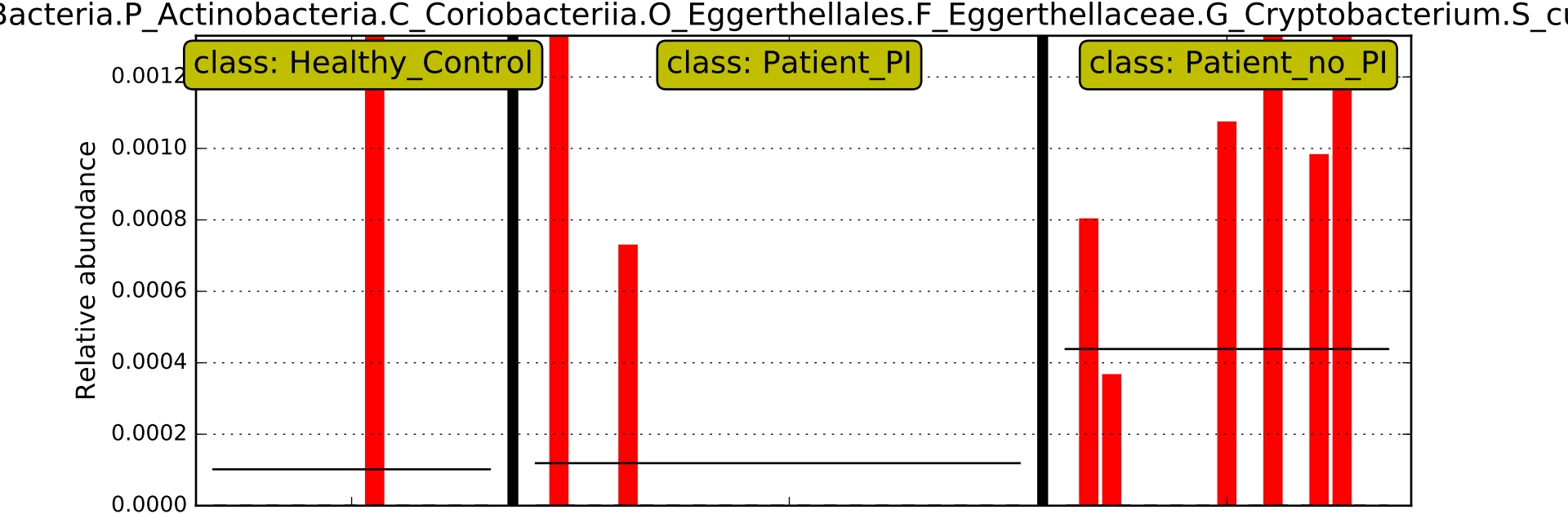
Bacteria.P_Actinobacteria.C_Coriobacteriia.O_Eggerthellales.F_Eggerthellaceae.G_Cryptobacterium.S_cu
class: Healthy_Control
class: Patient_PI
class: Patient_no_PI
Relative abundance
0.0012
0.0010
0.0008
0.0006
0.0004
0.0002
0.0000