

TACGTAGGTGGCAAGCGTTGTCCGGATTATTGGGCGTAAAGCGAGCGCA
GGCGGAAAAATAAGTCTAATGTGAAAGCCCTCGGCTTAAC

TAXONOMY: Assigned Greengenes taxonomy & most common taxonomy of 100% ID matches to RDP.

Greengenes lineage: k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Lactobacillaceae; g__Lactobacillus; s__

RDP lineage: d__Bacteria; k__; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Lactobacillaceae; g__Lactobacillus (165/165)

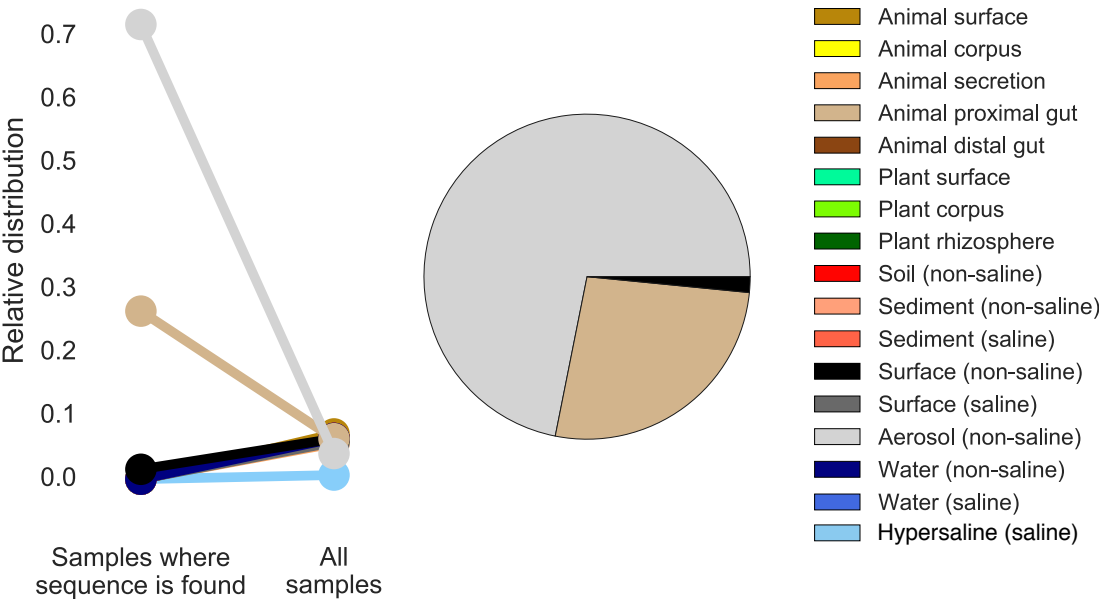
RDP species: Lactobacillus sp. (87/165) Lactobacillus amylovorus (64/165) Lactobacillus acidophilus (8/165)

PREVALENCE: Found in 3.20% of samples, rank #1511 out of 153302 tag sequences.

ABUNDANCE: Composes 0.035% of observations, rank #412 out of 153302 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 2000 samples evenly distributed by habitat & study.

PREVALENCE BY HABITAT:



ABUNDANCE BY ENVIRONMENTAL PARAMETERS:

