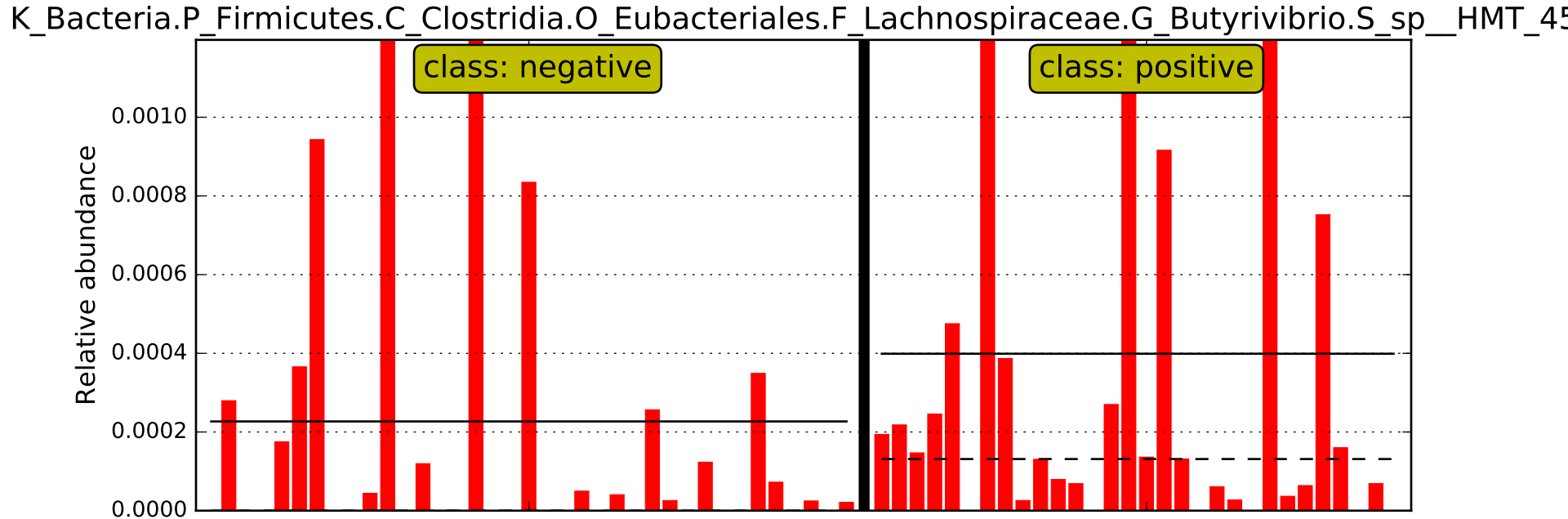
K_Bacteria.P_Firmicutes.C_Clostridia.O_Eubacteriales.F_Lachnospiraceae.G_Butyrivibrio.S_sp__HMT_45
class: negative
class: positive
Relative abundance
0.0010
0.0008
0.0006
0.0004
0.0002
0.0000