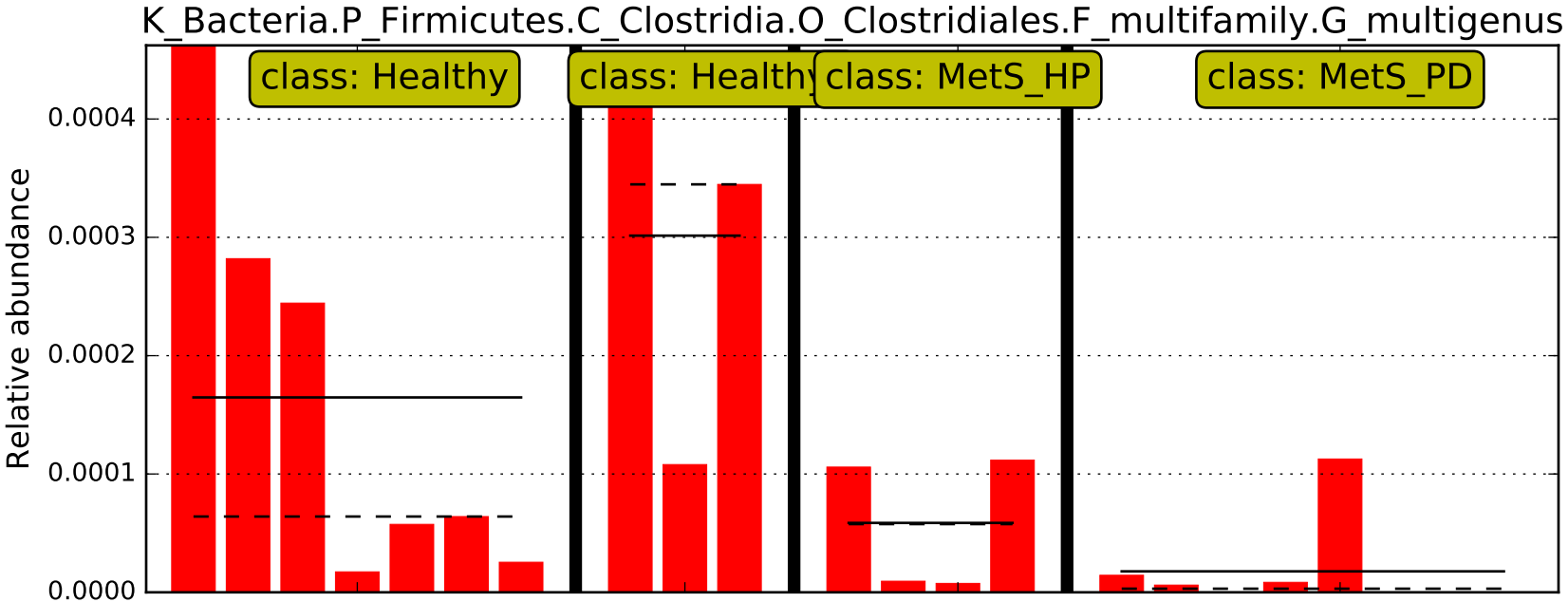
K_Bacteria.P_Firmicutes.C_Clostridia.O_Clostridiales.F_multifamily.G_multigenus
class: Healthy
class: Healthy
class: MetS_HP
class: MetS_PD
Relative abundance
0.0004
0.0003
0.0002
0.0001
0.0000