

Positive control ASV

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94efb59163996601f8a20a8f9b0ac0d9: d_Bacteria; p_Firmicutes; c_Bacilli; o_Bacillales;
  _Bacillaceae; g__Bacillus
b08e8a180eba5a20a70b190fbf5f9ad2: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria;
o__Enterobacterales; f__Enterobacteriaceae
fdaab5e761766ddd1297bd5f49d1cfb3: d Bacteria: p Proteobacteria: c Gammaproteobacteria:
o Enterobacterales
f0090fc348a4c45135f858d776e30dd0: d__Bacteria; p__Firmicutes; c__Bacilli;
o__Lactobacillales; f__Lactobacillaceae; g__Lactobacillus
456f366888eac2ce0809986d3ed1b237: d__Bacteria; p__Firmicutes; c__Bacilli;
o__Lactobacillales
d02f4c8904ba636c08bbdf8b65b5d7d0: d__Bacteria; p__Firmicutes; c__Bacilli;
o__Lactobacillales; f__Listeriaceae; g__Listeria
d3962c483c45f0765b9b7656ec7f43a0: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
3019073bed0ceecaece9f446607c003a: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria;
o__Enterobacterales; f__Enterobacteriaceae; g__Salmonella
f9eed79f2bd69b0da2385144b13658a7: d_Bacteria; p_Firmicutes; c_Bacilli;
o__Staphylococcales; f__Staphylococcaceae
5c9b89d5acf8e0020d9de19b53d21663: d_Bacteria; p_Firmicutes; c_Bacilli;
o__Staphylococcales; f__Staphylococcaceae; g__Staphylococcus
Others
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Negative Control ASV

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19d57d86ee1d5ea49fe9f4256bfa9d22: d Bacteria; p Bacteroidota; c Bacteroidia;
o Flavobacteriales; f__Flavobacteriaceae; g__Flavobacterium
1c34565a5c5d3d9fa3953518e9a4b59c: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria;
o__Burkholderiales; f__Oxalobacteraceae
33731ac0a1051571d808de3e91b11356: d Bacteria; p Proteobacteria; c Alphaproteobacteria;
o__Rhizobiales; f__Beijerinckiaceae; g__Methylobacterium-Methylorubrum;
s__Methylobacterium_jeotgali
7a6da1a5e2be4eedd87d67705eb68f4a: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria;
o__Alteromonadales; f__Alteromonadaceae; g__Rheinheimera
a64d8357b6a2010ba13a1ebfc7df7c9b: d__Bacteria; p__Proteobacteria; c__Alphaproteobacteria;
o__Rhodobacterales; f__Rhodobacteraceae; g__Amaricoccus; s__Amaricoccus_kaplicensis
d23c811ea38d6cb704bb13445f43d599: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
e1a72c614544de9ecfc0b0cd7cc24738: d__Bacteria; p__Bacteroidota; c__Bacteroidia;
o__Flavobacteriales; f__Flavobacteriaceae; g__Flavobacterium
e6242a18b930fae98ab395c927393aa3: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria;
o__Burkholderiales; f__Comamonadaceae
ebb95142c2d5fb1d9a1e0e779348b6f1: d_Bacteria; p_Bacteroidota; c_Bacteroidia;
o__Flavobacteriales; f__Flavobacteriaceae; g__Flavobacterium
f0af66950b4ed3e76bd78e09b28e1f02: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria;
o__Pseudomonadales; f__Moraxellaceae; g__Acinetobacter
Others
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