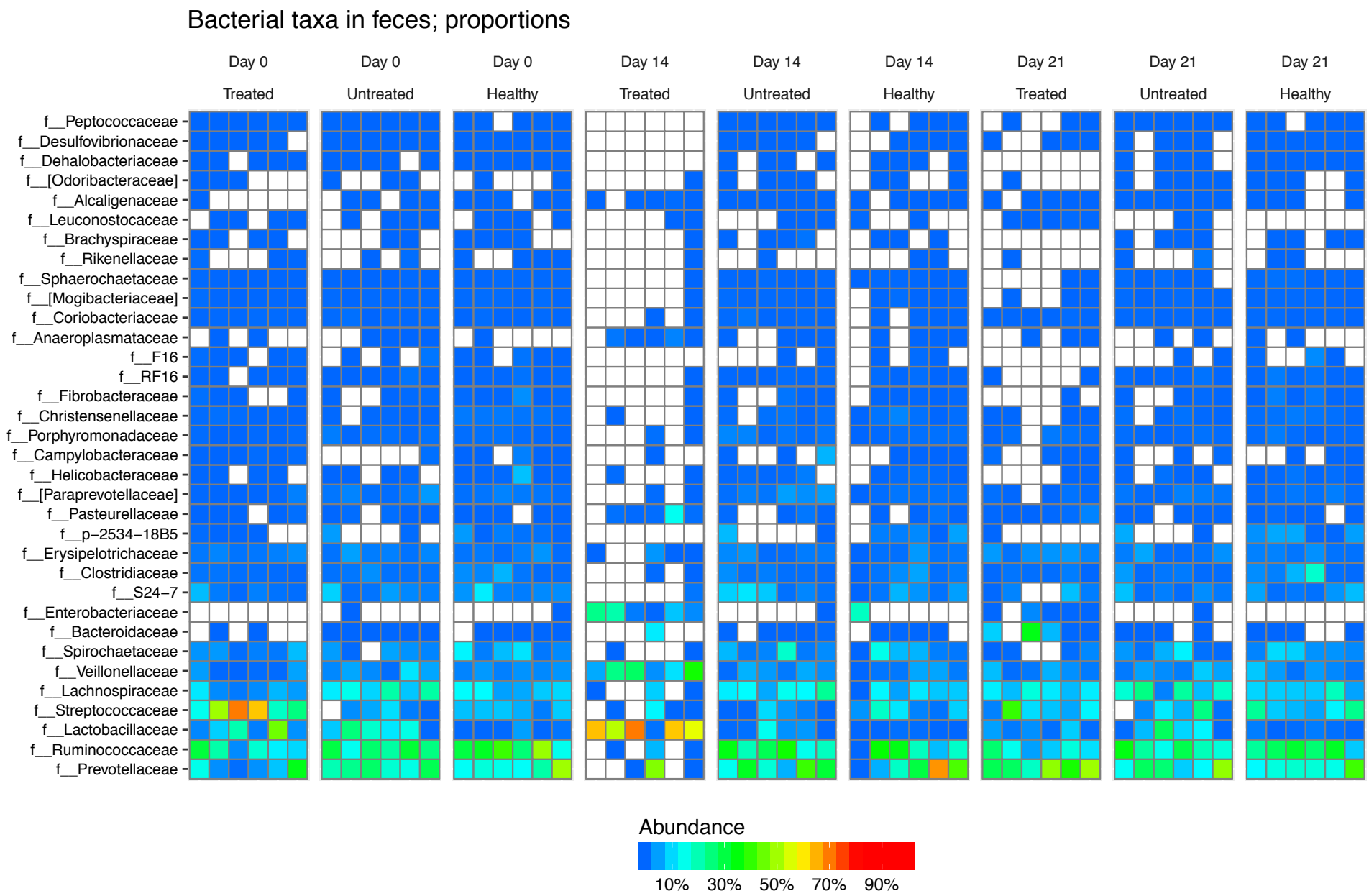
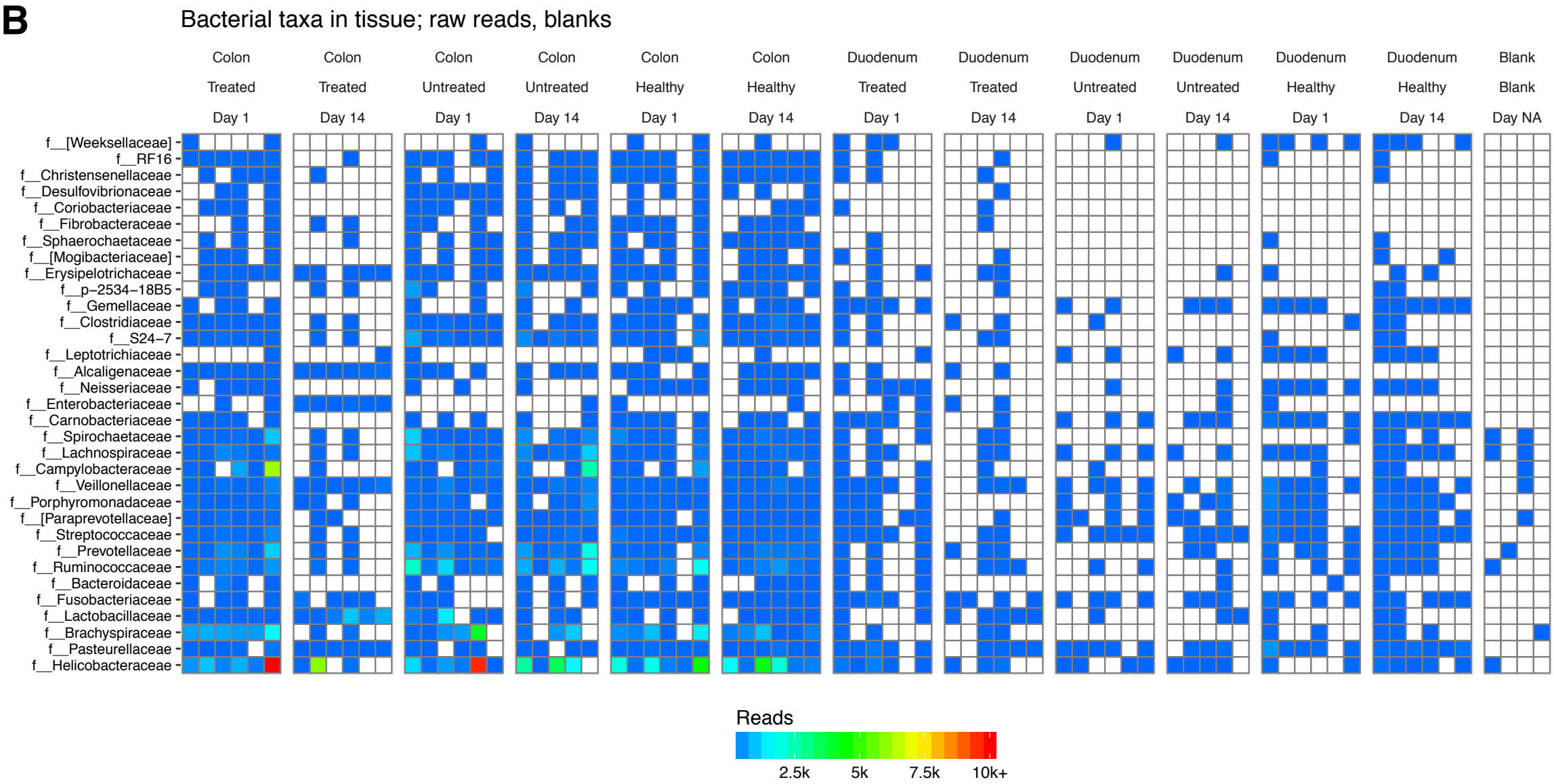
A


Bacterial taxa in feces; proportions
Day 0 Treated
Day 0 Untreated
Day 0 Healthy
Day 14 Treated
Day 14 Untreated
Day 14 Healthy
Day 21 Treated
Day 21 Untreated
Day 21 Healthy
f__Peptococcaceae
f__Desulfovibrionaceae
f__Dehalobacteriaceae
f__[Odoribacteraceae]
f__Alcaligenaceae
f__Leuconostocaceae
f__Brachyspiraceae
f__Rikenellaceae
f__Sphaerochaetaceae
f__[Mogibacteriaceae]
f__Coriobacteriaceae
f__Anaeroplasmataceae
f__F16
f__RF16
f__Fibrobacteraceae
f__Christensenellaceae
f__Porphyromonadaceae
f__Campylobacteraceae
f__Helicobacteraceae
f__[Paraprevotellaceae]
f__Pasteurellaceae
f__p−2534−18B5
f__Erysipelotrichaceae
f__Clostridiaceae
f__S24−7
f__Enterobacteriaceae
f__Bacteroidaceae
f__Spirochaetaceae
f__Veillonellaceae
f__Lachnospiraceae
f__Streptococcaceae
f__Lactobacillaceae
f__Ruminococcaceae
f__Prevotellaceae
Abundance
10%
30%
50%
70%
90%

B


Bacterial taxa in tissue; raw reads, blanks
Colon Treated Day 1
Colon Treated Day 14
Colon Untreated Day 1
Colon Untreated Day 14
Colon Healthy Day 1
Colon Healthy Day 14
Duodenum Treated Day 1
Duodenum Treated Day 14
Duodenum Untreated Day 1
Duodenum Untreated Day 14
Duodenum Healthy Day 1
Duodenum Healthy Day 14
Blank Blank Day NA
f__[Weeksellaceae]
f__RF16
f__Christensenellaceae
f__Desulfovibrionaceae
f__Coriobacteriaceae
f__Fibrobacteraceae
f__Sphaerochaetaceae
f__[Mogibacteriaceae]
f__Erysipelotrichaceae
f__p−2534−18B5
f__Gemellaceae
f__Clostridiaceae
f__S24−7
f__Leptotrichiaceae
f__Alcaligenaceae
f__Neisseriaceae
f__Enterobacteriaceae
f__Carnobacteriaceae
f__Spirochaetaceae
f__Lachnospiraceae
f__Campylobacteraceae
f__Veillonellaceae
f__Porphyromonadaceae
f__[Paraprevotellaceae]
f__Streptococcaceae
f__Prevotellaceae
f__Ruminococcaceae
f__Bacteroidaceae
f__Fusobacteriaceae
f__Lactobacillaceae
f__Brachyspiraceae
f__Pasteurellaceae
f__Helicobacteraceae
Reads
2.5k
5k
7.5k
10k+