Top drivers: community type Escherichia-Shigella-Ruminococcaceae UCG-002-Christensenellaceae R-7 group-Ruminococcaceae UCG-005-Ruminococcaceae UCG-010-[Eubacterium] coprostanoligenes group-Family-XIII AD3011 group-Ruminococcaceae NK4A214 group-Methanobrevibacter-Hydrogenoanaerobacterium-Defluviitaleaceae UCG-011-GCA-900066225 Ruminococcaceae UCG-014-Candidatus Soleaferrea-Ruminococcaceae V9D2013 group-Sanguibacteroides-Catabacter-Anaerotruncus-Akkermansia-CAG-352 0.3 0.9 0.0 Value