

**Supplementary Table 1.** Additional information on the relative amplicon-sequence-variants (ASVs)

<b>Genus</b>	<b>found_in_neg_ctls</b>	<b>number_of_ASVs_in_neg_ctls</b>	<b>number_of_ASVs_total</b>	<b>post_clean_mean_pct</b>	<b>num_samples_pos_after_cleaning</b>
<i>Comamonadaceae</i>	Y	1	4	4.61	14
<i>Staphylococcus</i>	Y	1	13	4.04	11
<i>Propionibacterium</i>	Y	1	3	3.75	9
<i>Pseudomonas</i>	Y	1	4	1.54	12
<i>Methylosula</i>	Y	1	1	1.16	1
<i>Acidimicrobiaceae</i>	Y	1	4	1.12	2
<i>Bacillus</i>	Y	1	15	0.89	8
<i>Microbacterium</i>	Y	2	10	0.53	3
<i>Acidibacter</i>	Y	1	2	0.34	2
<i>Lachnospiraceae</i>	Y	1	11	0.17	2
<i>Methylobacterium</i>	Y	1	3	0.14	4
<i>Belnapia</i>	Y	1	1	0.00	0
<i>Vasilyevaea</i>	Y	1	1	0.00	0
<i>Geodermatophilaceae</i>	Y	1	1	0.00	0
<i>Oligoflexales</i>	Y	1	1	0.00	0

Abbreviations: ctls, controls; neg, negative; num, number; pct, percentage; pos, positive