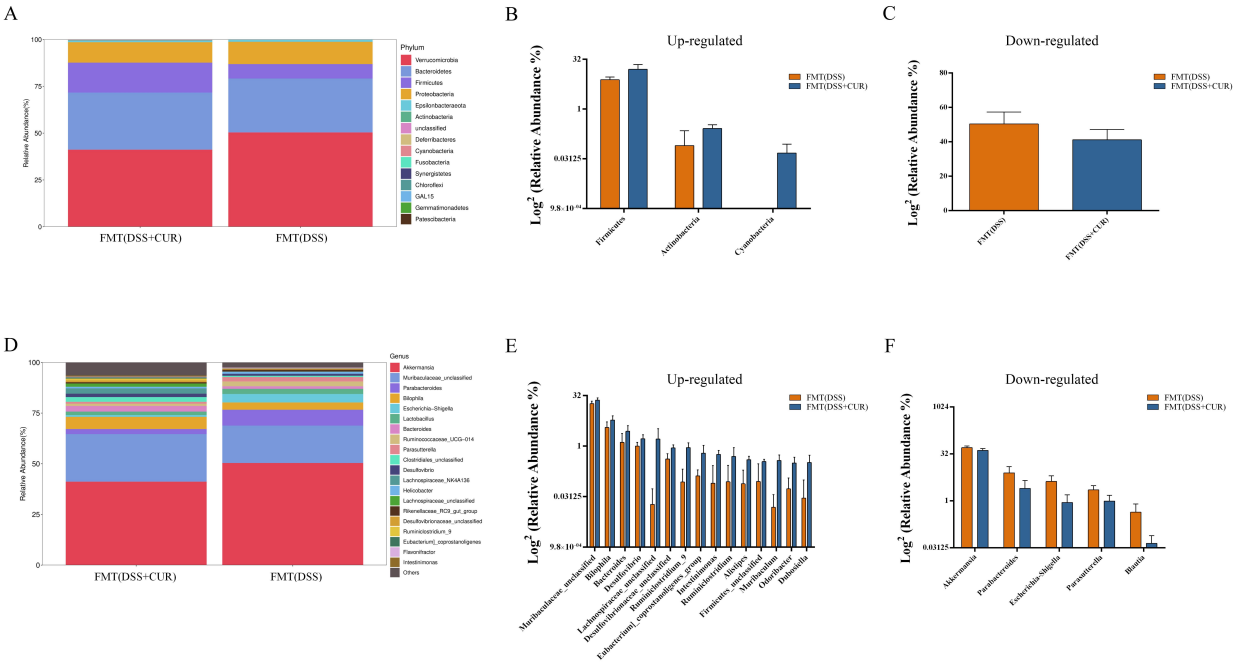
A
Relative Abundance(%)
FMT(DSS+CUR)
FMT(DSS)
Phylum
Verrucomicrobia
Bacteroidetes
Firmicutes
Proteobacteria
Epsilonbacteraeota
Actinobacteria
unclassified
Deferribacteres
Cyanobacteria
Fusobacteria
Synergistetes
Chloroflexi
GAL15
Gemmatimonadetes
Patescibacteria
B
Up-regulated
Log2 (Relative Abundance %)
FMT(DSS)
FMT(DSS+CUR)
Firmicutes
Actinobacteria
Cyanobacteria
C
Down-regulated
Log2 (Relative Abundance %)
FMT(DSS)
FMT(DSS+CUR)
D
Relative Abundance(%)
FMT(DSS+CUR)
FMT(DSS)
Genus
Akkermansia
Muribaculaceae_unclassified
Parabacteroides
Bilophila
Escherichia-Shigella
Lactobacillus
Bacteroides
Ruminococcaceae_UCG-014
Parasutterella
Clostridiales_unclassified
Desulfovibrio
Lachnospiraceae_NK4A136
Helicobacter
Lachnospiraceae_unclassified
Rikenellaceae_RC9_gut_group
Desulfovibrionaceae_unclassified
Ruminiclostridium_9
Eubacterium[_coprostanoligenes
Flavonifractor
Intestinimonas
Others
E
Up-regulated
Log2 (Relative Abundance %)
FMT(DSS)
FMT(DSS+CUR)
Marinifilaceae_unclassified
Bilophila
Bacteroides
Desulfovibrio
Lachnospiraceae_unclassified
Desulfovibrionaceae_unclassified
Ruminiclostridium_9
Eubacterium[_coprostanoligenes_group
Intestinimonas
Ruminiclostridium
Alistipes
Firmicutes_unclassified
Marinifilum
Odoribacter
Dubosiella
F
Down-regulated
Log2 (Relative Abundance %)
FMT(DSS)
FMT(DSS+CUR)
Akkermansia
Parabacteroides
Escherichia-Shigella
Parasutterella
Blautia