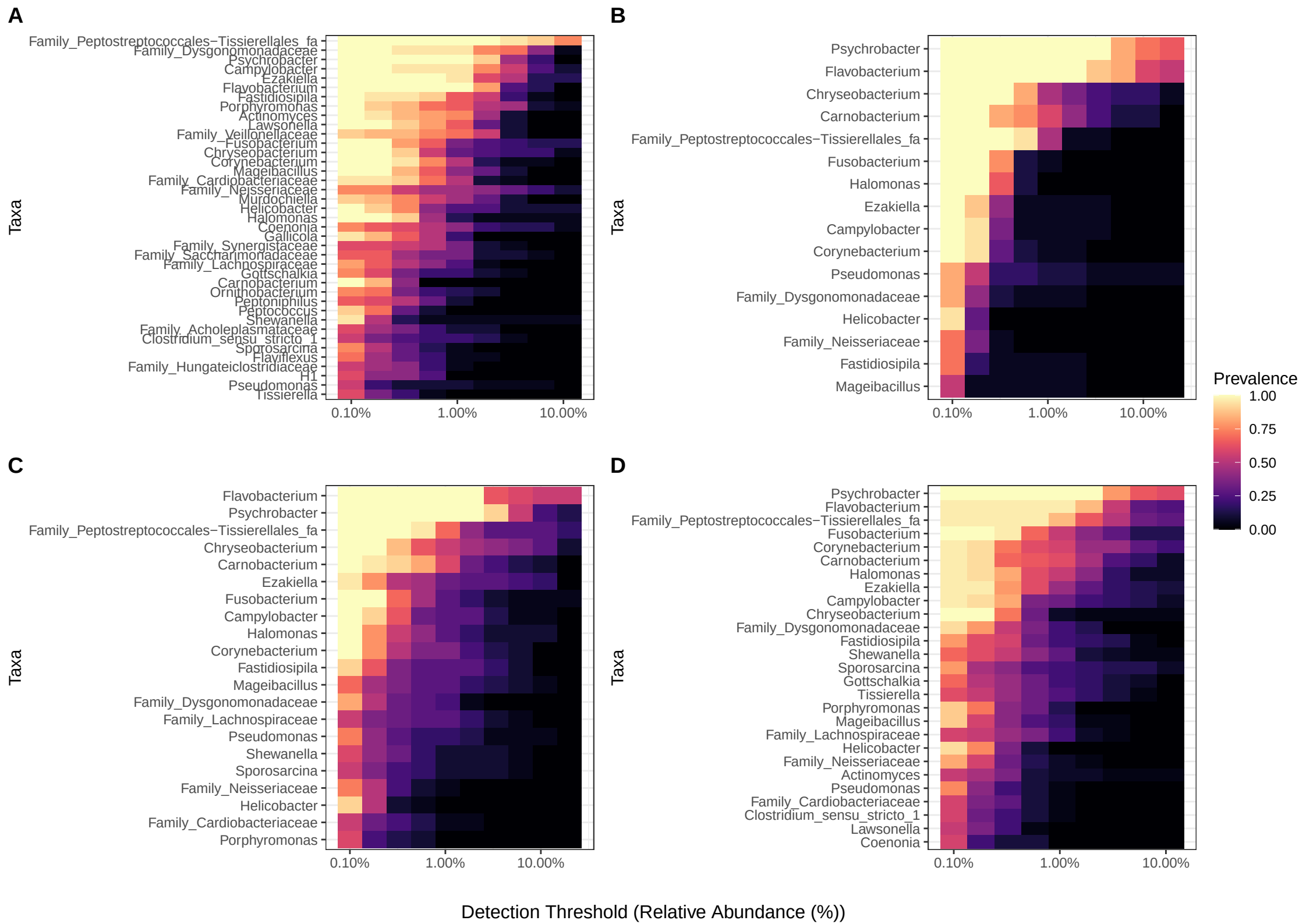

A
Family_Peptostreptococcales−Tissierellales_fa
Family_Dysgonomonadaceae
Psychrobacter
Campylobacter
Ezakiella
Flavobacterium
Fastidiosipila
Porphyromonas
Actinomyces
Lawsonella
Family_Veillonellaceae
Fusobacterium
Chryseobacterium
Corynebacterium
Mageibacillus
Family_Cardiobacteriaceae
Family_Neisseriaceae
Murdochiella
Helicobacter
Halomonas
Coenonia
Gallicola
Family_Synergistaceae
Family_Saccharimonadaceae
Family_Lachnospiraceae
Gottschalkia
Carnobacterium
Ornithobacterium
Peptoniphilus
Peptococcus
Shewanella
Family_Acholeplasmataceae
Clostridium_sensu_stricto_1
Sporosarcina
Flaviflexus
Family_Hungateiclostridiaceae
H1
Pseudomonas
Tissierella
Taxa
0.10%
1.00%
10.00%
B
Psychrobacter
Flavobacterium
Chryseobacterium
Carnobacterium
Family_Peptostreptococcales−Tissierellales_fa
Fusobacterium
Halomonas
Ezakiella
Campylobacter
Corynebacterium
Pseudomonas
Family_Dysgonomonadaceae
Helicobacter
Family_Neisseriaceae
Fastidiosipila
Mageibacillus
Taxa
0.10%
1.00%
10.00%
Prevalence
1.00
0.75
0.50
0.25
0.00
C
Flavobacterium
Psychrobacter
Family_Peptostreptococcales−Tissierellales_fa
Chryseobacterium
Carnobacterium
Ezakiella
Fusobacterium
Campylobacter
Halomonas
Corynebacterium
Fastidiosipila
Mageibacillus
Family_Dysgonomonadaceae
Family_Lachnospiraceae
Pseudomonas
Shewanella
Sporosarcina
Family_Neisseriaceae
Helicobacter
Family_Cardiobacteriaceae
Porphyromonas
Taxa
0.10%
1.00%
10.00%
D
Psychrobacter
Flavobacterium
Family_Peptostreptococcales−Tissierellales_fa
Fusobacterium
Corynebacterium
Carnobacterium
Halomonas
Ezakiella
Campylobacter
Chryseobacterium
Family_Dysgonomonadaceae
Fastidiosipila
Shewanella
Sporosarcina
Gottschalkia
Tissierella
Porphyromonas
Mageibacillus
Family_Lachnospiraceae
Helicobacter
Family_Neisseriaceae
Actinomyces
Pseudomonas
Family_Cardiobacteriaceae
Clostridium_sensu_stricto_1
Lawsonella
Coenonia
Taxa
0.10%
1.00%
10.00%
Detection Threshold (Relative Abundance (%))