

k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Aeromonadales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Alteromonadales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Cardiobacteriales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Chromatiales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Enterobacteriales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_HTCC2188
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Legionellales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Methylococcales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Oceanospirillales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Pasteurellales
k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Pseudomonadales

k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_RCP1-48
 k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Thiohalorhabdales
 k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Thiotrichales
 k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Vibrionales
 k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_Xanthomonadales
 k_Bacteria; p_Proteobacteria; c_Gammaproteobacteria; o_[Marinicellales]
 k_Bacteria; p_Proteobacteria; c_Zetaproteobacteria; o_Mariprofundales

k_Bacteria; p_Spirochaetes; c_Spirochaetes; o_Sphaerochaetales
 k_Bacteria; p_Spirochaetes; c_Spirochaetes; o_Spirochaetales
 k_Bacteria; p_Synergistetes; c_Synergistia; o_Synergistales

k_Bacteria; p_Tenericutes; c_Mollicutes; o_Acholeplasmatales
 k_Bacteria; p_Tenericutes; c_Mollicutes; o_Anaeroplasmatales
 k_Bacteria; p_Tenericutes; c_Mollicutes; o_Mycoplasmatales

k_Bacteria; p_Thermotogae; c_Thermotogae; o_Thermotogales
 k_Bacteria; p_Verrucomicrobia; c_Opitutae; o_Opitutales
 k_Bacteria; p_Verrucomicrobia; c_Opitutae; o_Puniceicoccales
 k_Bacteria; p_Verrucomicrobia; c_Opitutae; o_[Cerasicoccales]
 k_Bacteria; p_Verrucomicrobia; c_Verruco-5; o_SS1-B-03-39
 k_Bacteria; p_Verrucomicrobia; c_Verruco-5; o_WCHB1-41

k_Bacteria; p_Verrucomicrobia; c_Verrucomicrobiae; o_Verrucomicrobiales k_Bacteria; p_Verrucomicrobia; c_[Methylacidiphilae]; o_Methylacidiphilales

k_Bacteria; p_Verrucomicrobia; c_[Pedosphaerae]; o_[Pedosphaerales] k_Bacteria; p_Verrucomicrobia; c_[Spartobacteria]; o_[Chthoniobacterales]

k_Bacteria; p_SBR1093; c_; o_k_Bacteria; p_SR1; c_; o_

k_Bacteria; p_TM6; c_SBRH58; o_
k_Bacteria; p_TM6; c_SJA-4; o_
k_Bacteria; p_TM7; c_; o_
k_Bacteria; p_TM7; c_TM7-1; o_
k_Bacteria; p_TM7; c_TM7-3; o_

k_Bacteria; p_TM7; c_TM7-3; o_CW040
k_Bacteria; p_TM7; c_TM7-3; o_I025
k_Bacteria; p_Tenericutes; c_Mollicutes; o_

k_Bacteria; p_Tenericutes; c_Mollicutes; o_RF39 k_Bacteria; p_Tenericutes; c_RF3; o_ML615J-28

k_Bacteria; p_Verrucomicrobia; c_[Pedosphaerae]; o_

k_Bacteria; p_WS3; c_PRR-12; o_Sediment-1

k_Bacteria; p_[Caldithrix]; c_KSB1; o_Ucn15732

k_Bacteria; p_WPS-2; c_; o_

k_Bacteria; p_WS4; c_; o_ k_Bacteria; p_ZB3; c_BS119; o_