

1 **Supplementary Table 1**

2 The expected and observed relative abundance of bacterial species in the DNA extraction  
3 and the PCR step control samples.

<b>Bacterial taxon</b>	<b>Expected abundance</b>	<b>Obtained abundance in the PCR step control sample</b>	<b>Obtained abundance in the DNA extraction control sample</b>
<i>Pseudomonas</i>	12.5	7.229	10.439
<i>Enterobacteriaceae</i> (combination of <i>Escherichia</i> and <i>Salmonella</i> )	25.0	19.551	31.620
<i>Lactobacillus</i>	12.5	12.153	22.332
<i>Enterococcus</i>	12.5	8.202	4.976
<i>Staphylococcus</i>	12.5	19.613	10.535
<i>Listeria</i>	12.5	13.007	0.746
<i>Bacillus</i>	12.5	19.250	17.887

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