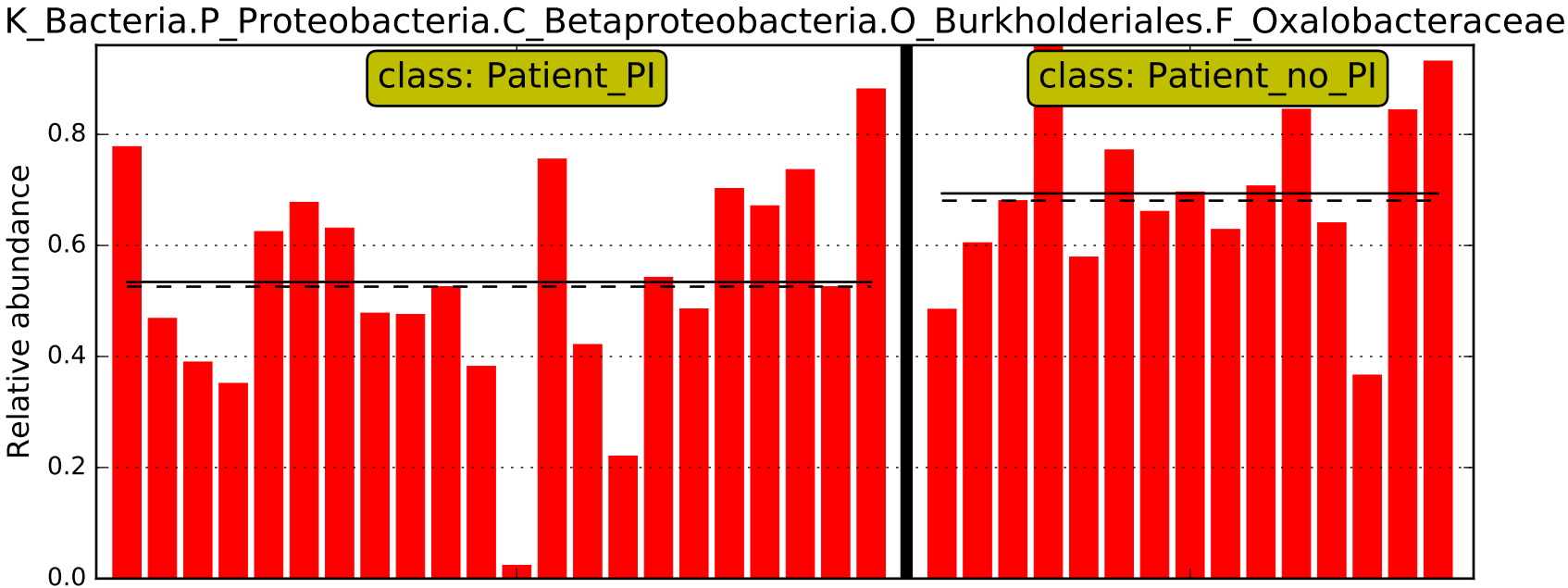
K_Bacteria.P_Proteobacteria.C_Betaproteobacteria.O_Burkholderiales.F_Oxalobacteraceae
class: Patient_PI
class: Patient_no_PI
Relative abundance
0.0
0.2
0.4
0.6
0.8