

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

**Ednrb mice - HSCR Dataset** OTU-ID **Taxon** Statistical Test Type **Facility** Association Log<sub>2</sub> FC 549756^ G: Lactobacillus G-Test Fecal P07-WT -0.29Boston Laramie **P07-KO** +2.4 Kruskal-Wallis **P07-KO** n/a G: Streptococcus Fecal Boston +10 P07-WT -2.5 Laramie n/a P: Actinobacteria Kruskal-Wallis Boston **P20-KO** +10 Laramie **P20-WT** -6.7 Colon Nonparametric T Boston **P20-KO** +10 **P20-WT** -6.7 Laramie n/a F: S24-7<sup>^</sup> Kruskal-Wallis Boston **P20-KO** +1.4 **P20-WT** -0.94 Laramie Fecal Nonparametric T Boston **P20-KO** +1.4 Laramie **P20-WT** -0.94 F: Ruminococcaceae Kruskal-Wallis Fecal Boston **P24-KO** n/a +2.7Laramie **P24-WT** -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.