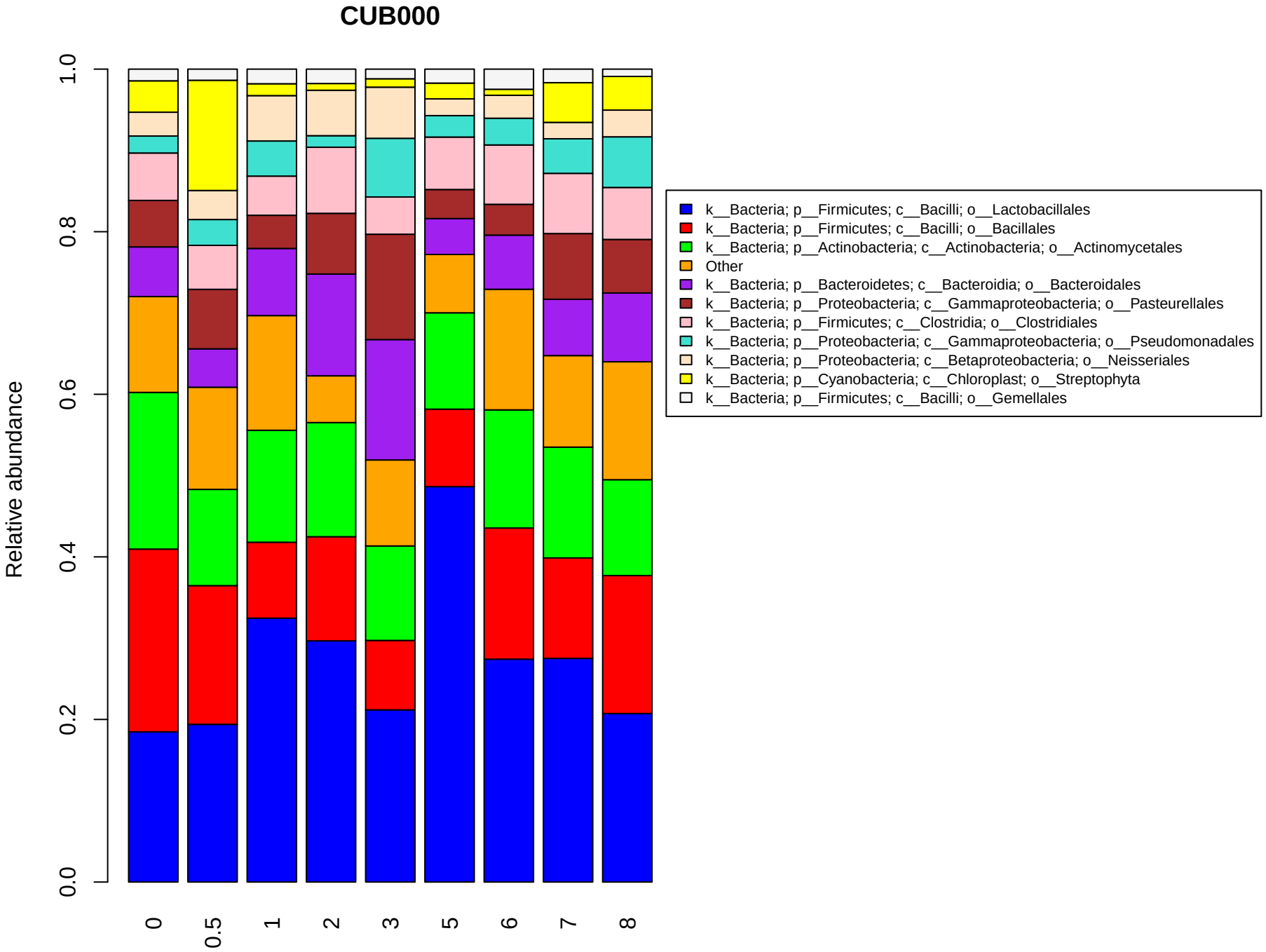
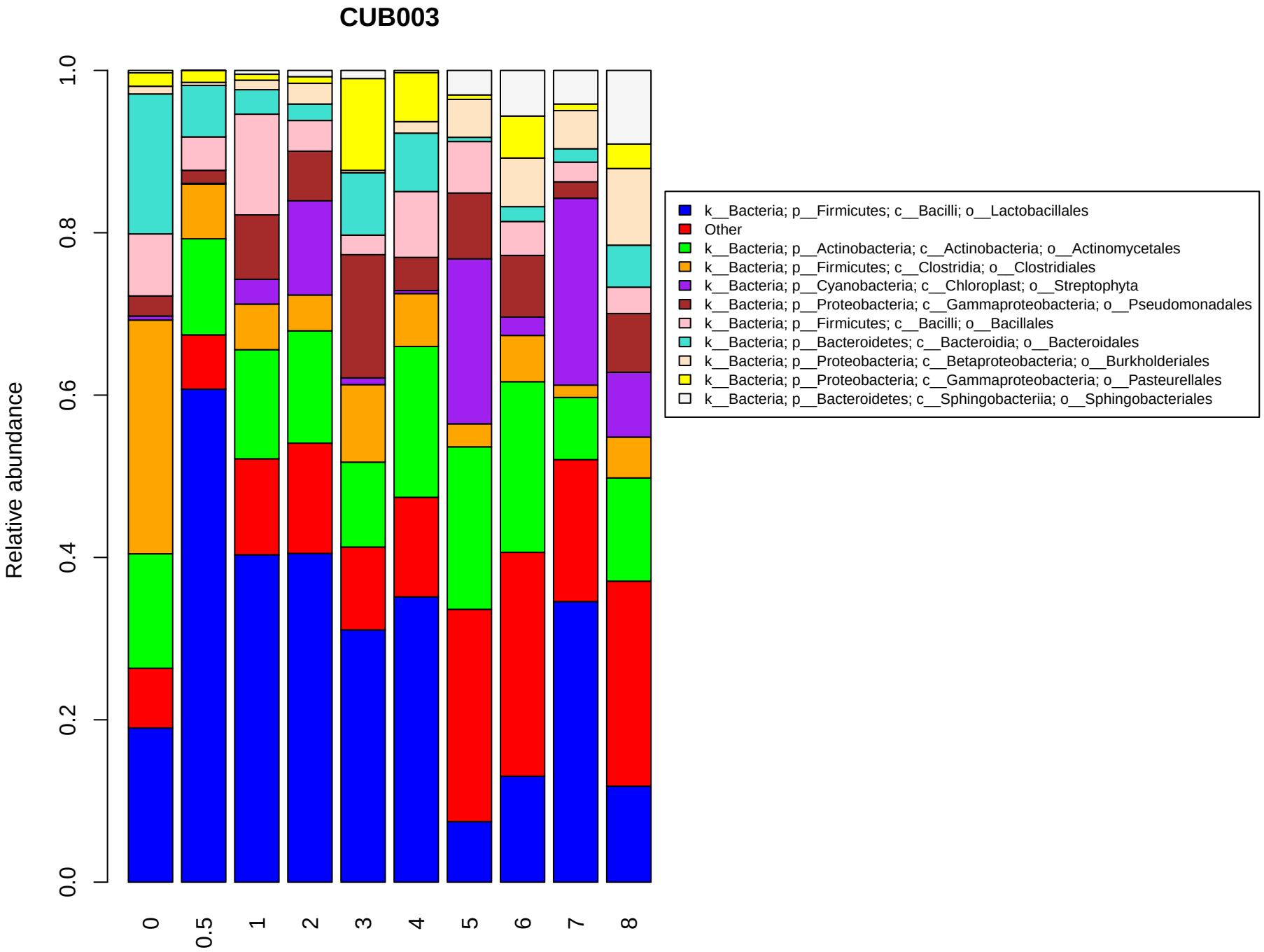

CUB000
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales


CUB003
Relative abundance
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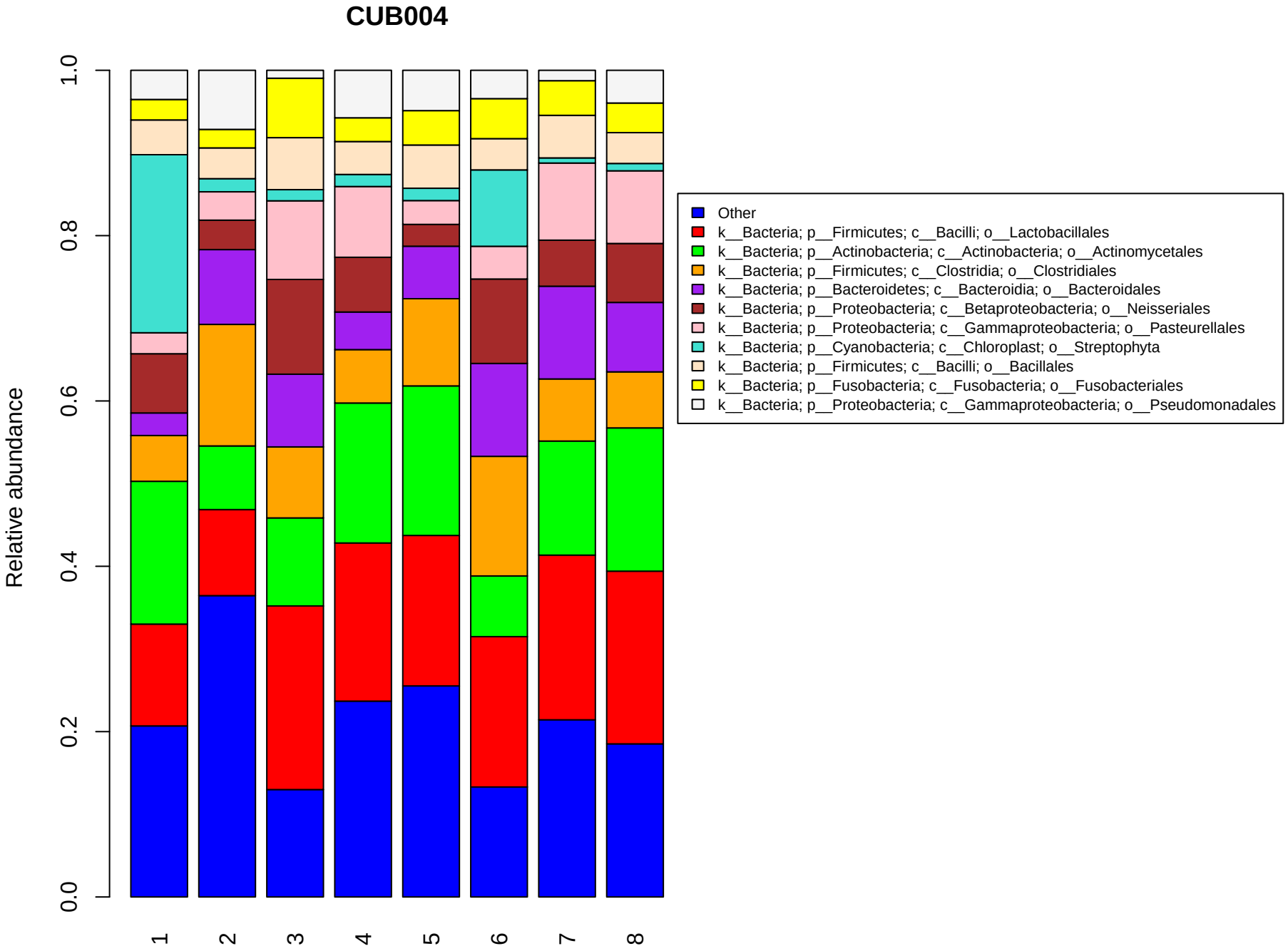
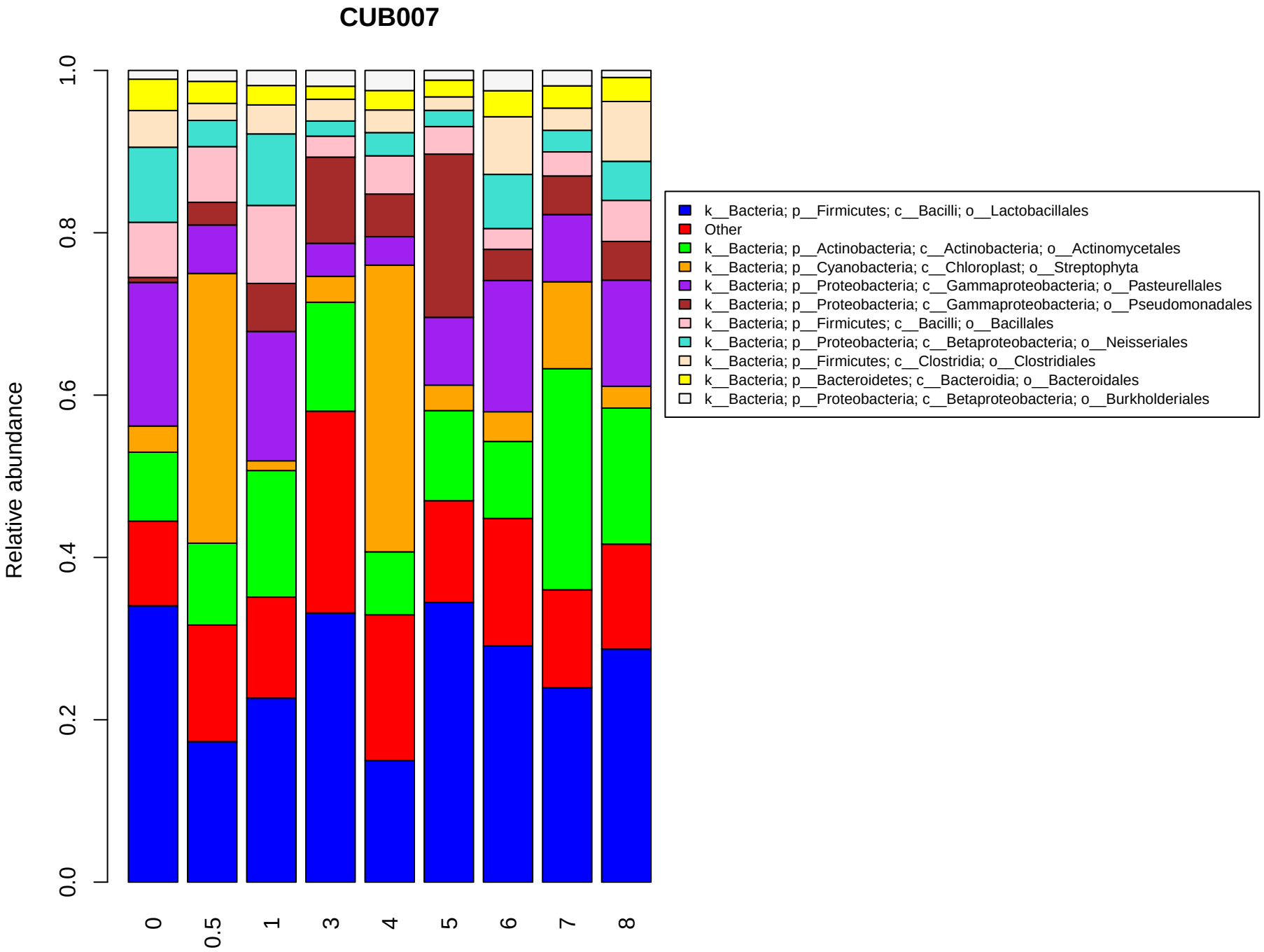
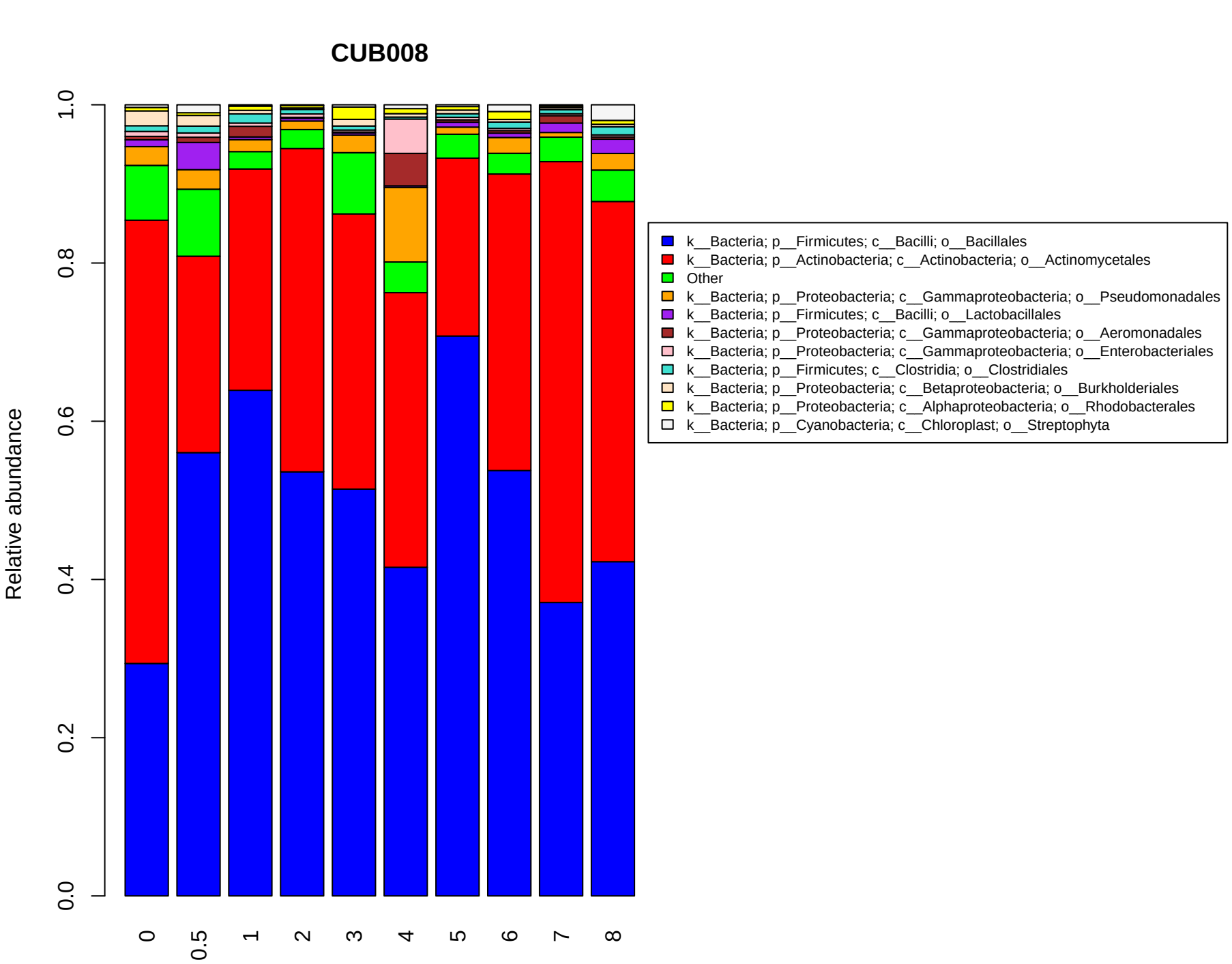
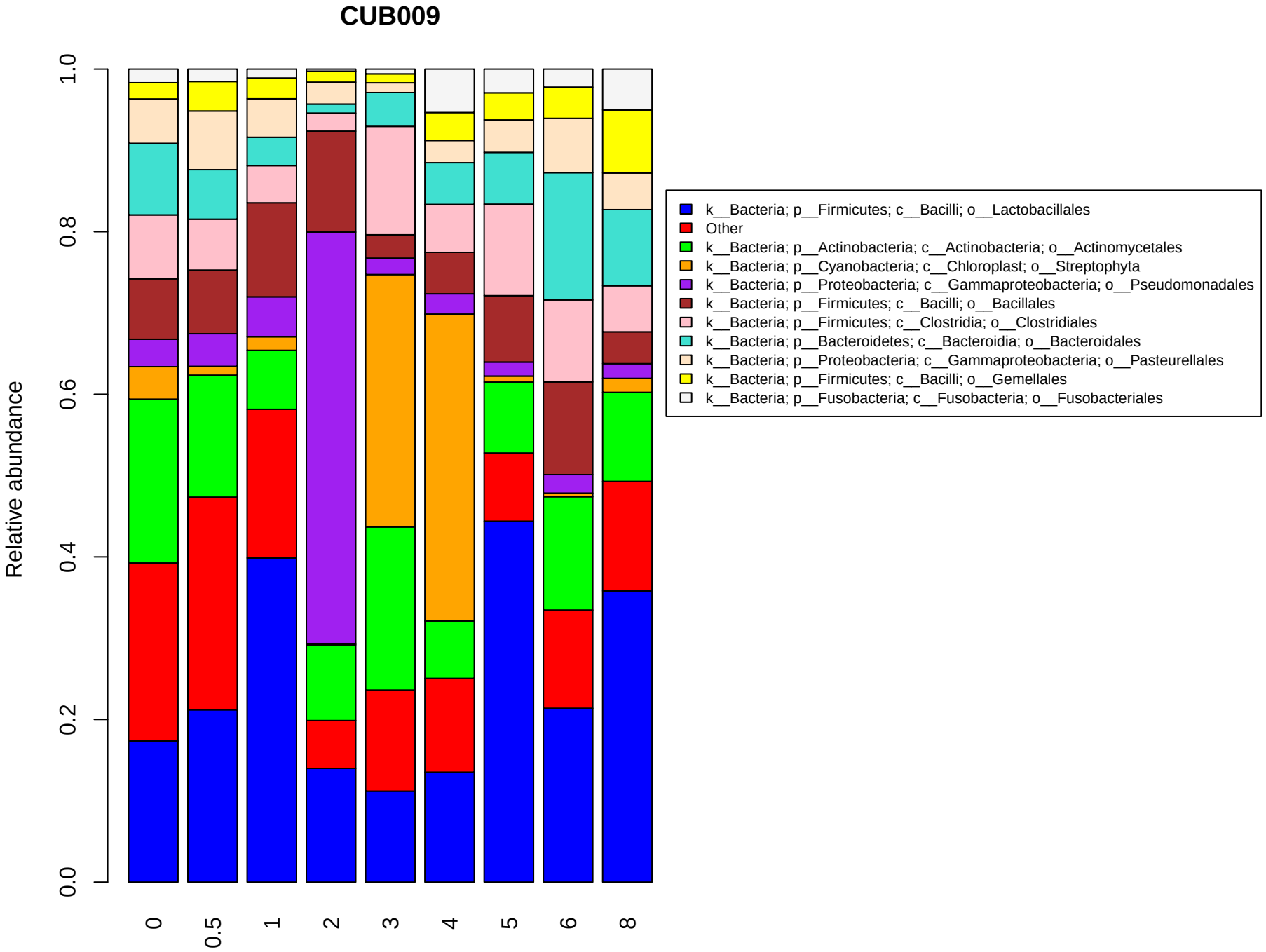
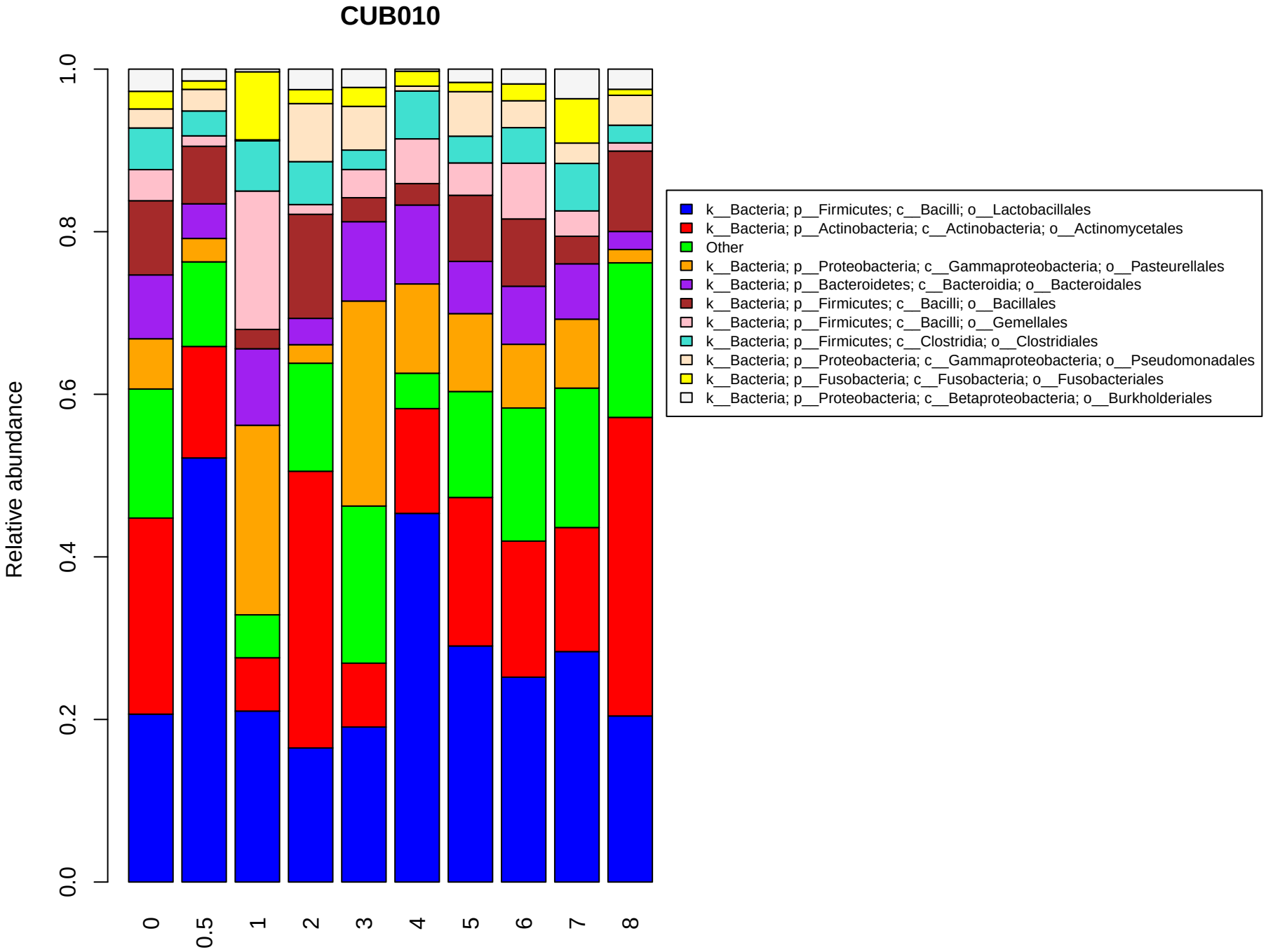
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales


CUB004
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
1
2
3
4
5
6
7
8
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales


CUB007
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


CUB008
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodobacterales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta


CUB009
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales


CUB010
Relative abundance
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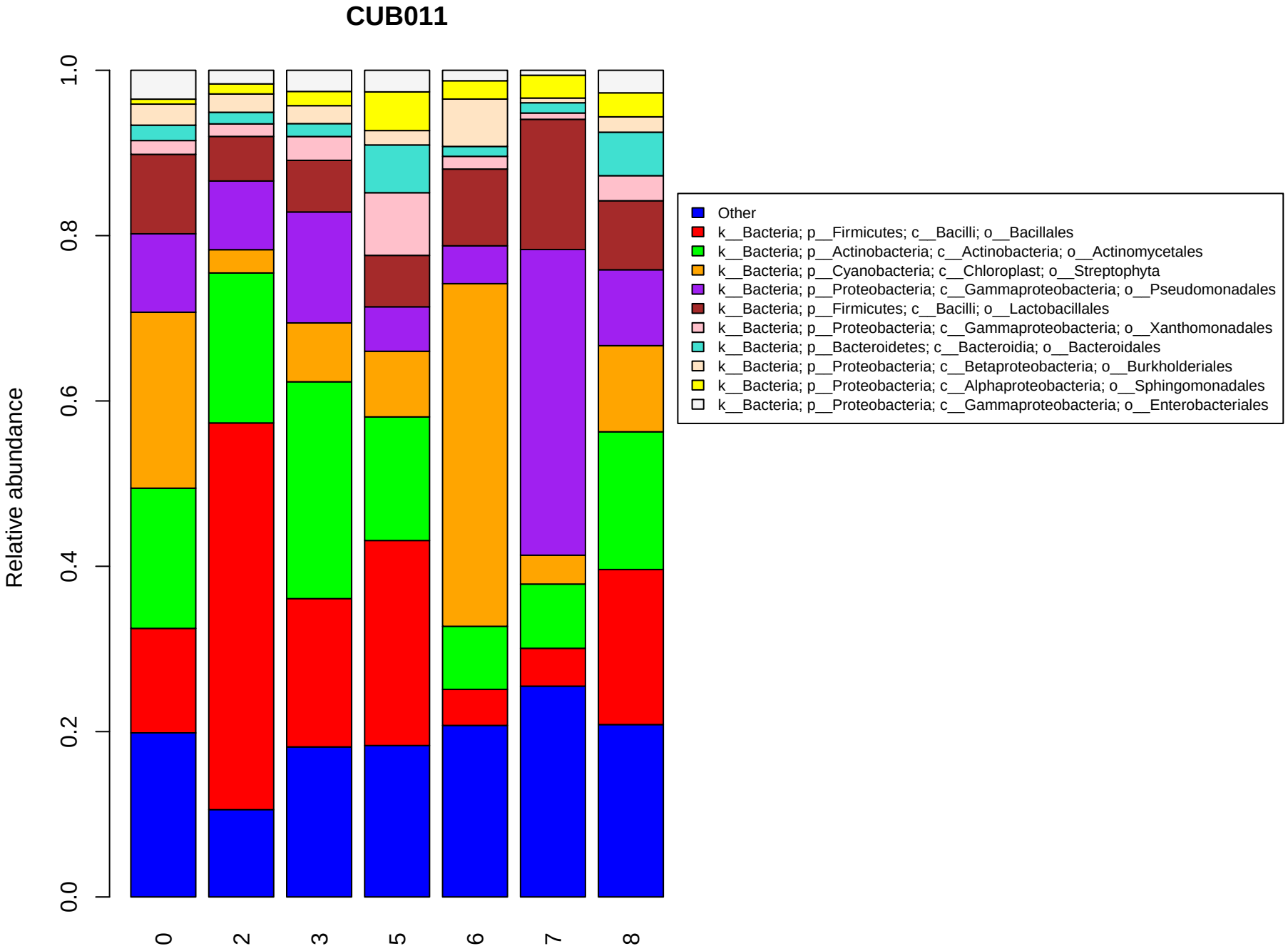
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


CUB011
Relative abundance
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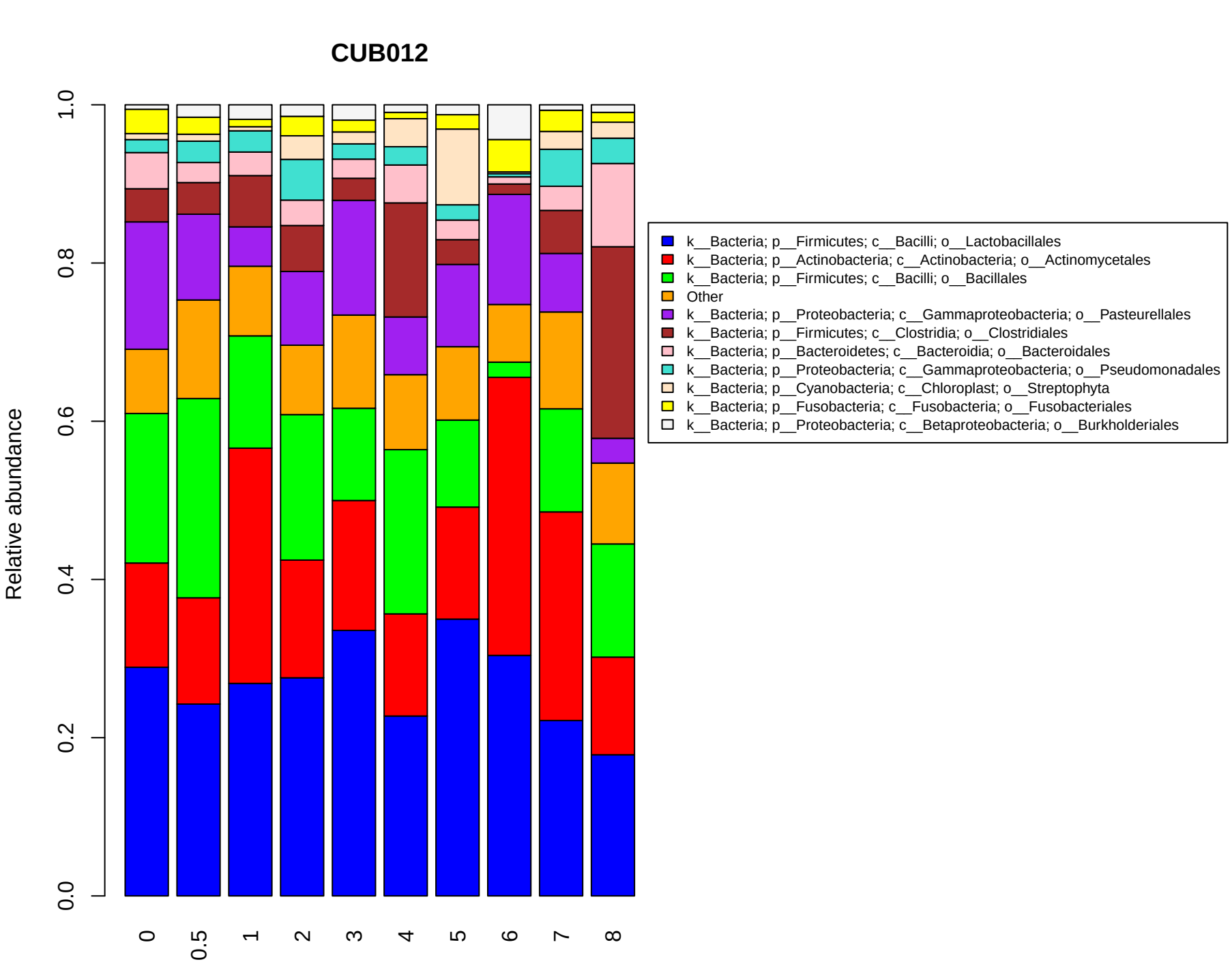
0.2
0.4
0.6
0.8
1.0
0
2
3
5
6
7
8
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales


CUB012
Relative abundance
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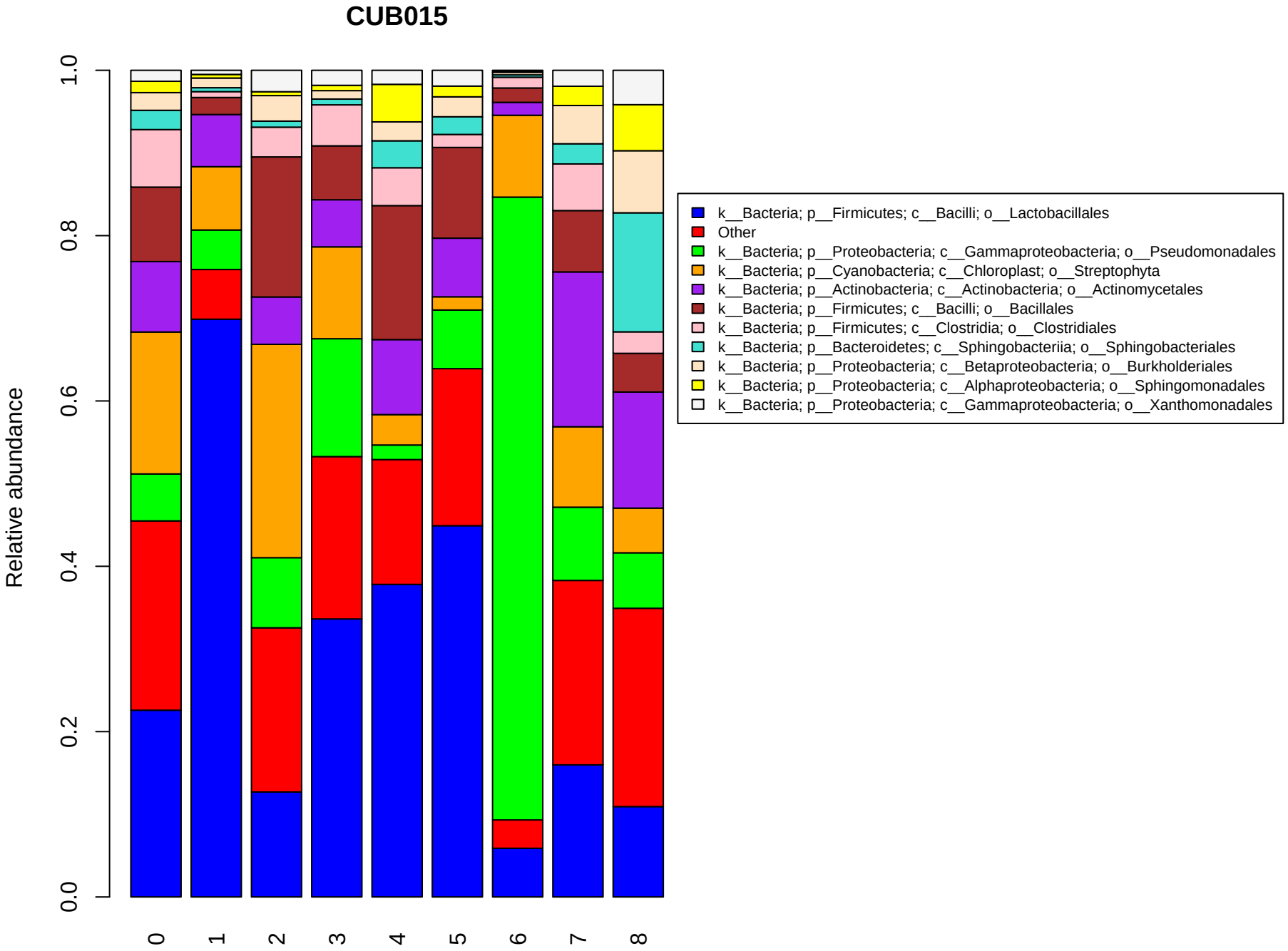
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


CUB015
Relative abundance
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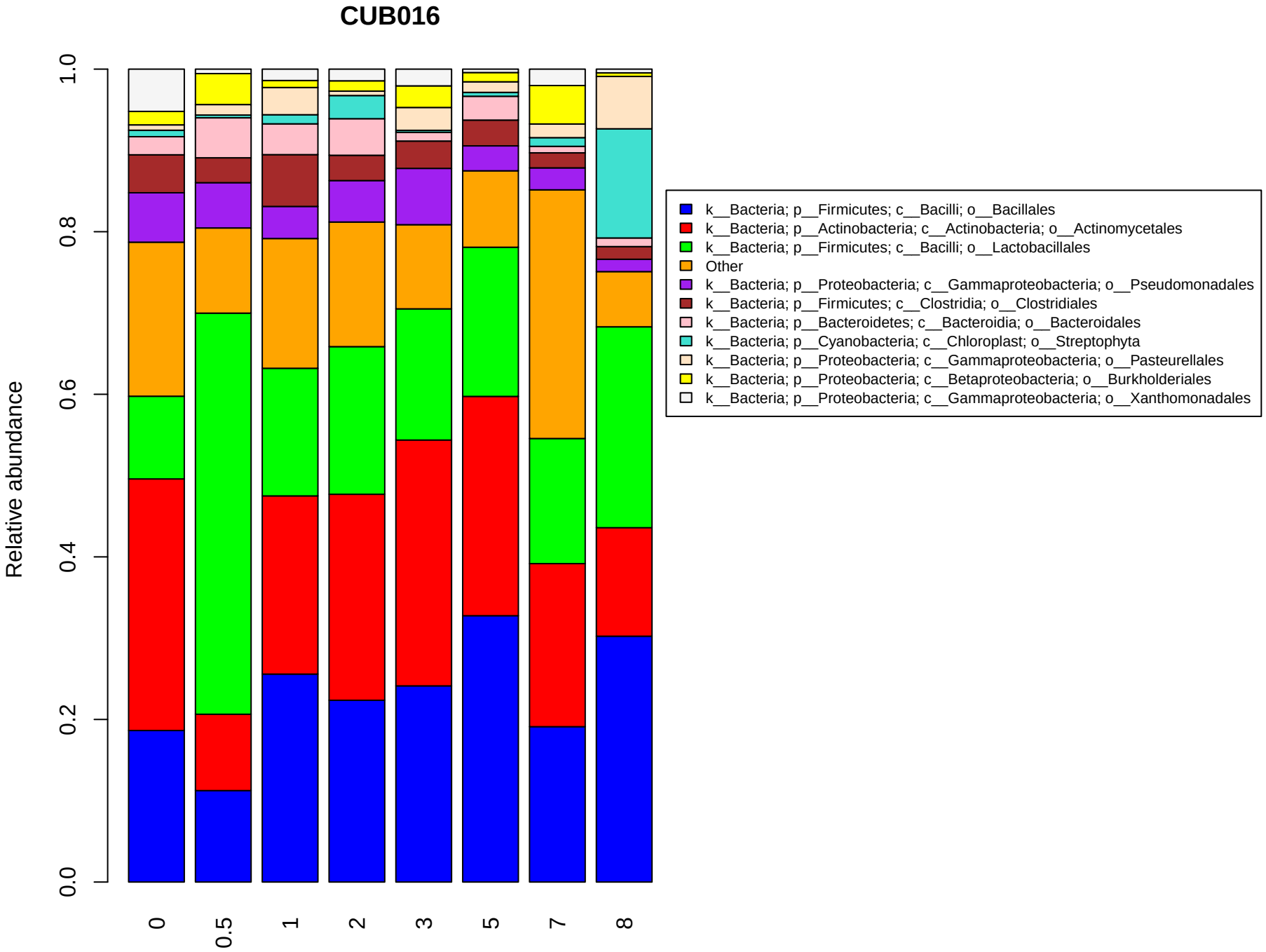
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales


CUB016
Relative abundance
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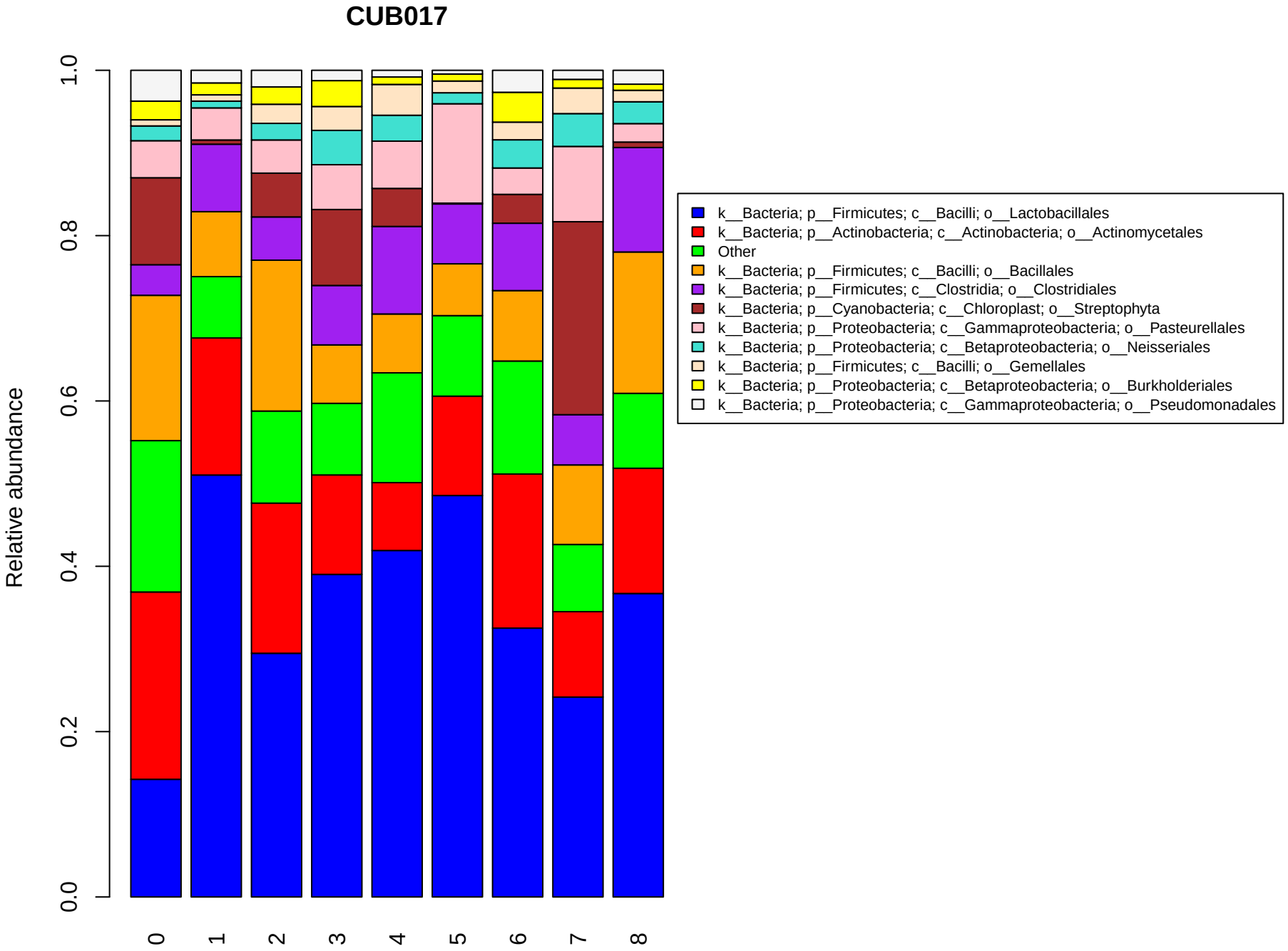
0.6
0.8
1.0
0
0.5
1
2
3
5
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales


CUB017
Relative abundance
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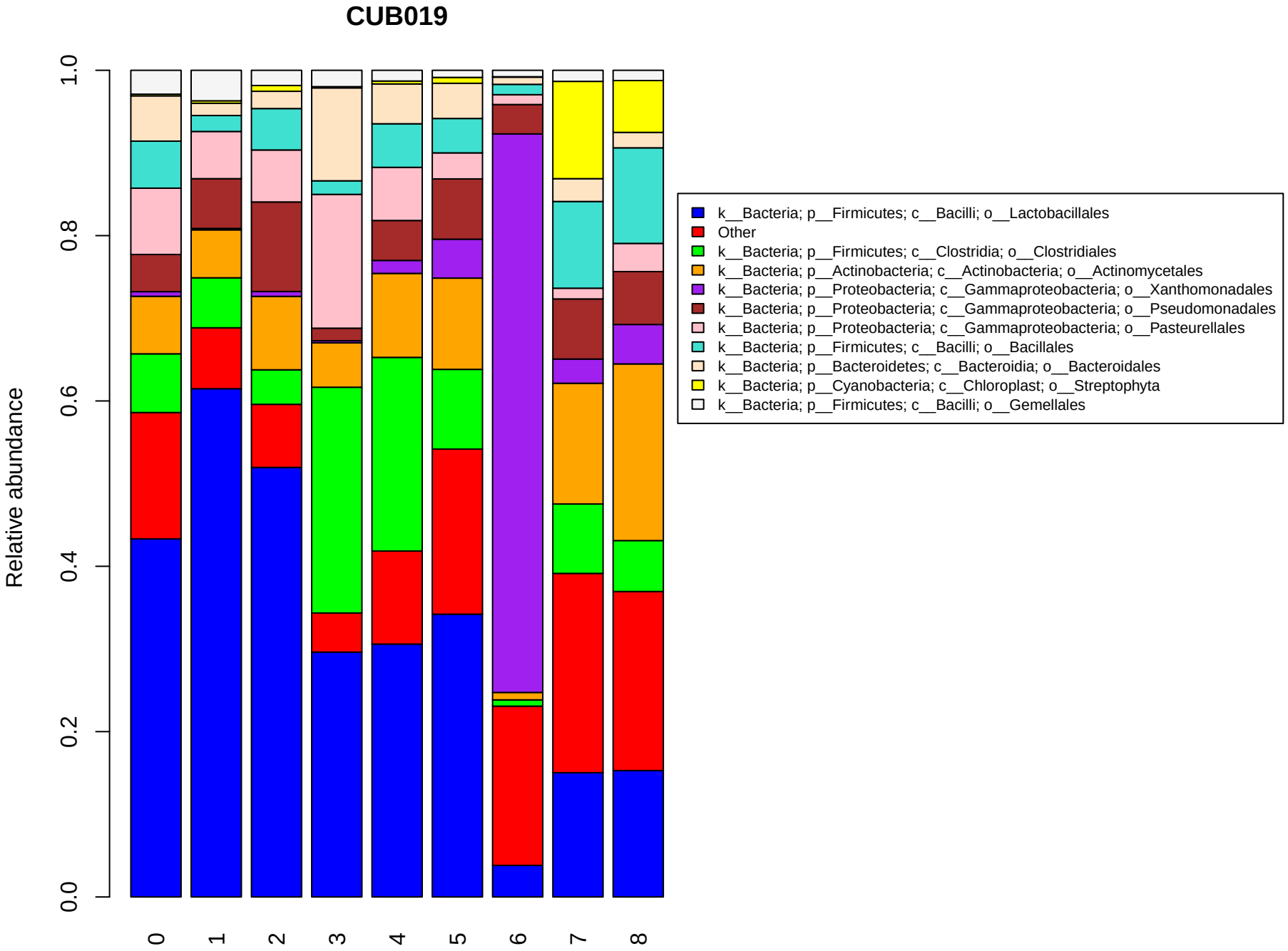
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales


CUB019
Relative abundance
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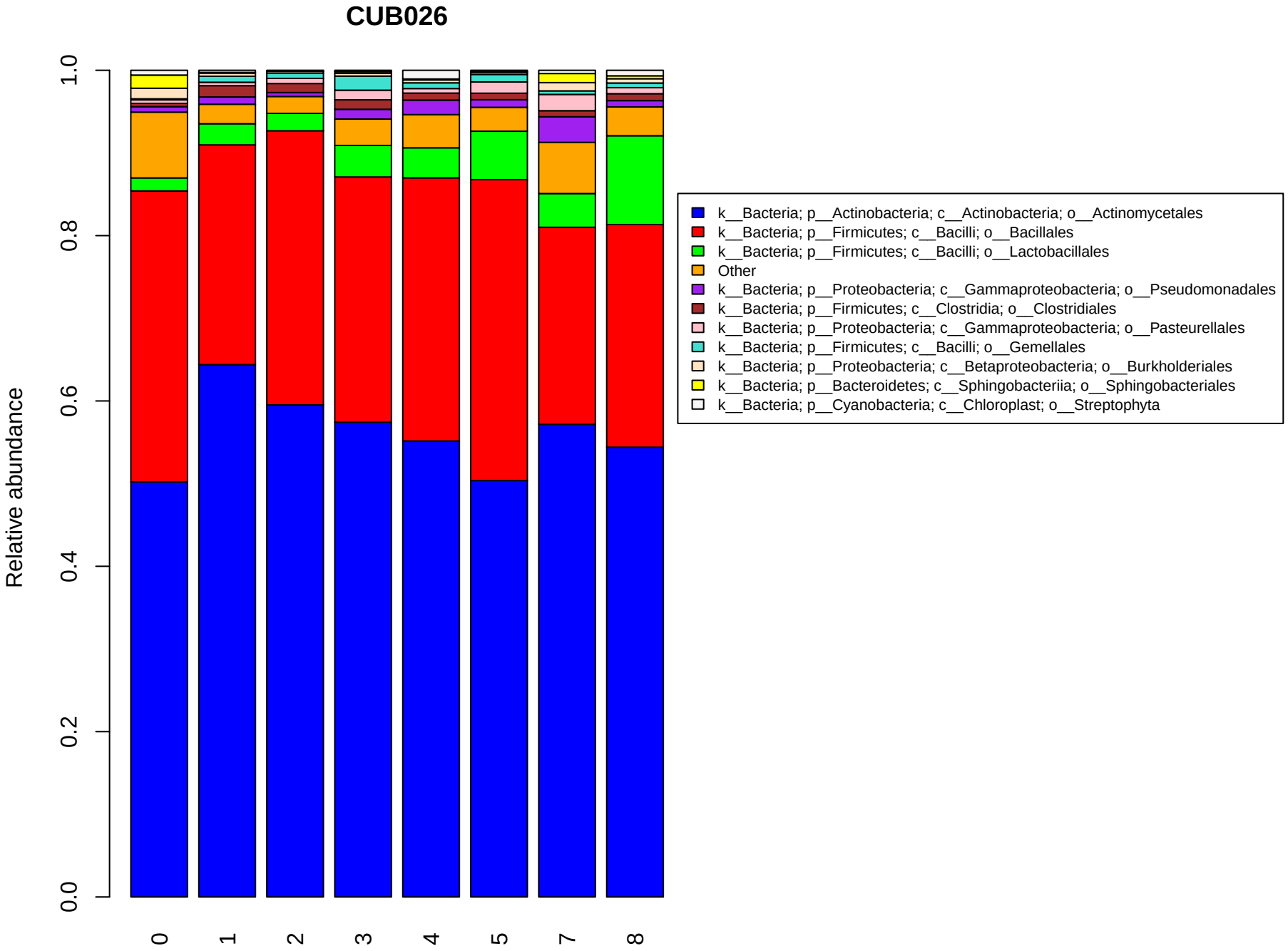
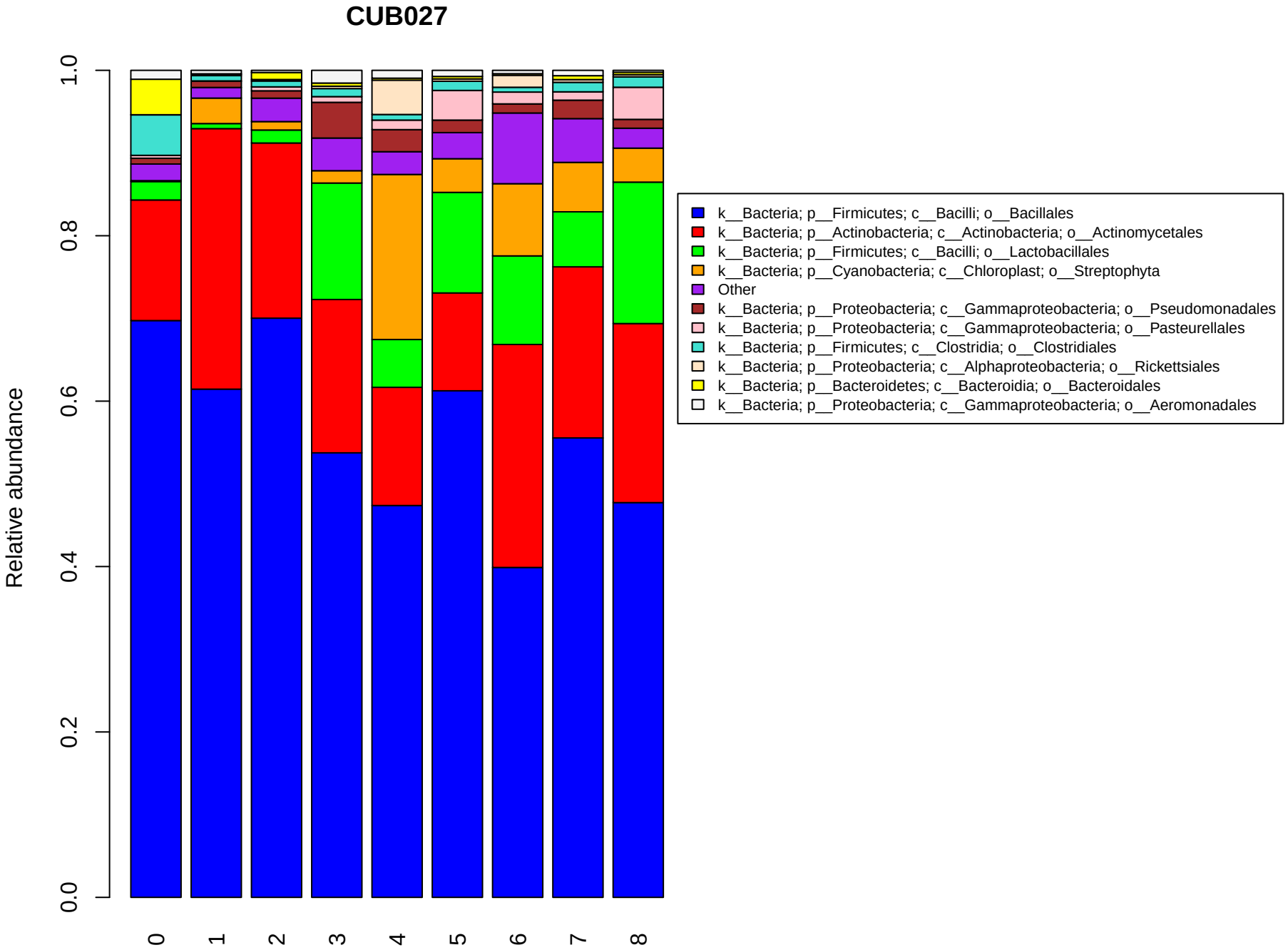
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales


CUB026
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
7
8
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta


CUB027
Relative abundance
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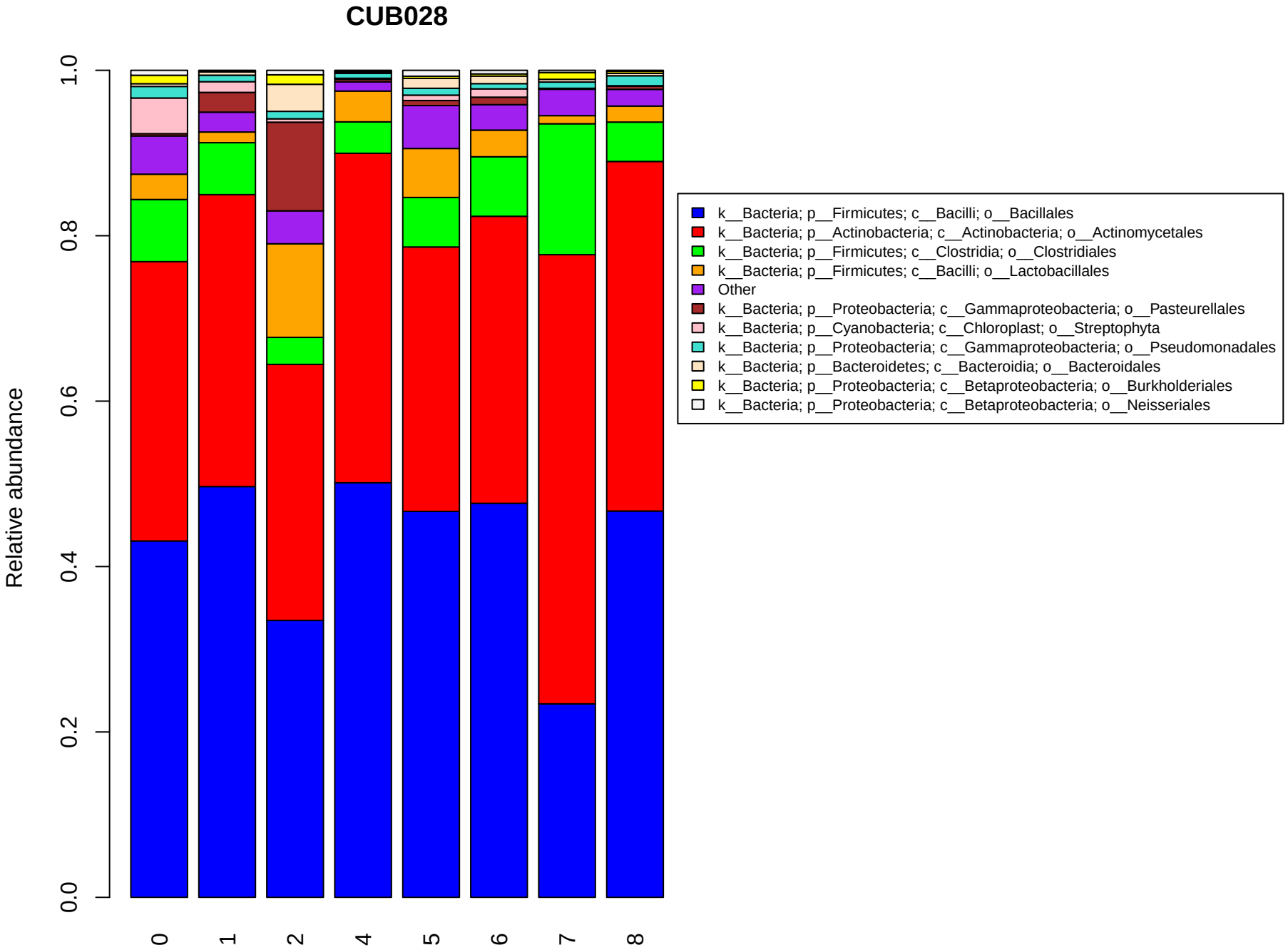
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rickettsiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales


CUB028
Relative abundance
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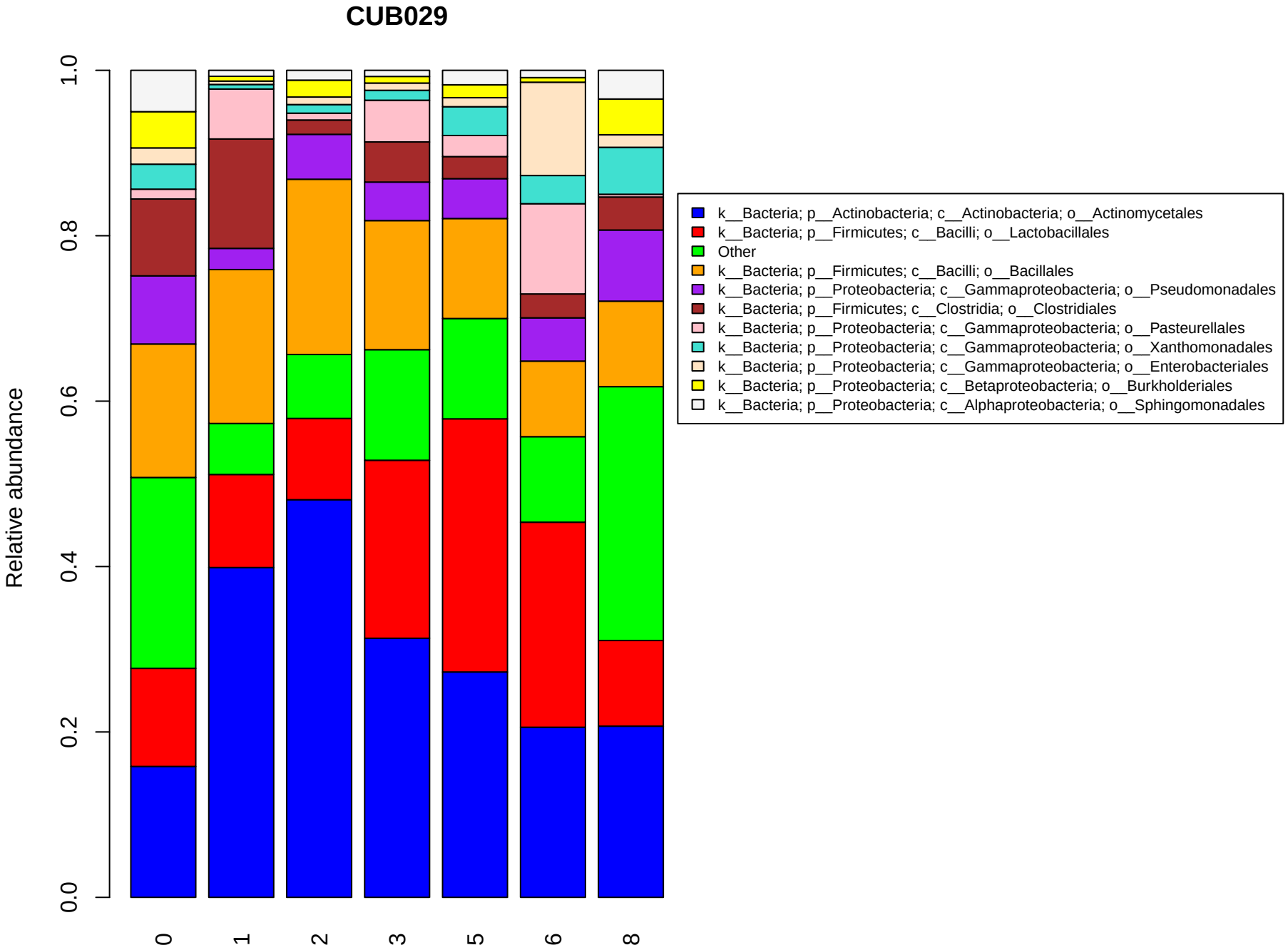
0.4
0.6
0.8
1.0
0
1
2
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales


CUB029
Relative abundance
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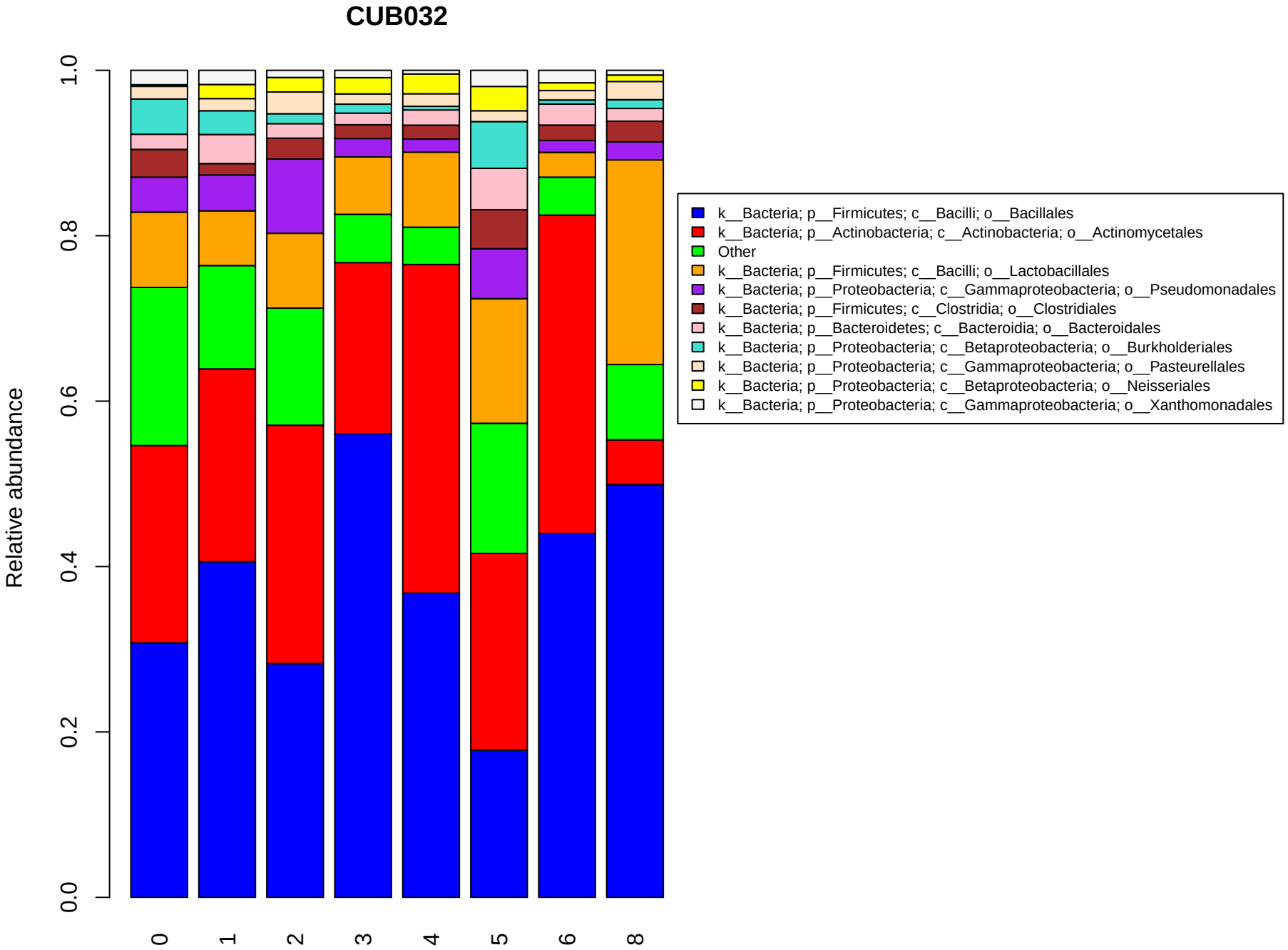
0.4
0.6
0.8
1.0
0
1
2
3
5
6
8
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales


CUB032
Relative abundance
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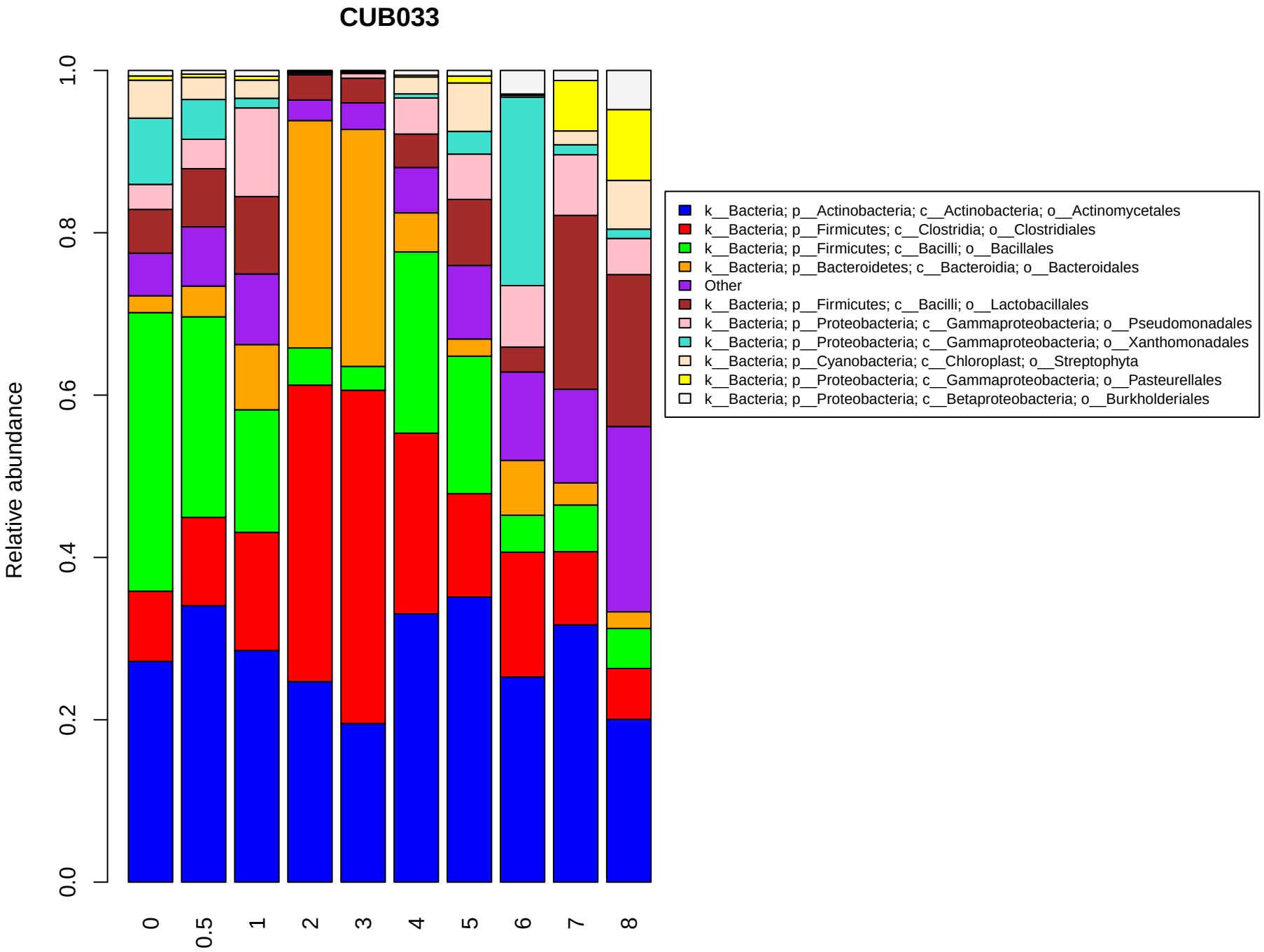
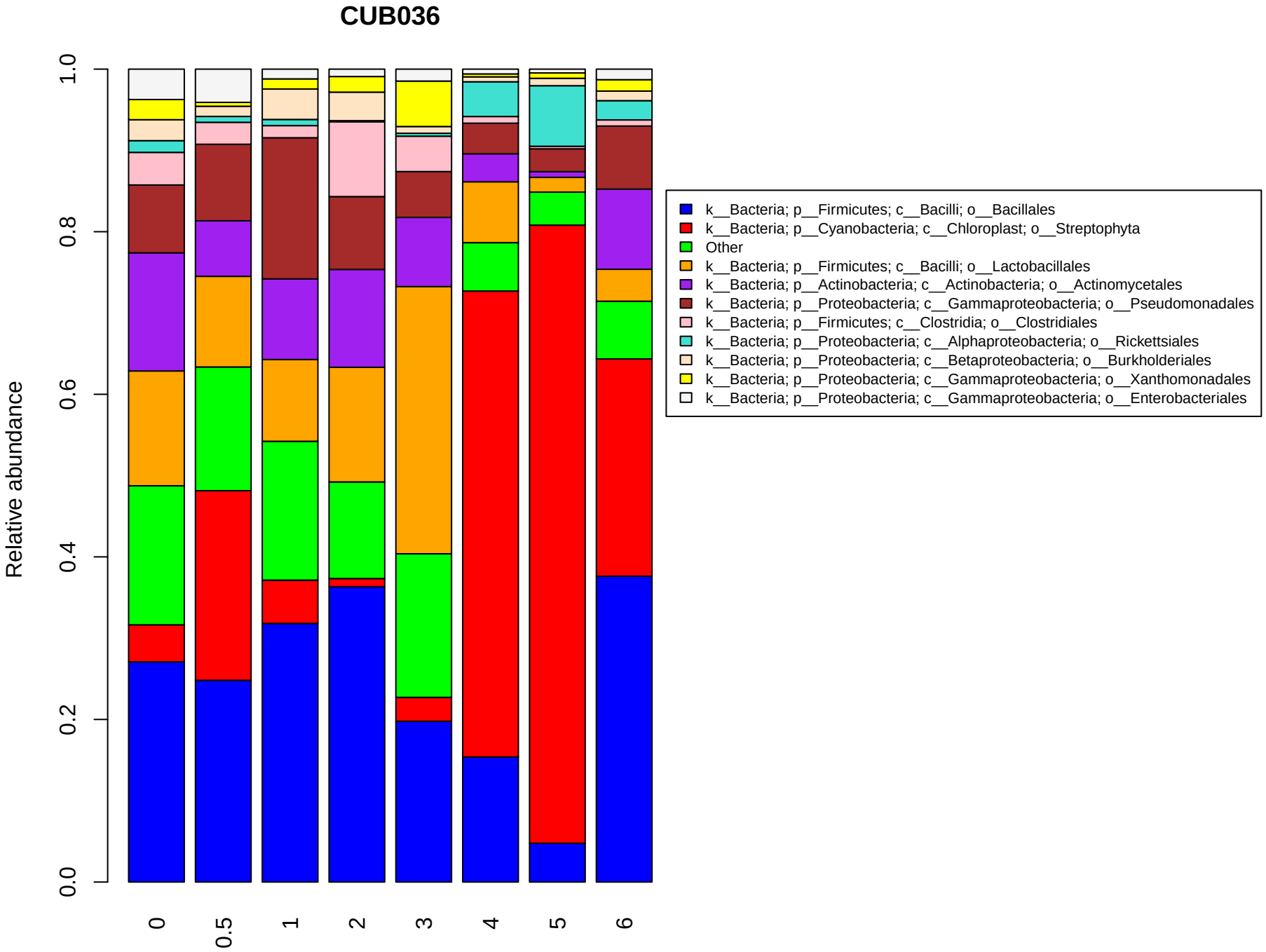
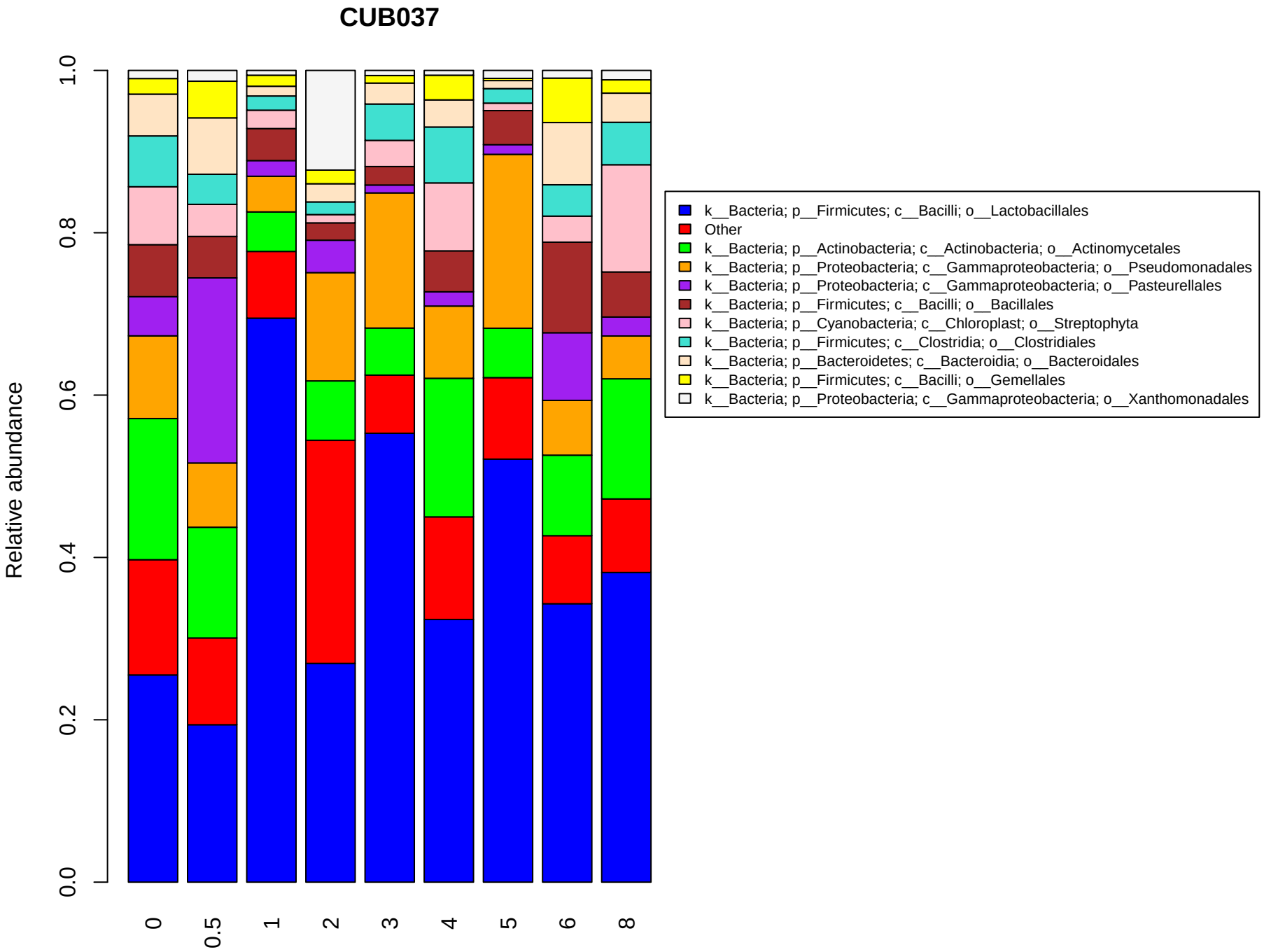
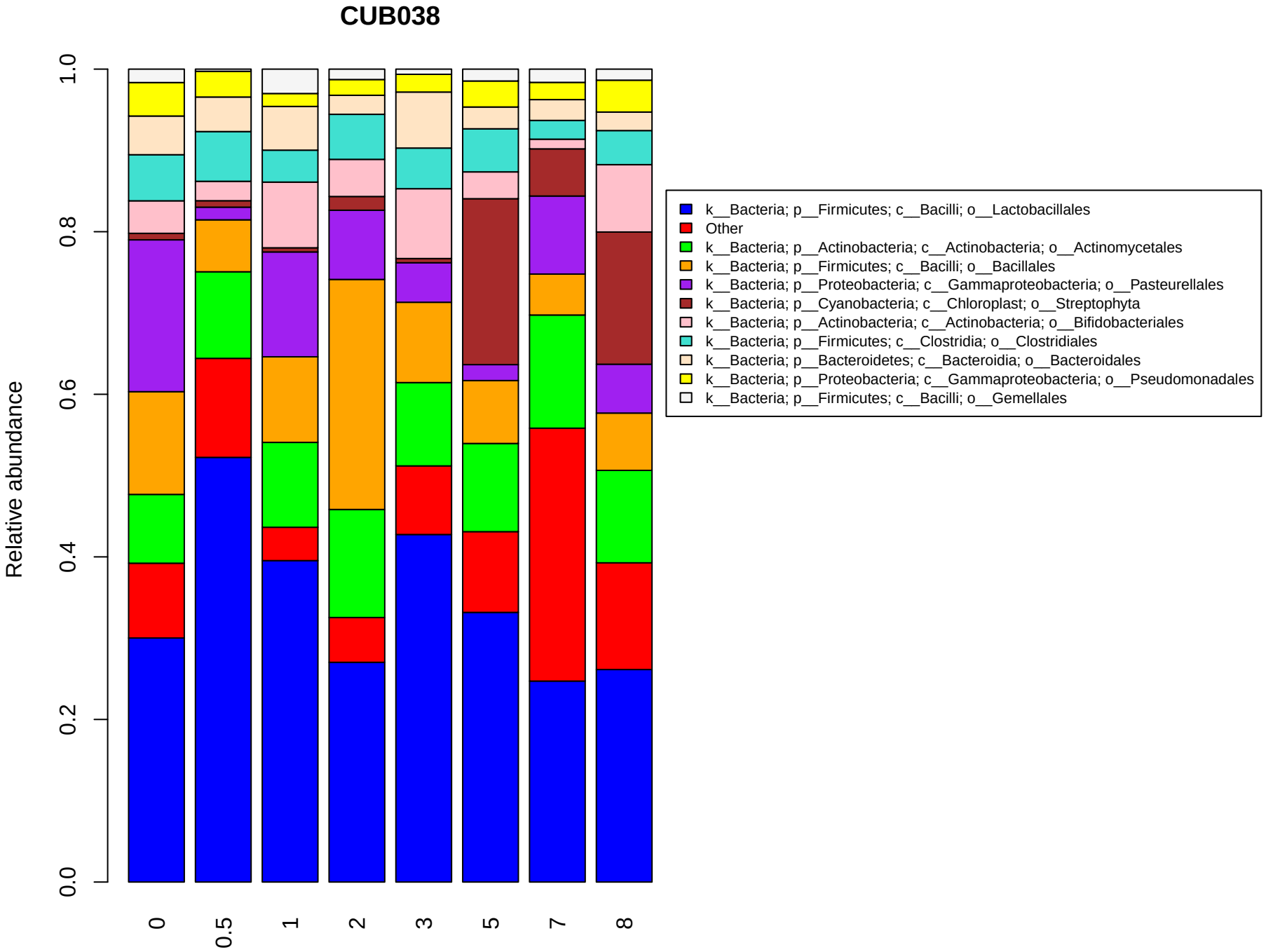
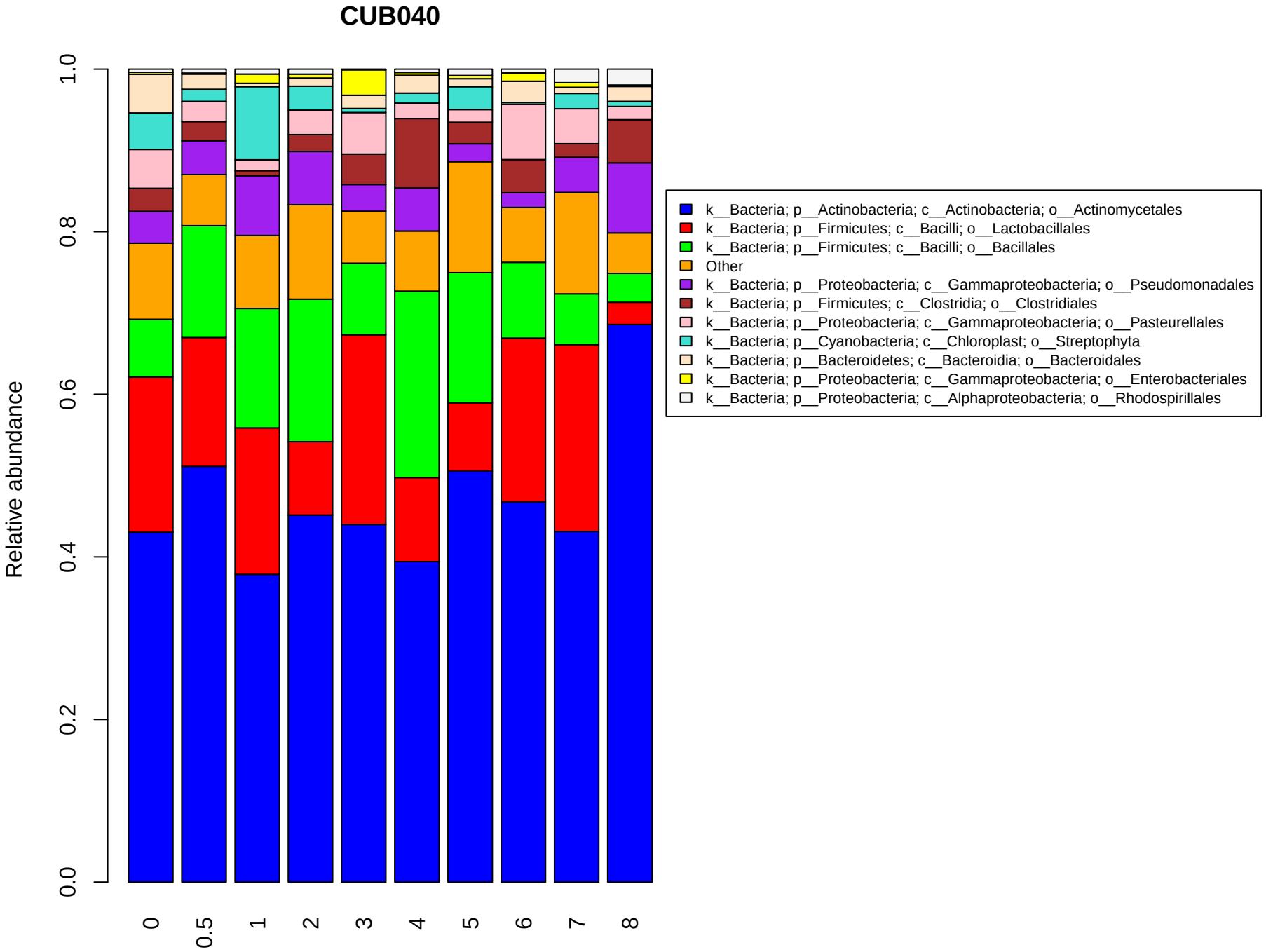
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales


CUB033
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


CUB036
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rickettsiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales


CUB037
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales


CUB038
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
5
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Bifidobacteriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales


CUB040
Relative abundance
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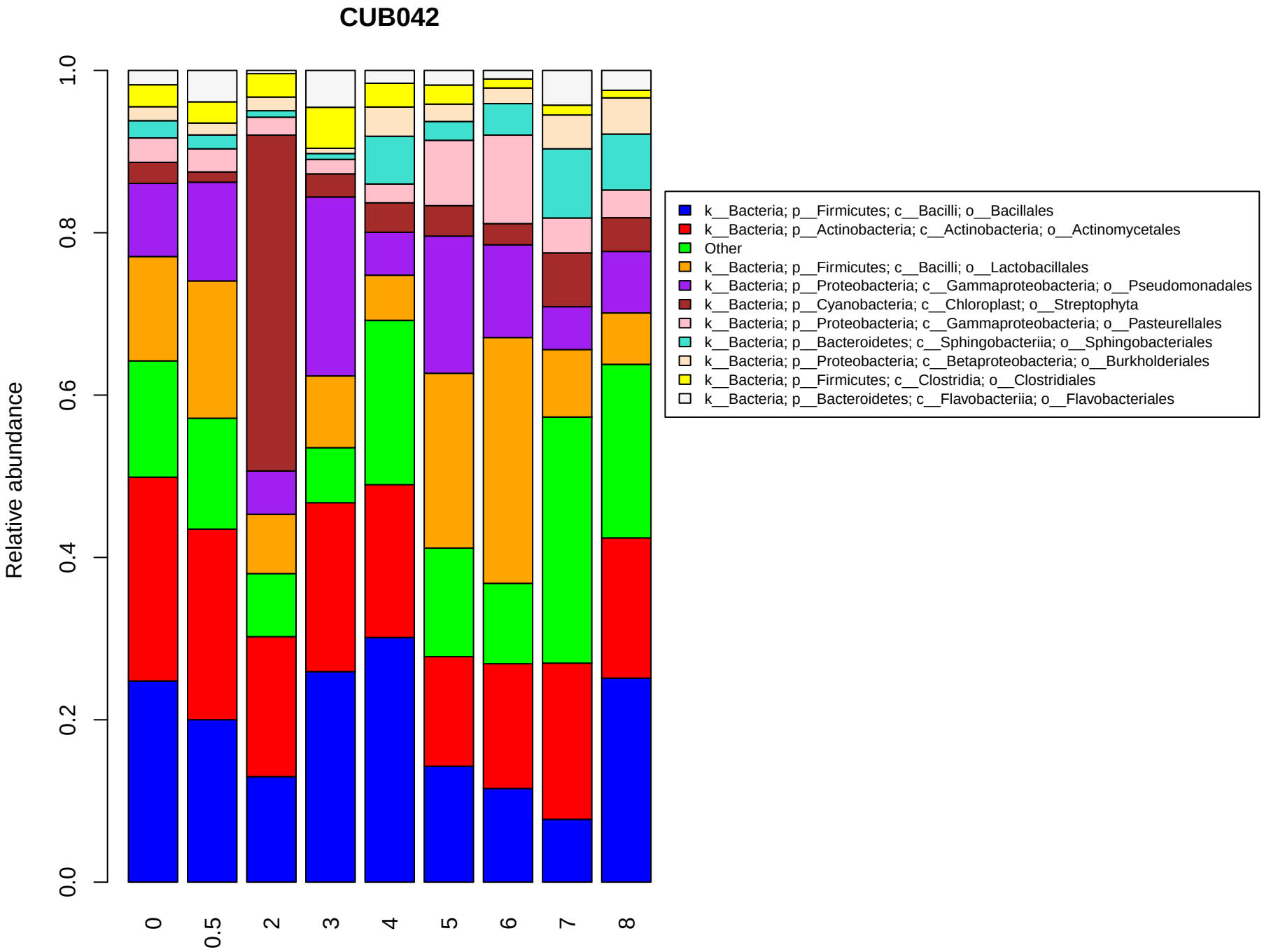
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodospirillales


CUB042
Relative abundance
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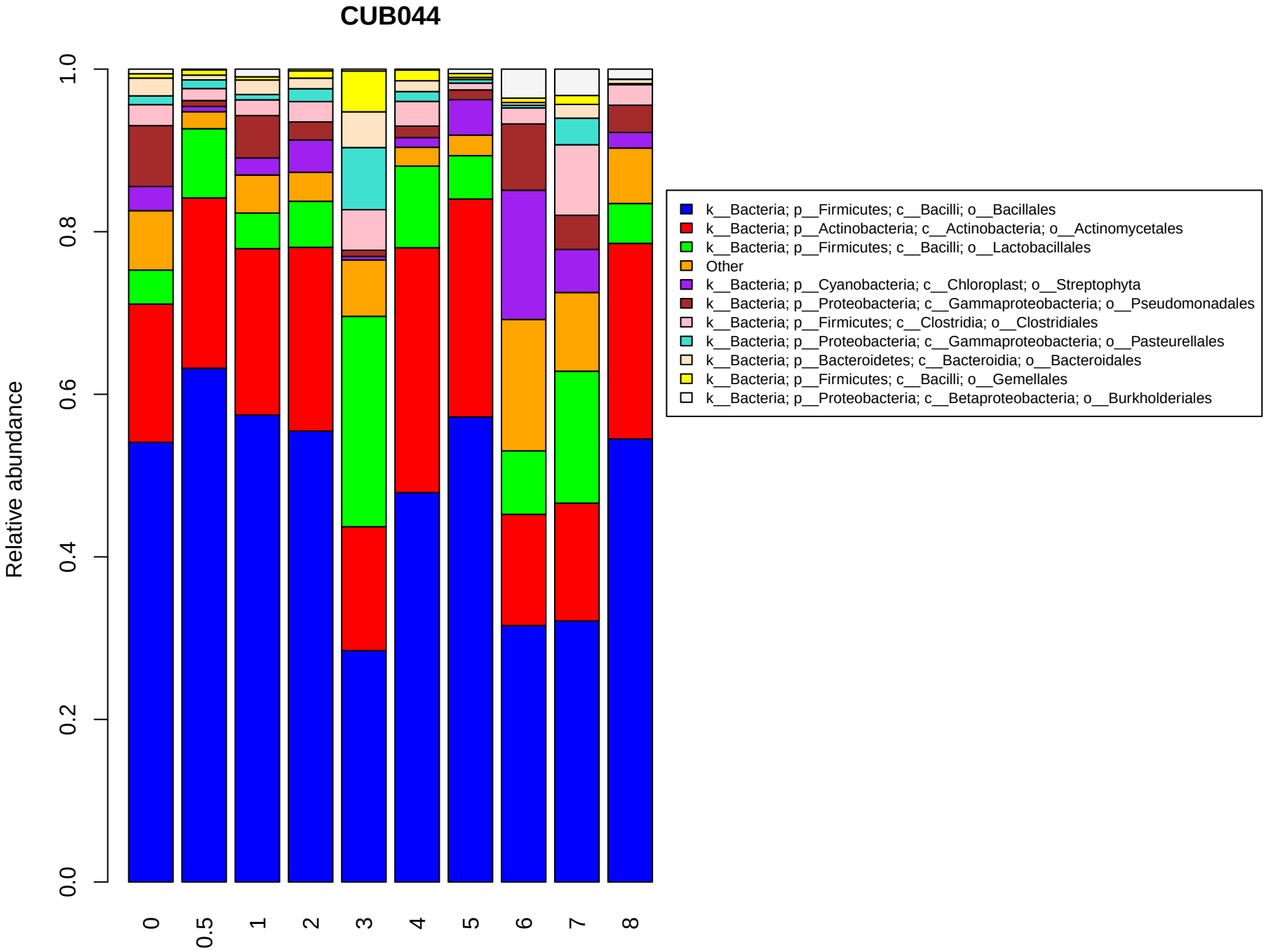
0.2
0.4
0.6
0.8
1.0
0
0.5
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales


CUB044
Relative abundance
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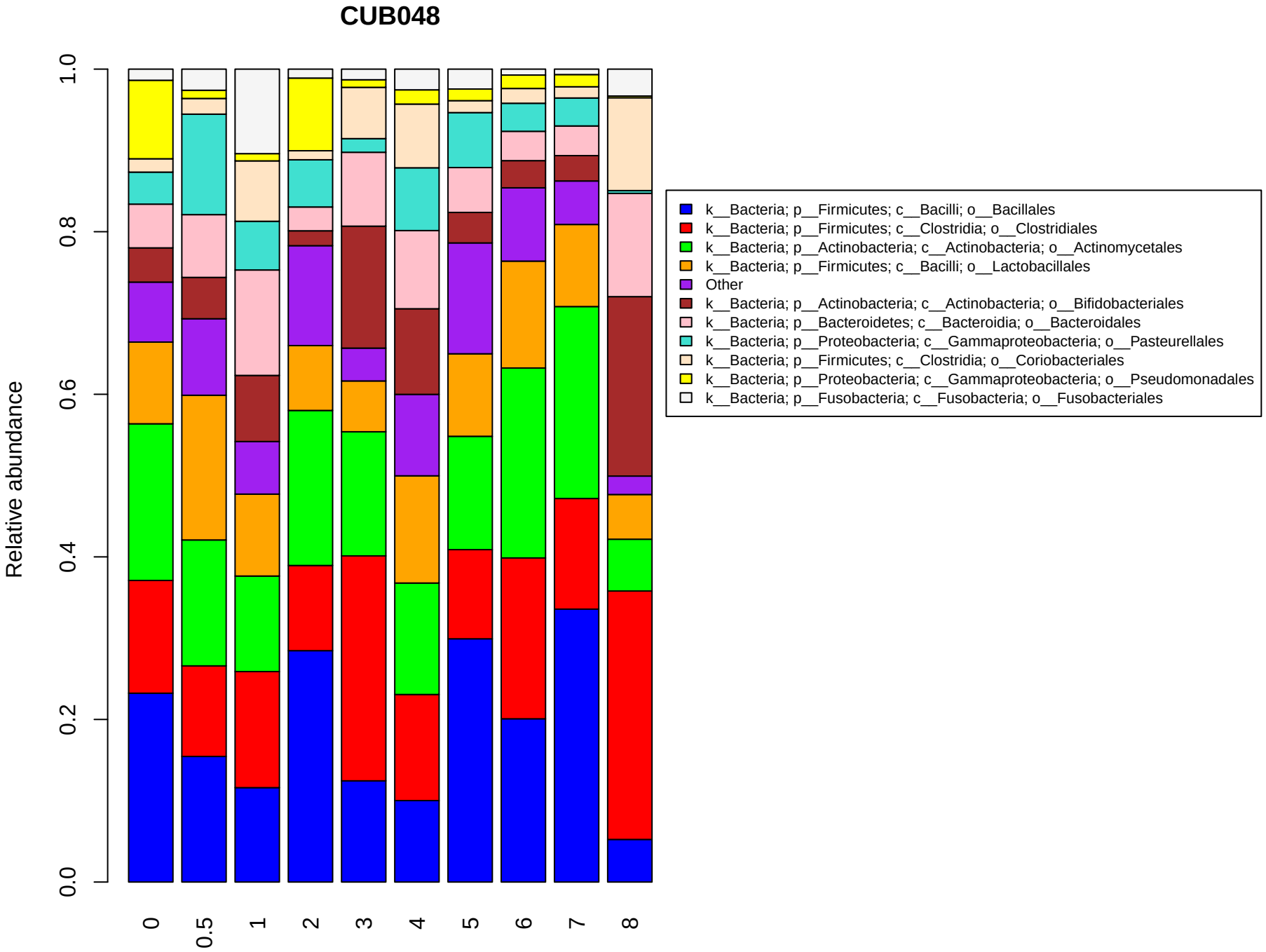
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


CUB048
Relative abundance
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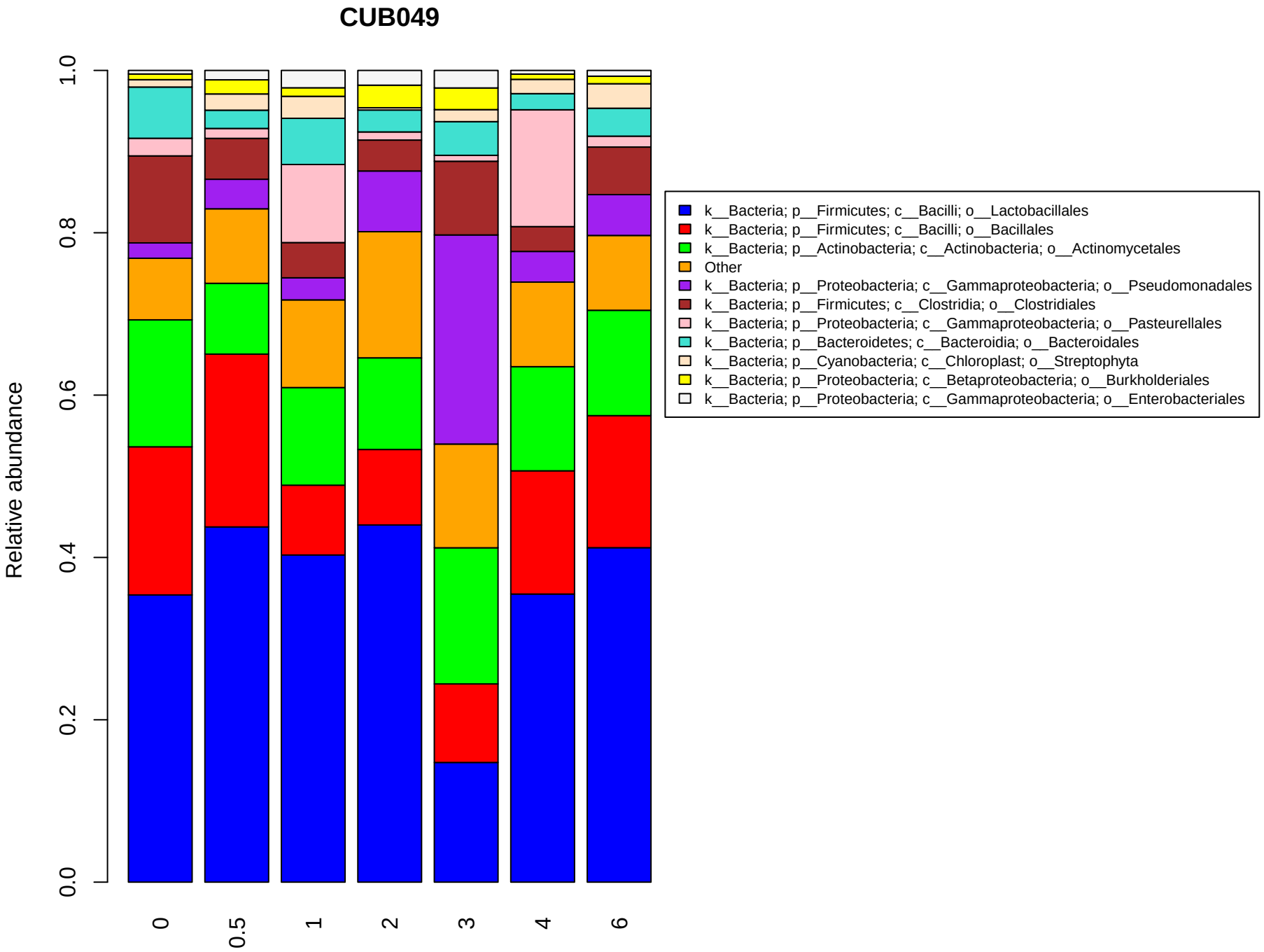
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Bifidobacteriales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Coriobacteriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales


CUB049
Relative abundance
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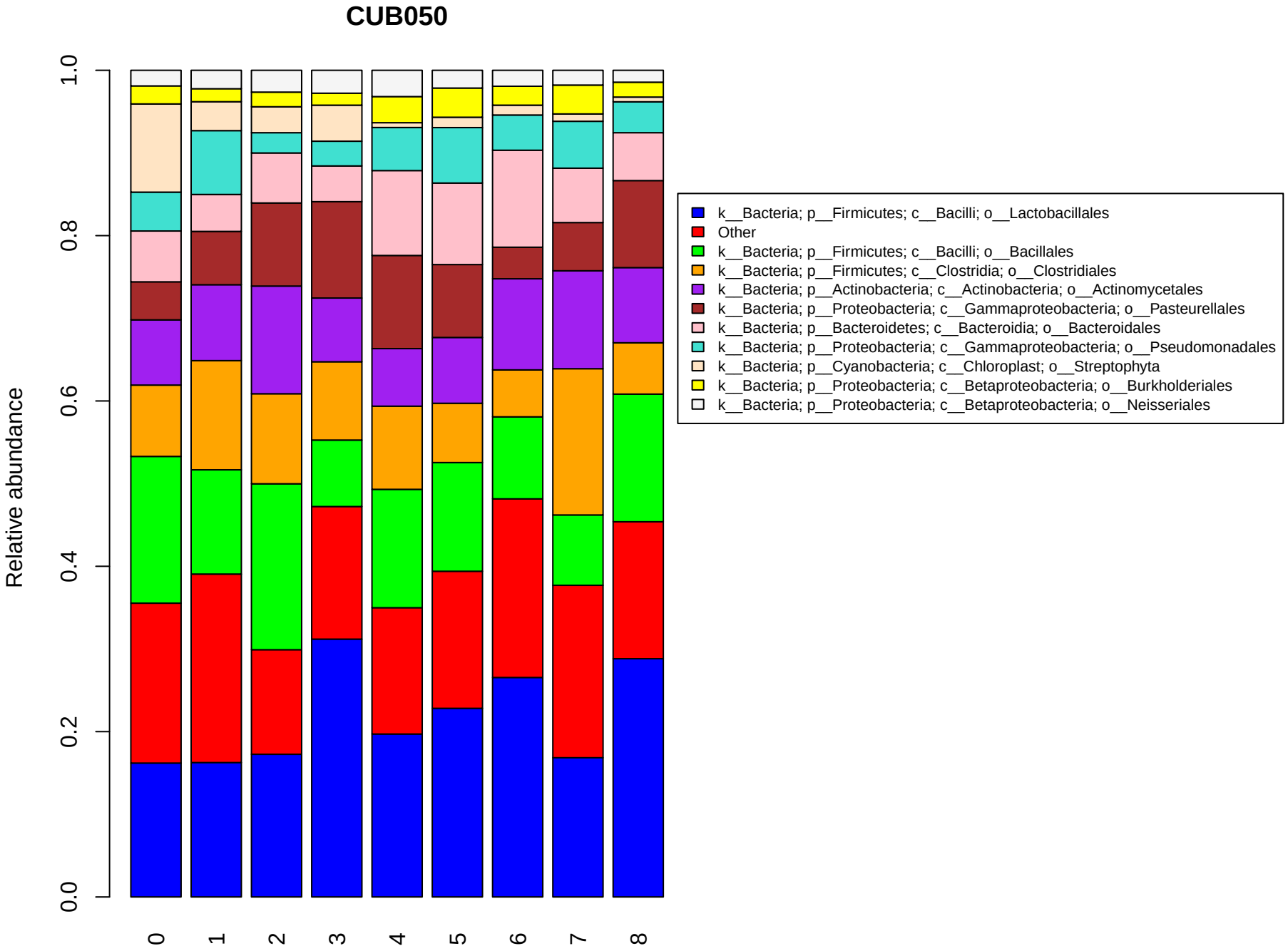
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
6
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales


CUB050
Relative abundance
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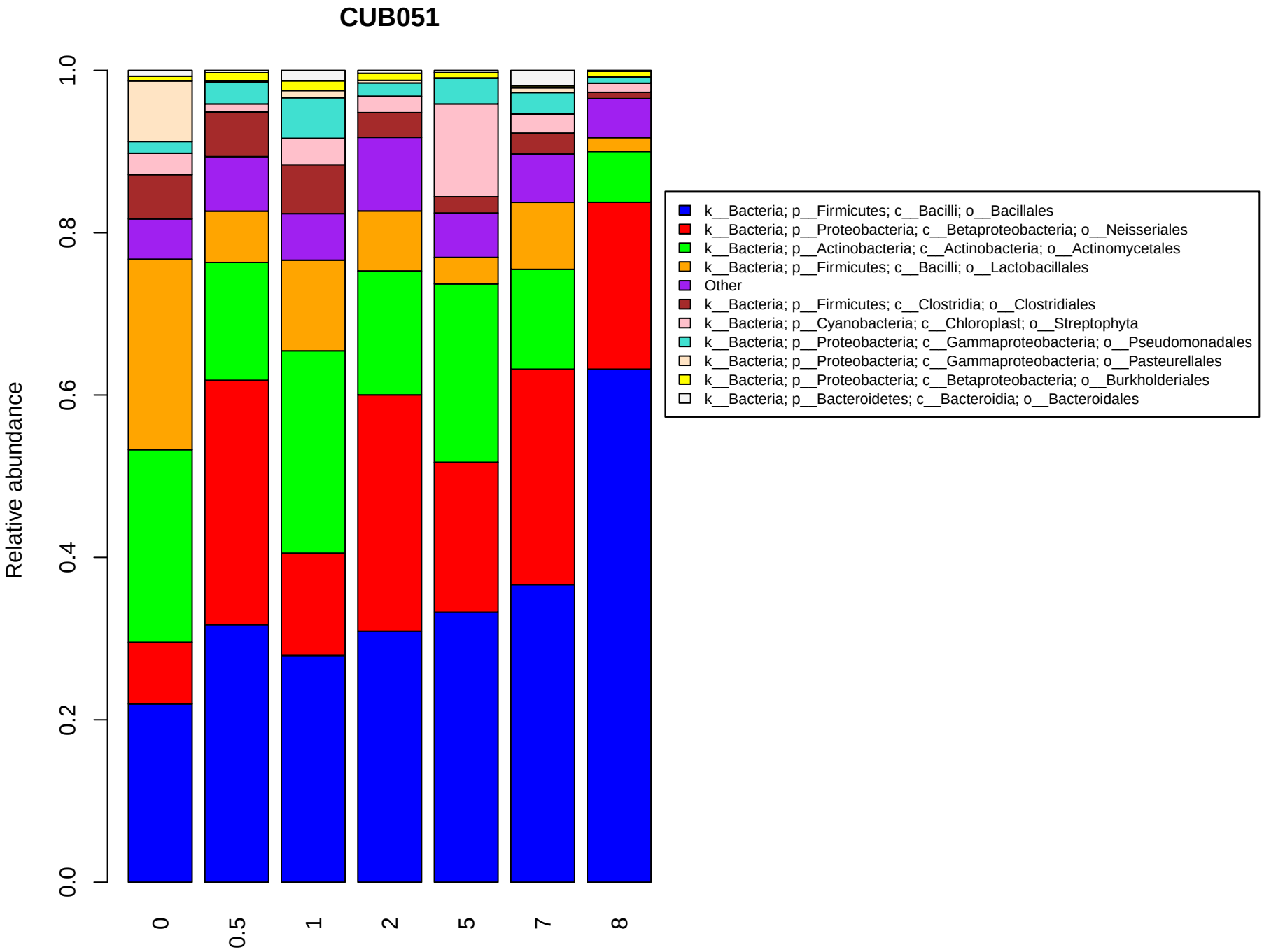
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales


CUB051
Relative abundance
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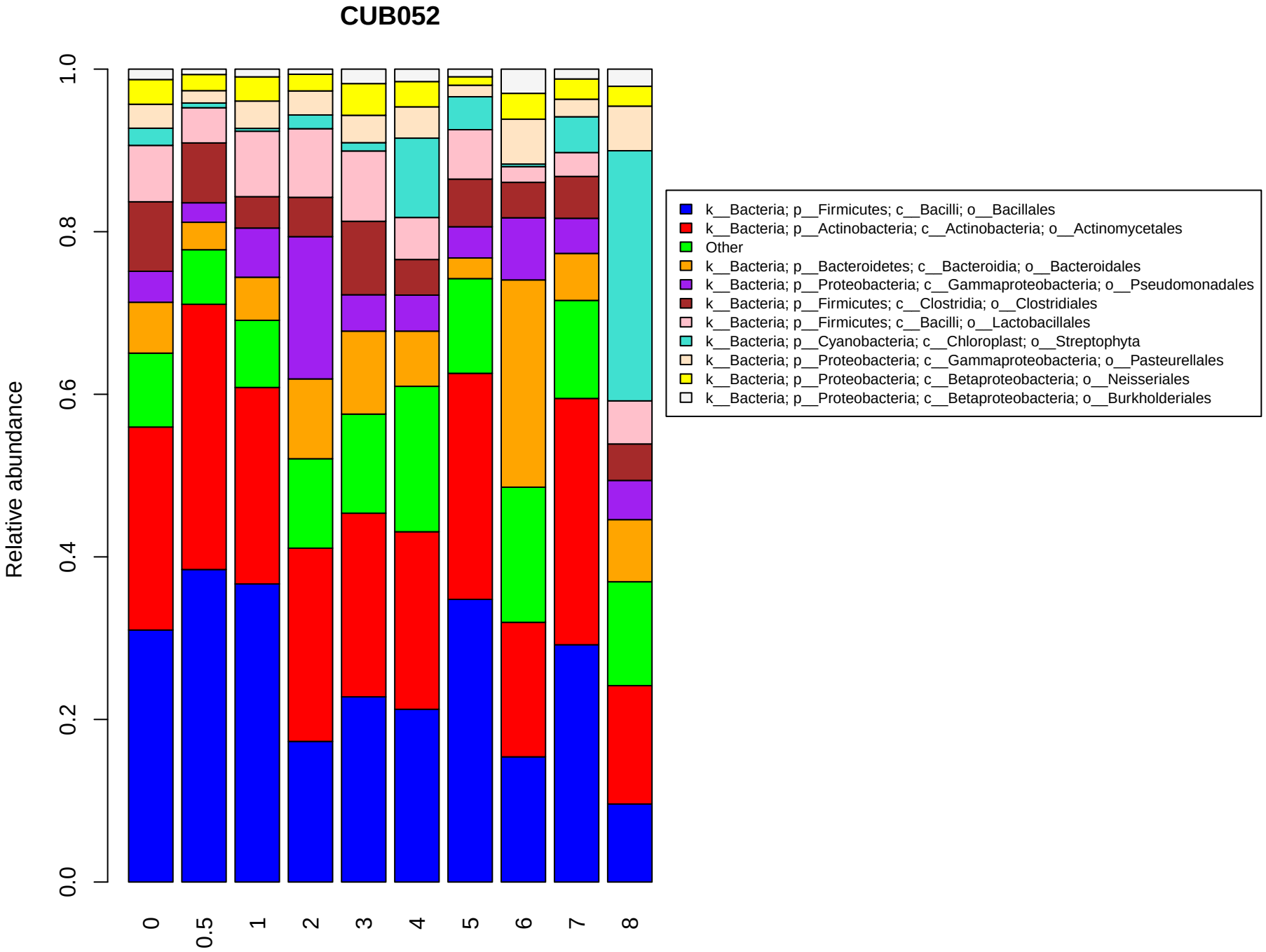
0.4
0.6
0.8
1.0
0
0.5
1
2
5
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales


CUB052
Relative abundance
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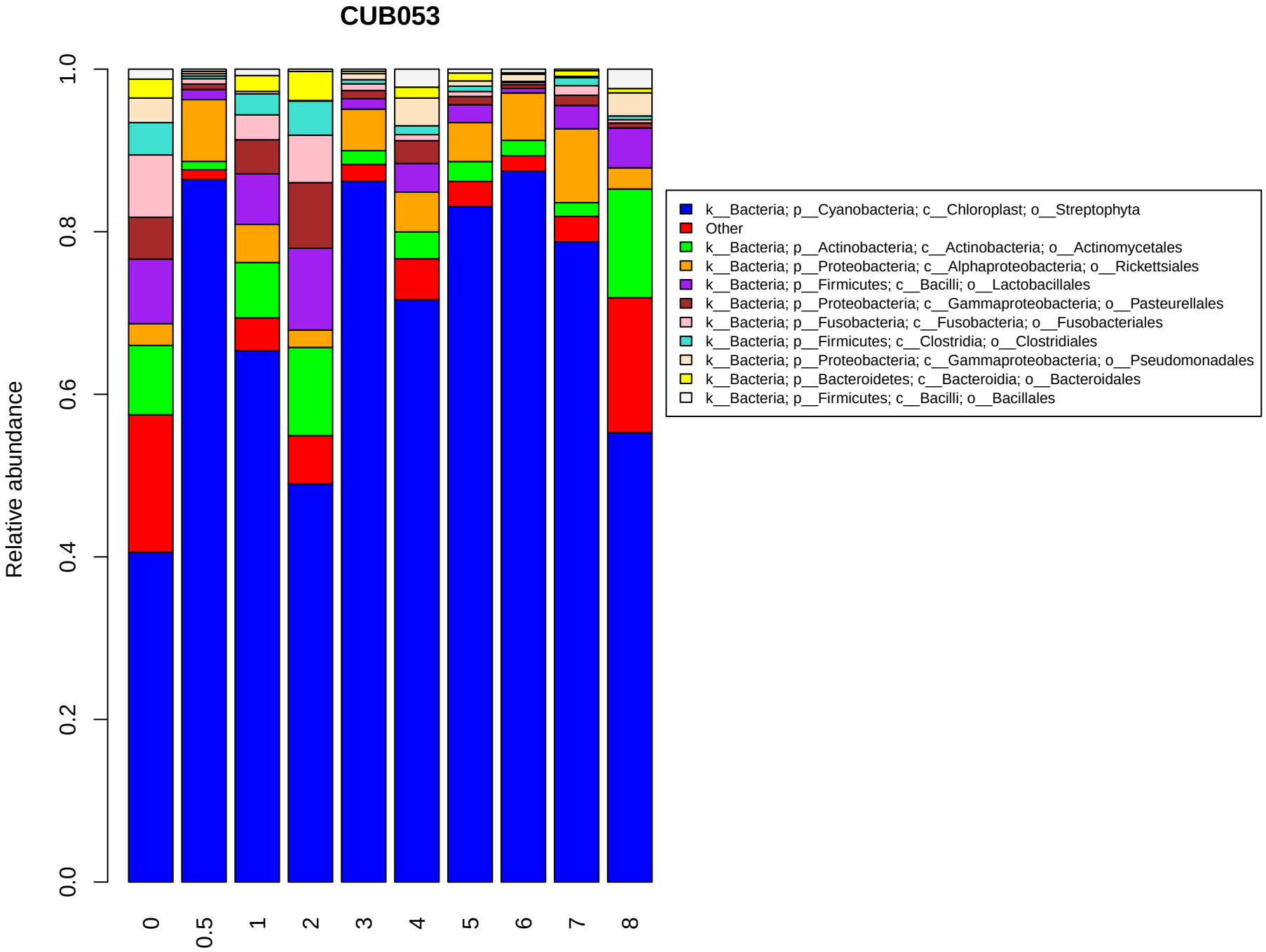
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


CUB053
Relative abundance
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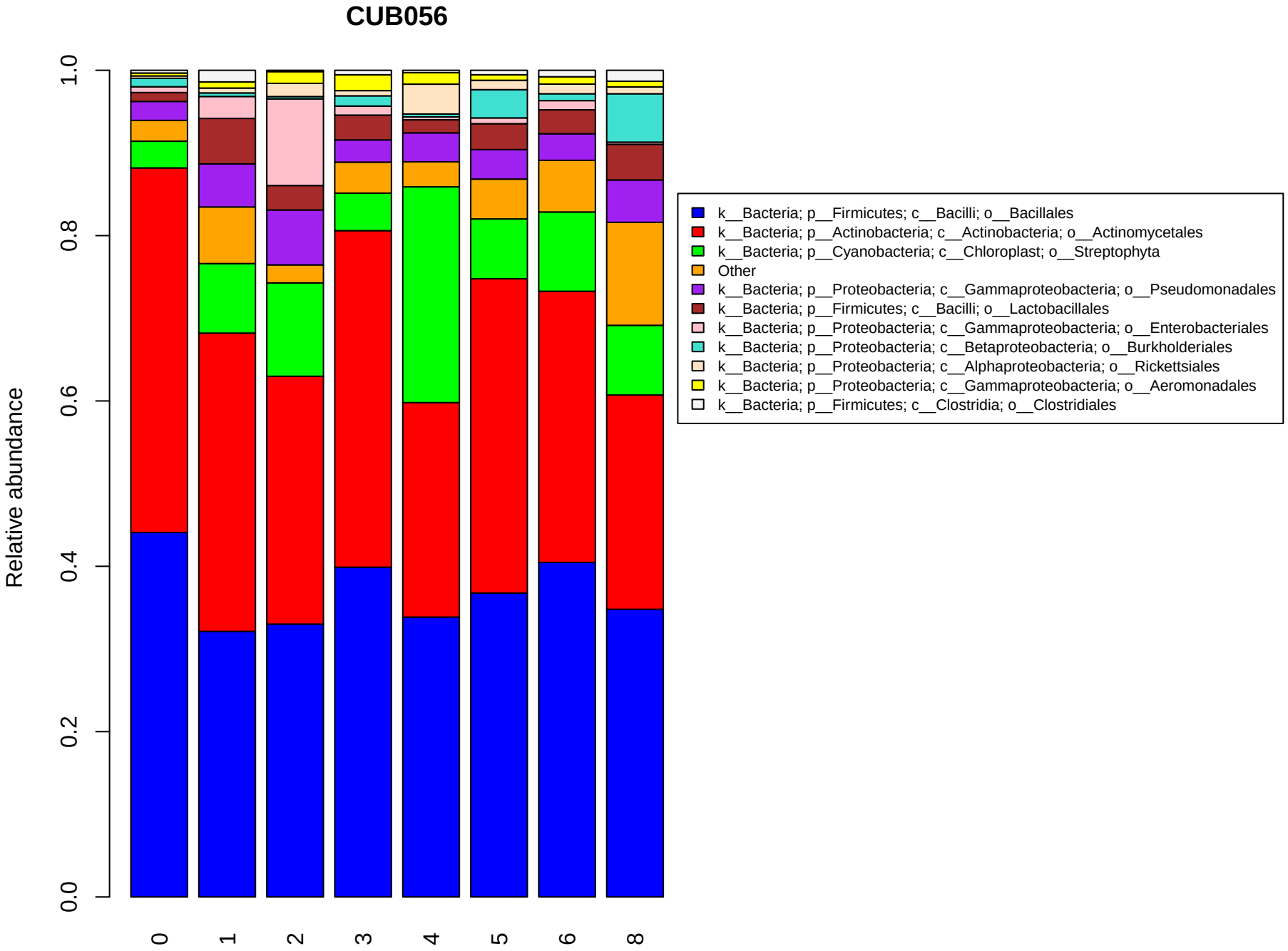
0.2
0.4
0.6
0.8
1.0
0
0.5
1
2
3
4
5
6
7
8
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rickettsiales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales


CUB056
Relative abundance
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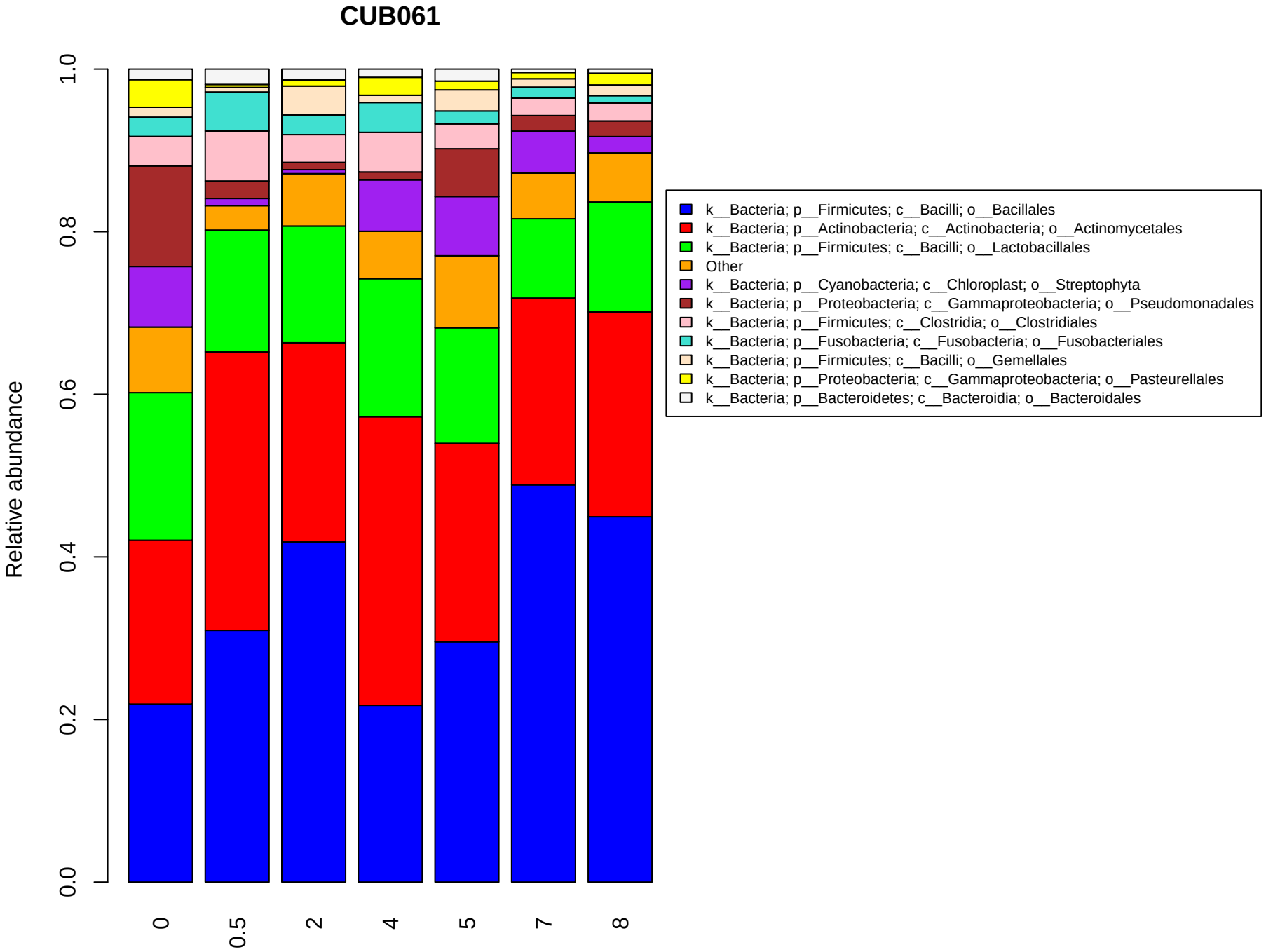
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rickettsiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales


CUB061
Relative abundance
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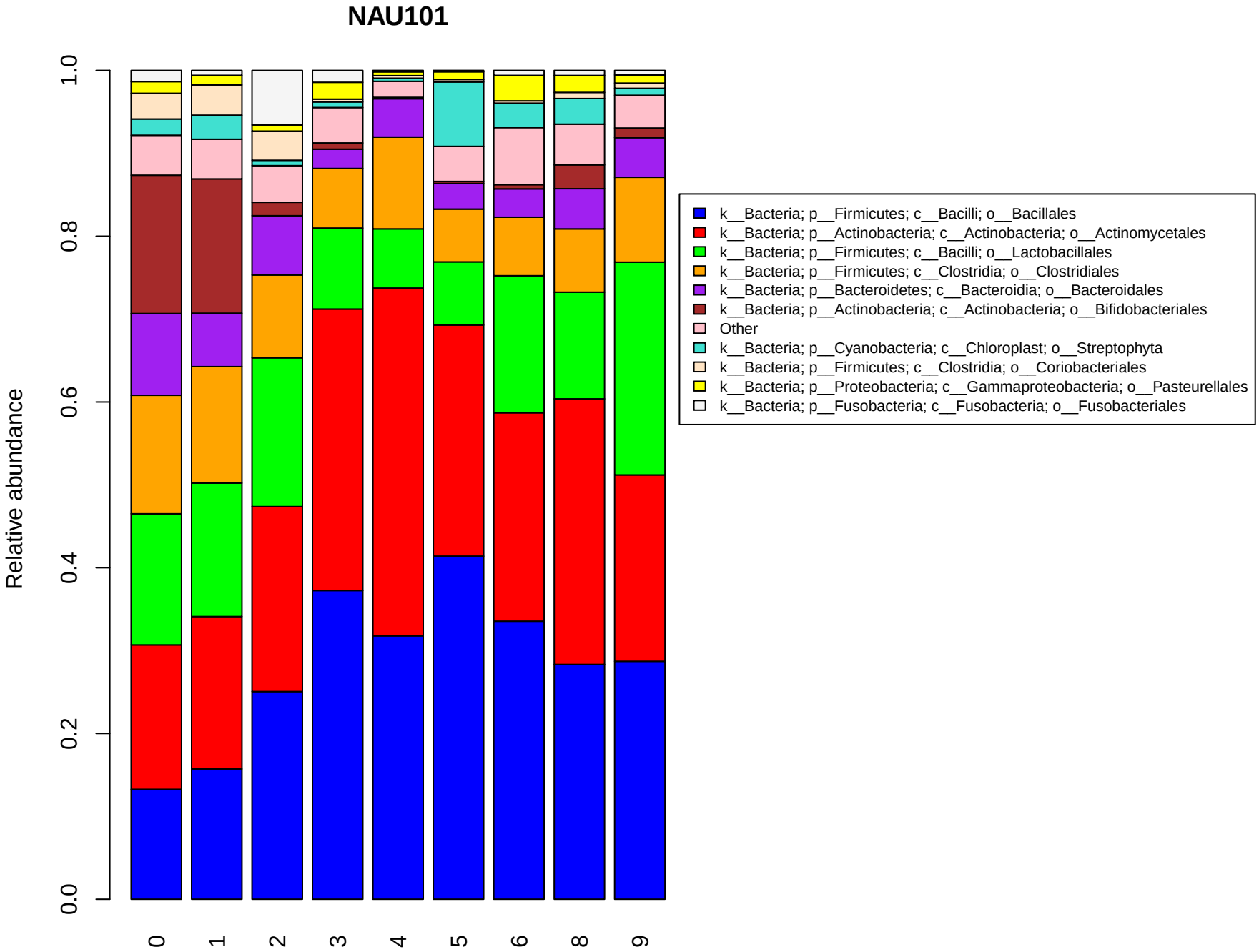
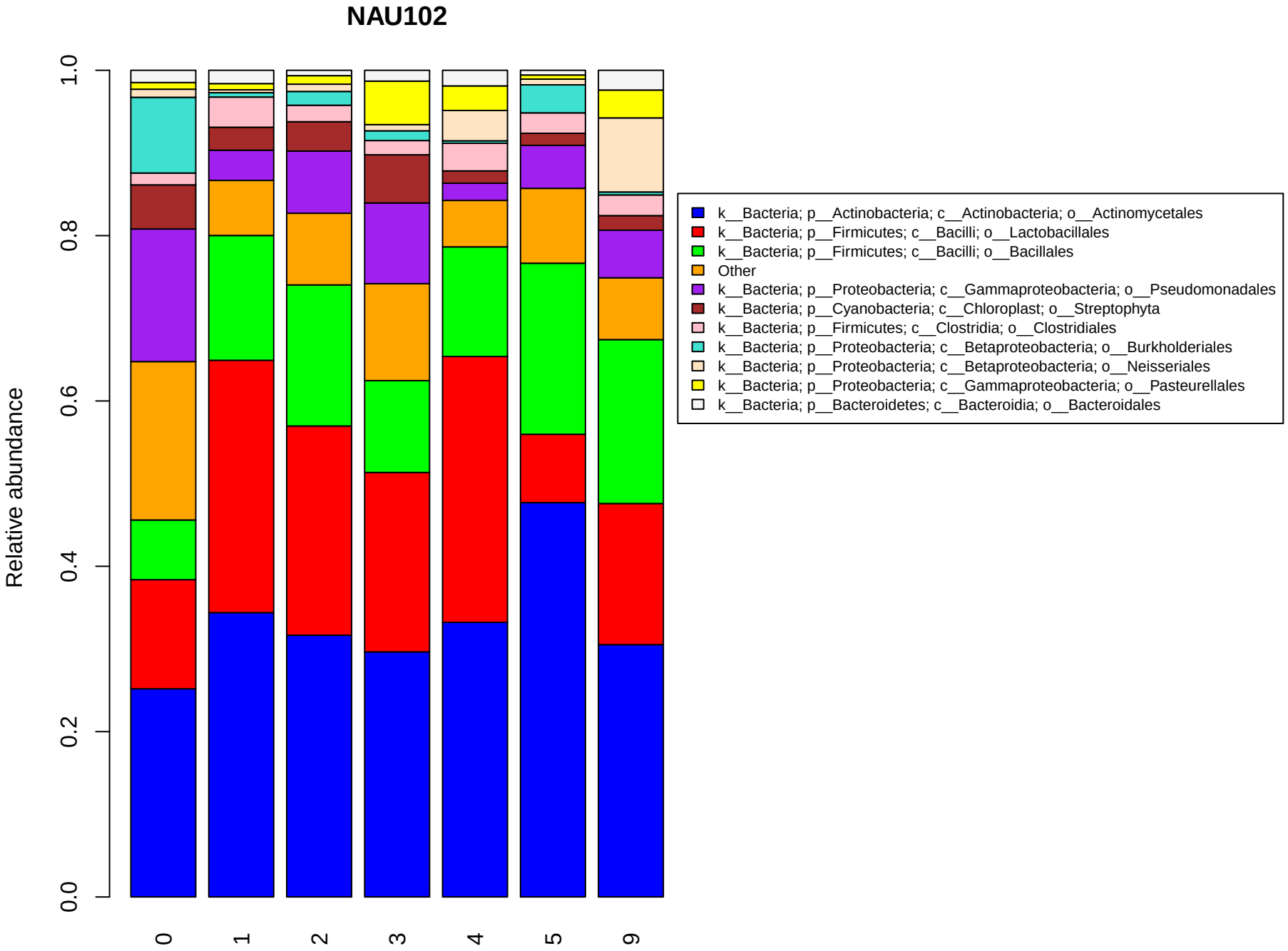
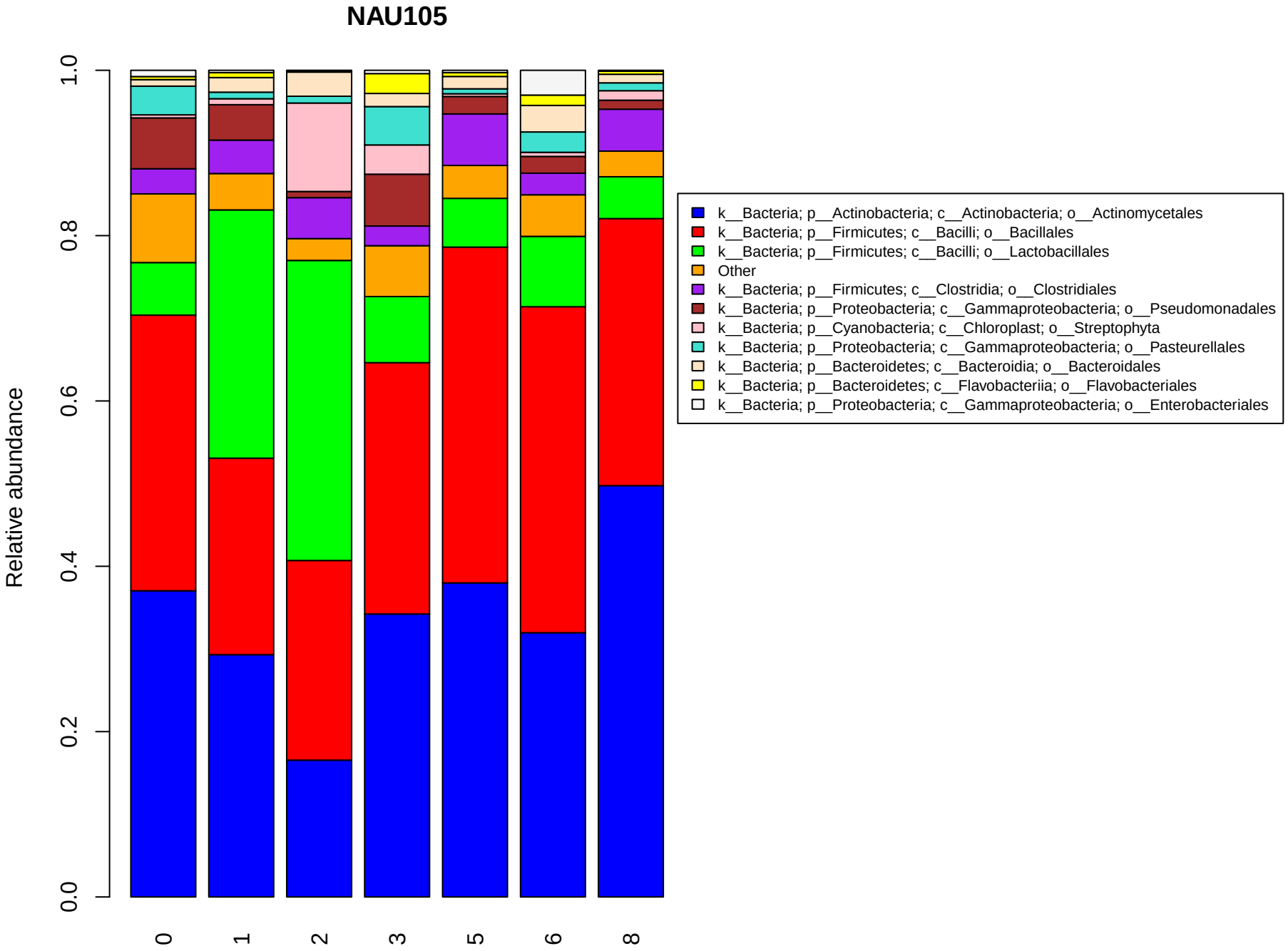
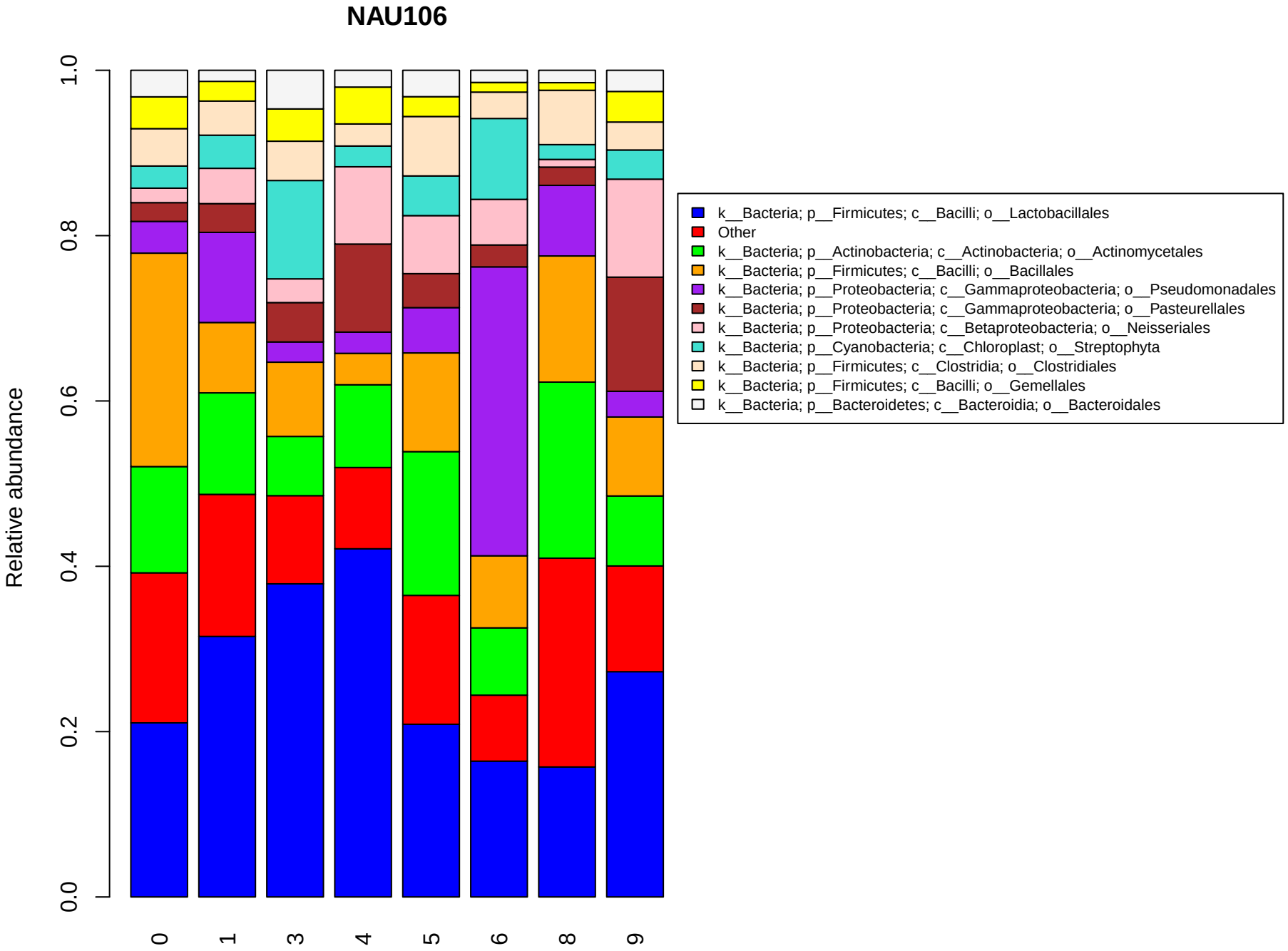
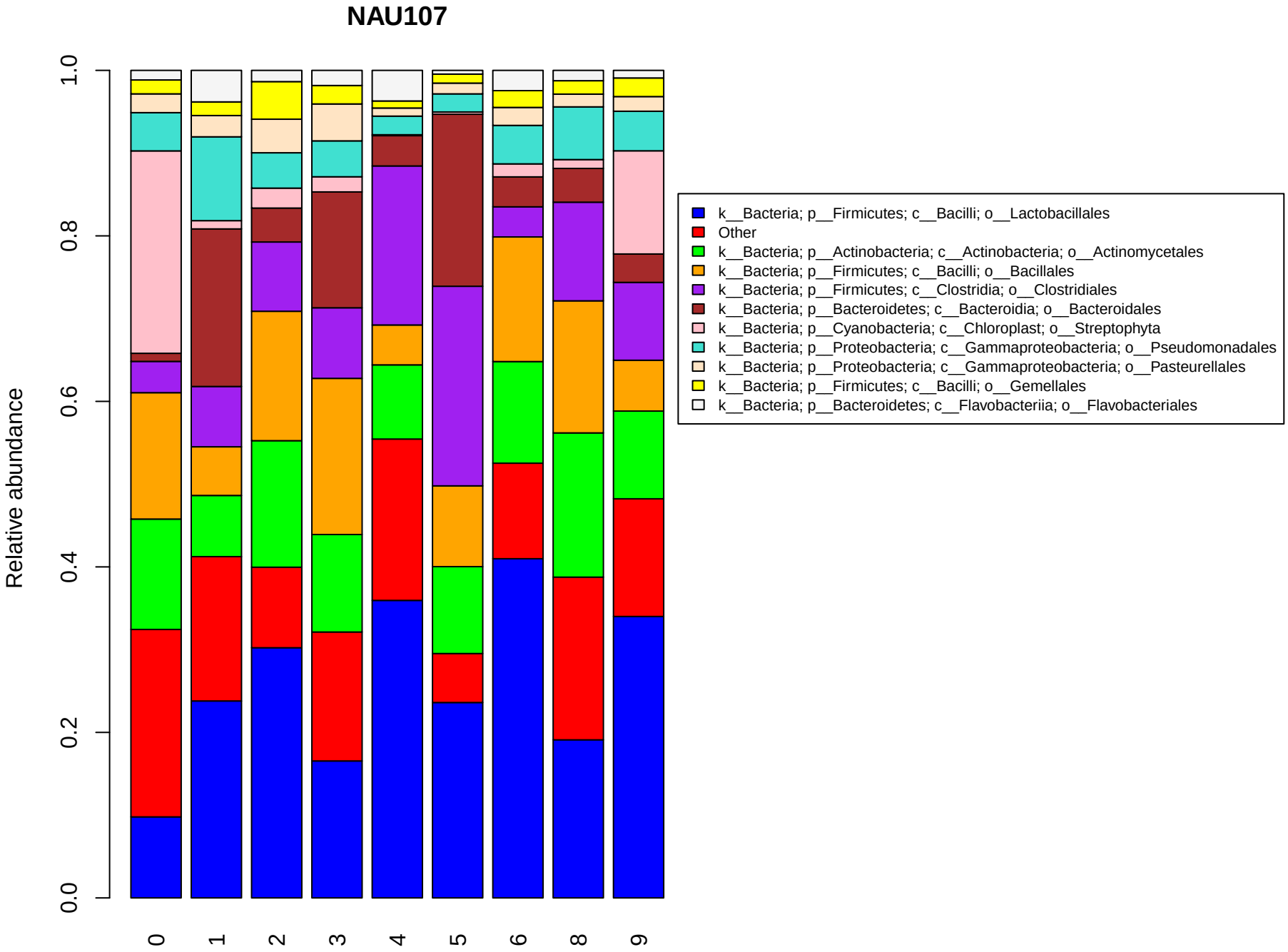
0.4
0.6
0.8
1.0
0
0.5
2
4
5
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales

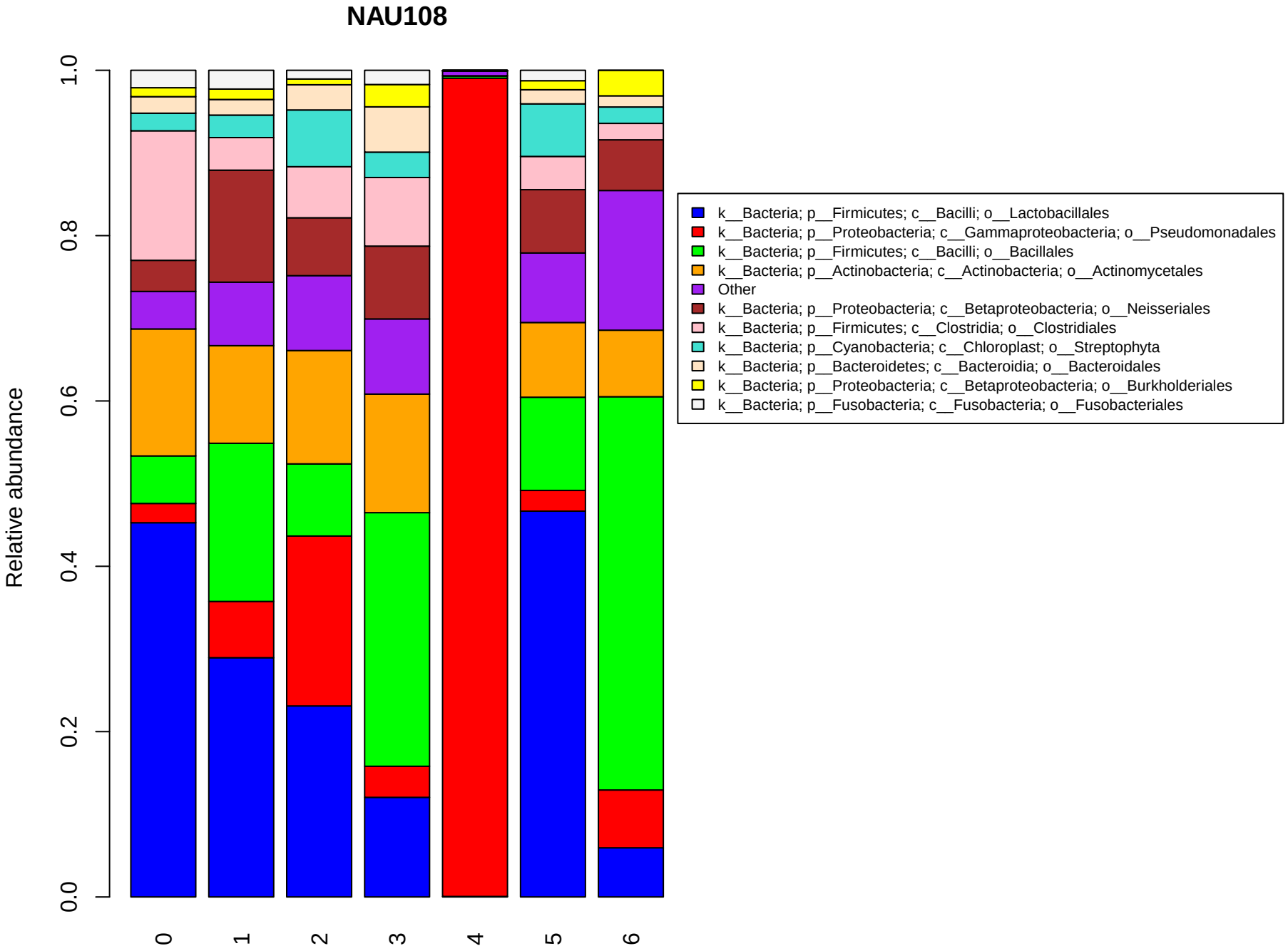

NAU101
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Bifidobacteriales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Clostridia; o__Coriobacteriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales

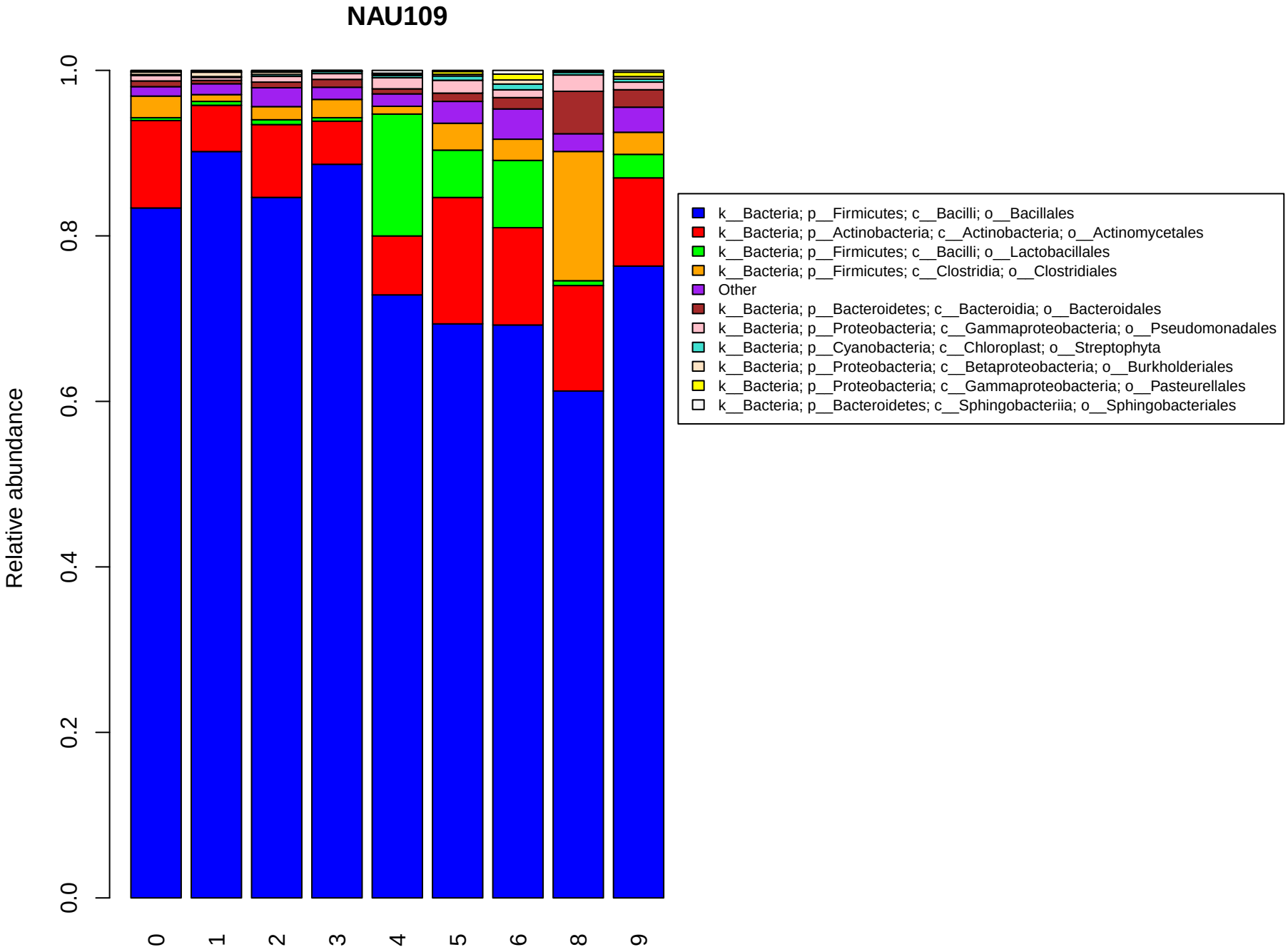

NAU102
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales

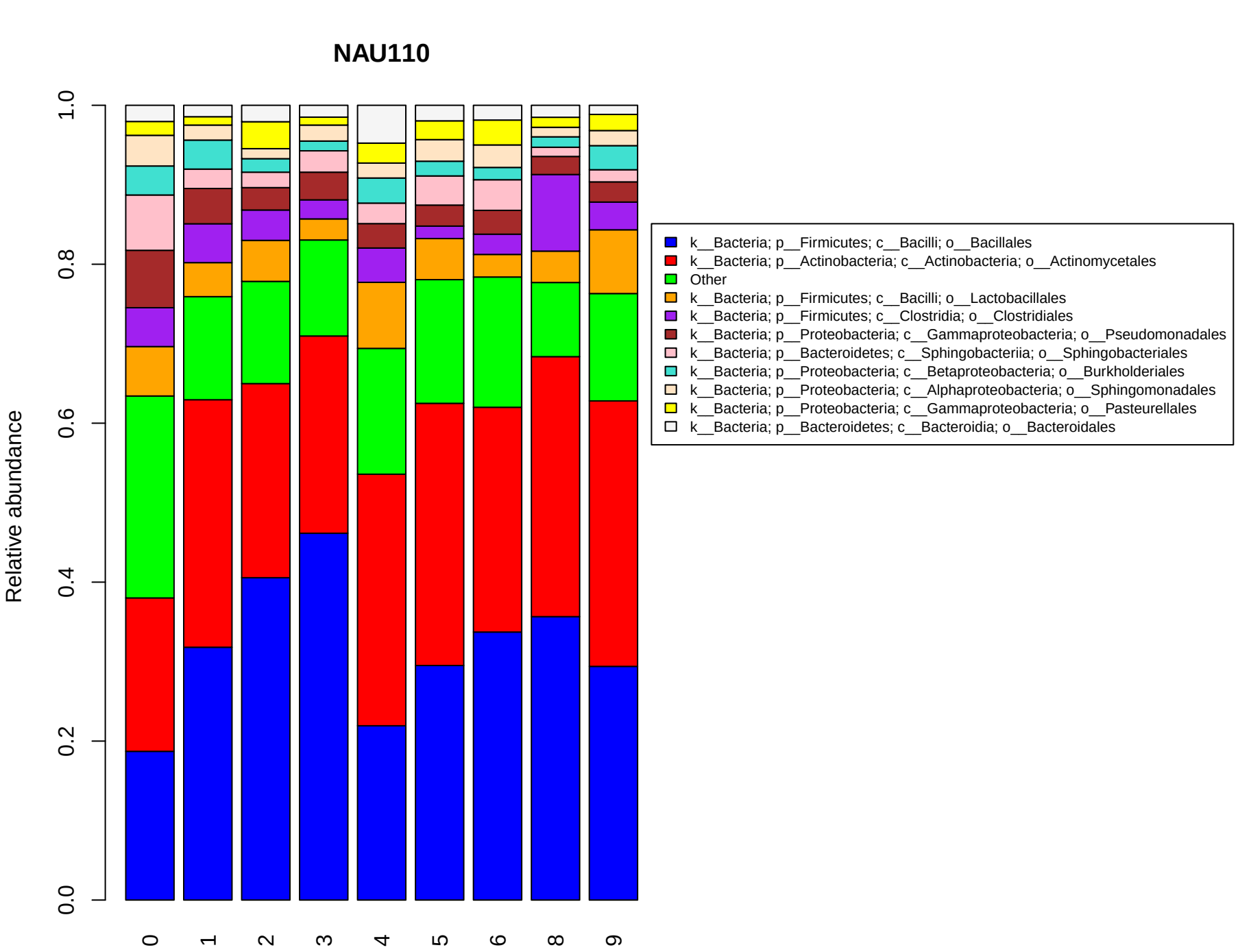

NAU105
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
8
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales

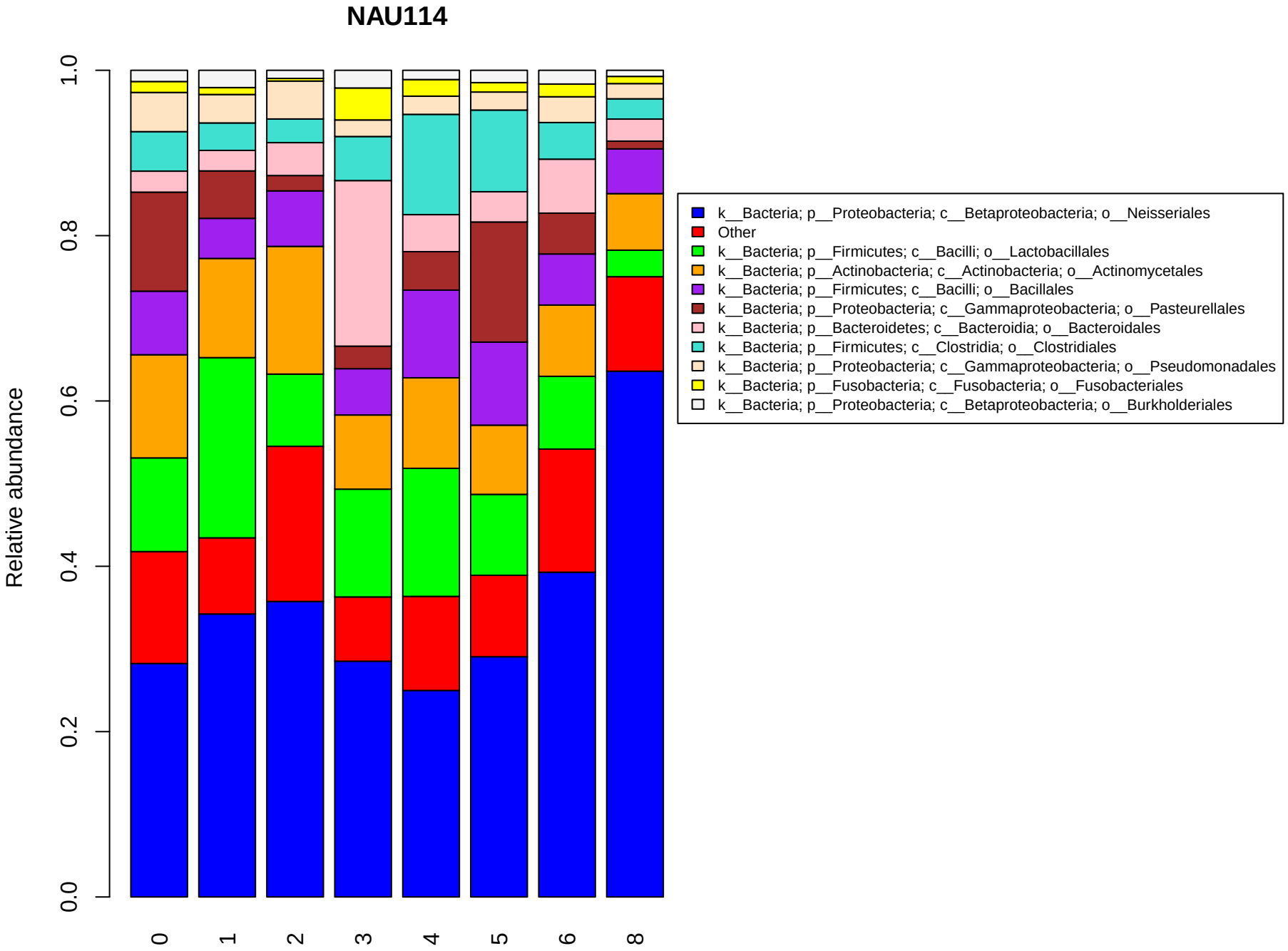

NAU106
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales

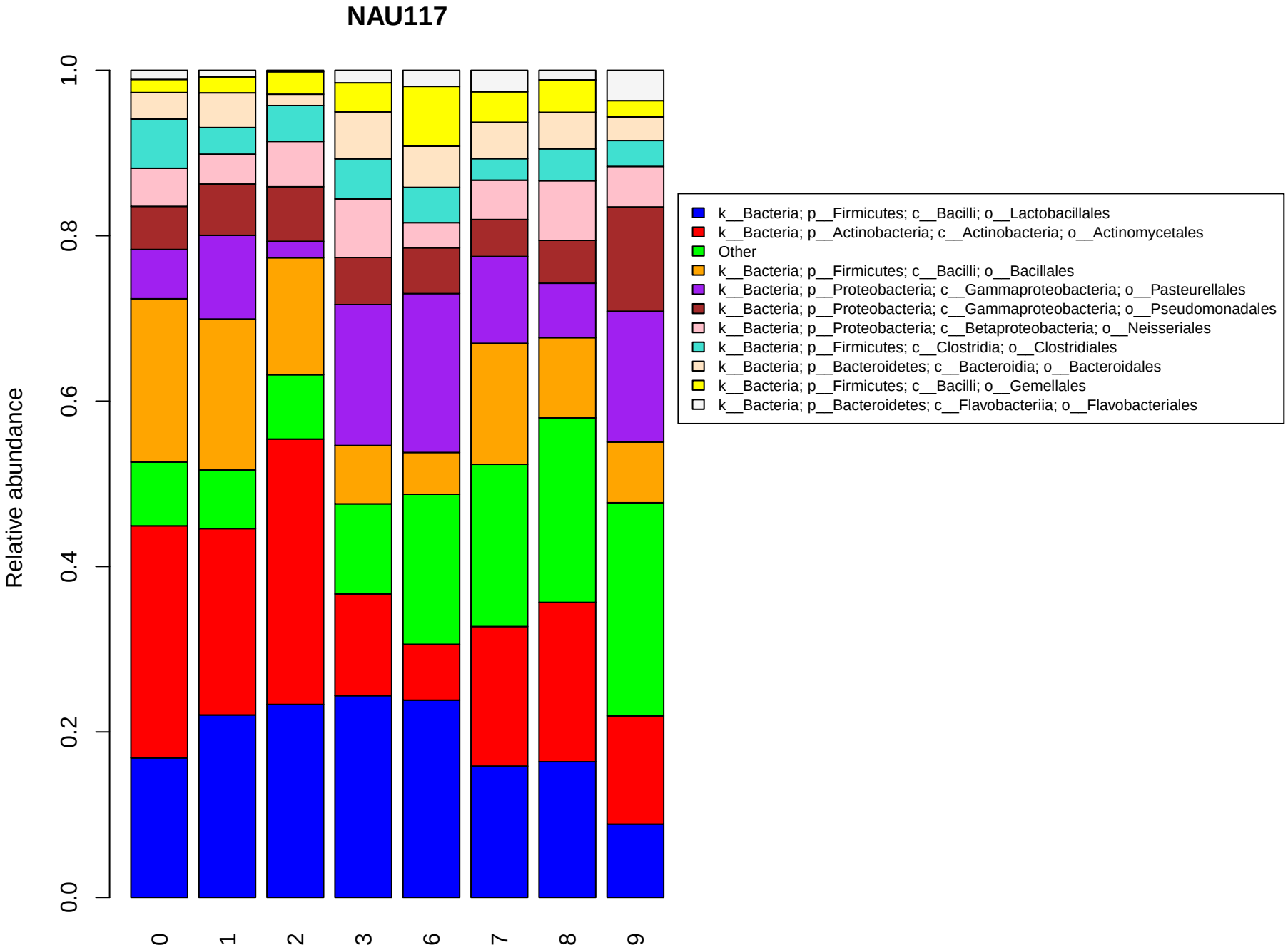

NAU107
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales

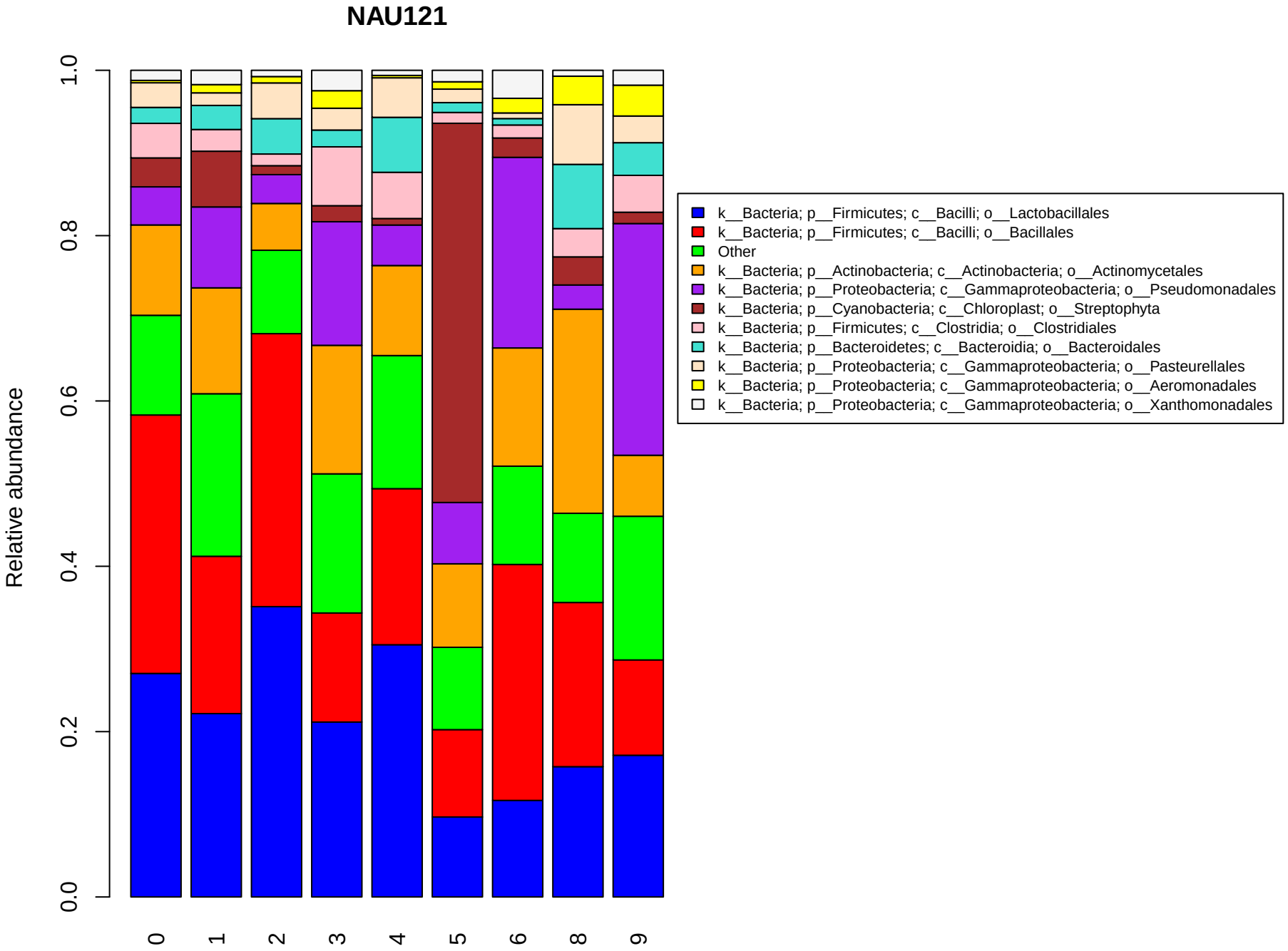

NAU108
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales

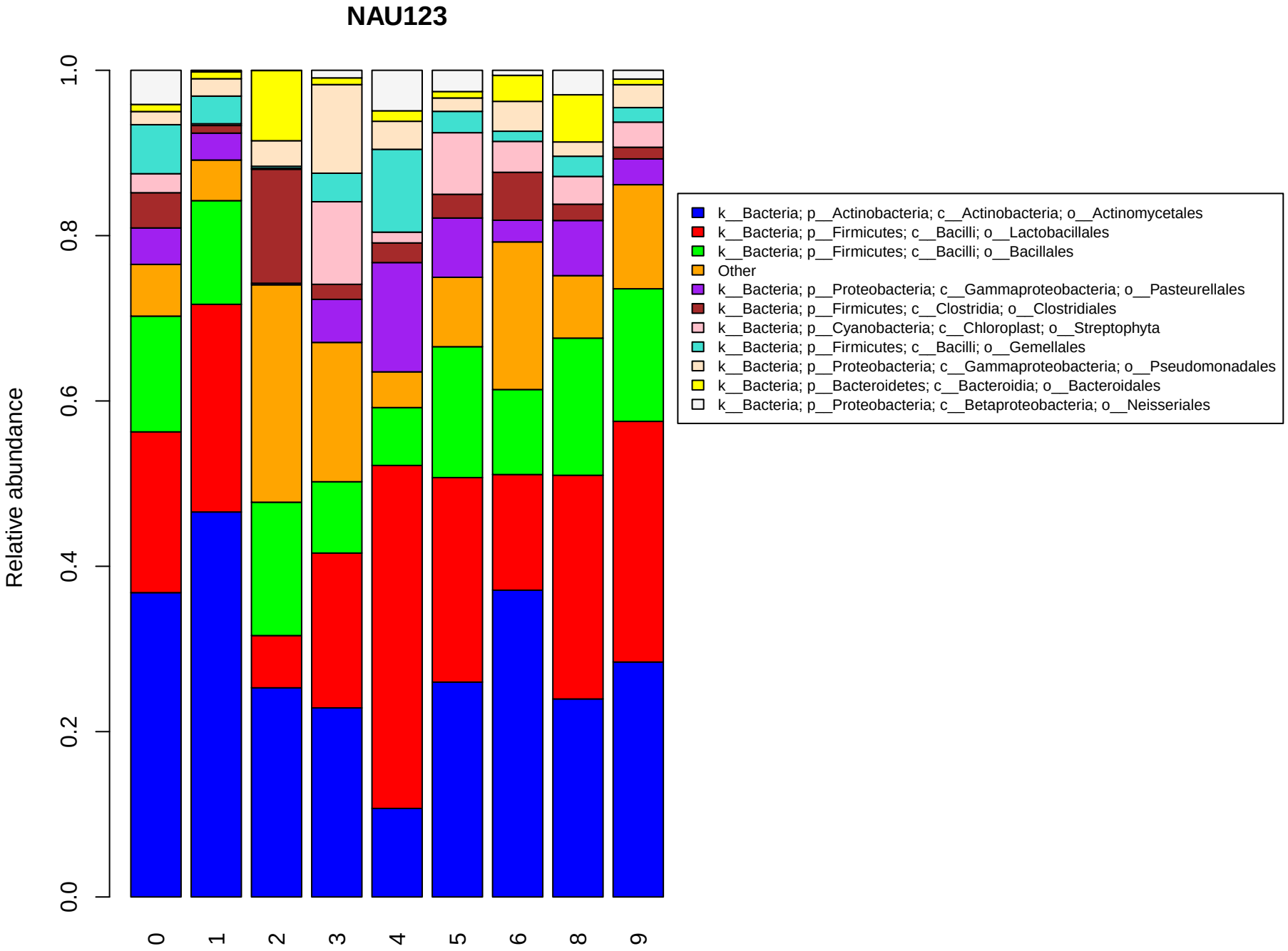

NAU109
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
Other
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales

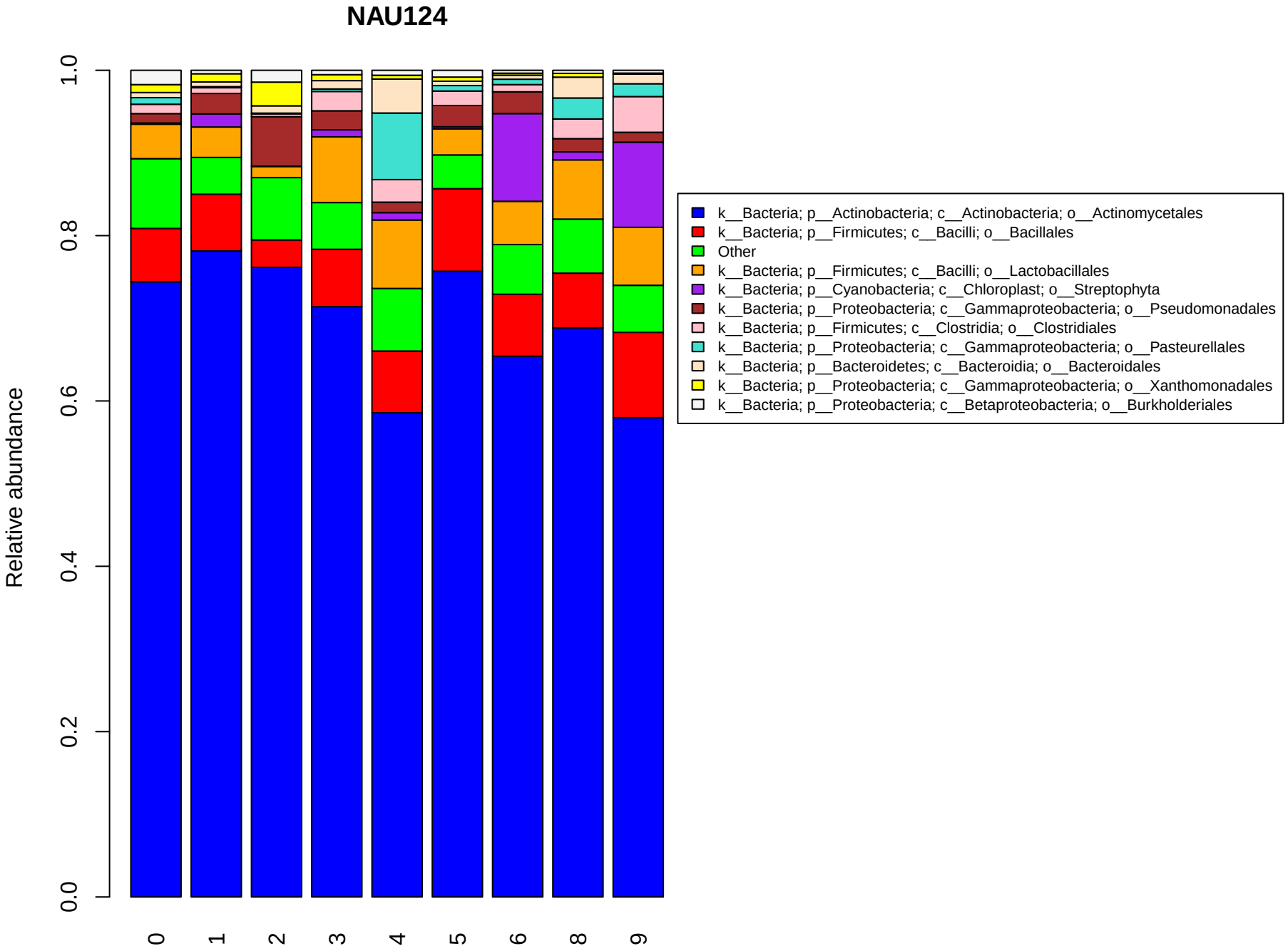

NAU110
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales

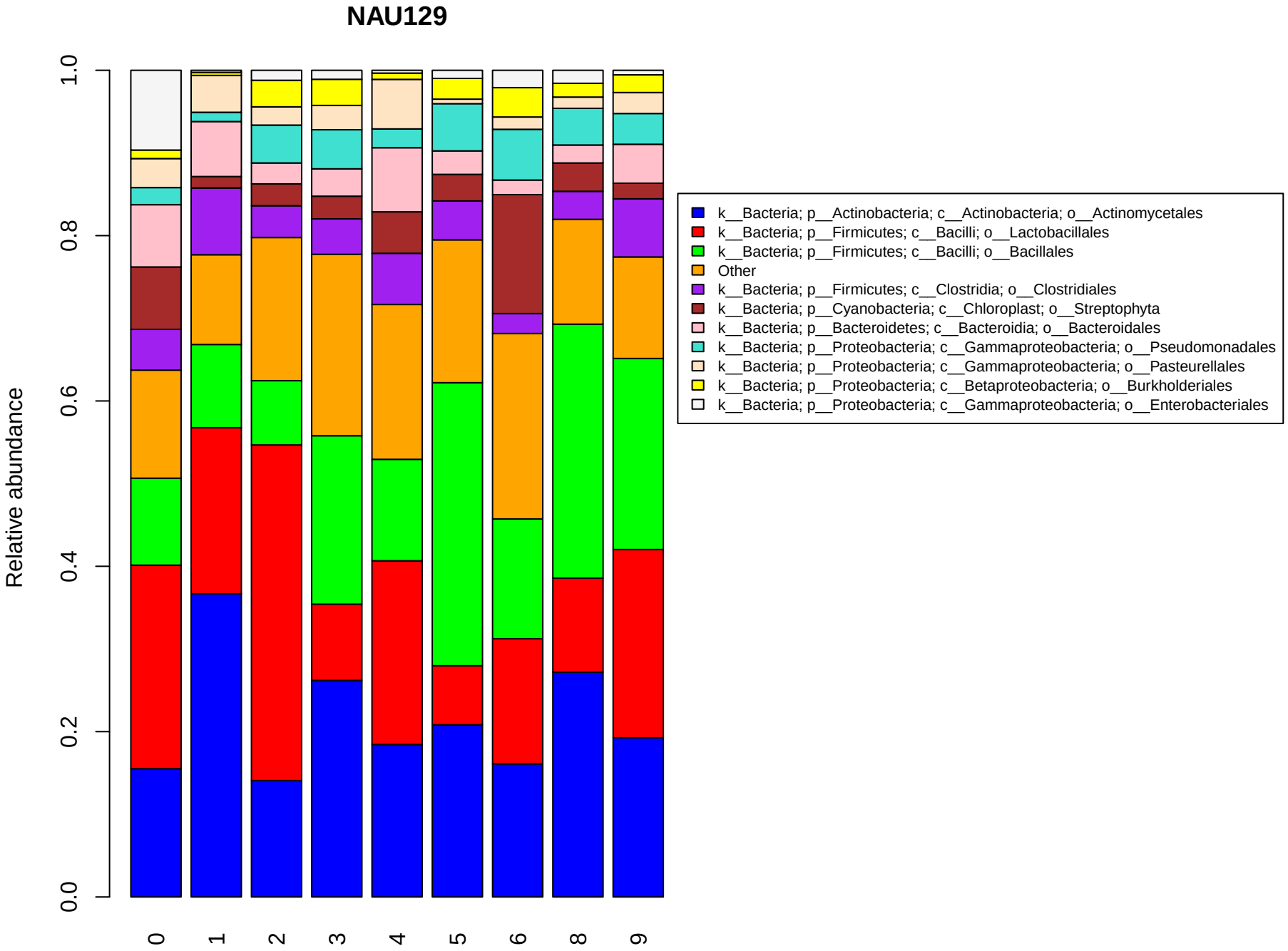

NAU114
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales

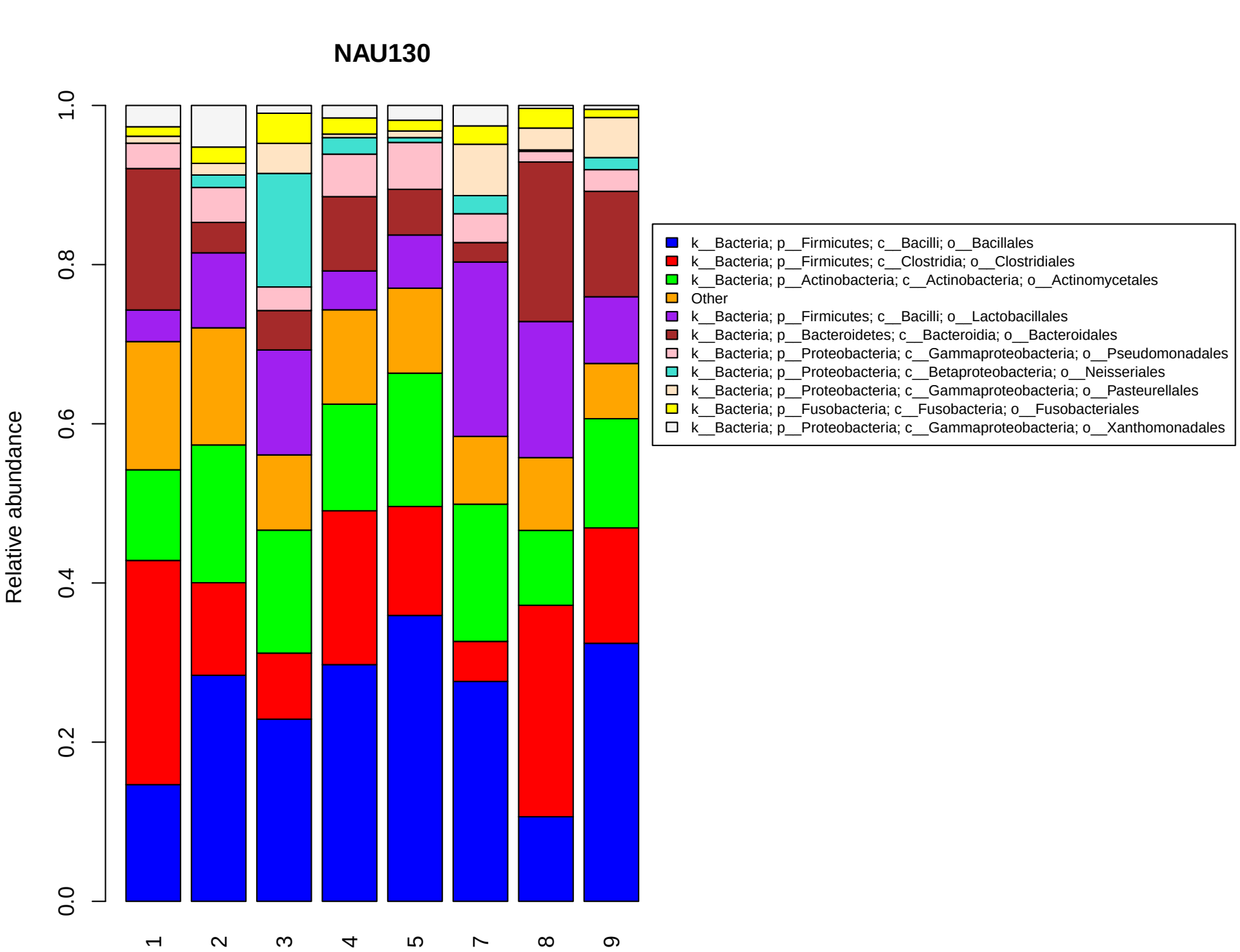

NAU117
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
6
7
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales

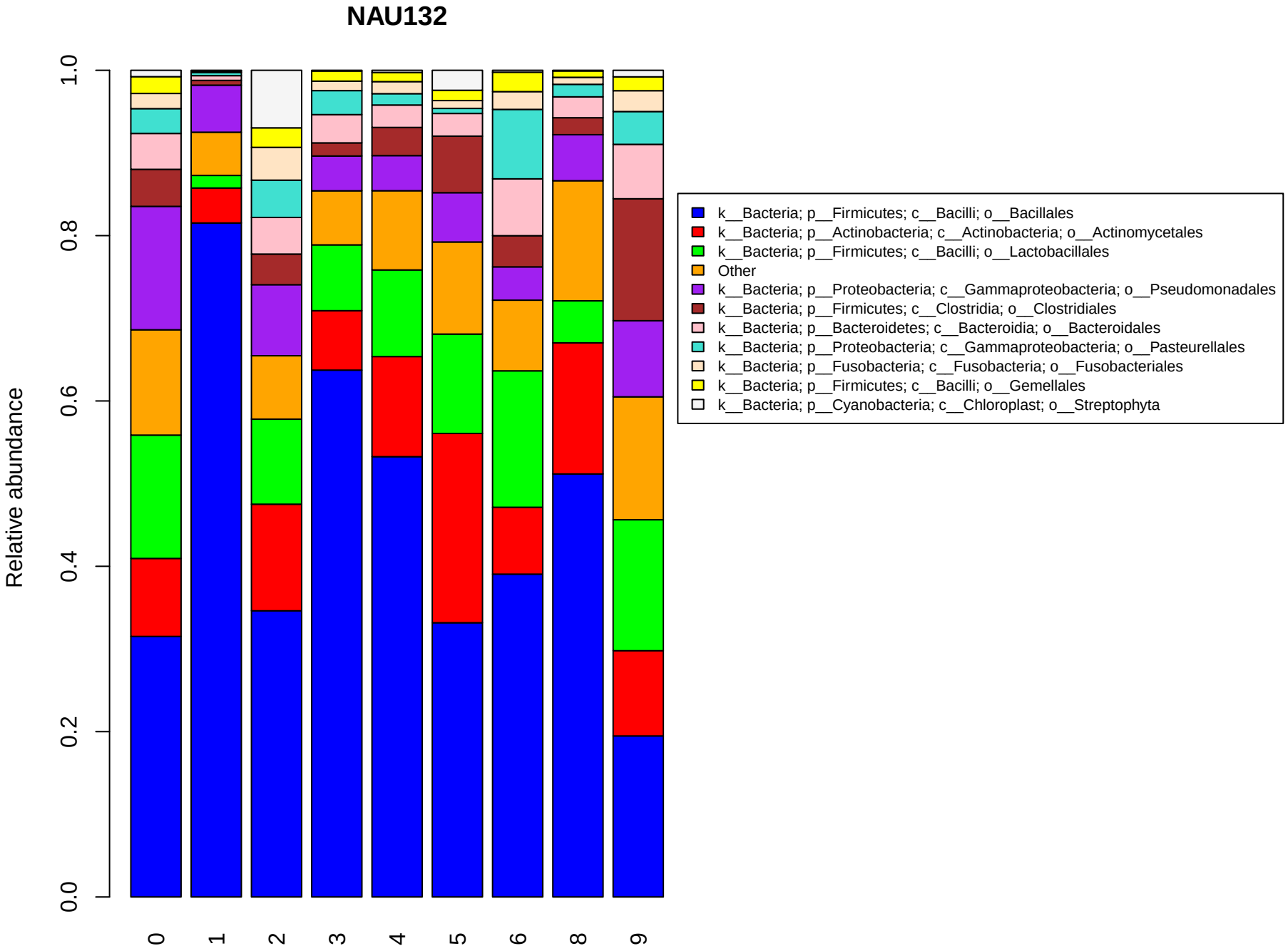

NAU121
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales

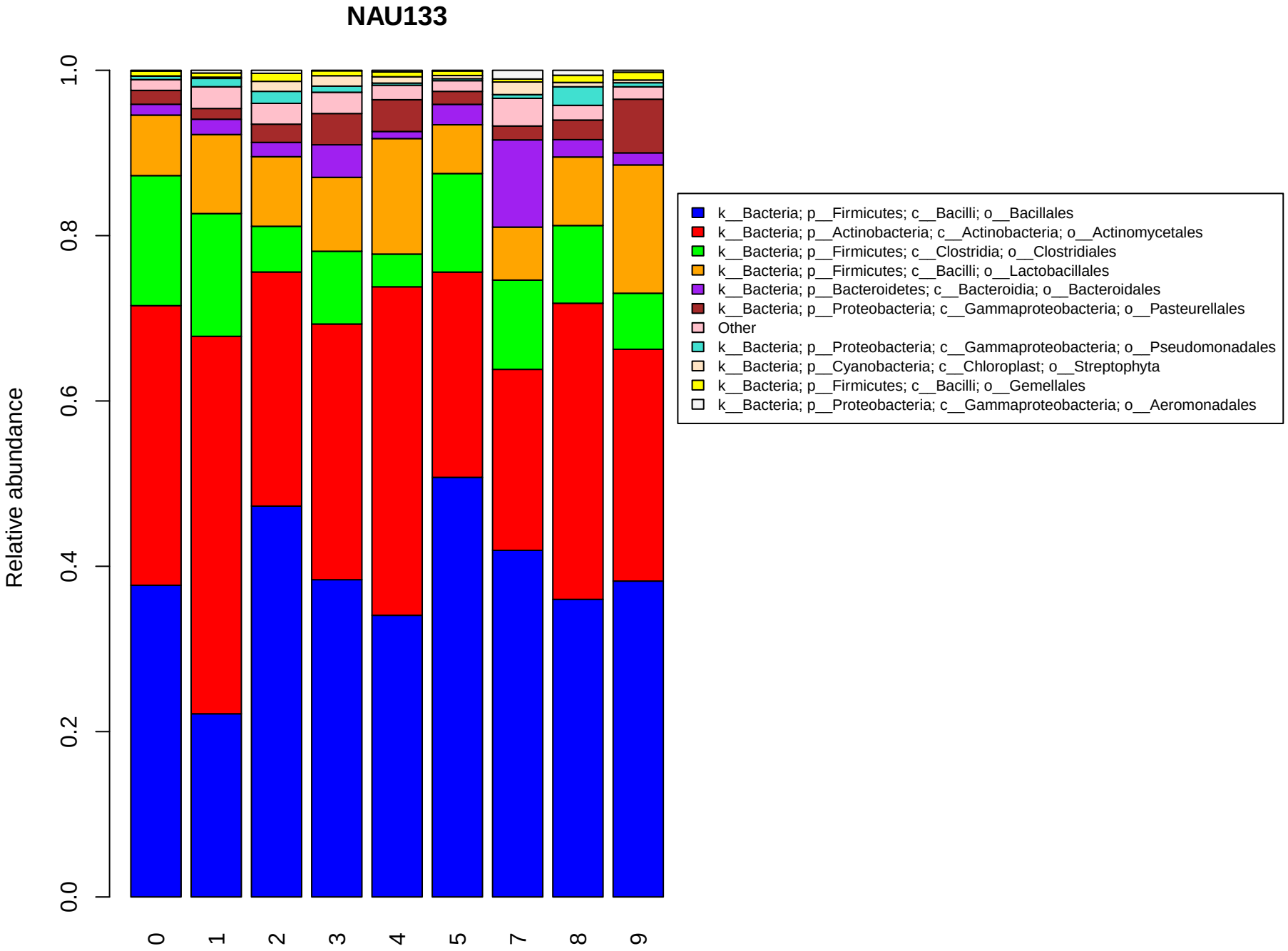

NAU123
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales

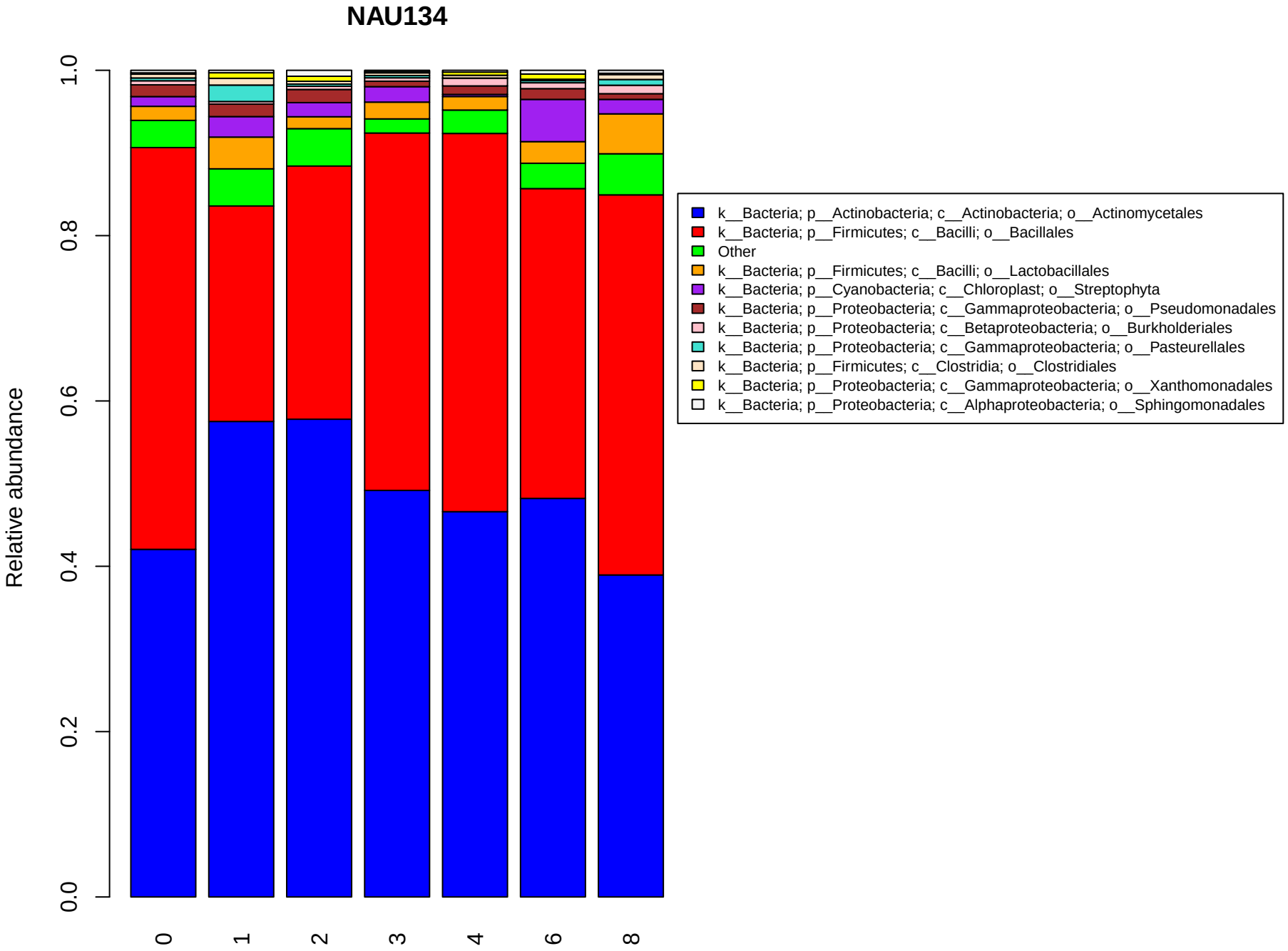

NAU124
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales

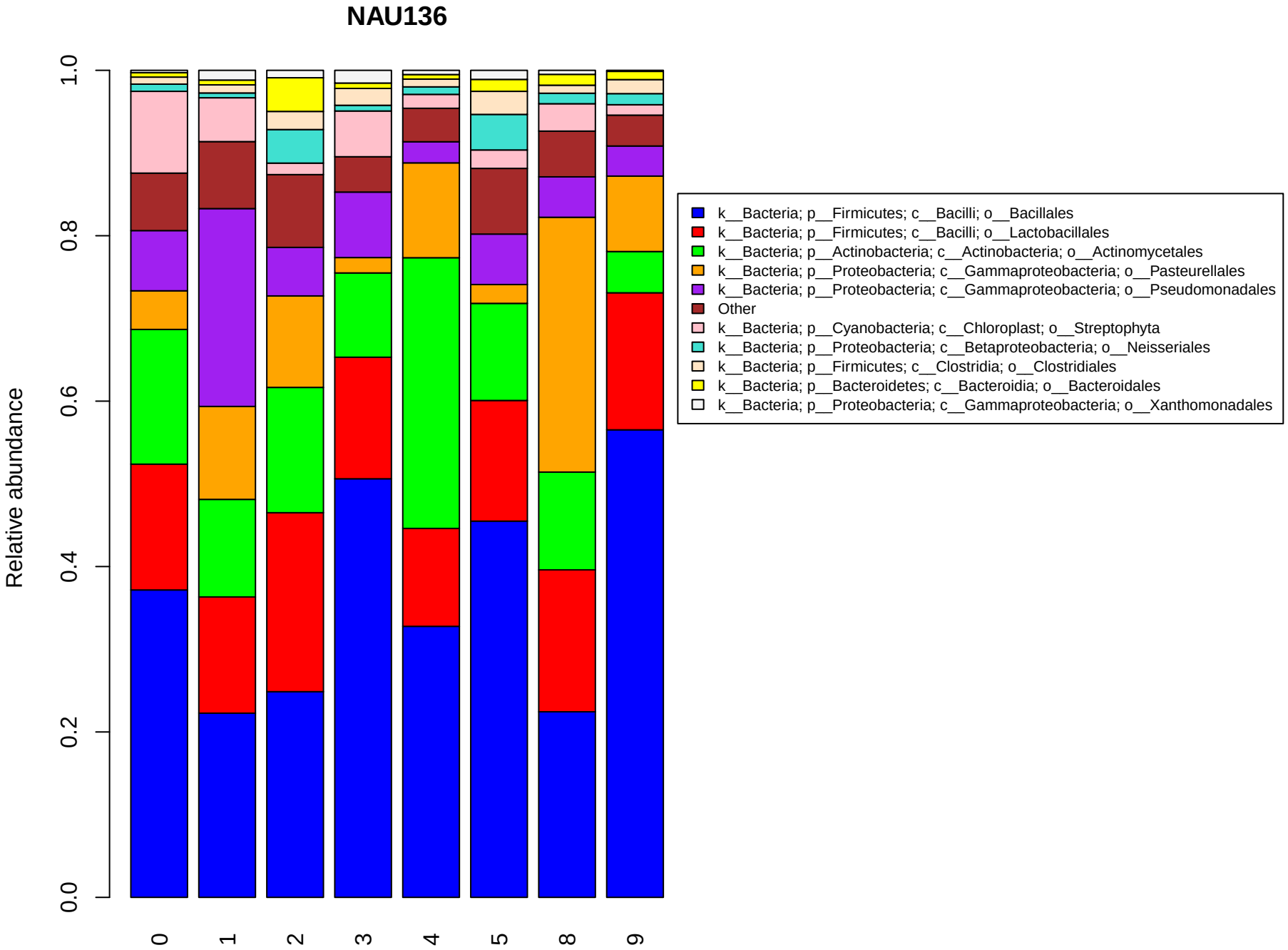

NAU129
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales

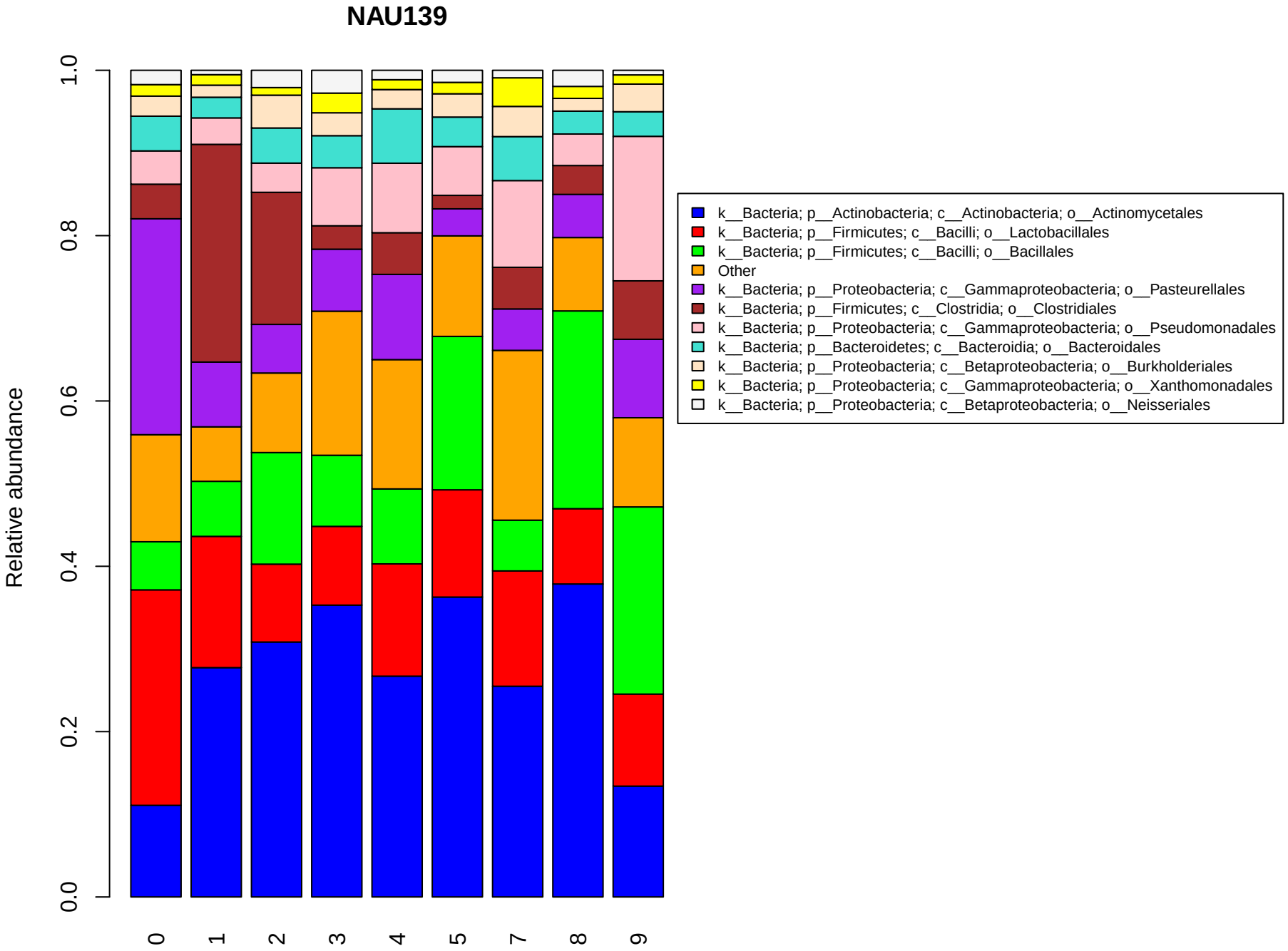

NAU130
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
1
2
3
4
5
7
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales

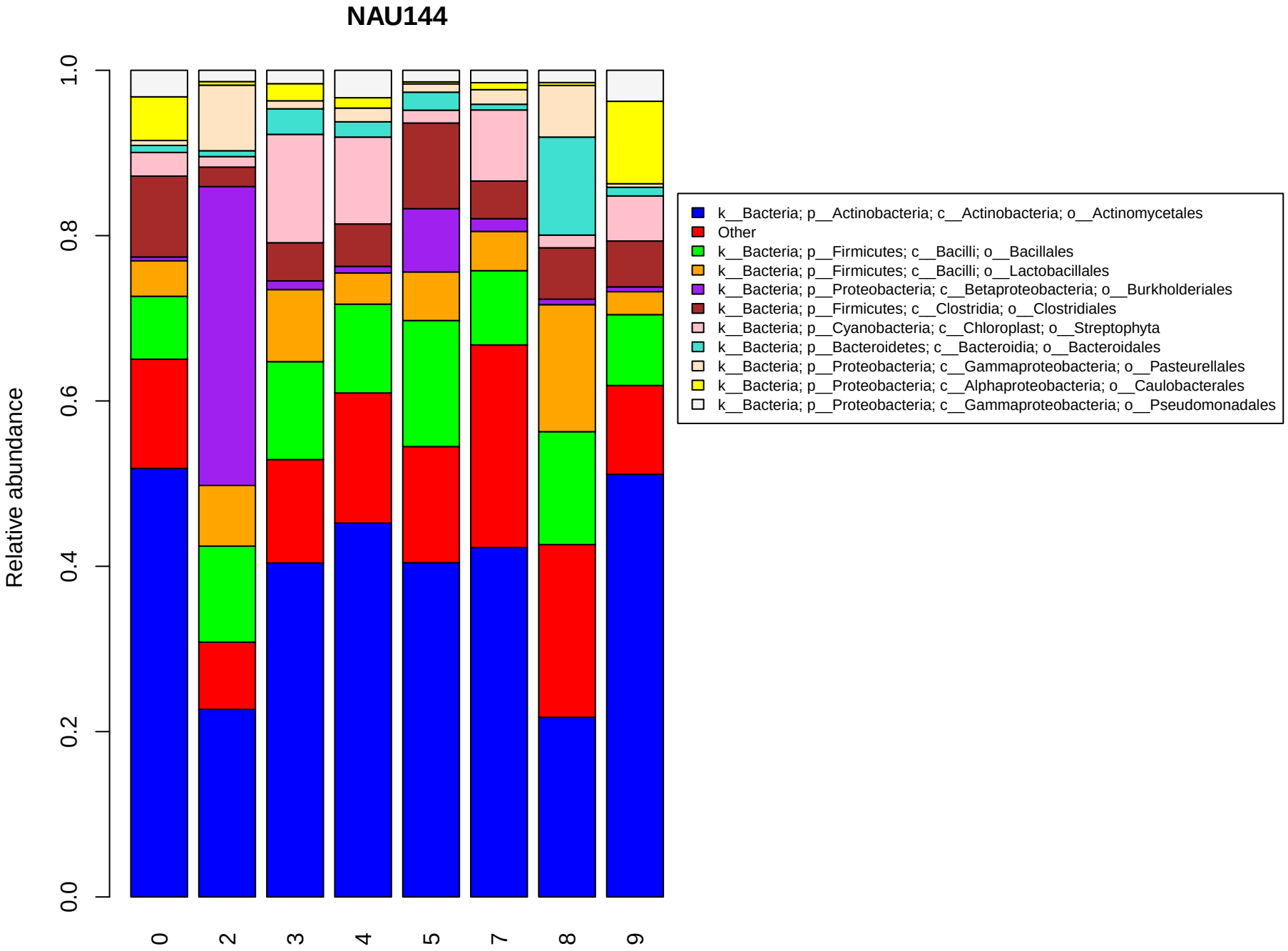

NAU132
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta

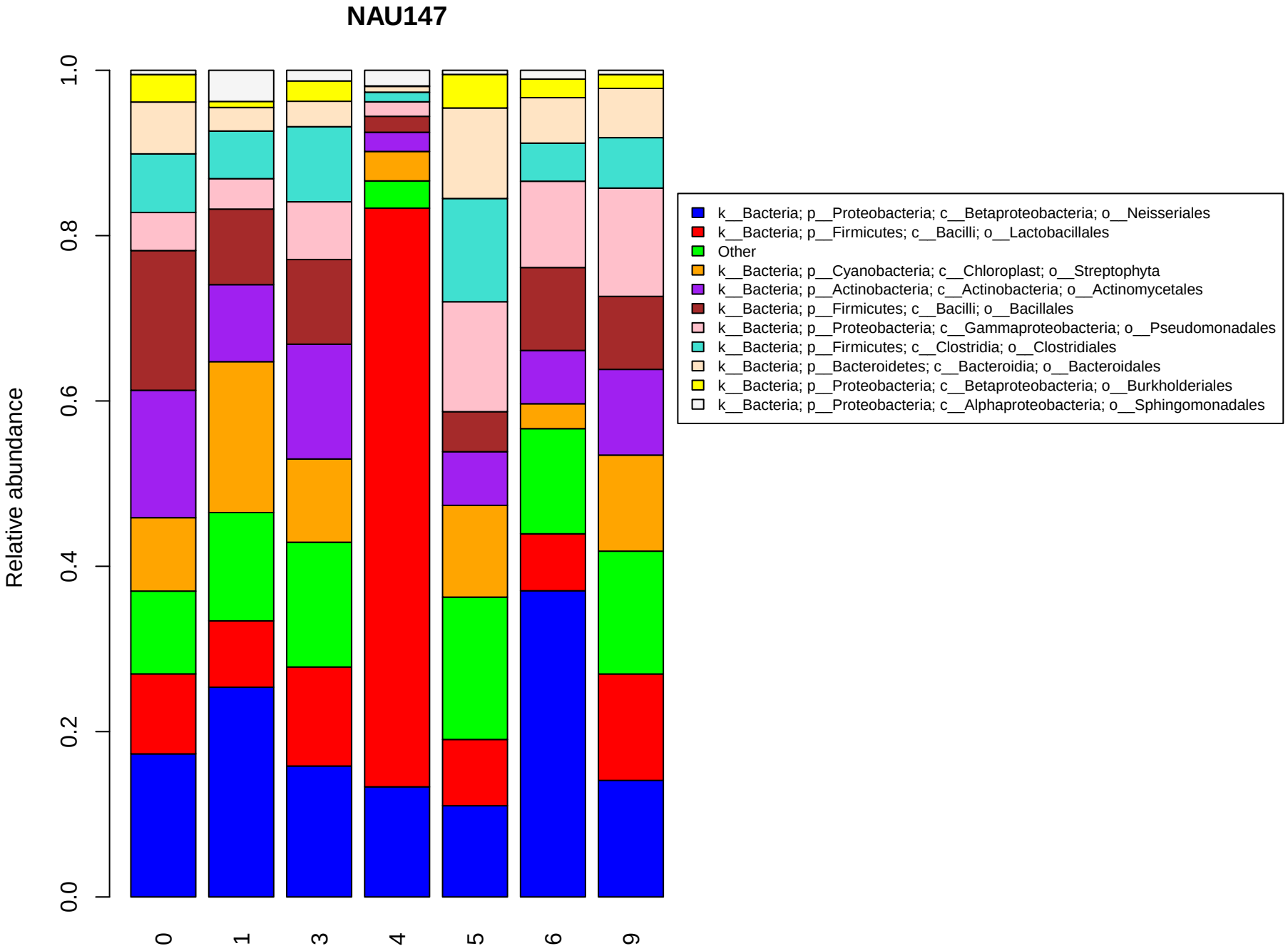

NAU133
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
7
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales

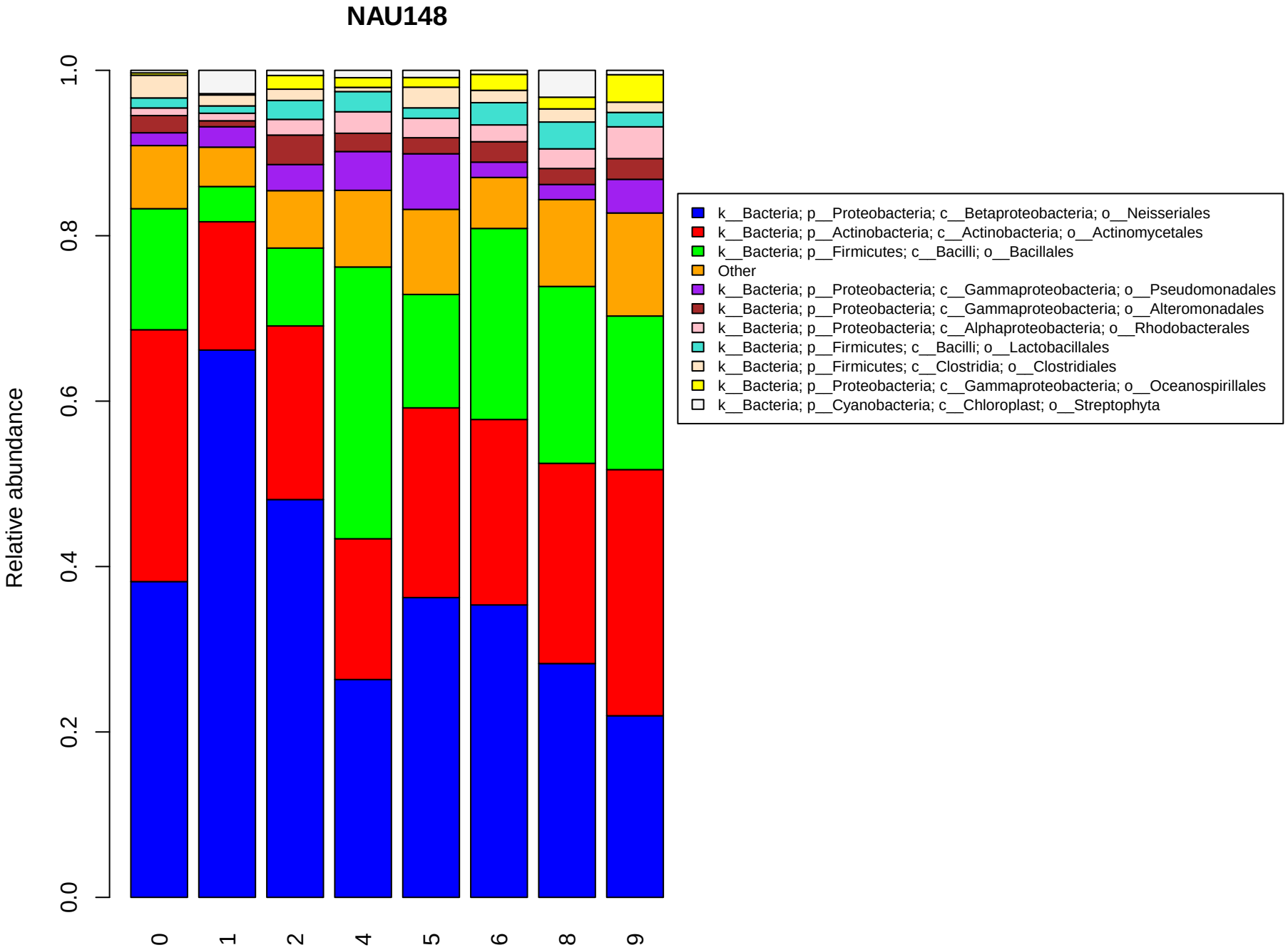

NAU134
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
6
8
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales

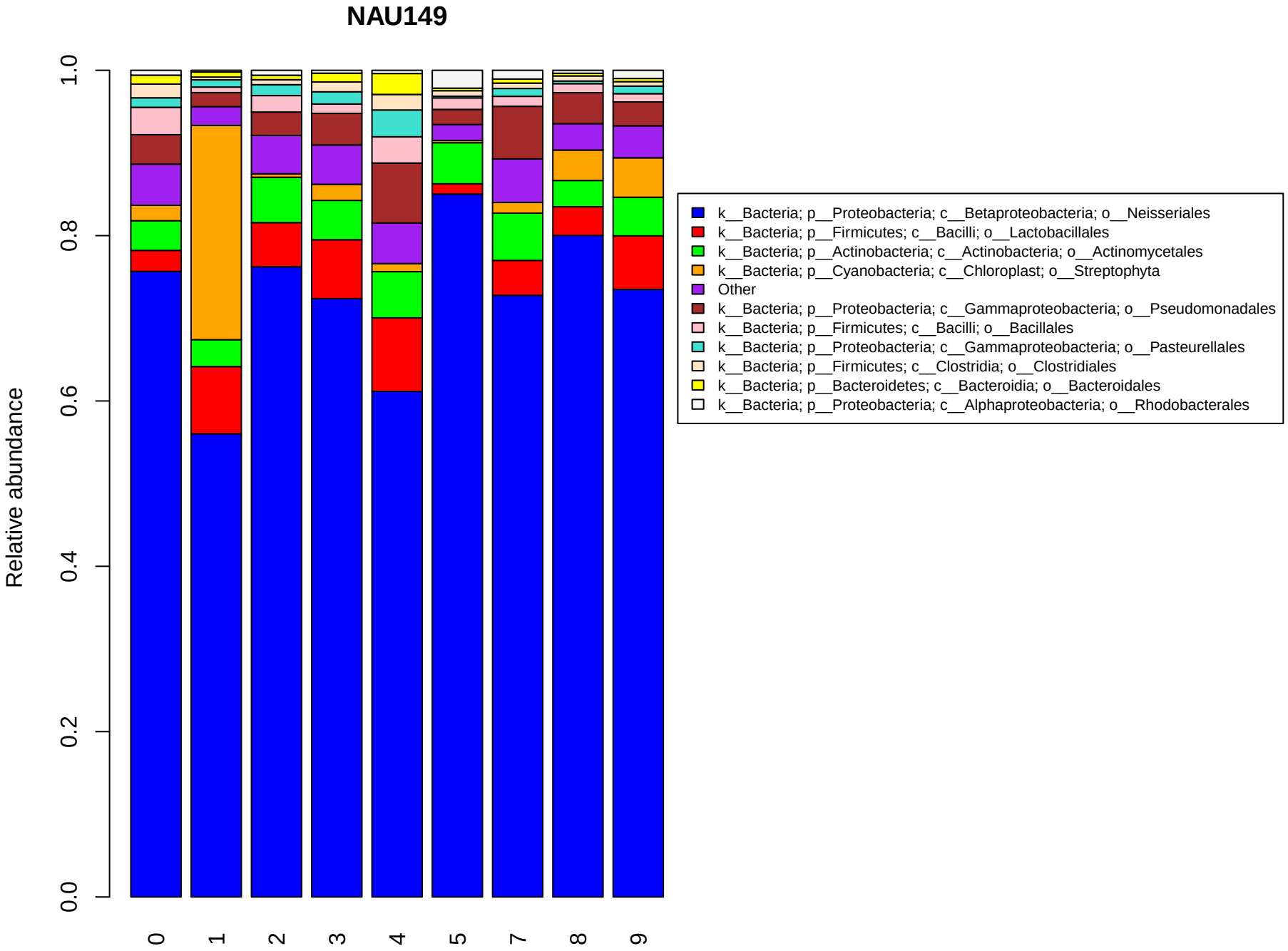

NAU136
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales

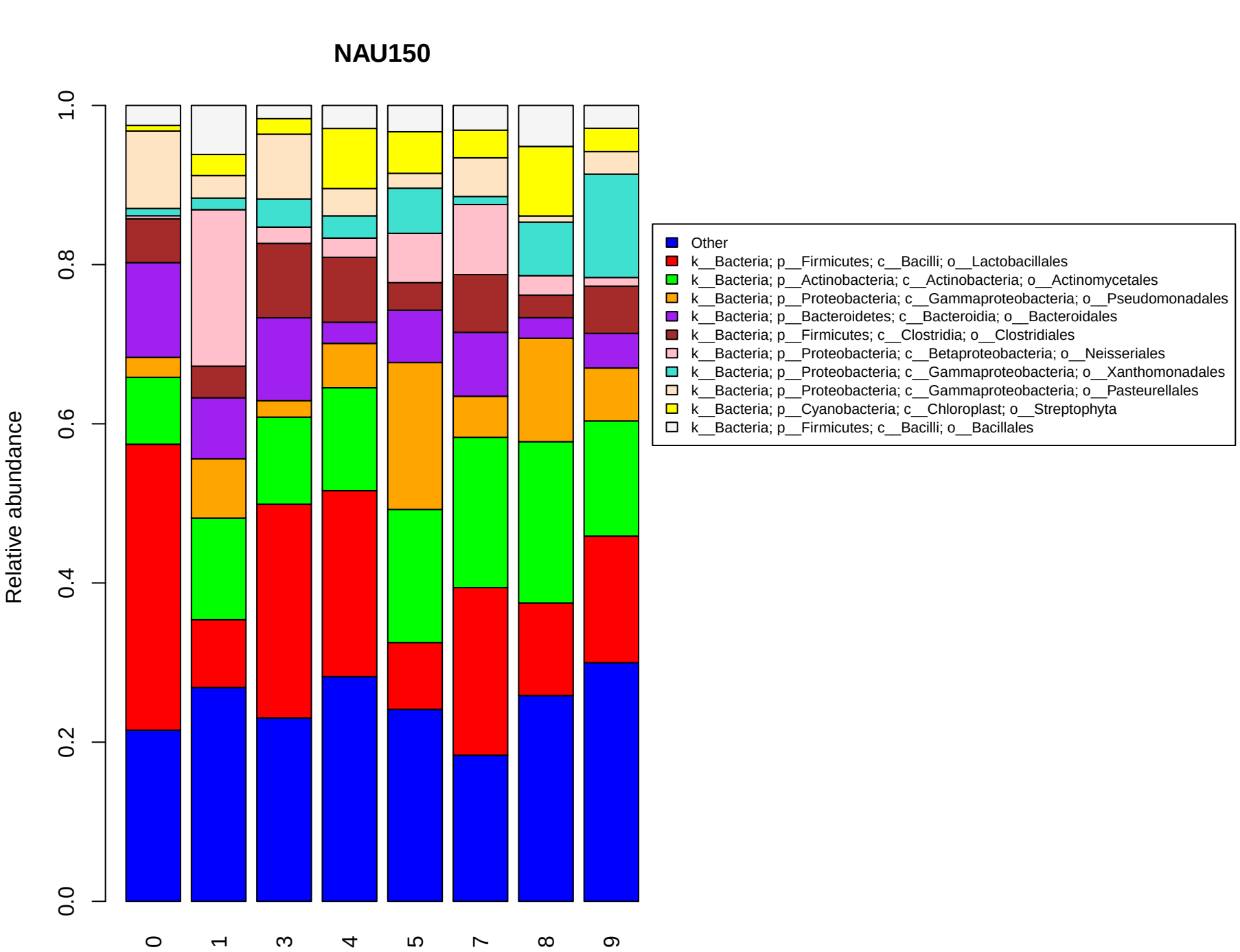

NAU139
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
7
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales

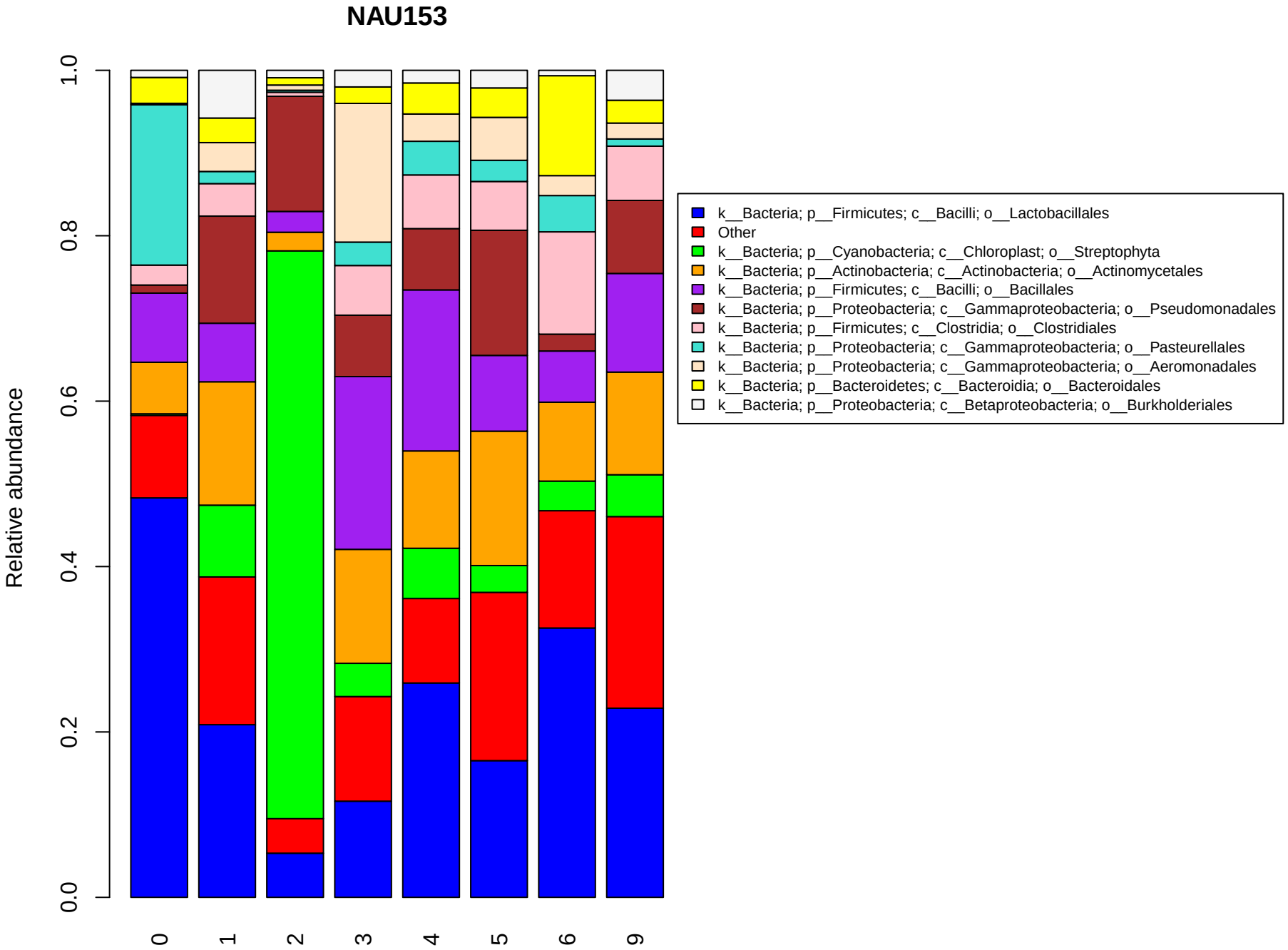

NAU144
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
2
3
4
5
7
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Caulobacterales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales

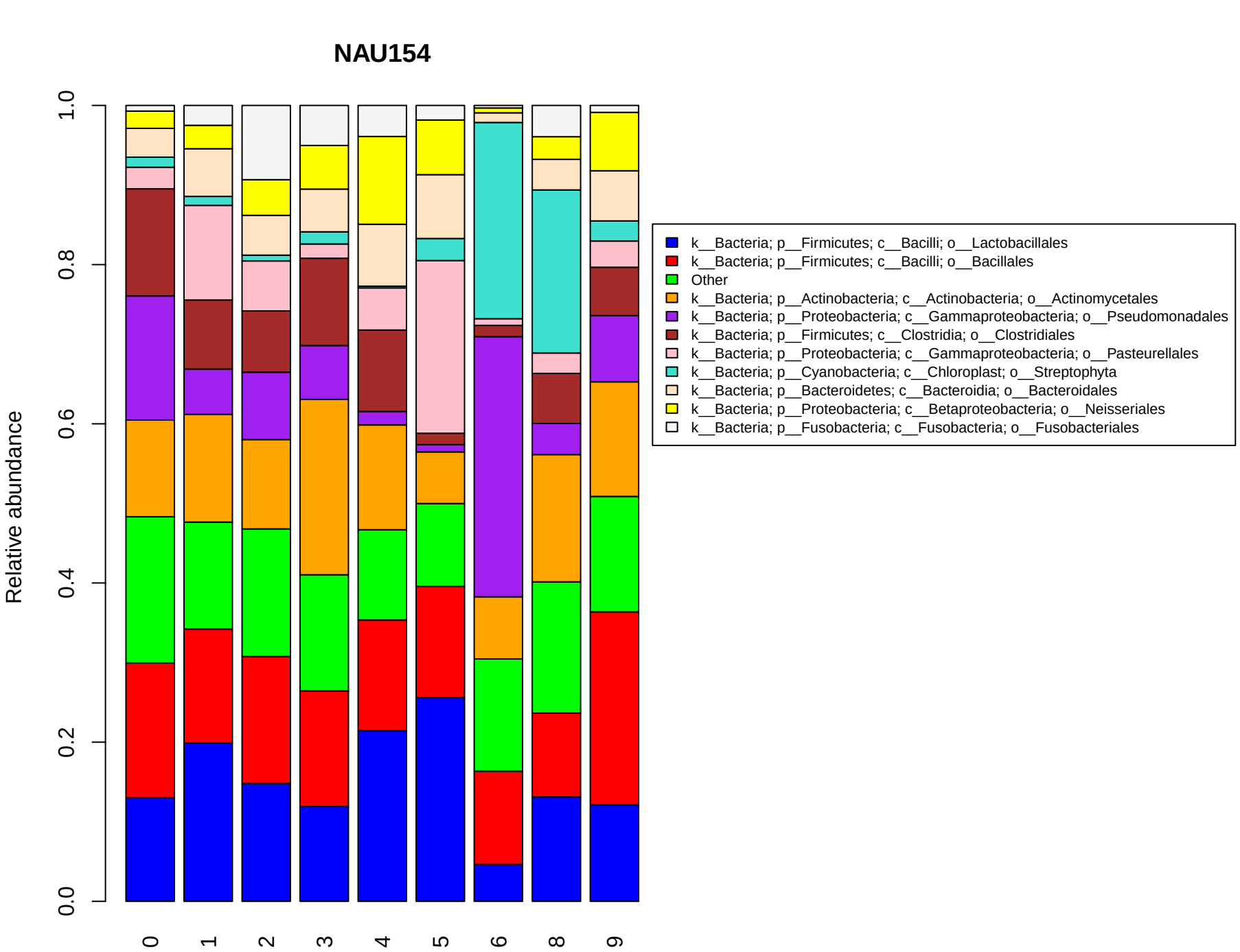

NAU147
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
3
4
5
6
9
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales

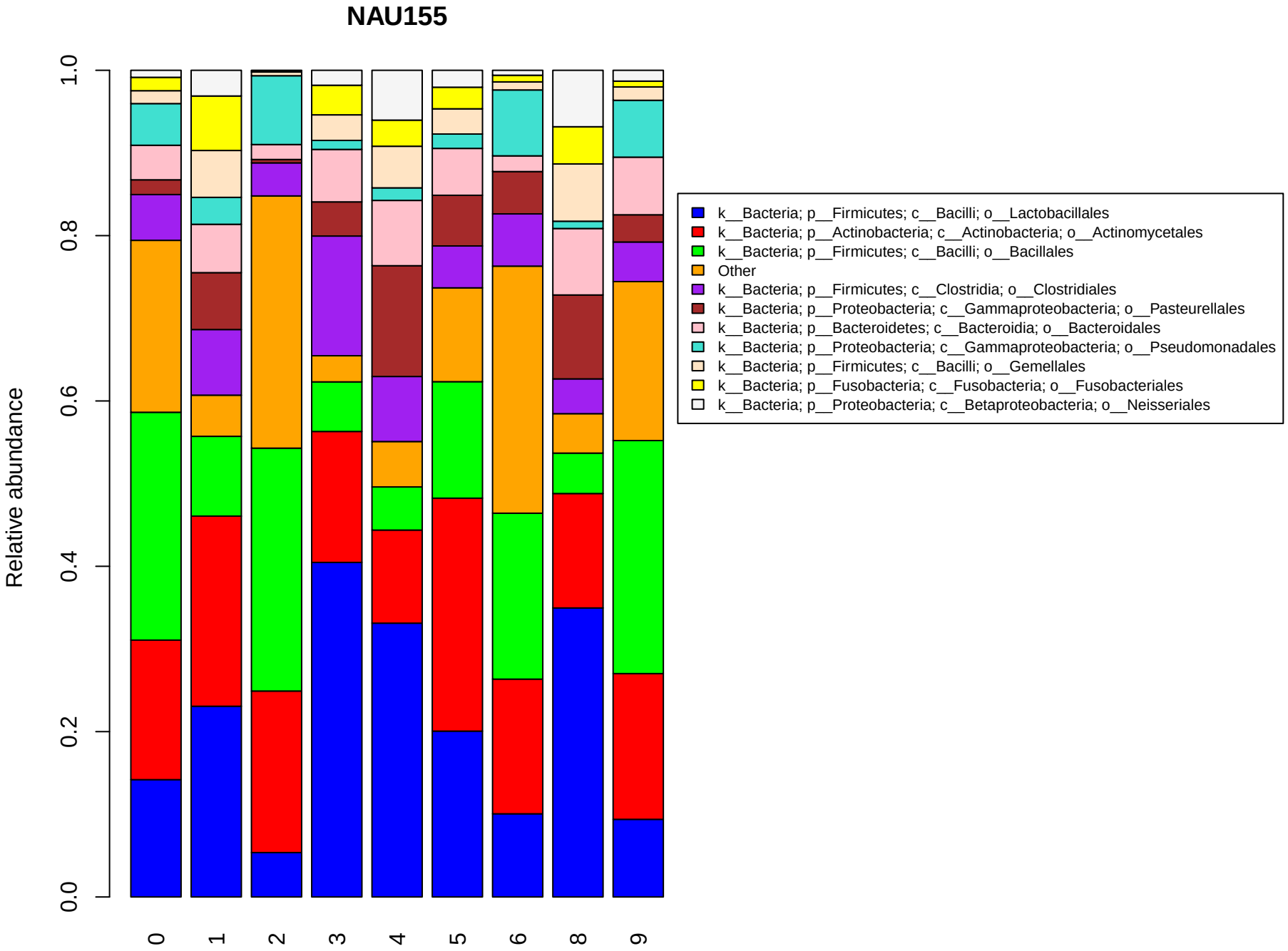

NAU148
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
4
5
6
8
9
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Alteromonadales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodobacterales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Oceanospirillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta

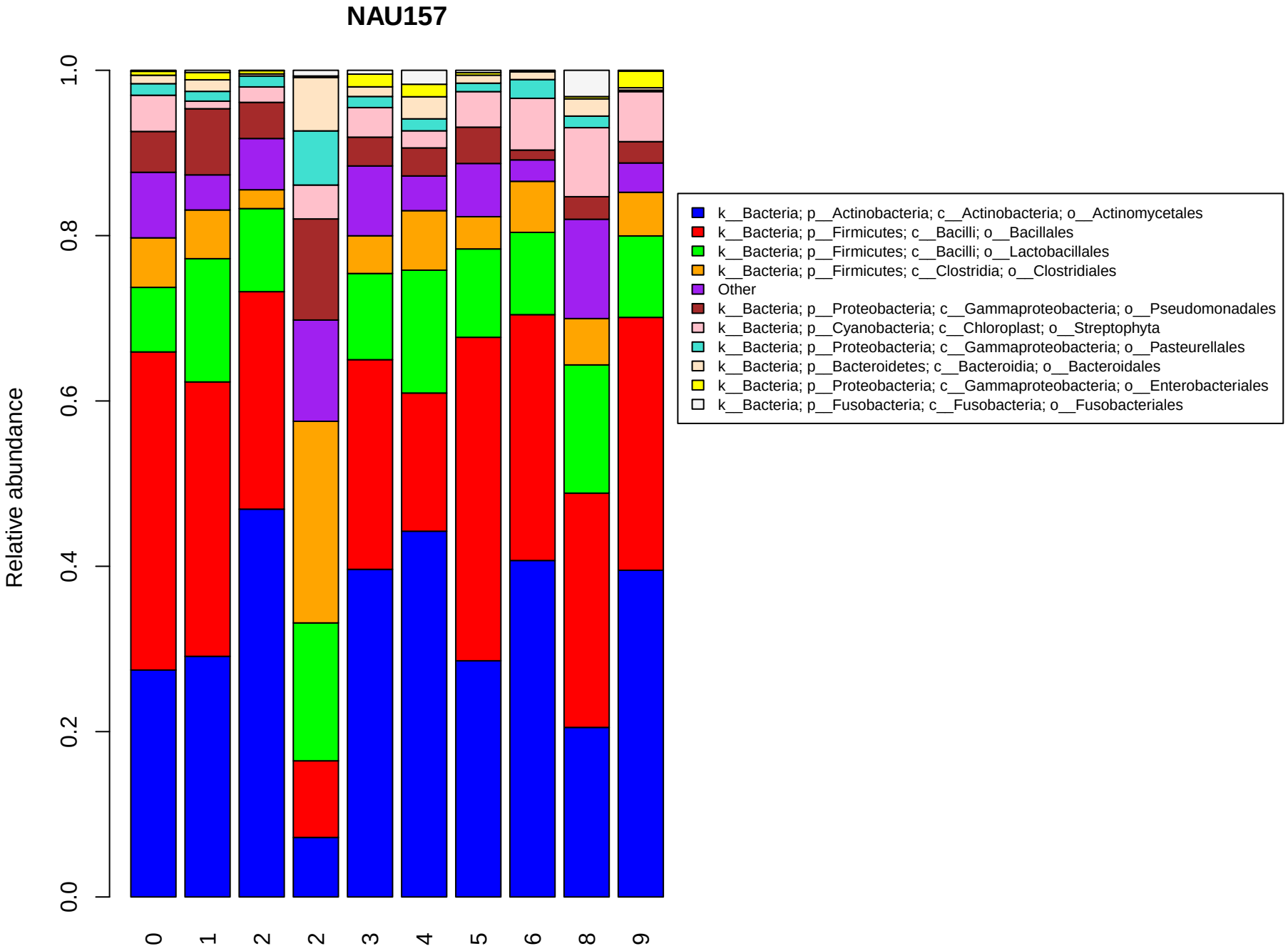

NAU149
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
7
8
9
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodobacterales

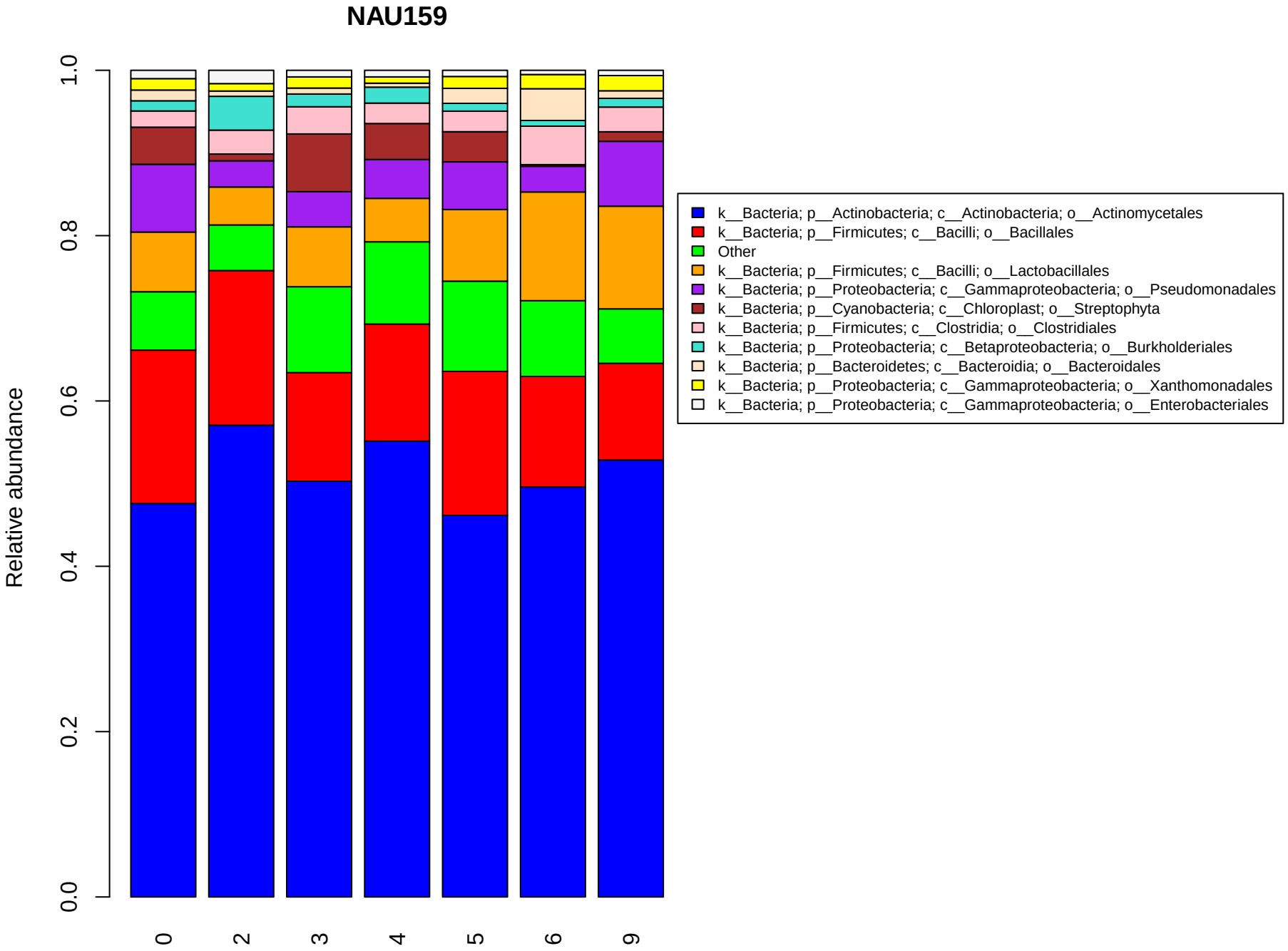

NAU150
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
3
4
5
7
8
9
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales

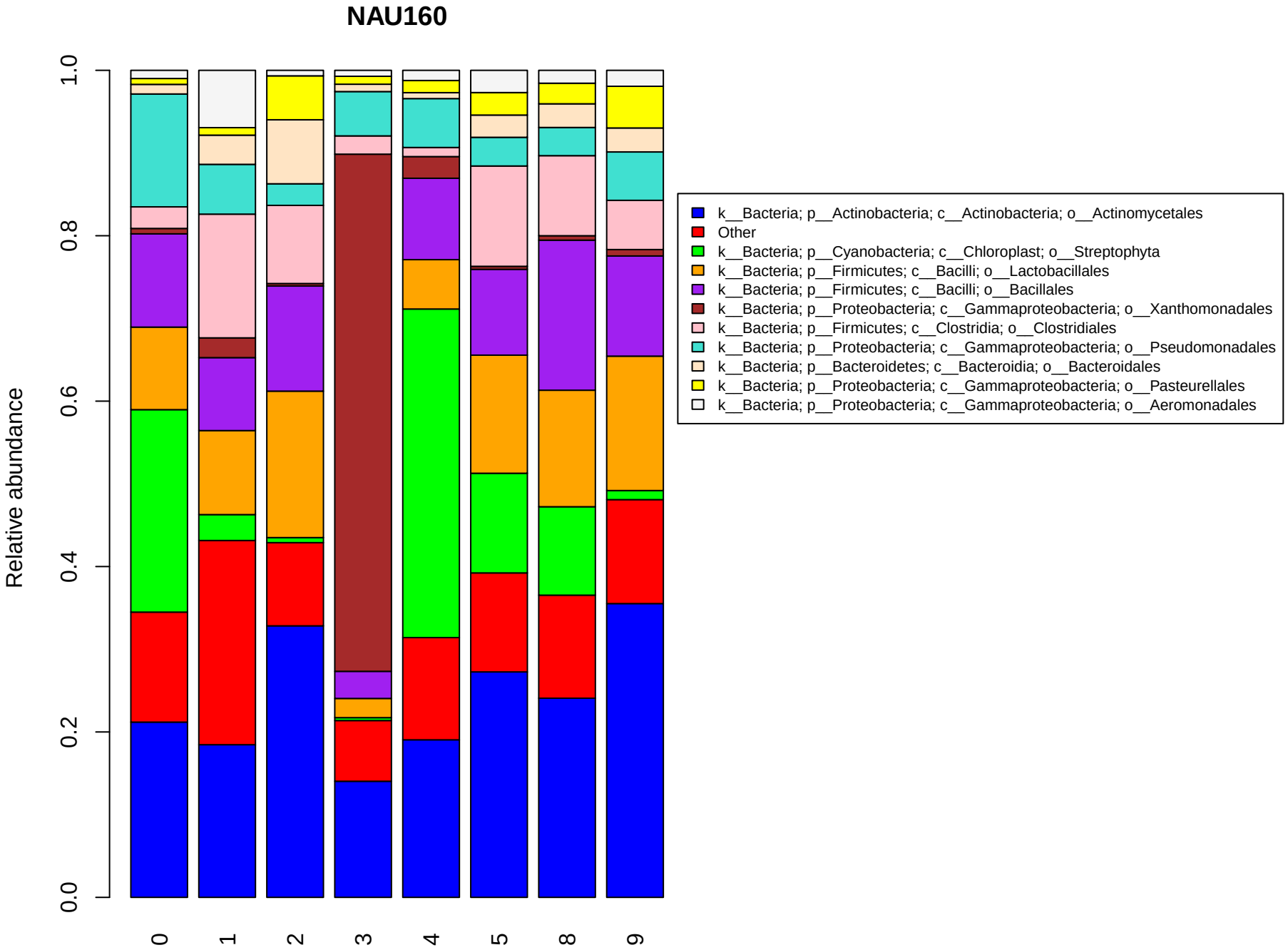

NAU153
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales

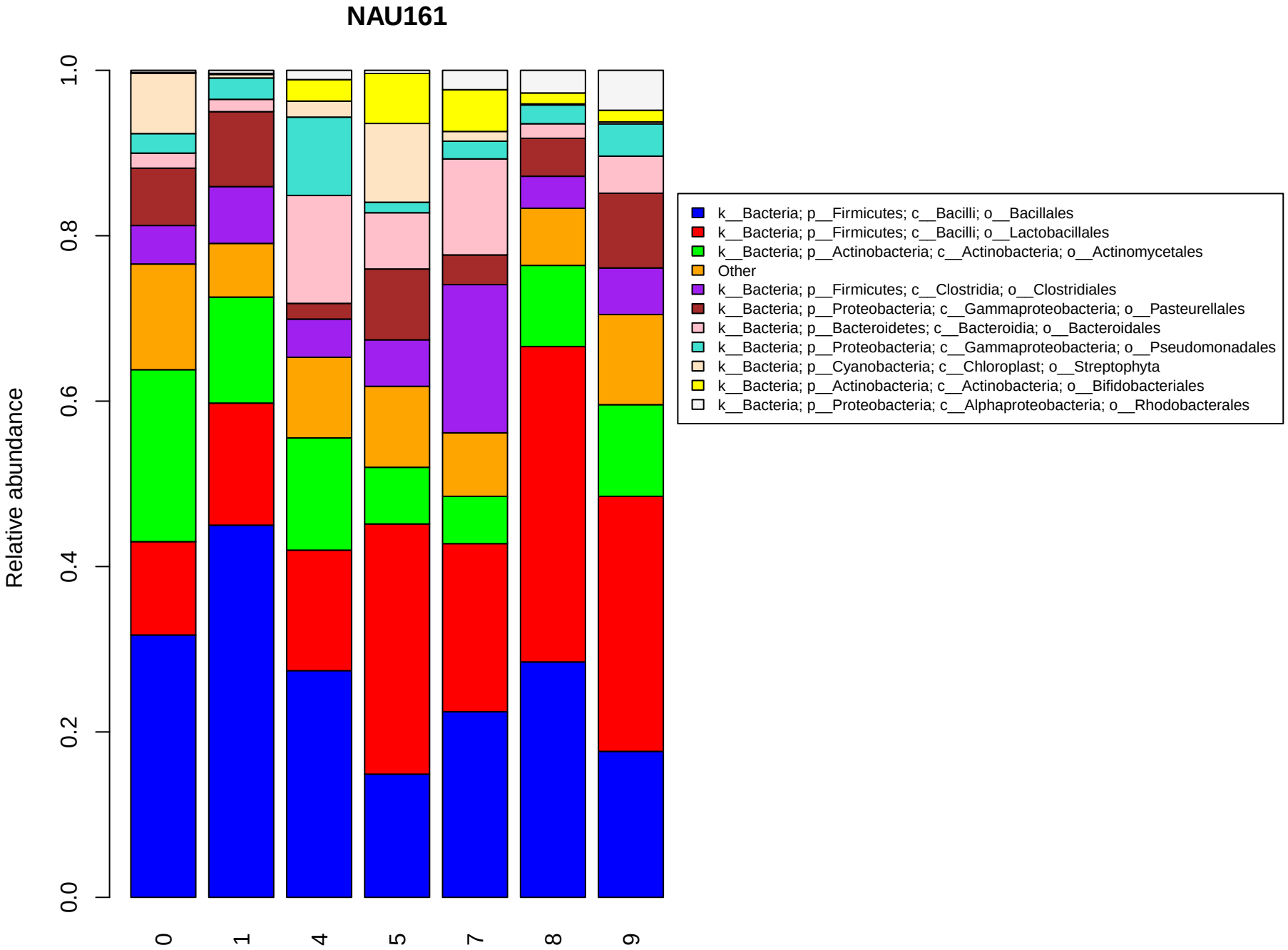

NAU154
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales

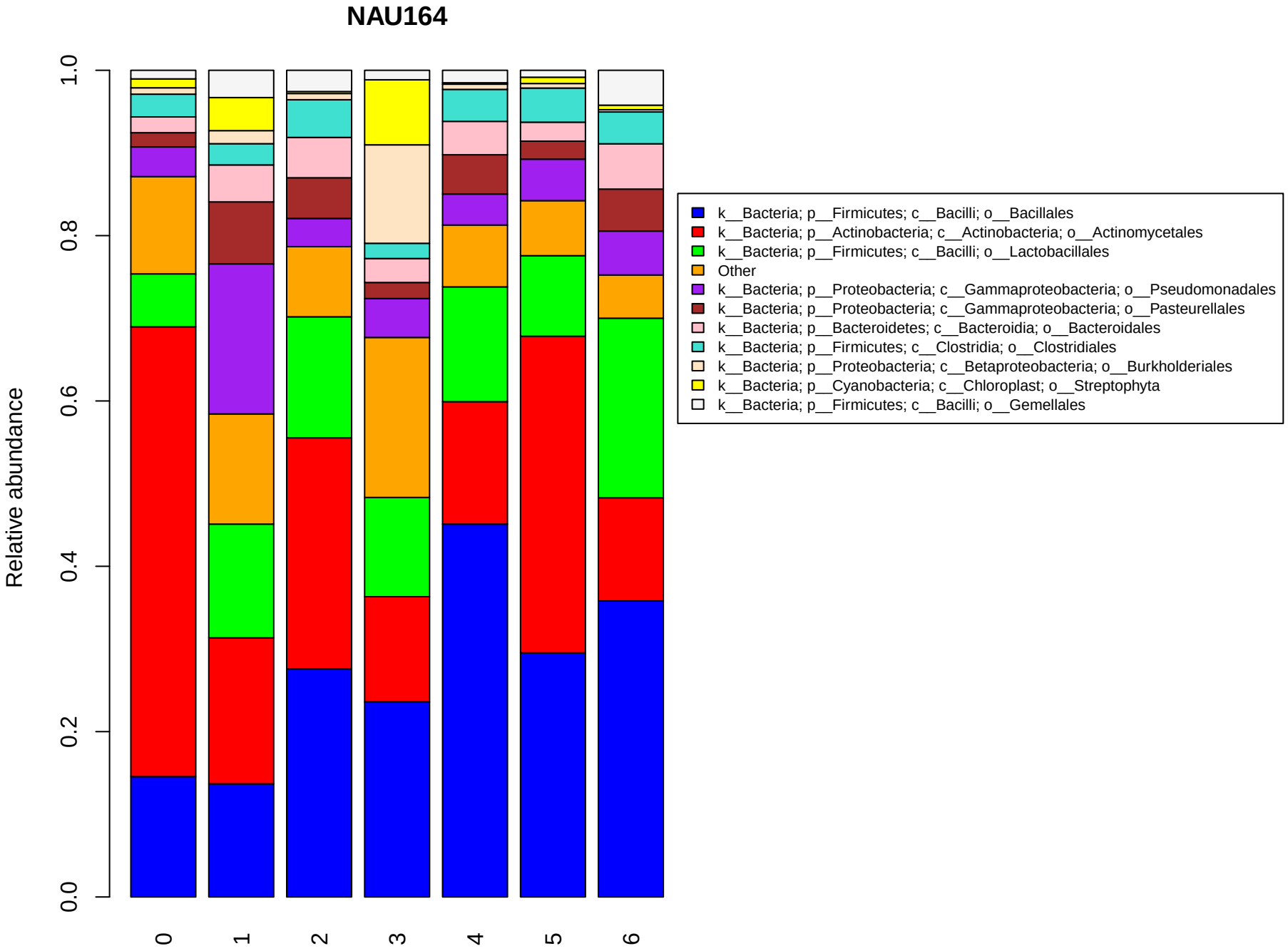

NAU155
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales

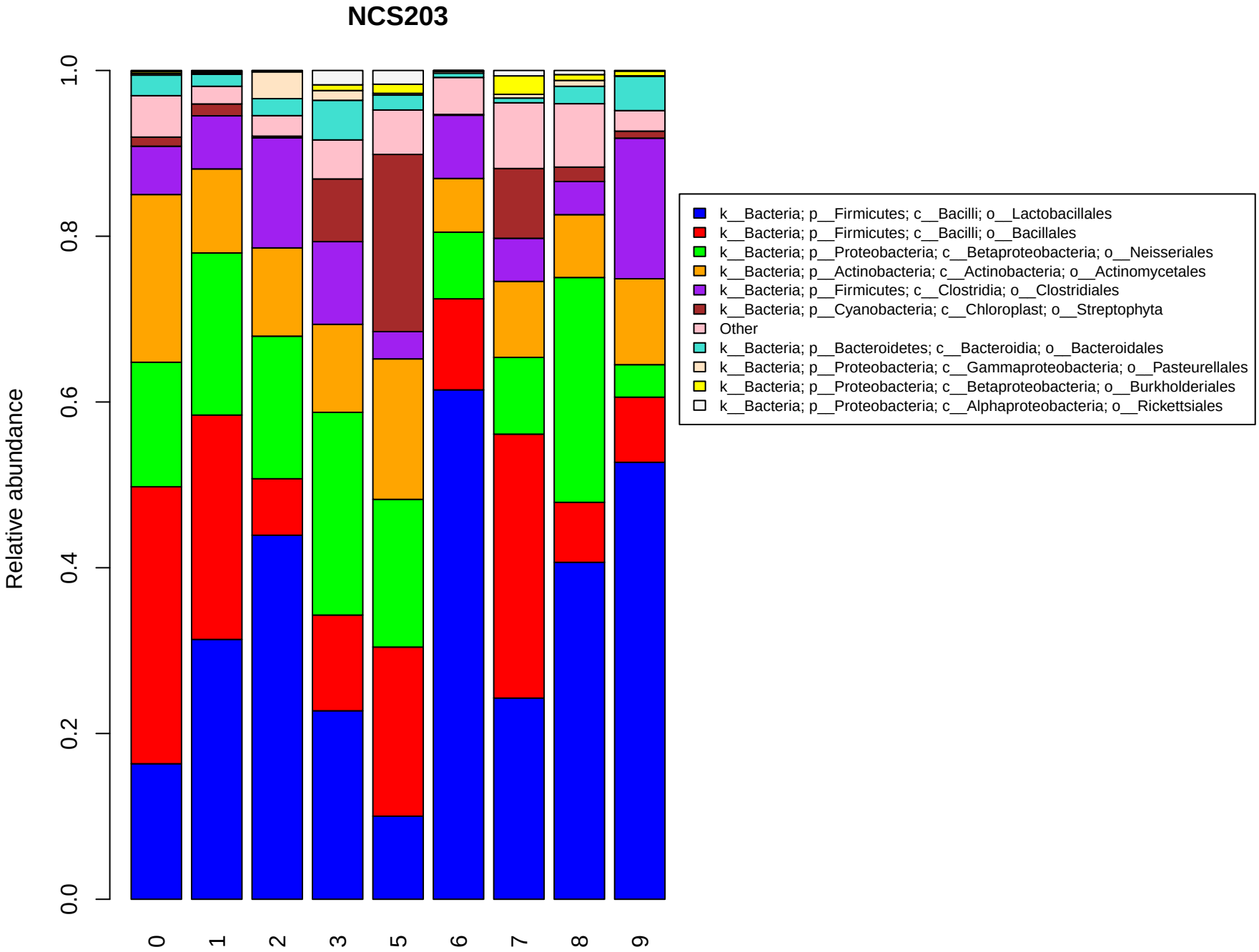

NAU157
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
2
3
4
5
6
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales

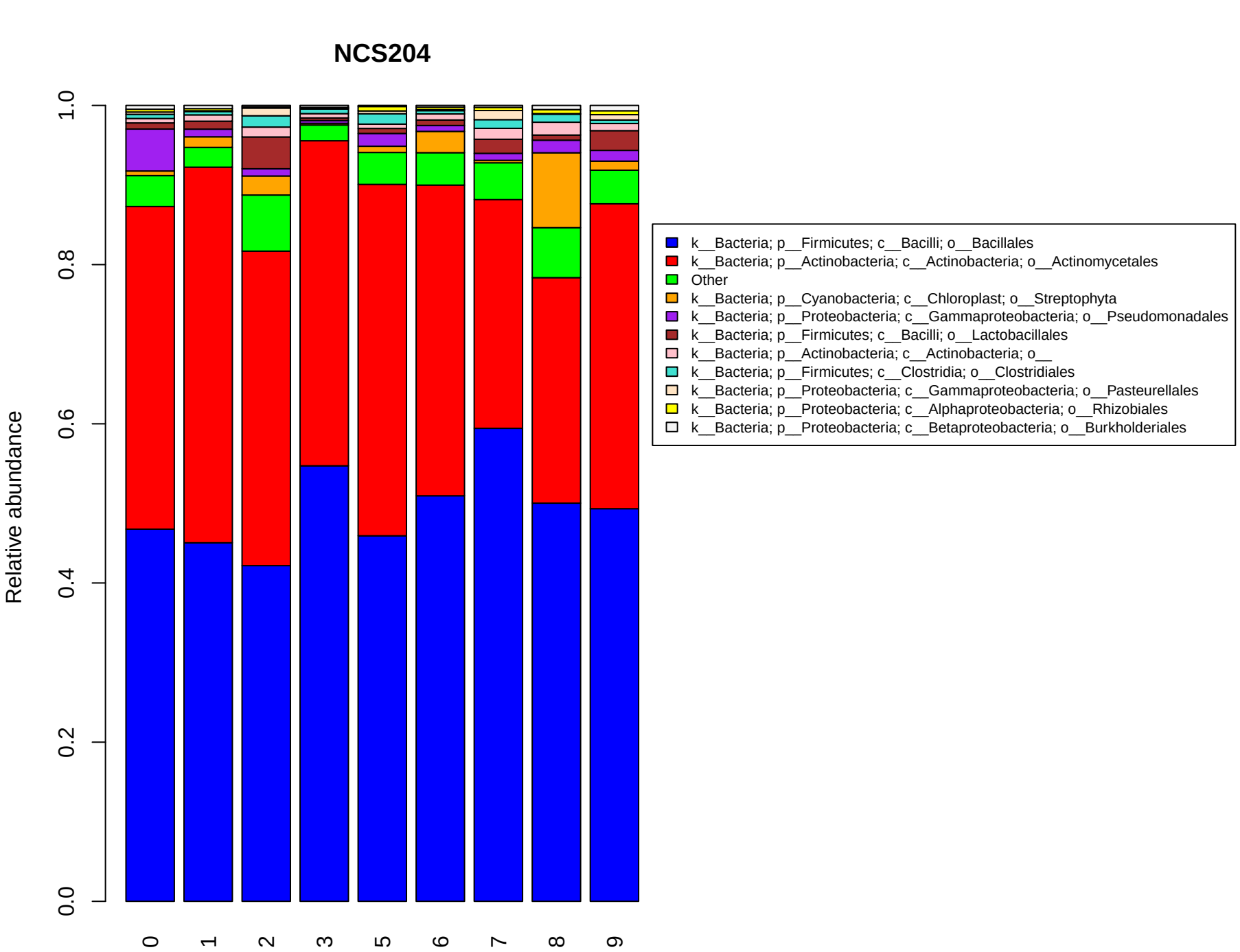

NAU159
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
2
3
4
5
6
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacteriales

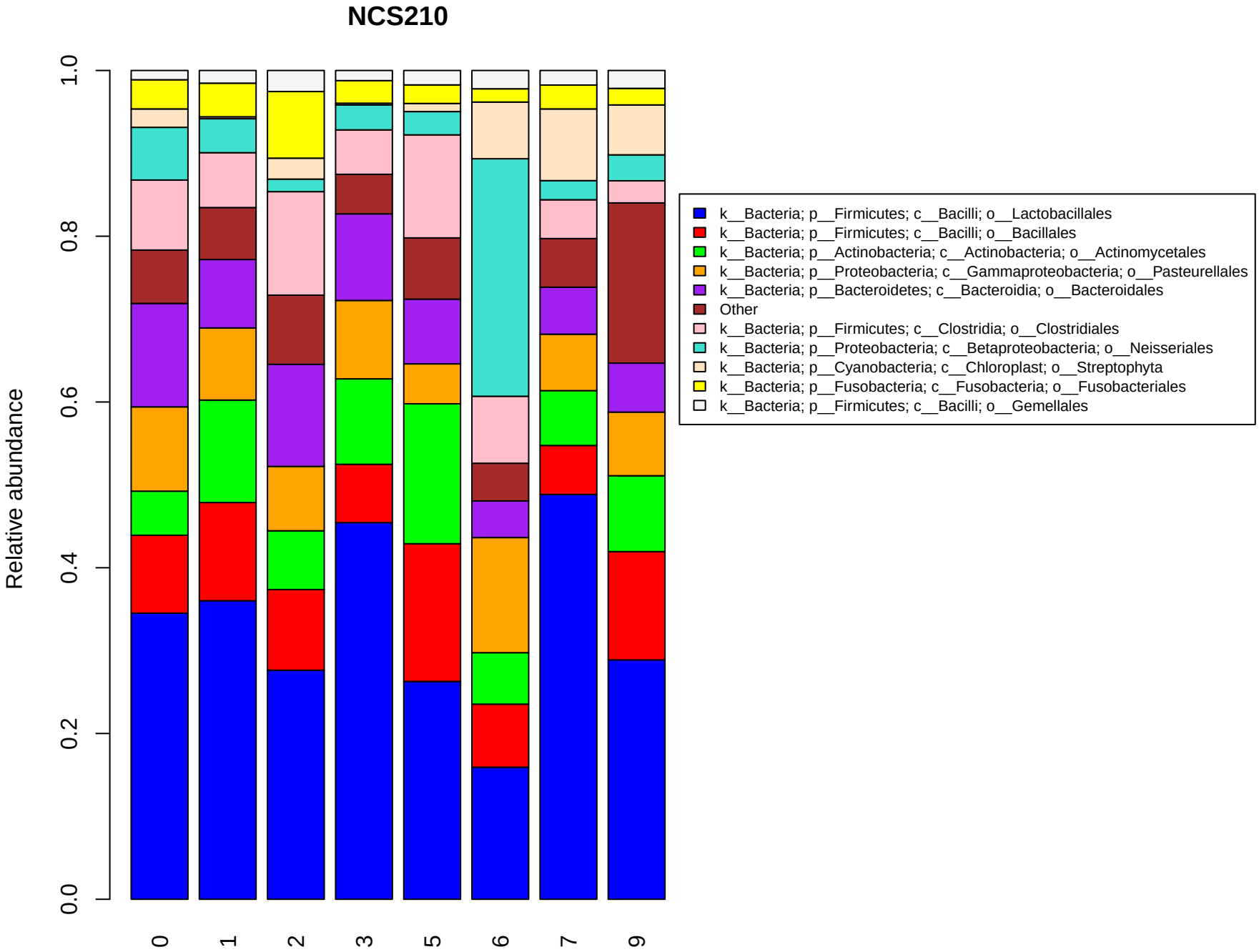

NAU160
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Aeromonadales

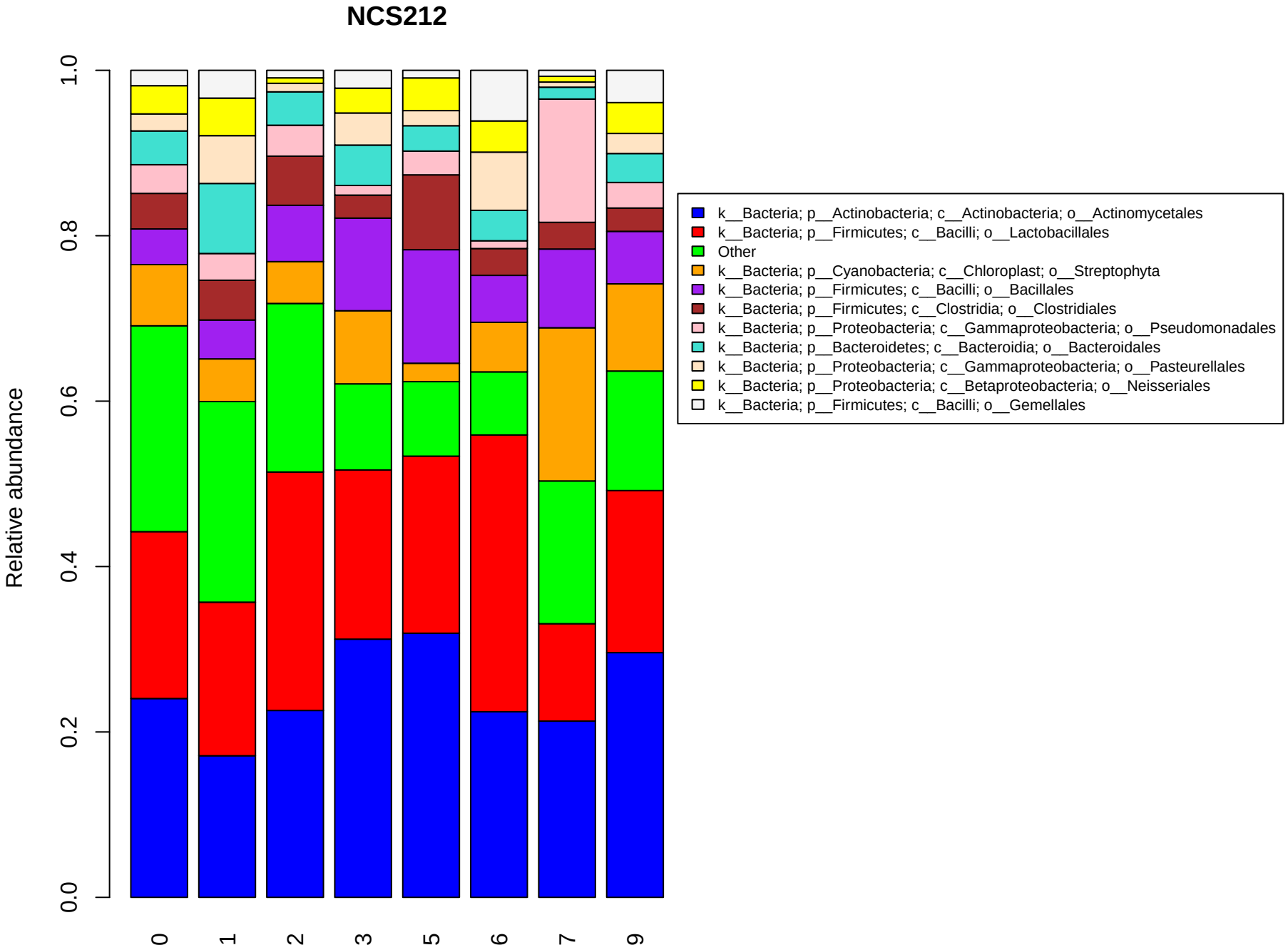

NAU161
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
4
5
7
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Bifidobacteriales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodobacterales

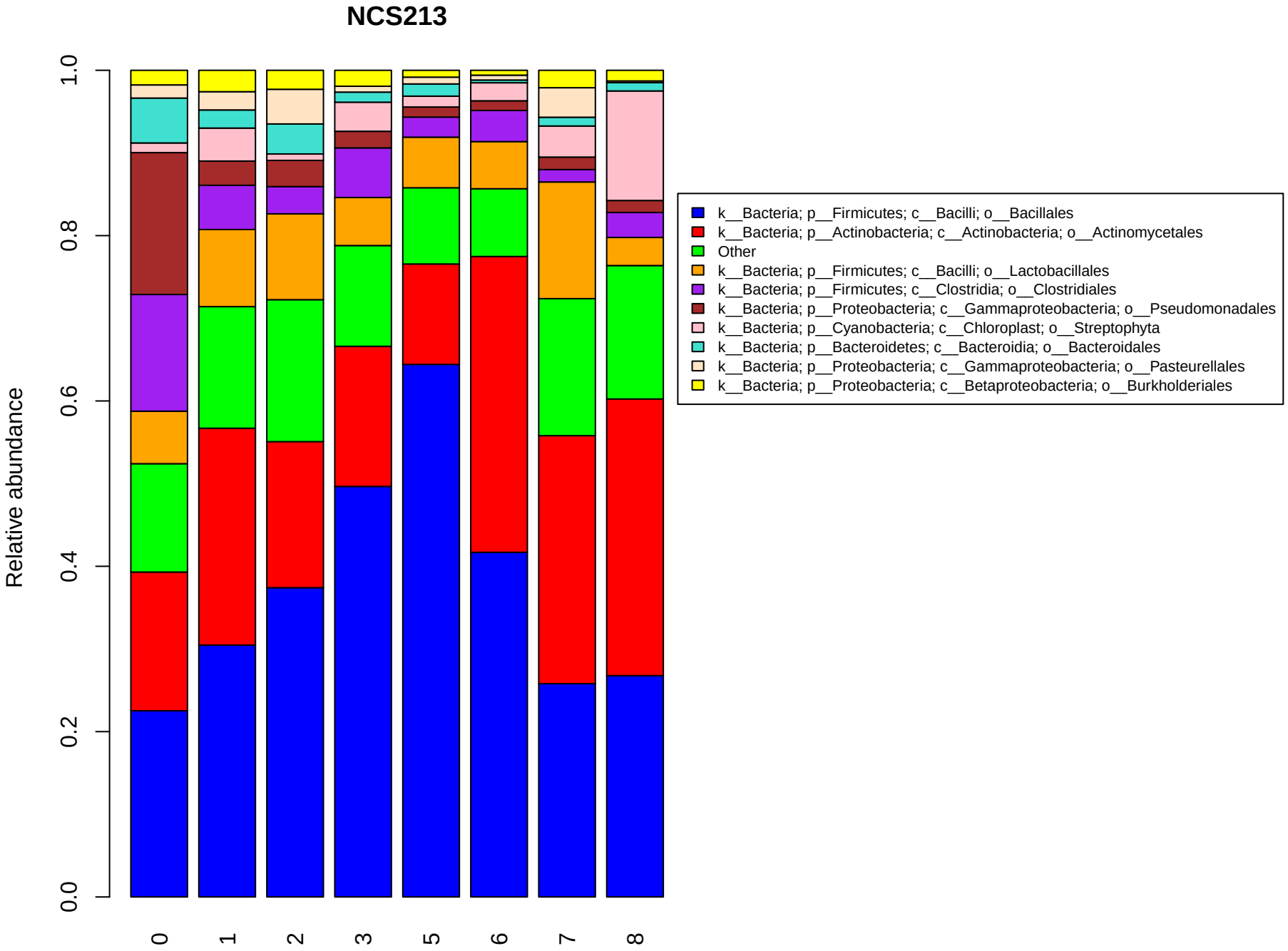

NAU164
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4
5
6
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales

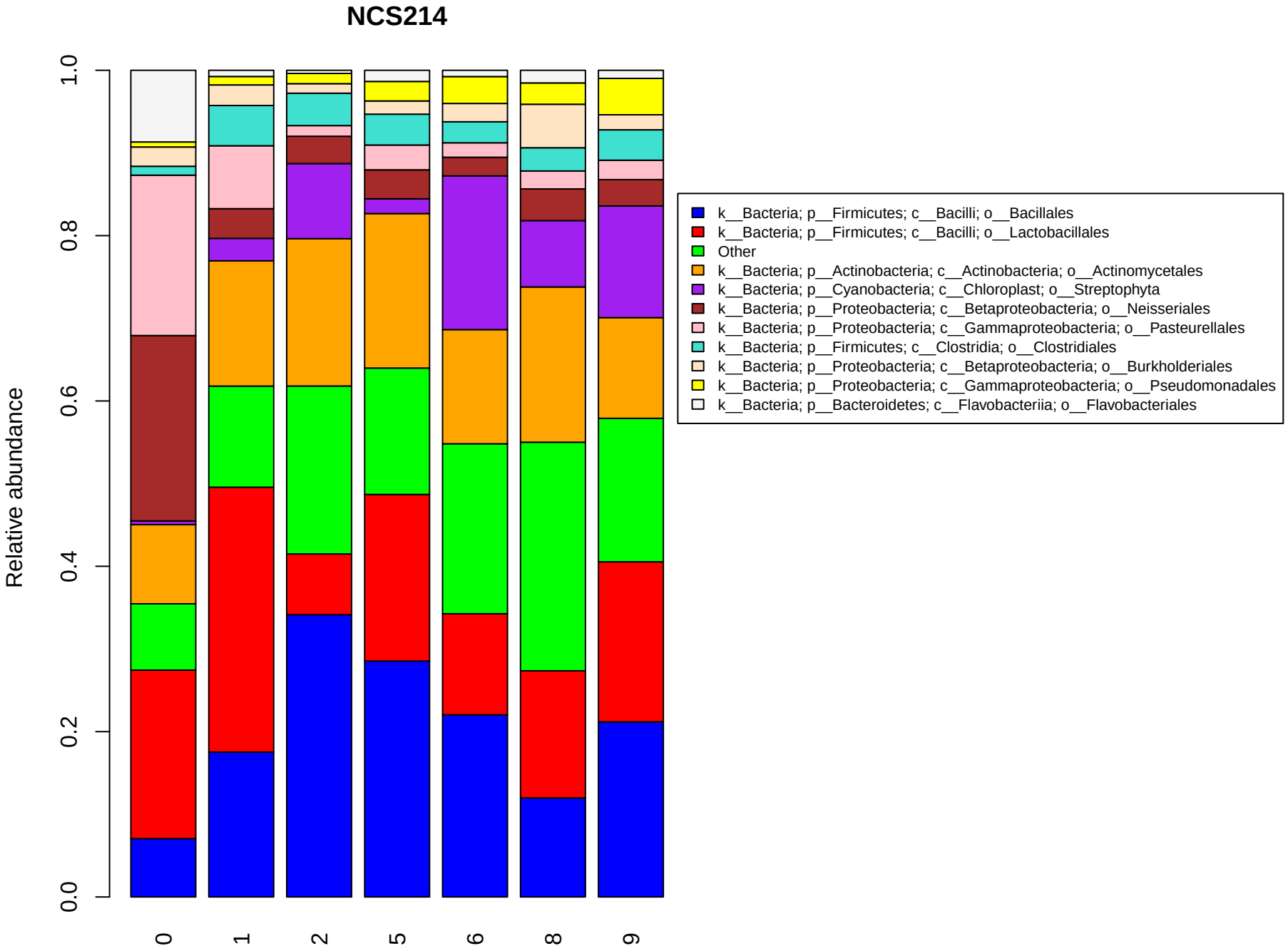

NCS203
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
Other
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rickettsiales

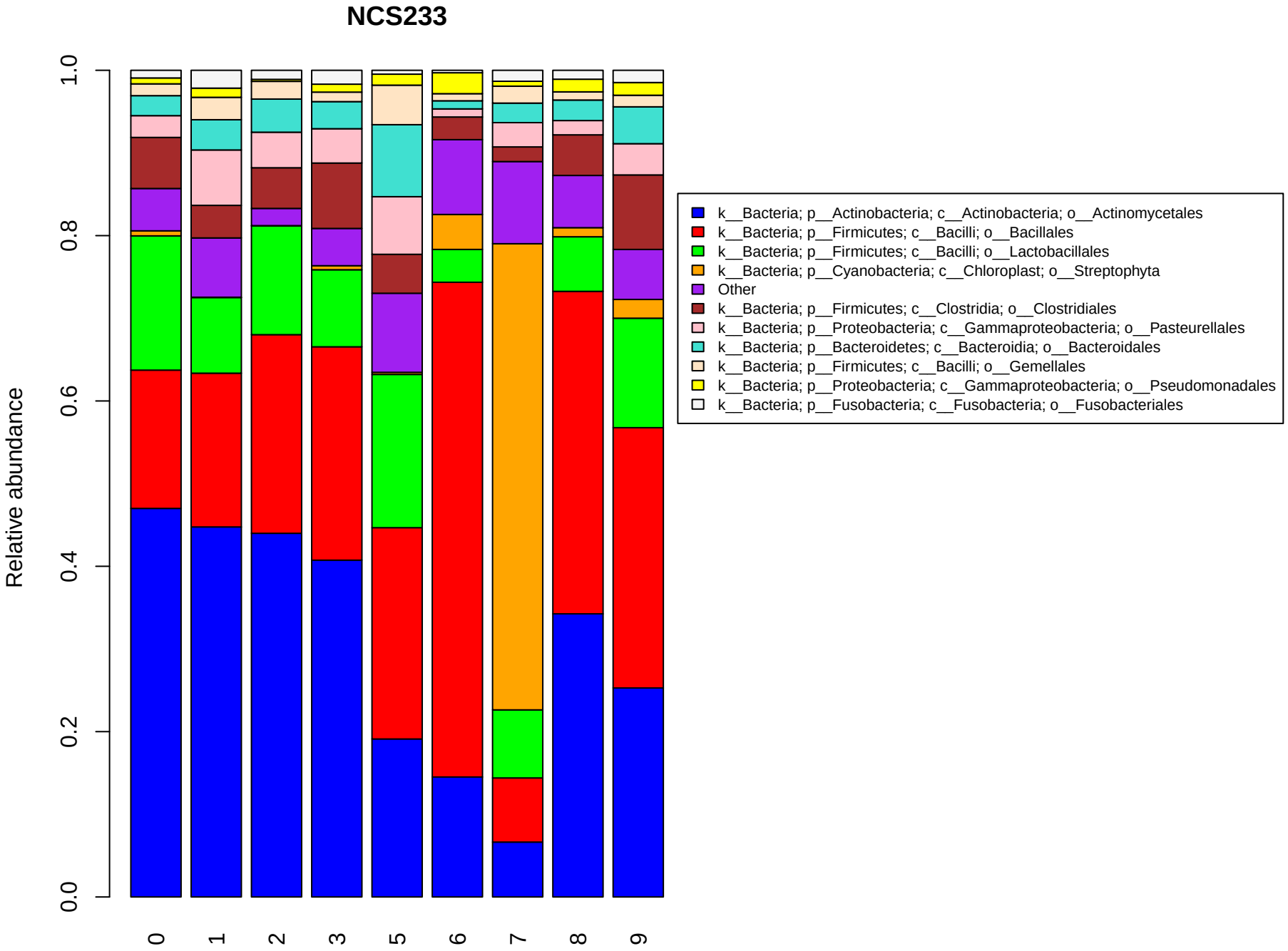

NCS204
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhizobiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales

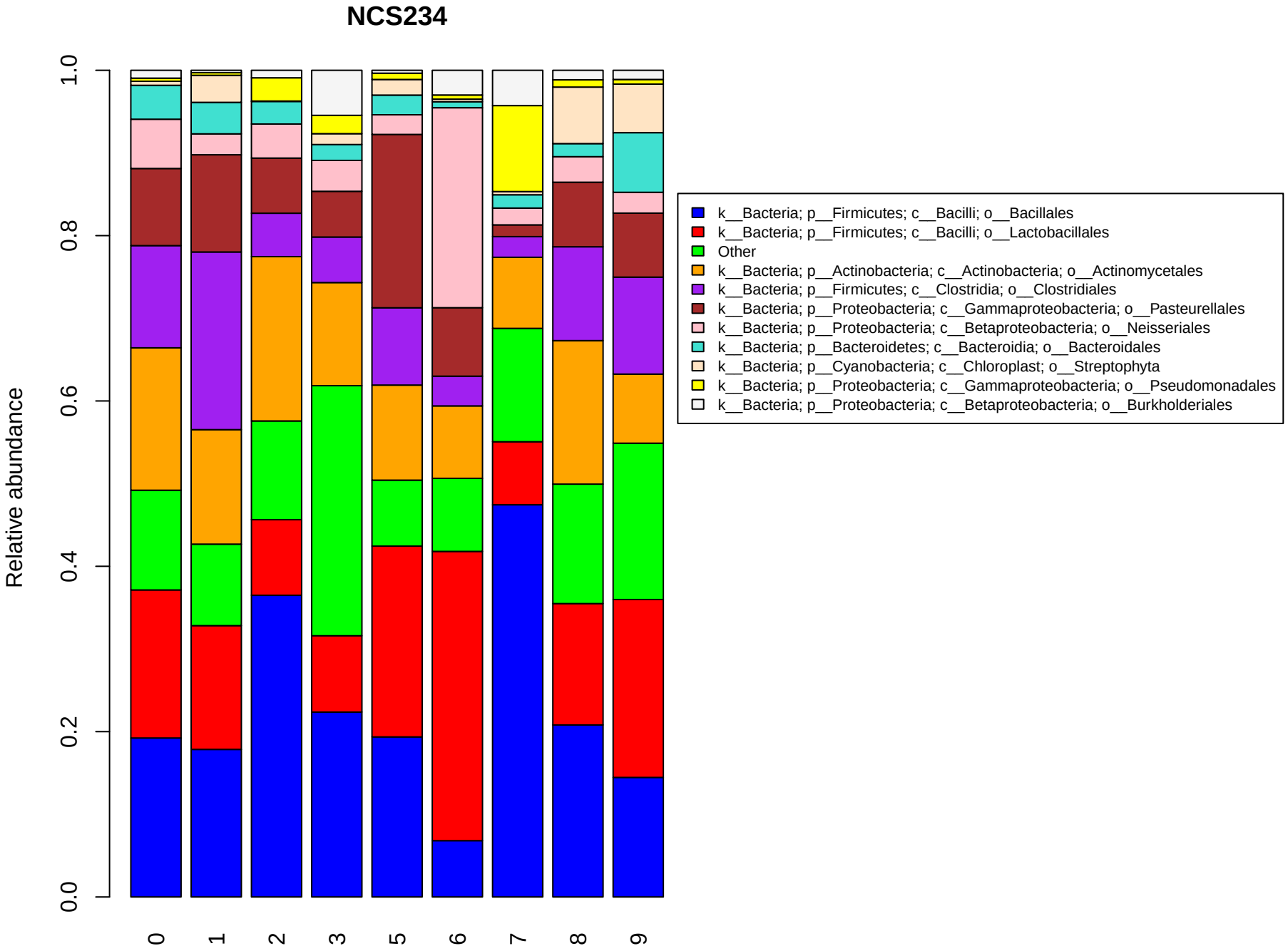

NCS210
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales

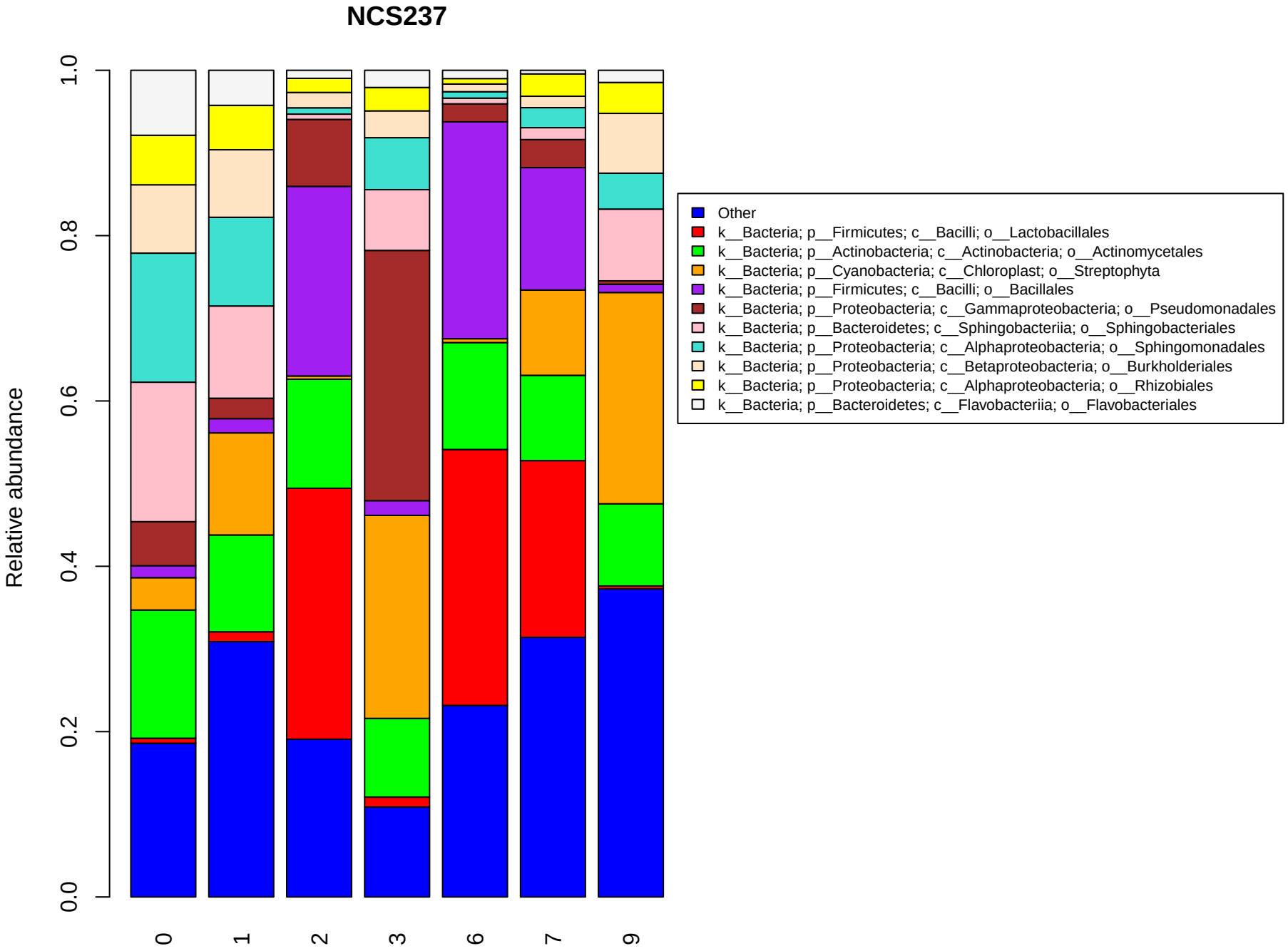

NCS212
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales

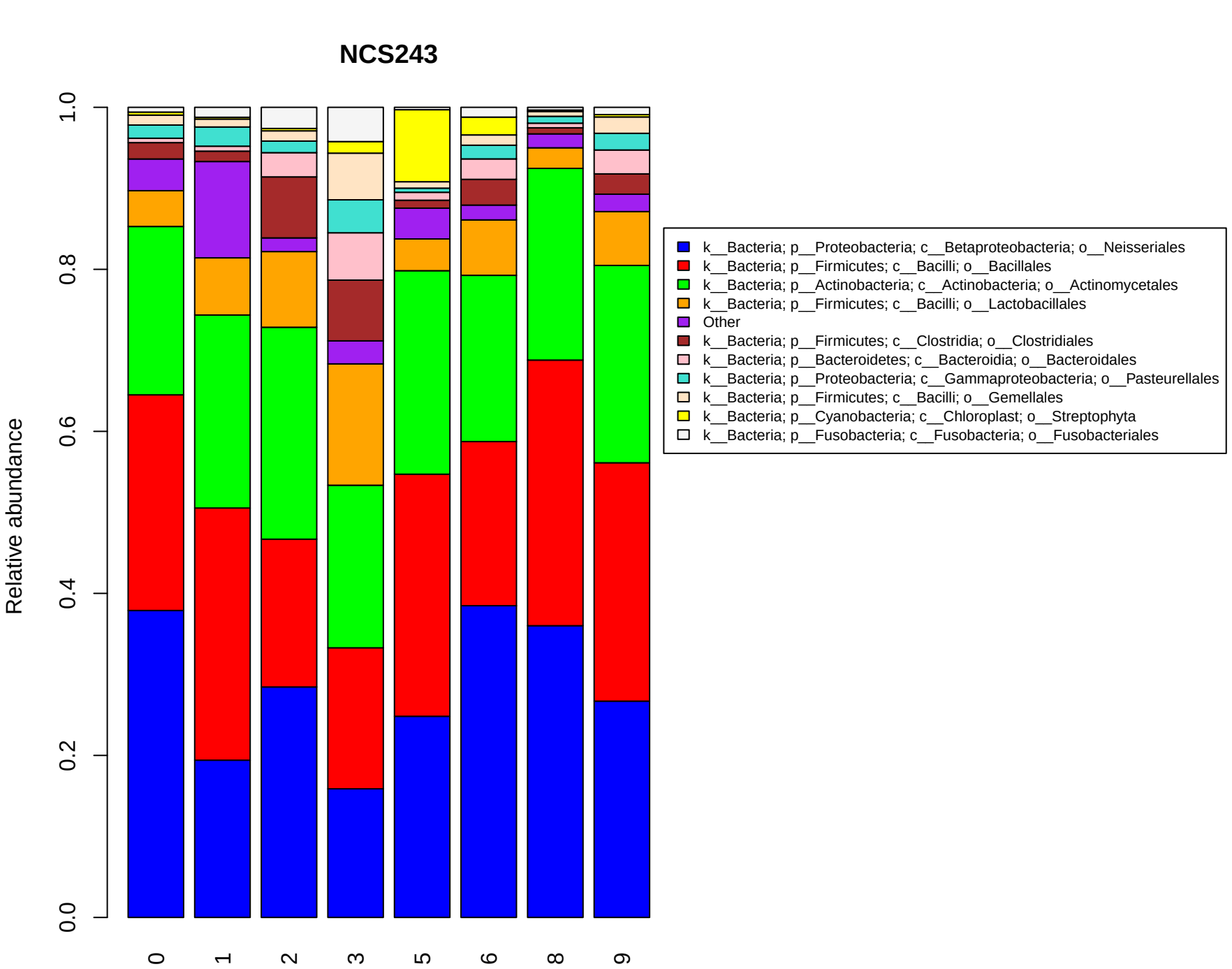

NCS213
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
8
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales

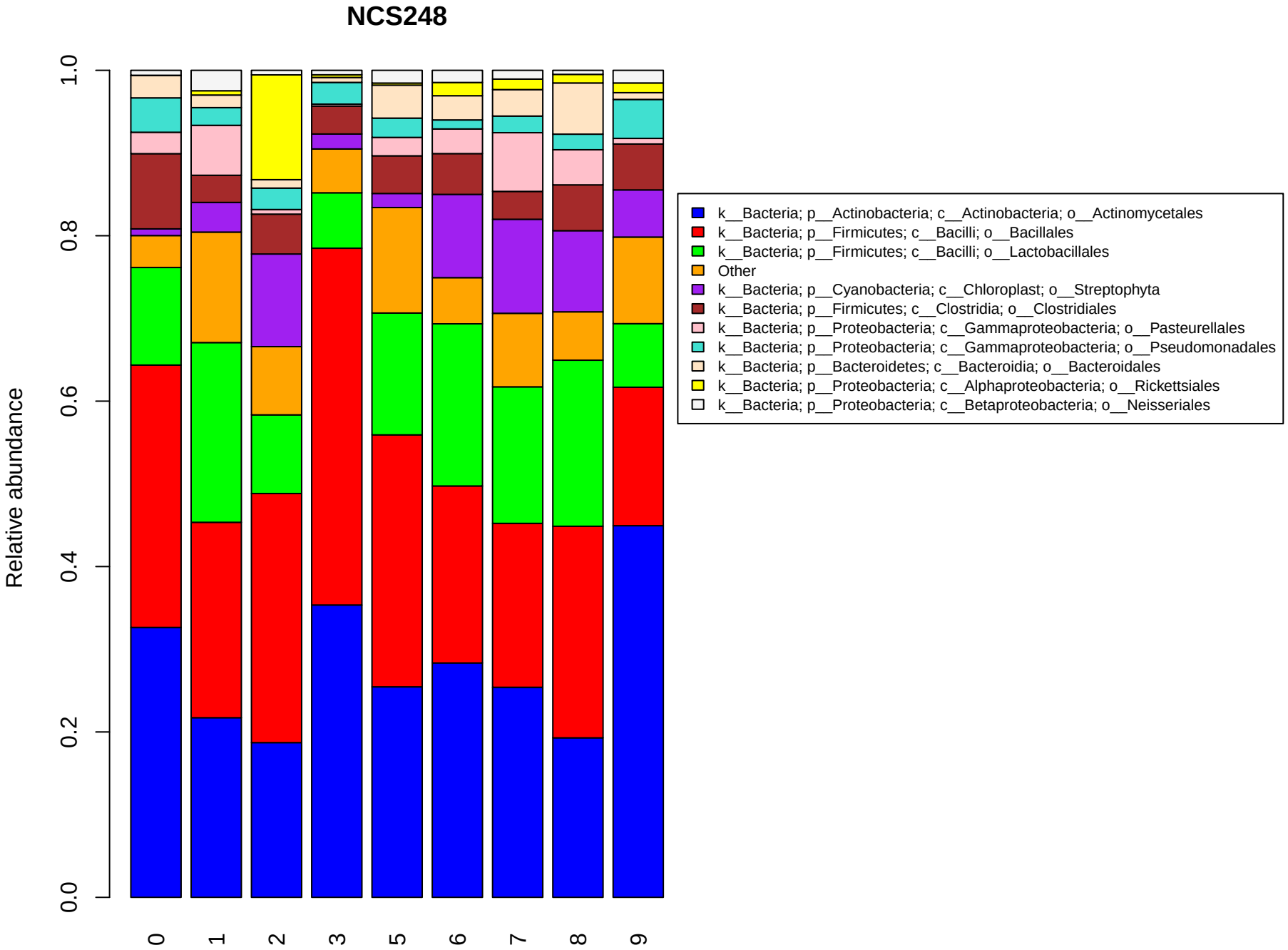

NCS214
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
5
6
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales

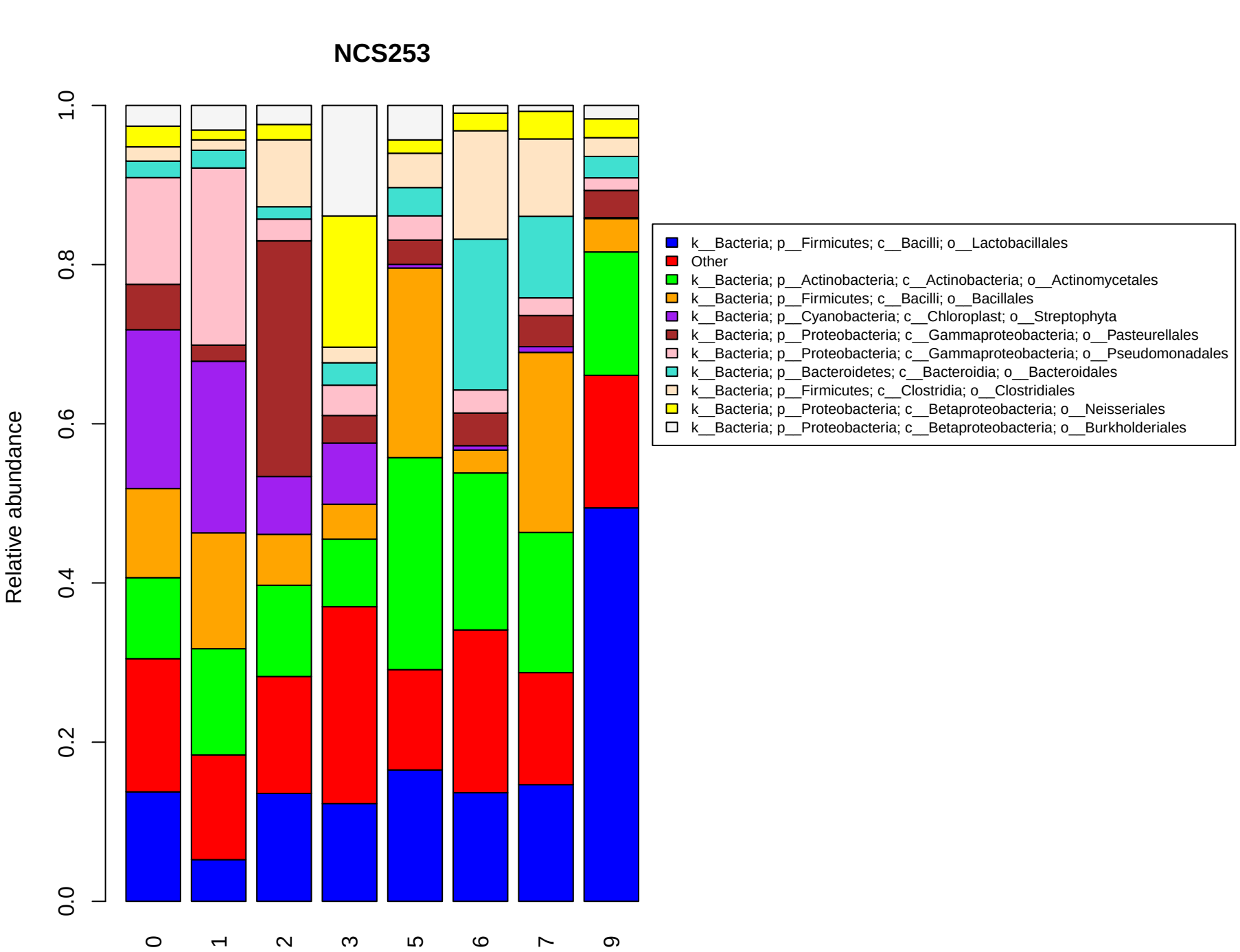

NCS233
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales

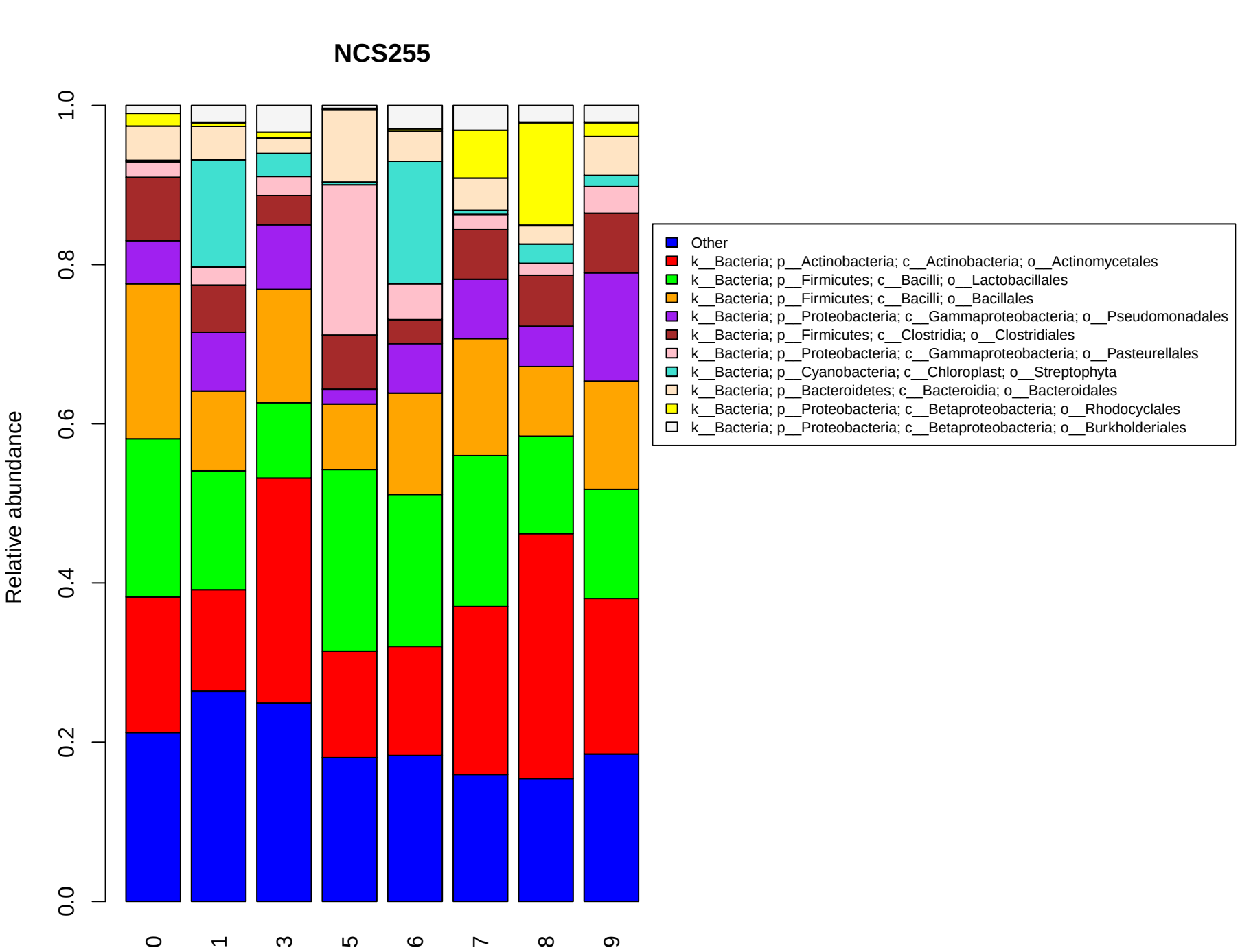

NCS234
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
8
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales

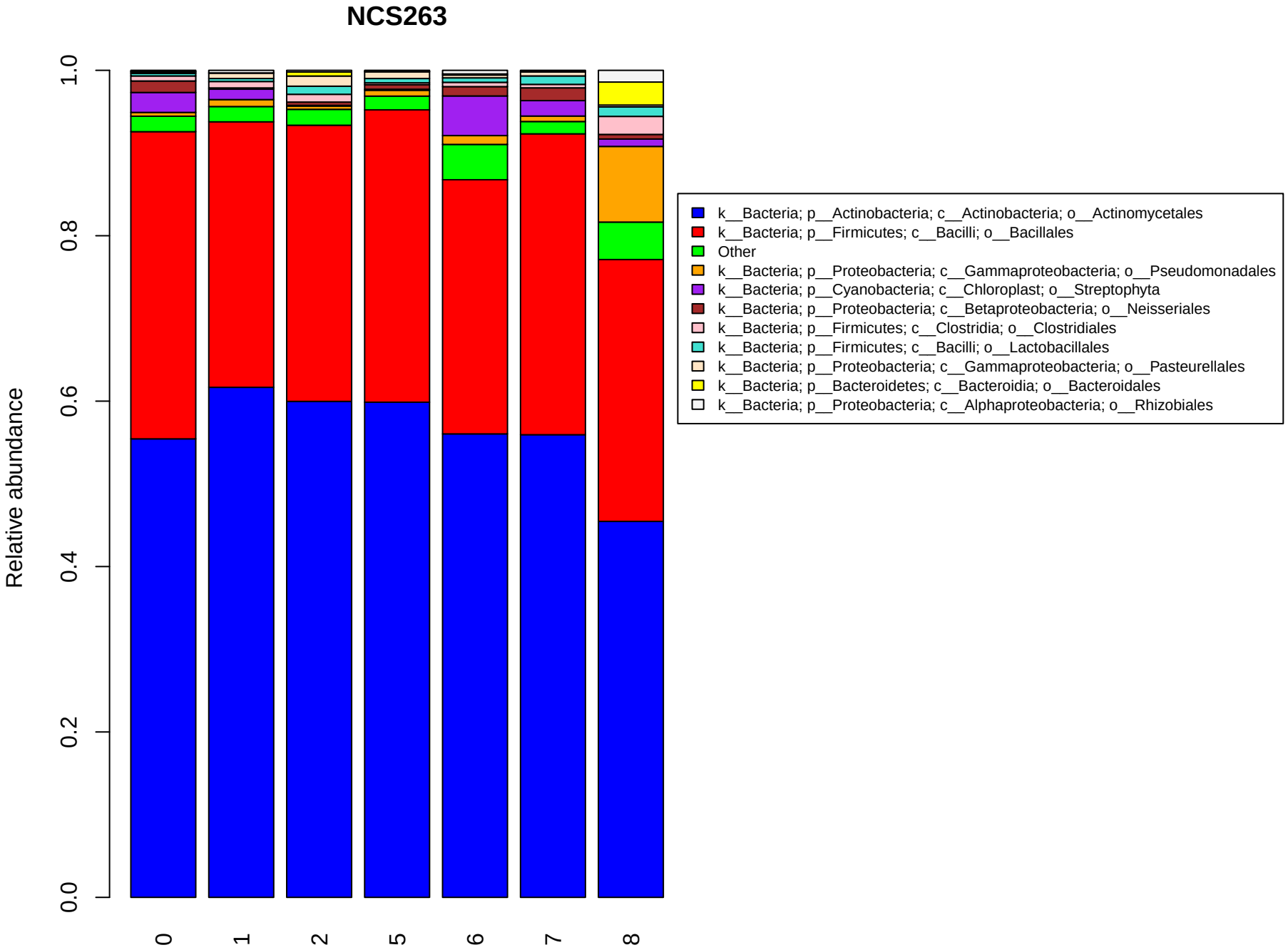

NCS237
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
6
7
9
Other
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Sphingobacteriia; o__Sphingobacteriales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Sphingomonadales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhizobiales
k__Bacteria; p__Bacteroidetes; c__Flavobacteriia; o__Flavobacteriales


NCS243
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
8
9
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Gemellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Fusobacteria; c__Fusobacteria; o__Fusobacteriales


NCS248
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
8
9
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rickettsiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales


NCS253
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
5
6
7
9
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


NCS255
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
3
5
6
7
8
9
Other
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Rhodocyclales
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Burkholderiales


NCS263
Relative abundance
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
5
6
7
8
k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales
Other
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales
k__Bacteria; p__Cyanobacteria; c__Chloroplast; o__Streptophyta
k__Bacteria; p__Proteobacteria; c__Betaproteobacteria; o__Neisseriales
k__Bacteria; p__Firmicutes; c__Clostridia; o__Clostridiales
k__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pasteurellales
k__Bacteria; p__Bacteroidetes; c__Bacteroidia; o__Bacteroidales
k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhizobiales