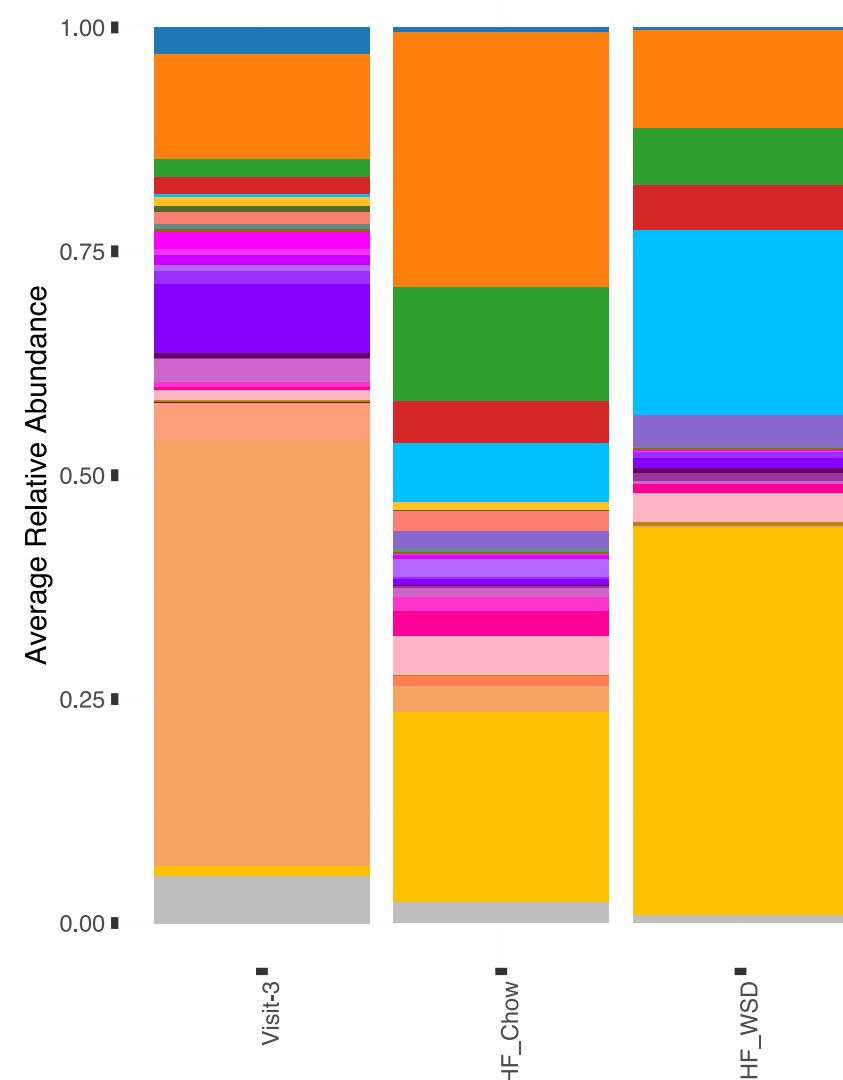


Alistipes  
 Bacteroides  
 Blautia  
 Lachnoclostridium  
 Akkermansia  
 Anaerostipes  
 Barnesiella  
 Bifidobacterium  
 Bilophila  
 Butyricoccus  
 Butyricimonas  
 Christensenellaceae\_R.7\_group  
 Colidextribacter  
 Coprobacter  
 Coprococcus  
 Erysipelatoclostridium  
 Flavonifractor

Gastranaerophilales  
 Incertae\_Sedis  
 Lachnospiraceae\_.\_\_\_  
 Oscillibacter  
 Oscillospiraceae.g\_\_uncultured  
 Parabacteroides  
 Paraprevotella  
 Parasutterella  
 Phascolarctobacterium  
 Ruminococcaceae\_.\_\_\_  
 Ruminococcus\_torques\_group  
 Sellimonas  
 Subdoligranulum  
 genera are among core microbiome (25%) of human-visit1 samples but not in core microbiome of mice samples  
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 Others



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