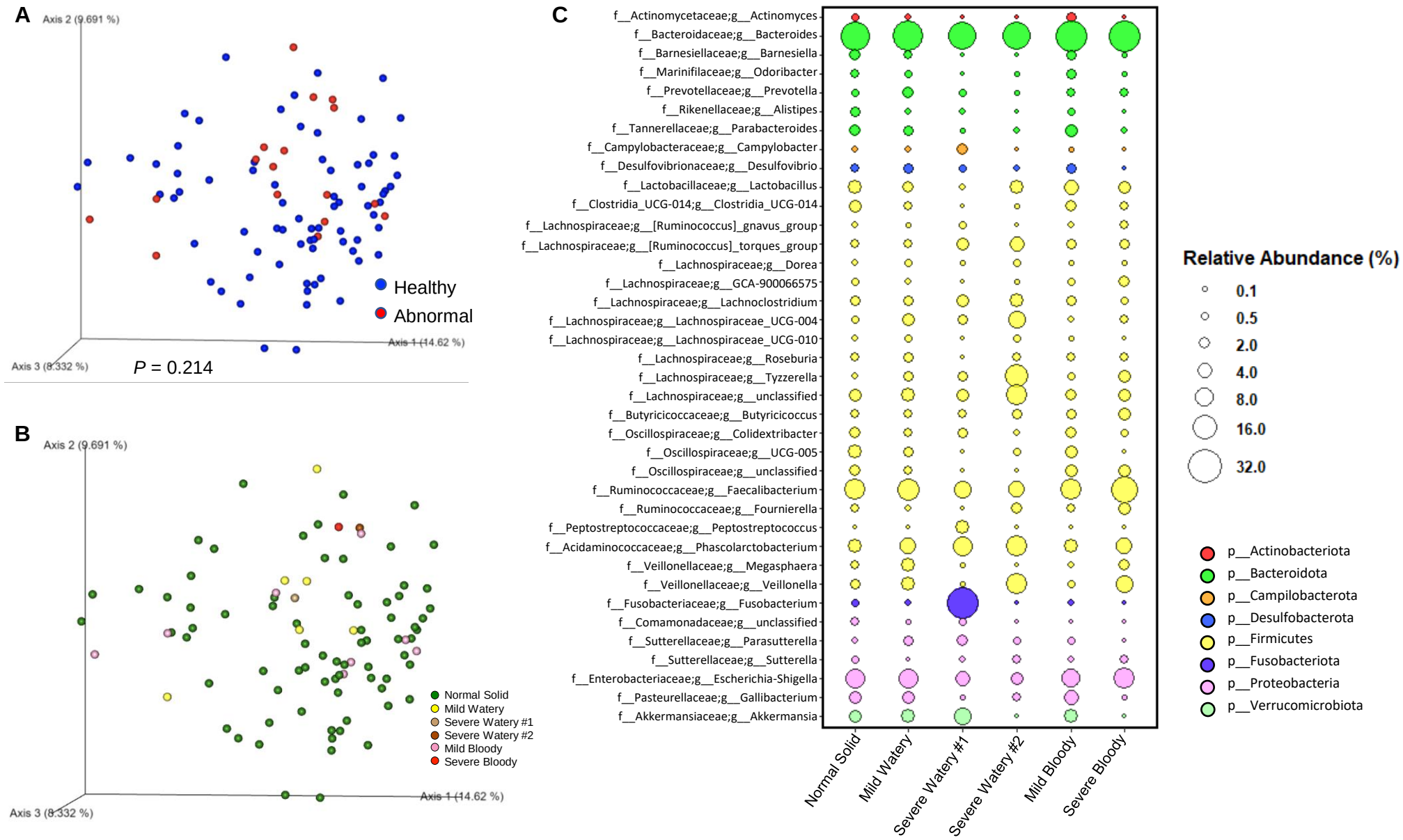
A
Axis 2 (9.691 %)
Axis 1 (14.62 %)
Axis 3 (8.332 %)
Healthy
Abnormal
P = 0.214
B
Axis 2 (9.691 %)
Axis 1 (14.62 %)
Axis 3 (8.332 %)
Normal Solid
Mild Watery
Severe Watery #1
Severe Watery #2
Mild Bloody
Severe Bloody
C
f__Actinomycetaceae;g__Actinomyces
f__Bacteroidaceae;g__Bacteroides
f__Barnesiellaceae;g__Barnesiella
f__Marinifilaceae;g__Odoribacter
f__Prevotellaceae;g__Prevotella
f__Rikenellaceae;g__Alistipes
f__Tannerellaceae;g__Parabacteroides
f__Campylobacteraceae;g__Campylobacter
f__Desulfovibrionaceae;g__Desulfovibrio
f__Lactobacillaceae;g__Lactobacillus
f__Clostridia_UCG-014;g__Clostridia_UCG-014
f__Lachnospiraceae;g__[Ruminococcus]_gnavus_group
f__Lachnospiraceae;g__[Ruminococcus]_torques_group
f__Lachnospiraceae;g__Dorea
f__Lachnospiraceae;g__GCA-900066575
f__Lachnospiraceae;g__Lachnoclostridium
f__Lachnospiraceae;g__Lachnospiraceae_UCG-004
f__Lachnospiraceae;g__Lachnospiraceae_UCG-010
f__Lachnospiraceae;g__Roseburia
f__Lachnospiraceae;g__Tyzzerella
f__Lachnospiraceae;g__unclassified
f__Butyricicoccaceae;g__Butyricicoccus
f__Oscillospiraceae;g__Colidextribacter
f__Oscillospiraceae;g__UCG-005
f__Oscillospiraceae;g__unclassified
f__Ruminococcaceae;g__Faecalibacterium
f__Ruminococcaceae;g__Fournierella
f__Peptostreptococcaceae;g__Peptostreptococcus
f__Acidaminococcaceae;g__Phascolarctobacterium
f__Veillonellaceae;g__Megasphaera
f__Veillonellaceae;g__Veillonella
f__Fusobacteriaceae;g__Fusobacterium
f__Comamonadaceae;g__unclassified
f__Sutterellaceae;g__Parasutterella
f__Sutterellaceae;g__Sutterella
f__Enterobacteriaceae;g__Escherichia-Shigella
f__Pasteurellaceae;g__Gallibacterium
f__Akkermansiaceae;g__Akkermansia
Normal Solid
Mild Watery
Severe Watery #1
Severe Watery #2
Mild Bloody
Severe Bloody
Relative Abundance (%)
0.1
0.5
2.0
4.0
8.0
16.0
32.0
p__Actinobacteriota
p__Bacteroidota
p__Campilobacterota
p__Desulfobacterota
p__Firmicutes
p__Fusobacteriota
p__Proteobacteria
p__Verrucomicrobiota