

S1 Table

<b>Sampling site</b>	<b>Taxon removed from AMOVA<sup>a</sup></b>	<b>P-value AMOVA</b>
<b>Cheek</b>	None	<0.001
	<i>Streptococcus salivarius</i> (a)	0.14
	<i>Streptococcus mitis</i> group (b)	0.09
	<i>Rothia</i> (c)	<0.001
	<i>Veillonella</i> (d)	<0.001
	<i>Actinomyces</i> (e)	<0.001
	<i>Gemella</i> (f)	<0.001
	<i>Corynebacterium</i> (g)	<0.001
	<i>Propionibacterium</i> (h)	0.001
<b>Dorsal forearm</b>	None	0.16
	<i>Propionibacterium</i> (a)	0.08
	<i>Streptococcus mitis</i> group (b)	0.36
	<i>Staphylococcus</i> (c)	0.17
	<i>Granulicatella</i> (d)	0.16
	<i>Rothia</i> (e)	0.17
	<i>Corynebacterium</i> (f)	0.13
	<i>Streptococcus salivarius</i> (g)	0.47
	<i>Prevotella</i> (h)	0.14
<b>Volar forearm</b>	None	0.16
	<i>Streptococcus mitis</i> group (a)	0.36
	<i>Gemella</i> (b)	0.16
	<i>Actinomyces</i> (c)	0.17
	<i>Rothia</i> (d)	0.16
	<i>Porphyromonas</i> (e)	0.15
	<i>Prevotella</i> (f)	0.14
	<i>Veillonella</i> (g)	0.18
	<i>Streptococcus salivarius</i> (h)	0.33

<sup>a</sup>Letters in parentheses indicate labels for taxa in figure 2

AMOVA, analysis of molecular variance.