

Note: (A) Bacteroides, (B) Muribaculaceae, (C) Akkermansia, (D) Lactobacillus, (E) Lachnoclostridium, (F) Phascolarctobacterium, (G) Lachnospiraceae_NK4A136_group, (H) Parabacteroides, (I) [Ruminococcus]_gnavus_group, (J) Anaerofilum, (K) Dorea, (L) Romboutsia, (M) Ruminococcaceae_UCG-014, (N) Hungatella, (O) Prevotellaceae_UCG-001, (P) Desulfovibrio, (Q) Ruminococcaceae_UCG-013, (R) Flavonifractor, (S) Blautia, (T) Eisenbergiella, (U) Parasutterella, (V) Bilophila, (W) Anaerostipes, (X) [Ruminococcus]_torques_group, (Y) [Eubacterium]_xylanophilum_group, (Z) Roseburia, (A1) Faecalibaculum, (B1) Ruminococcus_1, (C1) Ruminiclostridium_6, (D1) Prevotellaceae_NK3B31_group, (E1) [Eubacterium]_coprostanoligenes_group, (F1) Marvinbryantia, (G1) Ruminococcaceae_UCG-008, (H1) Oscillibacter, (I1) Fournierella, (J1) Ruminococcaceae_UCG-005, (K1) Turicibacter, (L1) Clostridium_sensu_stricto_1, (M1) Clostridioides, (N1) Ruminiclostridium_9, (O1) [Ruminococcus]_gavreaui_group, (P1) Ruminiclostridium_5, (Q1) Lachnospiraceae_UCG-006, (R1) UBA1819, (S1) Erysipelatoclostridium, (T1) Holdemania, (U1) Fusicatenibacter, (V1) Negativibacillus, (W1) Erysipelotrichaceae_UCG-003, (Y1) [Eubacterium]_nodatum_group