nus, Abundance)

na	Jeotgalicoccus	Lachnoclostridium	[Ruminococcus] torques group	
Saccharimonas	Acetatifactor	Desulfovibrio	Prevotellaceae UCG-001	
	Intestinimonas	Cladosporium	Paeniclostridium	
	Ruminococcus	[Ruminococcus] gnavus group	Sporosarcina	
rces	Roseburia	Blautia	Ureaplasma	
haeria	Staphylococcus	Oscillibacter	Butyricicoccus	
	UCG-003	Lachnospiraceae NK4A136 group	Clostridium sensu stricto 1	
	[Eubacterium] fissicatena group	Marvinbryantia	HT002	
bup	Lachnospiraceae UCG-006	Tuzzerella	Anaeroplasma	
	Enterorhabdus	Enterococcus	Eimeria	
	Limosilactobacillus	Odoribacter	Anaerostipes	
	Monoglobus	Parabacteroides	Mucispirillum	
n	Incertae Sedis	A2	Coriobacteriaceae UCG-002	
i] ventriosum group	Ligilactobacillus	Rikenella	Veillonella	
	GCA-900066575	Alistipes	Muribaculum	
ı] xylanophilum group	Lachnospiraceae UCG-001	Helicobacter	Erysipelatoclostridium	
ı] siraeum group	Anaerotruncus	Lachnospiraceae FCS020 group	Bacteroides	
	Romboutsia	Colidextribacter	Bilophila	
] oxidoreducens group	Paraprevotella	Terrisporobacter	Parasutterella	
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