



Genus	[Anaerorhabdus] furcosa group	Catenibacterium	Jeotgalicoccus	Oligoflexus	Shuttleworthia
	[Bacteroides] pectinophilus group	Catenisphaera	Kandleria	Olsenella	Solobacterium
	[Eubacterium] brachy group	Caviibacter	Lachnobacterium	Oribacterium	SP3-e08
	[Eubacterium] cellulosolvens group	Cellulosibacter	Lachnospiraceae AC2044 group	Oscillospira	Sphaerochaeta
	[Eubacterium] hallii group	Christensenellaceae R-7 group	Lachnospiraceae FCS020 group	p-1088-a5 gut group	Sphingobacterium
	[Eubacterium] nodatum group	Chryseobacterium	Lachnospiraceae FE2018 group	P3OB-42	Sphingobium
	[Eubacterium] oxidoreducens group	Clostridium sensu stricto 1	Lachnospiraceae ND3007 group	Papillibacter	Sphingomonas
	[Eubacterium] ruminantium group	Colidextribacter	Lachnospiraceae NK3A20 group	Pasteurella	Spirosoma
	[Eubacterium] saphenum group	Comamonas	Lachnospiraceae NK4A136 group	Pedobacter	Sporobacter
	[Eubacterium] siraeum group	Conchiformibius	Lachnospiraceae NK4B4 group	Peredibacter	Stenotrophomonas
	[Eubacterium] ventriosum group	Coprococcus	Lachnospiraceae UCG-001	Phenylobacterium	Streptococcus
	[Eubacterium] xylanophilum group	Coxiella	Lachnospiraceae UCG-002	Pirellula	Succiniclasticum
	[Ruminococcus] gauvreauii group	CPla-4 termite group	Lachnospiraceae UCG-006	Porphyromonas	Succinimonas
	Abditibacterium	Curtobacterium	Lachnospiraceae UCG-008	possible genus Sk003-Sk004	Succinivibrionaceae UCG-002
	Acetatifactor	Defluviitaleaceae UCG-011	Lachnospiraceae UCG-009	possible genus Sk018	Sutterella
	Acetitomaculum	Denitrobacterium	Lachnospiraceae UCG-010	Prevotella	Suttonella
	Acinetobacter	Desulfobulbus	Lachnospiraceae XPB1014 group	Prevotella_7	Synergistes
	Alloprevotella	Desulfovibrio	Lautropia	Prevotellaceae Ga6A1 group	Syntrophococcus
	Allorhizobium-Neorhizobium-Pararhizobium-Rhizobium	DEV114	M2PT2-76 termite group	Prevotellaceae NK3B31 group	Tardiphaga
	Altererythrobacter	Devosia	Mailhella	Prevotellaceae UCG-001	TM7a
	Alysiella	Dietzia	Mannheimia	Prevotellaceae UCG-003	Treponema
	Amnipila	DNF00809	Marvinbryantia	Prevotellaceae UCG-004	Trueperella
	Anaerobiospirillum	Dyadobacter	Massilia	Prevotellaceae YAB2003 group	Tundrisphaera
	Anaerofustis	Edaphobaculum	Methanobrevibacter	probable genus 10	Tyzzereella
	Anaeroplasma	Elusimicrobium	Methanomicrobium	Pseudobutyrvibrio	U29-B03
	Anaerosporobacter	EMP-G18	Methanosphaera	Pseudomonas	UCG-001
	Anaerovibrio	Endomicrobium	Methylobacterium-Methylorubrum	Psychroglaciecola	UCG-002
	Anaerovorax	Enterorhabdus	Microbacterium	Pyramidobacter	UCG-004
	Asteroleplasma	Erysipelotrichaceae UCG-009	Microvirga	Quinella	UCG-005
	Atopobium	Falsiporphyromonas	MN 122.2a	Raoultibacter	UCG-007
	Bacillus	Family XIII AD3011 group	Mogibacterium	Rhodococcus	UCG-009
	Bacteroides	Family XIII UCG-001	Monoglobus	Rhodopseudomonas	UCG-012
	Bergeyella	FD2005	Moraxella	Rikenellaceae RC9 gut group	V9D2013 group
	Bibersteinia	Fibrobacter	Moryella	Roseburia	Veillonellaceae UCG-001
	Blautia	Flavobacterium	Mucilaginibacter	Roseomonas	Verticella
	Brevundimonas	Flexilinea	Mycobacterium	Ruminiclostridium	Weissella
	Butyrivibrio	Fretibacterium	Mycoplasma	Ruminobacter	Xanthomonas
	CAG-352	Fusobacterium	NED5E9	Ruminococcus	XBB1006
	Campylobacter	hoa5-07d05 gut group	Neorhizobium	Saccharofermentans	Xylophilus
	Candidatus Methanomethylophilus	horsej-a03	NK4A214 group	Schwartzia	Z20
	Candidatus Saccharimonas	Howardella	Novosphingobium	Sediminispirochaeta	NA
	Candidatus Soleiferrea	Incertae Sedis	Nubsella	Selenomonas	