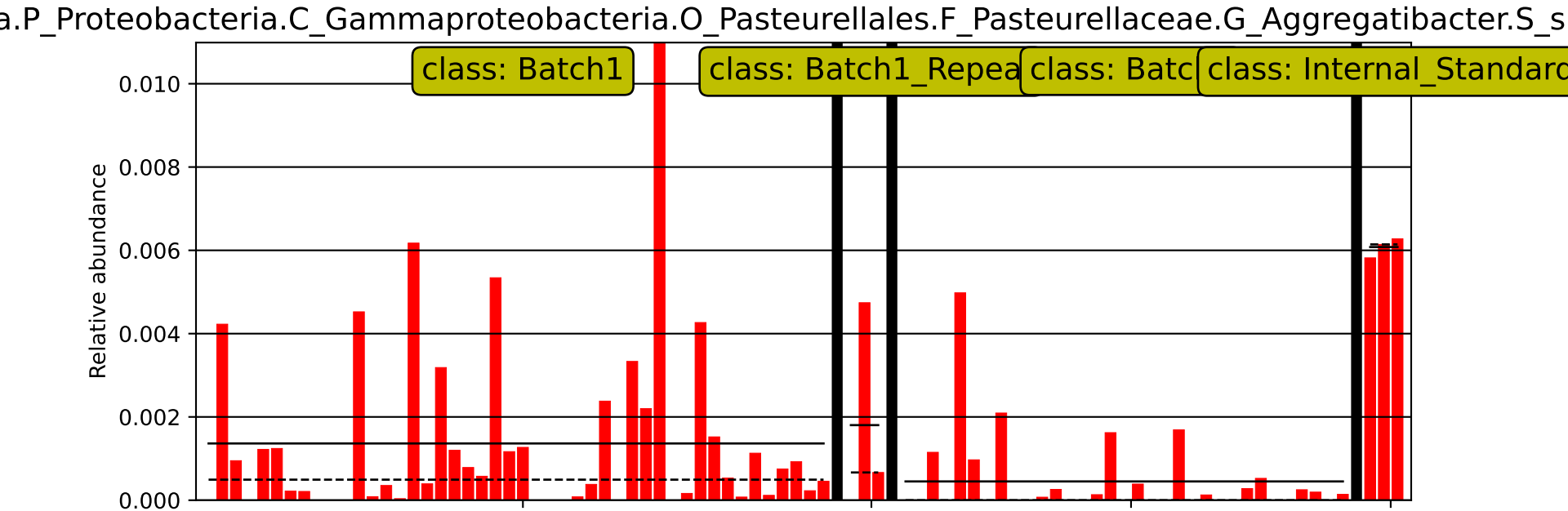
a.P_Proteobacteria.C_Gammaproteobacteria.O_Pasteurellales.F_Pasteurellaceae.G_Aggregatibacter.S_s
class: Batch1
class: Batch1_Repea
class: Batc
class: Internal_Standard
Relative abundance
0.010
0.008
0.006
0.004
0.002
0.000