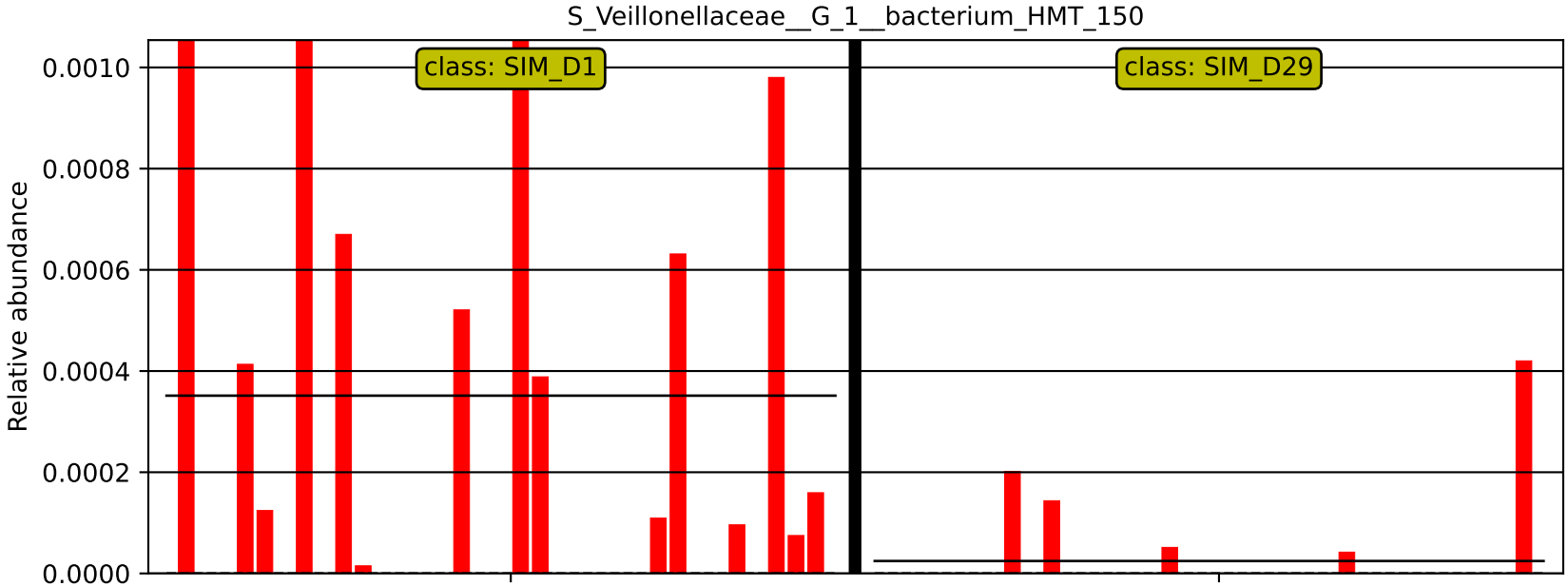
S_Veillonellaceae__G_1__bacterium_HMT_150
class: SIM_D1
class: SIM_D29
Relative abundance
0.0010
0.0008
0.0006
0.0004
0.0002
0.0000