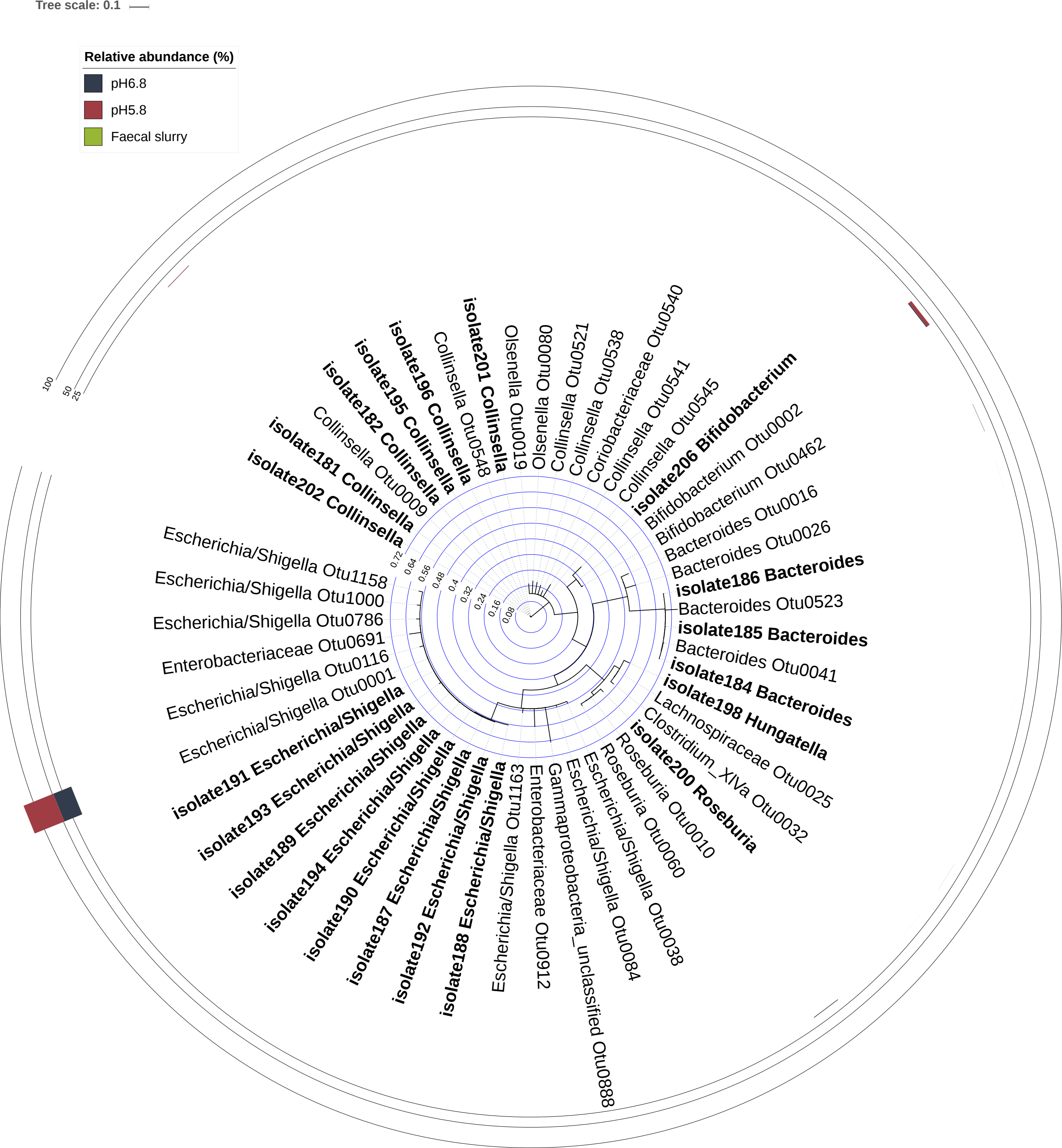
Tree scale: 0.1
Relative abundance (%)
pH6.8
pH5.8
Faecal slurry
100
50
25
0.08
0.16
0.24
0.32
0.4
0.48
0.56
0.64
0.72
Escherichia/Shigella Otu1158
Escherichia/Shigella Otu1000
Escherichia/Shigella Otu0786
Enterobacteriaceae Otu0691
Escherichia/Shigella Otu0116
Escherichia/Shigella Otu0001
isolate191 Escherichia/Shigella
isolate193 Escherichia/Shigella
isolate189 Escherichia/Shigella
isolate194 Escherichia/Shigella
isolate190 Escherichia/Shigella
isolate187 Escherichia/Shigella
isolate192 Escherichia/Shigella
isolate188 Escherichia/Shigella
Escherichia/Shigella Otu1163
Enterobacteriaceae Otu0912
Gammaproteobacteria_unclassified Otu0888
Escherichia/Shigella Otu0084
Escherichia/Shigella Otu0038
Roseburia Otu0060
Roseburia Otu0010
isolate200 Roseburia
Clostridium_XlVa Otu0032
Lachnospiraceae Otu0025
isolate198 Hungatella
isolate184 Bacteroides
Bacteroides Otu0041
isolate185 Bacteroides
Bacteroides Otu0523
isolate186 Bacteroides
Bacteroides Otu0026
Bacteroides Otu0016
Bifidobacterium Otu0462
Bifidobacterium Otu0002
isolate206 Bifidobacterium
Collinsella Otu0541
Coriobacteriaceae Otu0540
Collinsella Otu0538
Collinsella Otu0521
Olsenella Otu0080
Olsenella Otu0019
isolate201 Collinsella
Collinsella Otu0548
isolate196 Collinsella
isolate195 Collinsella
isolate182 Collinsella
Collinsella Otu0009
isolate181 Collinsella
isolate202 Collinsella