

Table S7 Differentially abundant non-candidate OTUs and taxa conserved between facilities

OTU-ID	Taxonomic Identity	Statistical Test	Sample Type	Association	n Facility	Log <sub>2</sub> FC
549756^	G: Lactobacillus	G-Test	Fecal	P07-WT P07-KO	Boston Laramie	-0.29 +2.4
n/a	G: Streptococcus	Kruskal-Wallis	Fecal	P07-KO P07-WT	Boston Laramie	+10 -2.5
n/a	P: Actinobacteria	Kruskal-Wallis	Colon	P20-KO P20-WT	Boston Laramie	+10 -6.7
		Nonparametric T		P20-KO P20-WT	Boston Laramie	+10 -6.7
n/a	F: S24-7^	Kruskal-Wallis	Fecal	P20-KO P20-WT	Boston Laramie	+1.4 -0.94
		Nonparametric T		P20-KO P20-WT	Boston Laramie	+1.4 -0.94
n/a 	F: Ruminococcaceae	Kruskal-Wallis	Fecal	P24-KO P24 <b>-</b> WT	Boston Laramie	+2.7 -4.8

<sup>^</sup> Indicates OTUs or taxa observed at multiple ages. P: represents Phylum. F: represents Family. G: represents Genus. Log<sub>2</sub> Fold Change (FC) was calculated KO/WT, therefore (+) indicates association with KO mice and (-) indicates association with WT mice.