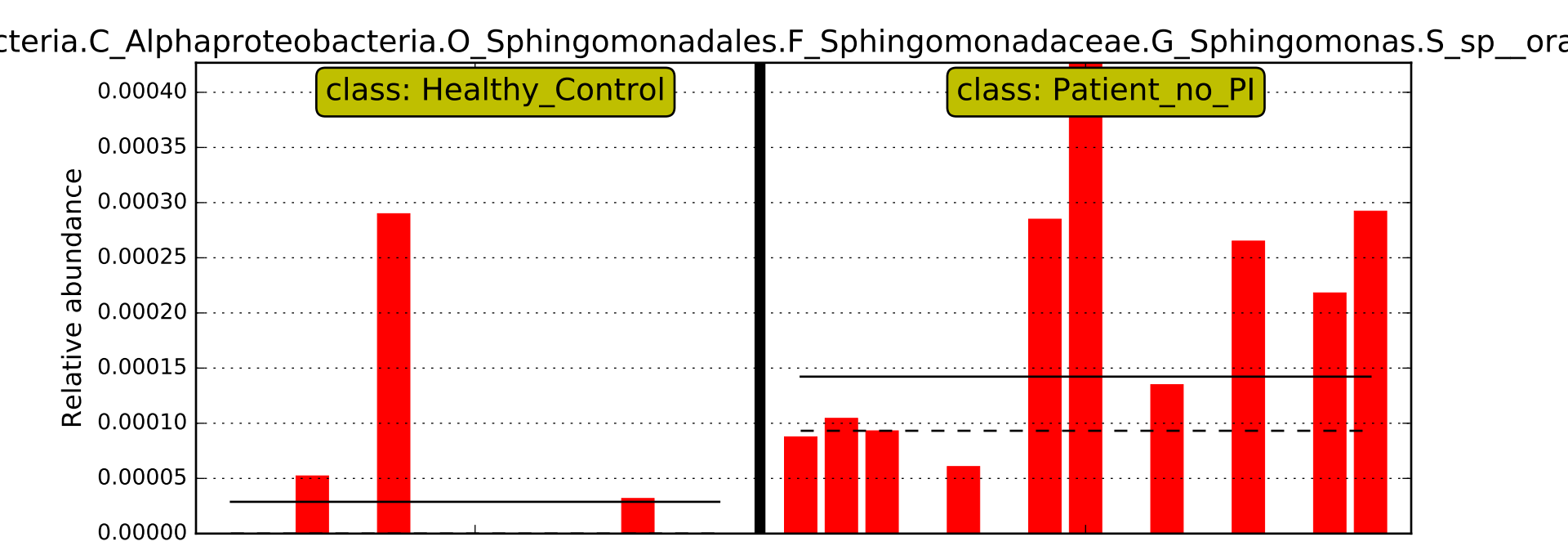
cteria.C_Alphaproteobacteria.O_Sphingomonadales.F_Sphingomonadaceae.G_Sphingomonas.S_sp__ora
class: Healthy_Control
class: Patient_no_PI
Relative abundance
0.00040
0.00035
0.00030
0.00025
0.00020
0.00015
0.00010
0.00005
0.00000