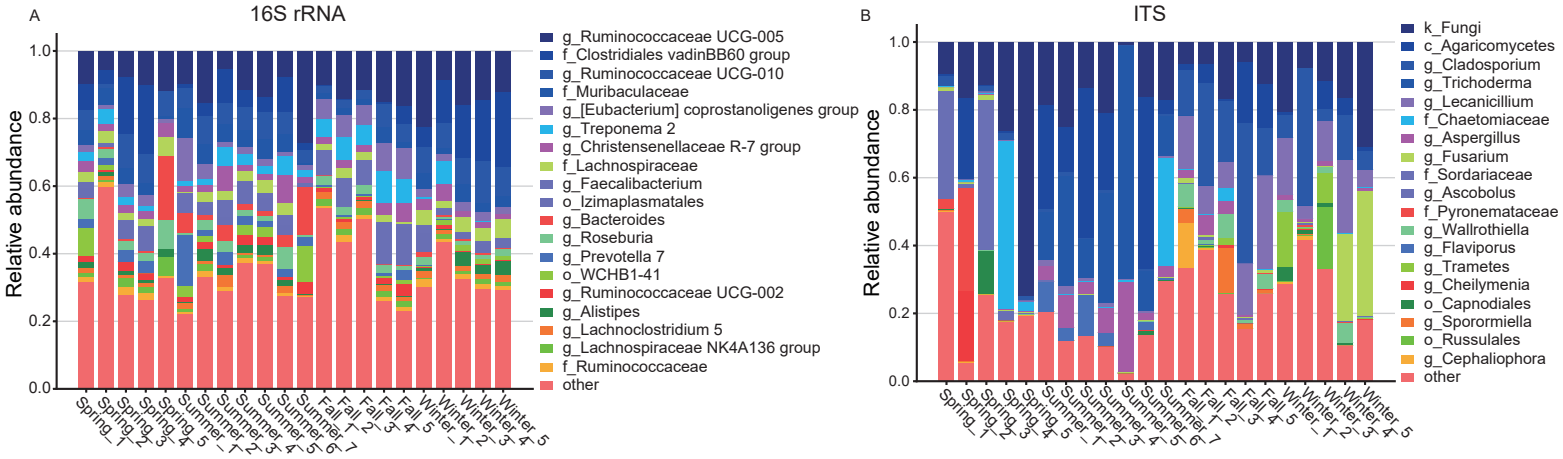
A
16S rRNA
Relative abundance
1.0
0.8
0.6
0.4
0.2
0.0
Spring_1
Spring_2
Spring_3
Spring_4
Summer_5
Summer_1
Summer_2
Summer_3
Summer_4
Summer_5
Summer_6
Fall_1
Fall_2
Fall_3
Fall_4
Fall_5
Winter_1
Winter_2
Winter_3
Winter_4
Winter_5
g_Ruminococcaceae UCG-005
f_Clostridiales vadinBB60 group
g_Ruminococcaceae UCG-010
f_Muribaculaceae
g_[Eubacterium] coprostanoligenes group
g_Treponema 2
g_Christensenellaceae R-7 group
f_Lachnospiraceae
g_Faecalibacterium
o_Izimaplasmatales
g_Bacteroides
g_Roseburia
g_Prevotella 7
o_WCHB1-41
g_Ruminococcaceae UCG-002
g_Alistipes
g_Lachnoclostridium 5
g_Lachnospiraceae NK4A136 group
f_Ruminococcaceae
other
B
ITS
Relative abundance
1.0
0.8
0.6
0.4
0.2
0.0
Spring_1
Spring_2
Spring_3
Spring_4
Summer_5
Summer_1
Summer_2
Summer_3
Summer_4
Summer_5
Summer_6
Fall_1
Fall_2
Fall_3
Fall_4
Fall_5
Winter_1
Winter_2
Winter_3
Winter_4
Winter_5
k_Fungi
c_Agaricomycetes
g_Cladosporium
g_Trichoderma
g_Lecanicillium
f_Chaetomiaceae
g_Aspergillus
g_Fusarium
f_Sordariaceae
g_Ascobolus
f_Pyronemataceae
g_Wallrothiella
g_Flaviporus
g_Trametes
g_Cheilymenia
o_Capnodiales
g_Sporormiella
o_Russulales
g_Cephaliophora
other