

Classifier: RDP Naive Bayesian rRNA Classifier Version 2.11,
September 2015

Taxonomical Hierarchy: RDP 16S rRNA training set 16

Query File: F9-F12 GenBank.fas

Submit Date: Wed Jul 10 07:20:23 EDT 2019

Confidence threshold (for classification to Root ONLY): 95%

Symbol +/- indicates predicted sequence orientation

F9.36;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;13%
F9.37;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;14%
F9.39;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;13%
F9.42;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;14%
F9.43;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;14%
F9.44;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;13%
F9.45;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;14%
F9.46;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;13%
F9.48;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;14%
F9.49;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
98%;"Flavobacteriales";98%;Flavobacteriaceae;77%;Kordia;14%
F9.7;+;Root;100%;Bacteria;100%;"Proteobacteria";
100%;Betaproteobacteria;100%;Burkholderiales;100%;Comamonadaceae;
100%;Variovorax;100%
F9.35;+;Root;100%;Bacteria;100%;Firmicutes;100%;Bacilli;
100%;Lactobacillales;100%;Streptococcaceae;100%;Streptococcus;100%
F9.2;+;Root;100%;Bacteria;100%;Firmicutes;100%;Bacilli;
100%;Lactobacillales;100%;Streptococcaceae;100%;Streptococcus;100%
F9.40;+;Root;100%;Bacteria;100%;"Proteobacteria";
100%;Alphaproteobacteria;100%;Rhodobacterales;100%;Rhodobacteraceae;
100%;Oceanicola;72%
F9.47;+;Root;100%;Bacteria;100%;"Proteobacteria";
100%;Alphaproteobacteria;100%;Rhodobacterales;100%;Rhodobacteraceae;
100%;Oceanicola;72%
F9.41;+;Root;100%;Bacteria;100%;"Proteobacteria";
100%;Alphaproteobacteria;100%;Rhodobacterales;100%;Rhodobacteraceae;
100%;Leisingera;100%
F9.53;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
100%;"Flavobacteriales";100%;Flavobacteriaceae;100%;Tenacibaculum;
93%
F9.64;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
100%;"Flavobacteriales";100%;Flavobacteriaceae;100%;Flavivirga;24%
F9.63;+;Root;100%;Bacteria;100%;"Bacteroidetes";100%;Flavobacteriia;
100%;"Flavobacteriales";100%;Flavobacteriaceae;100%;Arenitalea;100%
F9.38;+;Root;100%;Bacteria;100%;"Proteobacteria";
100%;Alphaproteobacteria;100%;Rhizobiales;100%;Phyllobacteriaceae;
100%;Pseudahrensia;74%

F9.54;+;Root;100%;Bacteria;100%;"Actinobacteria";
 100%;Actinobacteria;100%;Actinobacteridae;100%;Actinomycetales;
 100%;Propionibacterineae;100%;Propionibacteriaceae;
 100%;Propionibacterium;100%
 F9.59;+;Root;100%;Bacteria;100%;"Actinobacteria";
 100%;Actinobacteria;100%;Actinobacteridae;100%;Actinomycetales;
 100%;Propionibacterineae;100%;Propionibacteriaceae;
 100%;Propionibacterium;100%
 F9.55;+;Root;100%;Bacteria;100%;"Actinobacteria";
 100%;Actinobacteria;100%;Actinobacteridae;100%;Actinomycetales;
 100%;Propionibacterineae;100%;Propionibacteriaceae;
 100%;Propionibacterium;100%
 F9.56;+;Root;100%;Bacteria;100%;"Actinobacteria";
 100%;Actinobacteria;100%;Actinobacteridae;100%;Actinomycetales;
 100%;Propionibacterineae;100%;Propionibacteriaceae;
 100%;Propionibacterium;100%
 F9.57;+;Root;100%;Bacteria;100%;"Actinobacteria";
 100%;Actinobacteria;100%;Actinobacteridae;100%;Actinomycetales;
 100%;Propionibacterineae;100%;Propionibacteriaceae;
 100%;Propionibacterium;100%
 F9.58;+;Root;100%;Bacteria;100%;"Actinobacteria";
 100%;Actinobacteria;100%;Actinobacteridae;100%;Actinomycetales;
 100%;Propionibacterineae;100%;Propionibacteriaceae;
 100%;Propionibacterium;100%
 F9.21;+;Root;100%;Bacteria;100%;"Actinobacteria";
 100%;Actinobacteria;100%;Actinobacteridae;100%;Actinomycetales;
 100%;Micrococcineae;100%;Microbacteriaceae;100%;Rhodoluna;19%
 F9.20;+;Root;100%;Bacteria;100%;"Proteobacteria";
 100%;Betaproteobacteria;100%;Burkholderiales;100%;Burkholderiaceae;
 100%;Ralstonia;100%
 F9.1;+;Root;100%;Bacteria;100%;"Proteobacteria";
 100%;Betaproteobacteria;100%;Burkholderiales;100%;Burkholderiaceae;
 100%;Burkholderia;100%
 F9.3;+;Root;100%;Bacteria;100%;"Proteobacteria";
 100%;Betaproteobacteria;100%;Burkholderiales;100%;Burkholderiaceae;
 100%;Lautropia;100%
 F9.52;+;Root;100%;Bacteria;100%;"Proteobacteria";
 100%;Alphaproteobacteria;100%;Rhodobacterales;98%;Rhodobacteraceae;
 98%;Ahrensia;97%
 F9.25;+;Root;100%;Bacteria;100%;"Proteobacteria";
 100%;Alphaproteobacteria;100%;Rhodobacterales;100%;Rhodobacteraceae;
 100%;Oceanicola;72%
 F9.26;+;Root;100%;Bacteria;100%;"Proteobacteria";
 100%;Alphaproteobacteria;100%;Rhodospirillales;
 100%;Acetobacteraceae;100%;Acidisphaera;68%