Hindgut Relative Abundance Taxa 1.00-Other c__Clostridia; o__Clostridiales o__Clostridiales; f__Ruminococcaceae f__Moraxellaceae; g__Acinetobacter f_Ruminococcaceae; g_Ruminococcus o__Clostridiales; f__Lachnospiraceae 0.75 o Bacteroidales; f S24-7 Root Relative Abundance f_Lachnospiraceae; g_Butyrivibrio f_Prevotellaceae; g_Prevotella f_Spirochaetaceae; g_Treponema f_Ruminococcaceae; g_Oscillospira f_Bacteroidaceae; g_Bacteroides 0.50o__Enterobacteriales; f__Enterobacteriaceae f_Lachnospiraceae; g_Coprococcus o_Clostridiales; f_Christensenellaceae o_Coriobacteriales; f_Coriobacteriaceae c_Bacteroidia; o_Bacteroidales o_Bacillales; f_Planococcaceae 0.25 f__Lachnospiraceae; g__Dorea f__Lactobacillaceae; g__Lactobacillus c__Bacilli; o__Bacillales f__[Paraprevotellaceae]; g__CF231 f__Comamonadaceae; g__Comamonas 0.00 o__Xanthomonadales; f__Xanthomonadaceae o__Flavobacteriales; f__Flavobacteriaceae 2 5 6 Subject