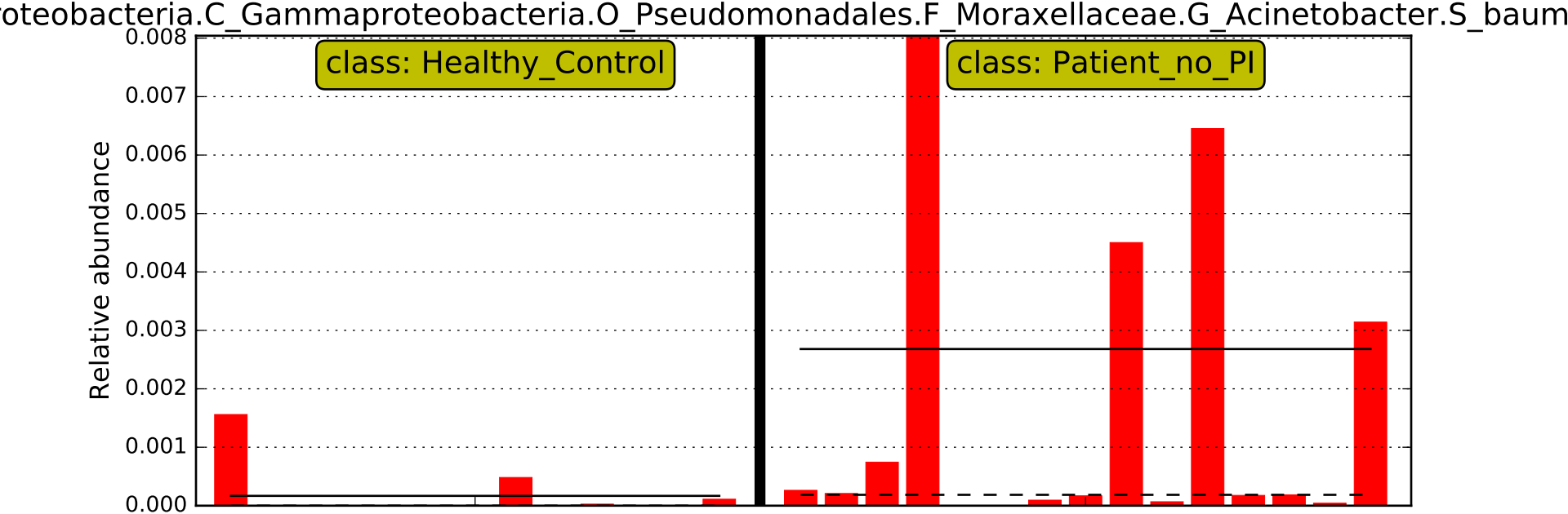
roteobacteria.C_Gammaproteobacteria.O_Pseudomonadales.F_Moraxellaceae.G_Acinetobacter.S_baum
class: Healthy_Control
class: Patient_no_PI
Relative abundance
0.008
0.007
0.006
0.005
0.004
0.003
0.002
0.001
0.000