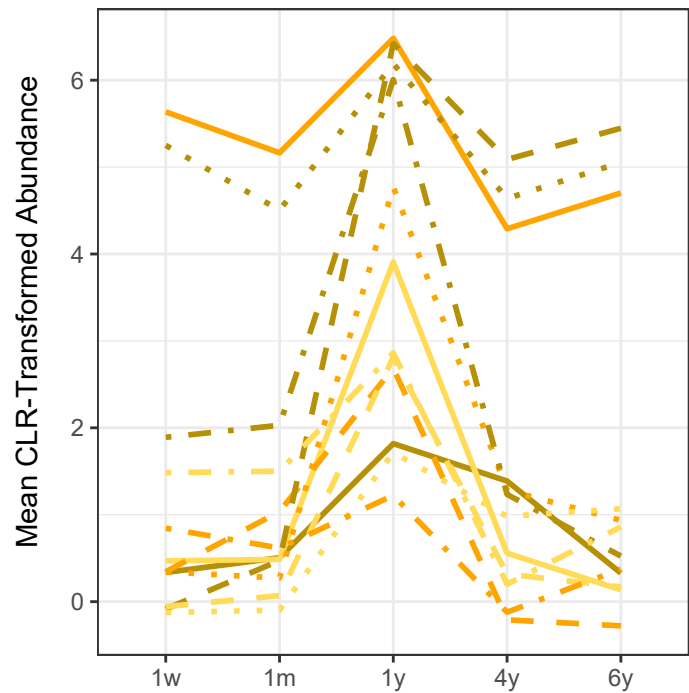
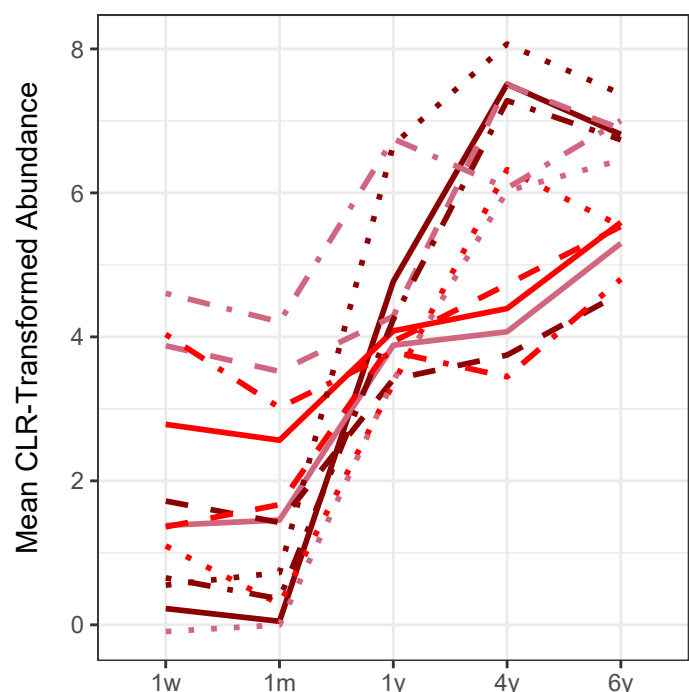


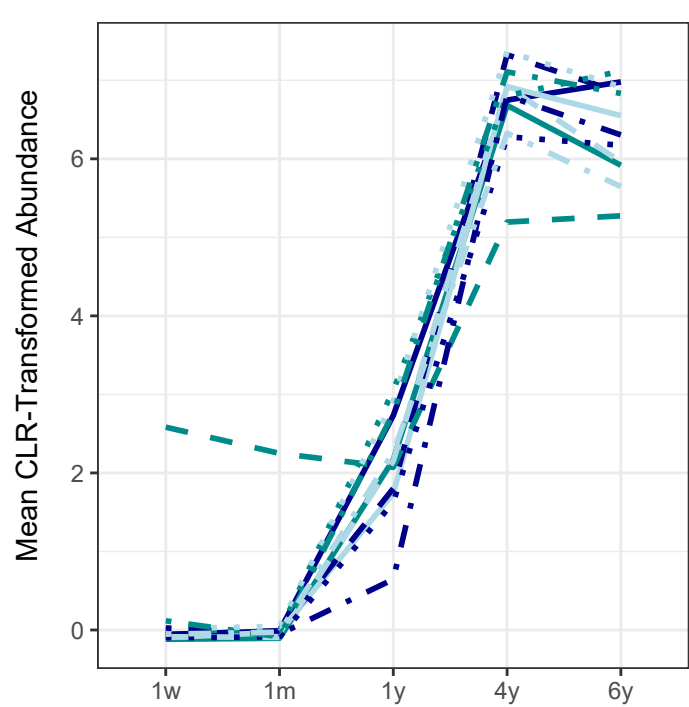
- Early ASVs**
- Bifidobacterium bifidum
 - Bifidobacterium breve/infantis
 - Clostridium_P perfringens
 - Escherichia coli/coli_C
 - Family_Enterobacteriaceae26
 - Family_Enterococcaceae
 - Staphylococcus argenteus/aureus
 - Streptococcus australis/koreensis
 - Streptococcus australis_A/chosunense
 - Streptococcus prausnitzii_F/salivarius
 - Veillonella dispar/dispar_A
 - Veillonella parvula_A



- Intermediate ASVs**
- Bacteroides fragilis/fragilis_A
 - Eggerthella lenta
 - Enterocloster bolteae/citroniae
 - Erysipelatoclostridium ramosum
 - Faecalibacterium sp.24
 - Faecalimonas phoceensis
 - Flavonifractor plautii/sp000508885
 - Haemophilus_D flavescens/parainfluenzae
 - Lachnospira eligens/eligens_A
 - Parabacteroides sp.31
 - Ruminococcus_B gnavus
 - Veillonella tobetsuensis



- Late ASVs**
- Anaerostipes hadrus/hadrus_A
 - Bacteroides faecichinchillae/faecis
 - Bacteroides sp.41
 - Bacteroides sp902362375
 - Bacteroides uniformis
 - Bifidobacterium callitrichidarum/catenulatum
 - Blautia_A wexlerae_B
 - Clostridium cuniculi/saudiense
 - Dialister invisus
 - Intestinibacter bartlettii
 - Phocaecicola dorei
 - Phocaecicola sp.22



- Latest ASVs**
- Agathobacter rectalis
 - Anaerobutyricum hallii
 - Blautia_A faecis
 - Blautia_A massiliensis/schinkii
 - CAG-110 sp900760325
 - Clostridium sp001916075
 - Dorea_A longicatena/longicatena_B
 - Faecalibacterium prausnitzii_C/prausnitzii_D
 - Fusicatenibacter saccharivorans
 - Gemmiger quicibialis
 - Romboutsia timonensis
 - Terrisporobacter sp902363255