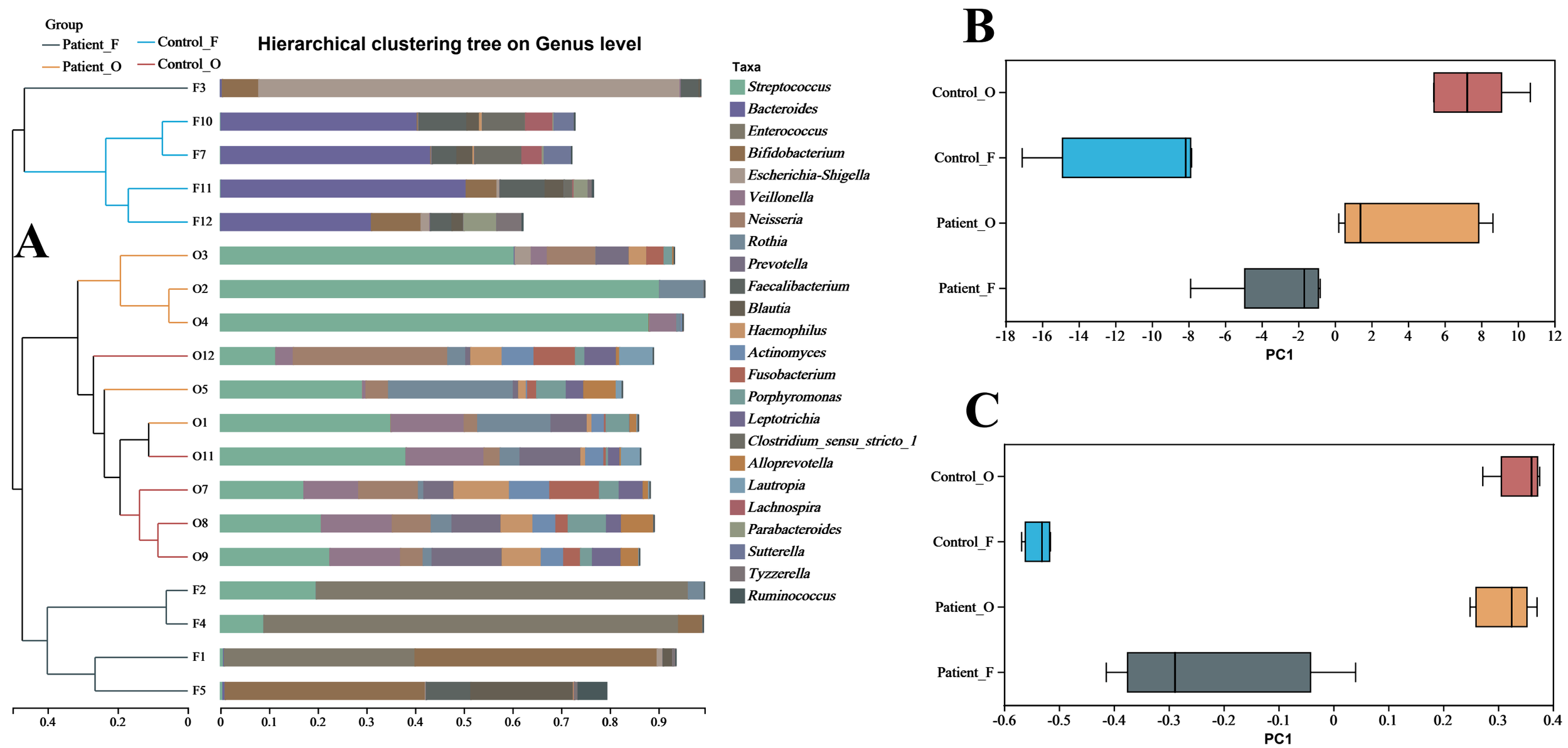
A
Group
Patient_F
Control_F
Patient_O
Control_O
Hierarchical clustering tree on Genus level
F3
F10
F7
F11
F12
O3
O2
O4
O12
O5
O1
O11
O7
O8
O9
F2
F4
F1
F5
0.4
0.2
0
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Taxa
Streptococcus
Bacteroides
Enterococcus
Bifidobacterium
Escherichia-Shigella
Veillonella
Neisseria
Rothia
Prevotella
Faecalibacterium
Blautia
Haemophilus
Actinomyces
Fusobacterium
Porphyromonas
Leptotrichia
Clostridium_sensu_stricto_1
Alloprevotella
Lautropia
Lachnospira
Parabacteroides
Sutterella
Tyzzerella
Ruminococcus
B
Control_O
Control_F
Patient_O
Patient_F
-18
-16
-14
-12
-10
-8
-6
-4
-2
0
2
4
6
8
10
12
PC1
C
Control_O
Control_F
Patient_O
Patient_F
-0.6
-0.5
-0.4
-0.3
-0.2
-0.1
0
0.1
0.2
0.3
0.4
PC1