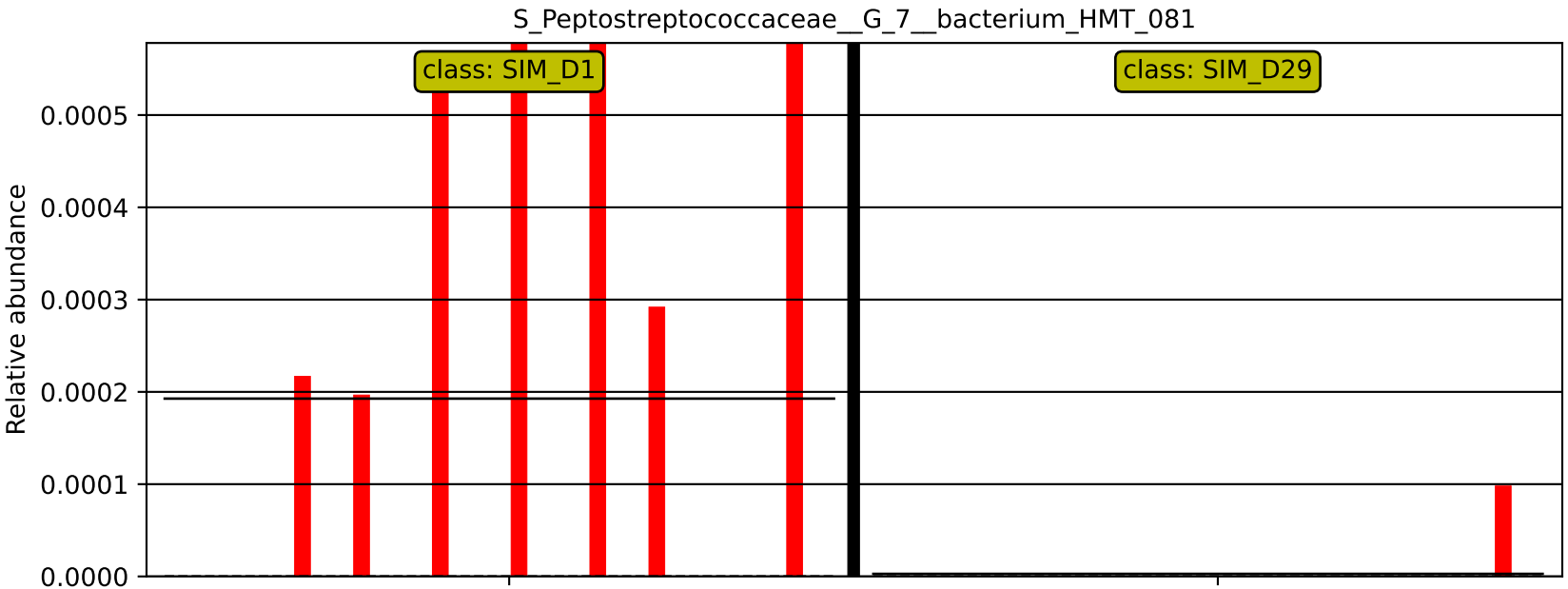
S_Peptostreptococcaceae__G_7__bacterium_HMT_081
class: SIM_D1
class: SIM_D29
Relative abundance
0.0005
0.0004
0.0003
0.0002
0.0001
0.0000