```
8.0
                k_;p_;c_;o_;f__
                k__; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales; f__Propionibacteriaceae
                 k ; p Actinobacteria; c Actinobacteria; o Bifidobacteriales; f Bifidobacteriaceae
                k__; p__Actinobacteria; c__Actinobacteria; o__Coriobacteriales; f__
0.6
                k ; p Actinobacteria; c Actinobacteria; o Coriobacteriales; f Coriobacteriaceae
                k__; p__Bacteroidetes; c__; o__; f__
                k ; p Bacteroidetes; c Bacteroidia; o Bacteroidales; f
                k ; p Bacteroidetes; c Bacteroidia; o Bacteroidales; f Bacteroidaceae
                k ; p Bacteroidetes; c Bacteroidia; o Bacteroidales; f Porphyromonadaceae
                k_; p_Bacteroidetes; c_Bacteroidia; o_Bacteroidales; f_Prevotellaceae
0.4
                k ; p Bacteroidetes; c Bacteroidia; o Bacteroidales; f Rikenellaceae
                k__; p__Cyanobacteria; c__4C0d-2; o__YS2; f__
                 k ; p Firmicutes; c ; o ; f
                 k ; p Firmicutes; c Bacilli; o Lactobacillales; f Enterococcaceae
                 k ; p Firmicutes; c Bacilli; o Lactobacillales; f Lactobacillaceae
                 k ; p Firmicutes; c Bacilli; o Lactobacillales; f Streptococcaceae
0.2
                 k ; p Firmicutes; c Bacilli; o Turicibacterales; f Turicibacteraceae
                 k ; p Firmicutes; c Clostridia; o ; f
                 k ; p Firmicutes; c Clostridia; o Clostridiales; f
                 k_; p_Firmicutes; c_Clostridia; o_Clostridiales; f_Catabacteriaceae
                k ; p Firmicutes; c Clostridia; o Clostridiales; f Clostridiaceae
                 k_; p_Firmicutes; c_Clostridia; o_Clostridiales; f_ClostridialesFamilyXI.IncertaeSedis
0.0
               k ; p Firmicutes; c Clostridia; o Clostridiales; f ClostridialesFamilyXIII.IncertaeSedis
                 k ; p Firmicutes; c Clostridia; o Clostridiales; f Dehalobacteriaceae
               k ; p Firmicutes; c Clostridia; o Clostridiales; f Lachnospiraceae
                k ; p Firmicutes; c Clostridia; o Clostridiales; f Peptococcaceae
   ethnicity
             k ; p Firmicutes; c Clostridia; o Clostridiales; f Ruminococcaceae
              k ; p Firmicutes; c Clostridia; o Clostridiales; f Veillonellaceae
             k_; p_Fusobacteria; c_Fusobacteria; o_Fusobacteriales; f_Fusobacteriaceae
             k_; p_Lentisphaerae; c_Lentisphaerae; o_Victivallales; f_Victivallaceae
             k_; p_Proteobacteria; c_Alphaproteobacteria; o_; f_
             k ; p Proteobacteria; c Betaproteobacteria; o ; f
             k_; p_Proteobacteria; c_Betaproteobacteria; o Burkholderiales; f
             k ; p Proteobacteria; c Betaproteobacteria; o Burkholderiales; f Alcaligenaceae
             k ; p Proteobacteria; c Betaproteobacteria; o Burkholderiales; f Oxalobacteraceae
             k ; p Proteobacteria; c Deltaproteobacteria; o Desulfovibrionales; f Desulfovibrionaceae
               k ; p Proteobacteria; c Epsilonproteobacteria; o Campylobacterales; f Campylobacteraceae
                k__; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales; f__Succinivibrionaceae
             k_; p_Proteobacteria; c_Gammaproteobacteria; o_Enterobacteriales; f_Enterobacteriaceae
               k_; p_Proteobacteria; c_Gammaproteobacteria; o_Pasteurellales; f_Pasteurellaceae
             k ; p Synergistetes; c Synergistia; o Synergistales; f Synergistaceae
               k ; p Tenericutes; c ; o ; f
             k__; p__Tenericutes; c__Erysipelotrichi; o__Erysipelotrichales; f__
               k__; p__Tenericutes; c__Erysipelotrichi; o__Erysipelotrichales; f__Erysipelotrichaceae
             k__; p__Tenericutes; c__Erysipelotrichi; o__Erysipelotrichales; f__vadinHA31
               k__; p__Tenericutes; c__ML615J-28; o__; f__
              k__; p__Tenericutes; c__Mollicutes; o__; f__
               k_; p_Tenericutes; c_Mollicutes; o_Anaeroplasmatales; f_Anaeroplasmataceae
              k__; p__Tenericutes; c__Mollicutes; o__RF39; f__
               k__; p__Verrucomicrobia; c__Opitutae; o__Puniceicoccales; f__Puniceicoccaceae
              k_; p__Verrucomicrobia; c__Verrucomicrobiae; o__Verrucomicrobiales; f__Verrucomicrobiaceae
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1.0