

ct_reorder(Genus, Abundance)

Vishniacozyma	Jeotgalicoccus	Lachnoclostridium	[Ruminococcus] torques group	Lactobacillus
Candidatus Saccharimonas	Acetatifactor	Desulfovibrio	Prevotellaceae UCG-001	Tritrichomonas
Aspergillus	Intestinimonas	Cladosporium	Paeniclostridium	Mycoplasma
Loliinae	Ruminococcus	[Ruminococcus] gnavus group	Sporosarcina	Debaryomyces
Sporobolomyces	Roseburia	Blautia	Ureaplasma	Candidatus Sto
Neophaeosphaeria	Staphylococcus	Oscillibacter	Butyricicoccus	Oxyurida
Spirurida	UCG-003	Lachnospiraceae NK4A136 group	Clostridium sensu stricto 1	[Eubacterium] n
Dorea	[Eubacterium] fissicatena group	Marvinbryantia	HT002	Rikenellaceae F
NK4A214 group	Lachnospiraceae UCG-006	Tuzzerella	Anaeroplasma	Alloprevotella
Helianthus	Enterorhabdus	Enterococcus	Eimeria	Oryza
Brassica	Limosilactobacillus	Odoribacter	Anaerostipes	Escherichia-Sh
Penicillium	Monoglobus	Parabacteroides	Mucispirillum	Saccharomyces
Microdochium	Incertae Sedis	A2	Coriobacteriaceae UCG-002	Mammalia
[Eubacterium] ventriosum group	Ligilactobacillus	Rikenella	Veillonella	[Clostridium] inn
UCG-009	GCA-900066575	Alistipes	Muribaculum	Vigna
[Eubacterium] xylanophilum group	Lachnospiraceae UCG-001	Helicobacter	Erysipelatoclostridium	Brachypodium
[Eubacterium] siraeum group	Anaerotruncus	Lachnospiraceae FCS020 group	Bacteroides	Akkermansia
ASF356	Romboutsia	Colidextribacter	Bilophila	Alloteropsis
[Eubacterium] oxidoreducens group	Paraprevotella	Terrisporobacter	Parasutterella	NA