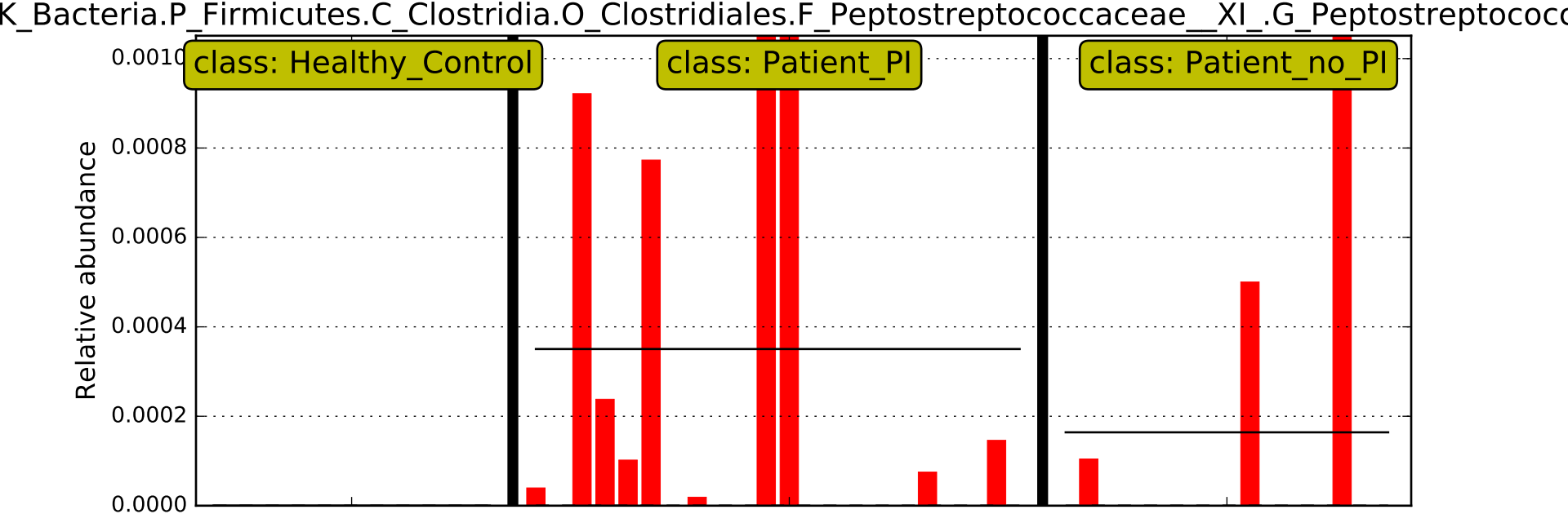
K_Bacteria.P_Firmicutes.C_Clostridia.O_Clostridiales.F_Peptostreptococcaceae__XI_.G_Peptostreptococ
class: Healthy_Control
class: Patient_PI
class: Patient_no_PI
Relative abundance
0.0010
0.0008
0.0006
0.0004
0.0002
0.0000