

K_Bacteria.P_Firmicutes.C_Negativicutes.O_Selenomonadales.F_Veillonellaceae.G_Selenomonas.S_nox

Relative abundance

class: Control_plate

class: Patient_sample

0.007
0.006
0.005
0.004
0.003
0.002
0.001
0.000

