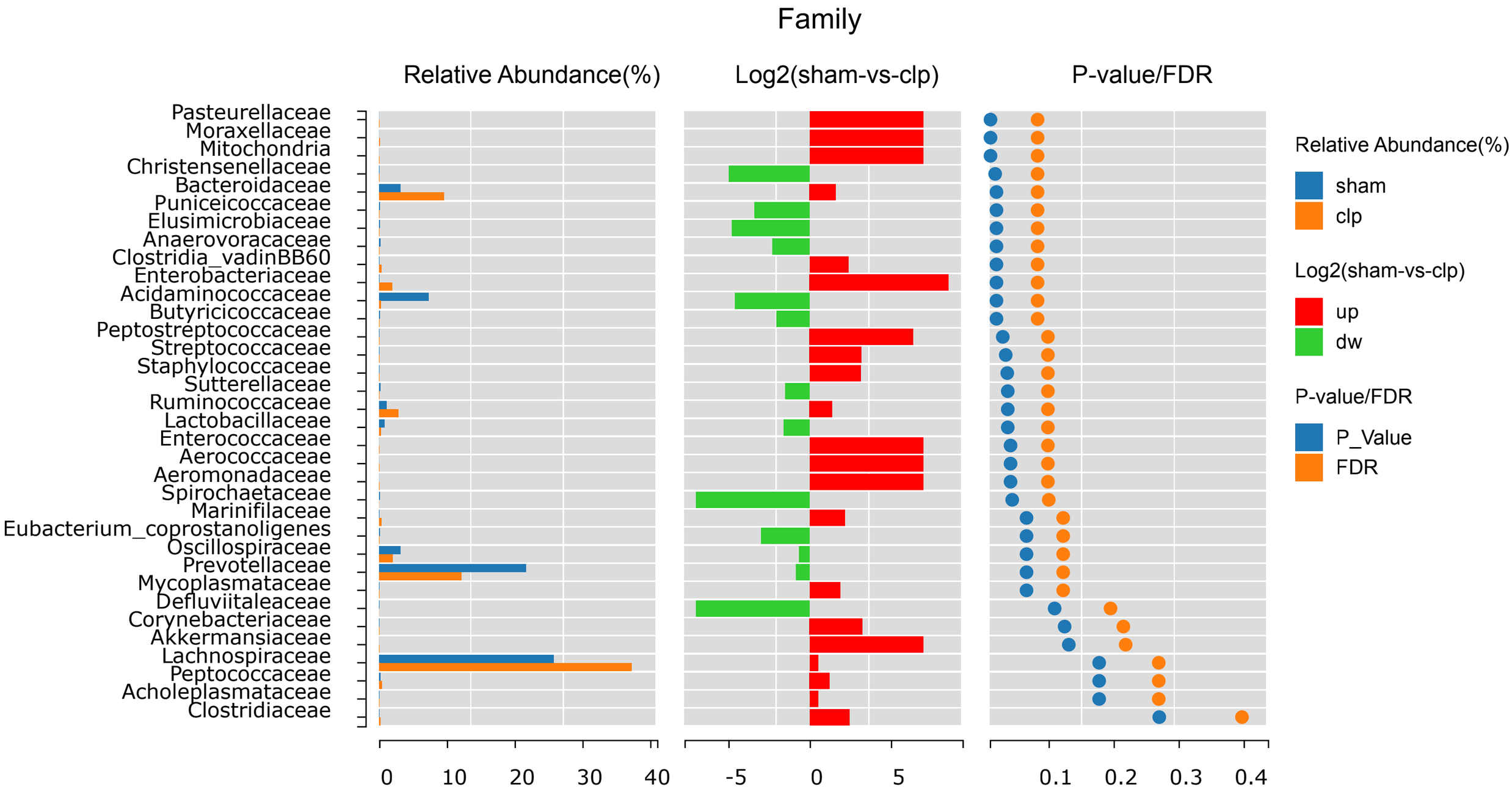
Family
Relative Abundance(%)
Log2(sham-vs-clp)
P-value/FDR
Pasteurellaceae
Moraxellaceae
Mitochondria
Christensenellaceae
Bacteroidaceae
Puniceicoccaceae
Elusimicrobiaceae
Anaerovoracaceae
Clostridia_vadinBB60
Enterobacteriaceae
Acidaminococcaceae
Butyricicoccaceae
Peptostreptococcaceae
Streptococcaceae
Staphylococcaceae
Sutterellaceae
Ruminococcaceae
Lactobacillaceae
Enterococcaceae
Aerococcaceae
Aeromonadaceae
Spirochaetaceae
Marinifilaceae
Eubacterium_coprostanoligenes
Oscillospiraceae
Prevotellaceae
Mycoplasmataceae
Defluviitaleaceae
Corynebacteriaceae
Akkermansiaceae
Lachnospiraceae
Peptococcaceae
Acholeplasmataceae
Clostridiaceae
0
10
20
30
40
-5
0
5
0.1
0.2
0.3
0.4
Relative Abundance(%)
sham
clp
Log2(sham-vs-clp)
up
dw
P-value/FDR
P_Value
FDR