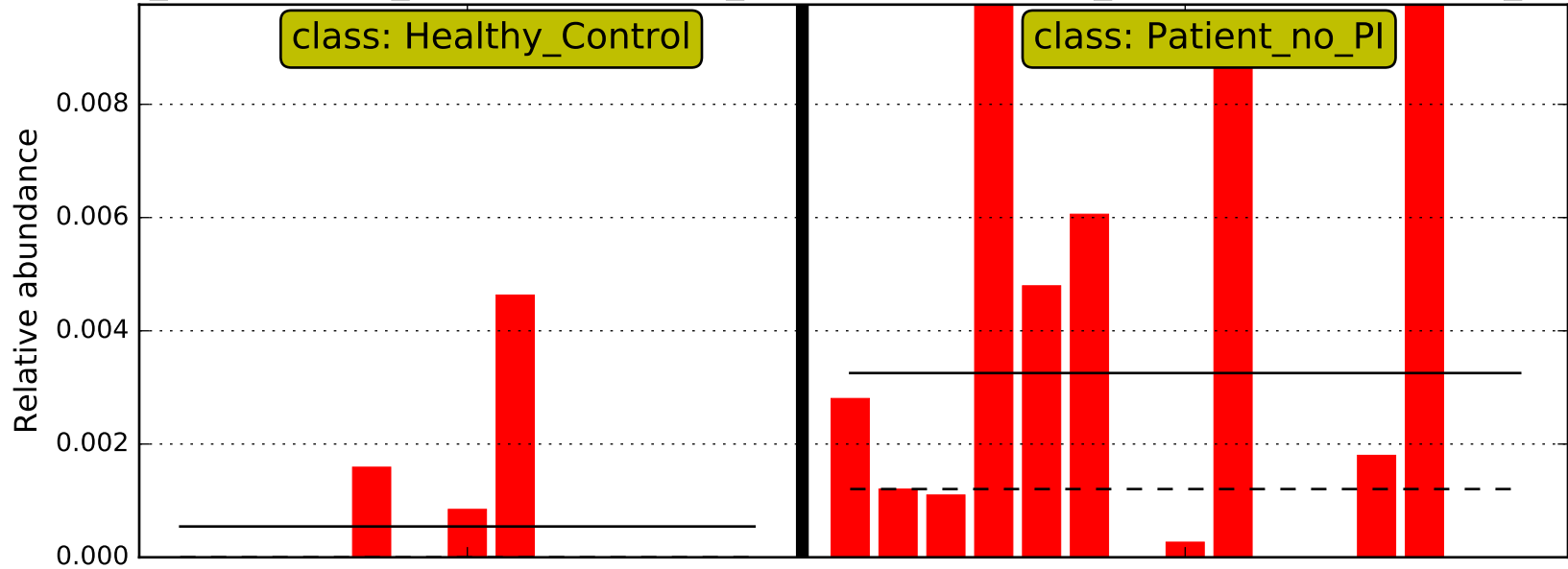


K_Bacteria.P_Firmicutes.C_Erysipelotrichia.O_Erysipelotrichales.F_Erysipelotrichaceae.G_Solobacterium



class: Healthy_Control

class: Patient_no_PI

Relative abundance

0.008
0.006
0.004
0.002
0.000