

TACGGGGGGGGCAAGCGTTGTTTCGGAATTACTGGGCGTAAAGGGCTCGTA
GGCGGCCAACTAAGTCAGACGTGAAATCCCTCGGCTTAAC

TAXONOMY:

Greengenes lineage: k__Bacteria; p__Acidobacteria; c__Acidobacteria-6; o__iii1-15; f__; g__; s__

RDP (100% ID) lineage: d__Bacteria; k__; p__Acidobacteria; c__Acidobacteria_Gp6; o__; f__; g__Gp6
(4/4)

RDP (100% ID) species: unidentified bacterium (2/4) agricultural soil (1/4) Acidobacteria bacterium (1/4)

PREVALENCE: Found in 23.95% of samples, rank #5 out of 155002 tag sequences.

ABUNDANCE: Composes 0.127% of observations, rank #83 out of 155002 tag sequences.

METHODS: Amplicon PCR with 16S rRNA V4 primers 515f–806rB. Sequencing with Illumina HiSeq 90/100-bp or MiSeq 150-bp single reads. Sequences error-checked with Deblur, trimmed to 90 bp & rarefied to 5000 obs./sample. Showing even sample type distribution subset of 2000 samples.

PREVALENCE BY SAMPLE TYPE:



