

Summary Table File: LimSC25.taxa.genus.NoControls.N500.summary\_table.tsv  
Factor File: 0137\_Lim\_Metadata\_20171019.tsv  
Output File Root: sickle.genus.tail.t20

Diversity Index: tail

Summary Table:  
Num Samples: 49  
Num Categories: 199

Factor Table:  
Num Samples: 58  
Num Factors: 27

Shared Samples: 49

Number of Top Categories from each sample to summarize: 20

Samples exclusive to Summary Table:  
character(0)

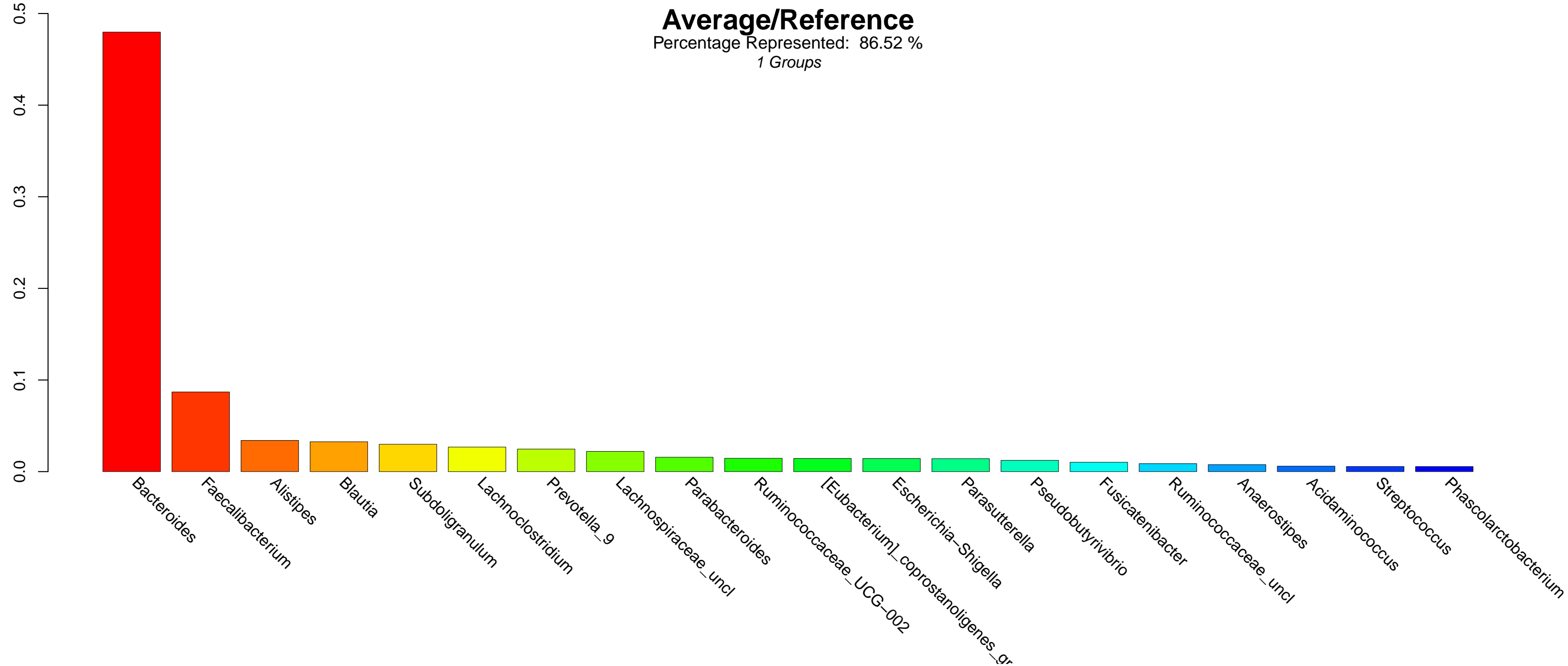
Samples exclusive to Factor Table:  
[1] "0137.BC000299.01.ST" "0137.10022176929.01.ST" "0000.0385.20170823"  
[4] "0000.0386.20170824" "0000.0387.20170825" "0000.0388.20170828"  
[7] "0000.0389.Ext.N" "0001.0390.20170913" "0002.0391.20170913"

# Mean abundances across all samples

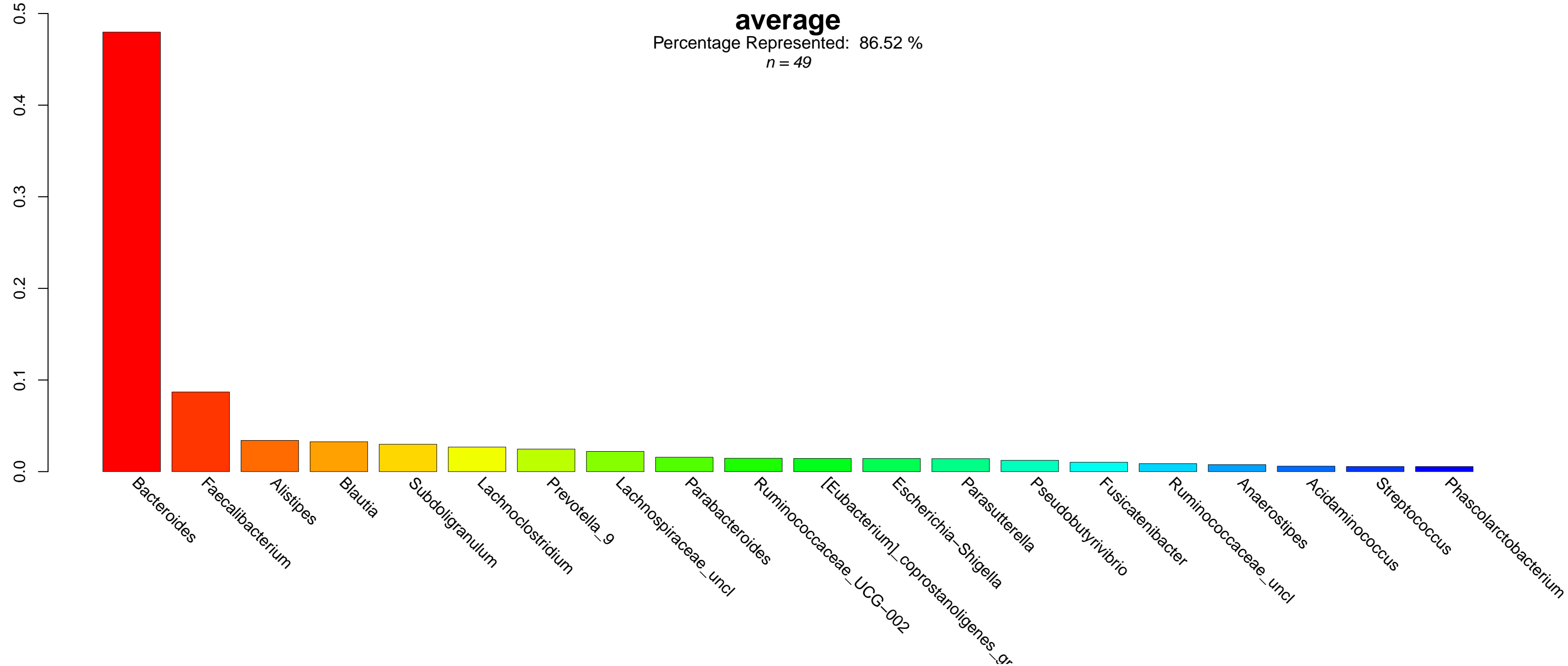
**Average/Reference**

Percentage Represented: 86.52 %

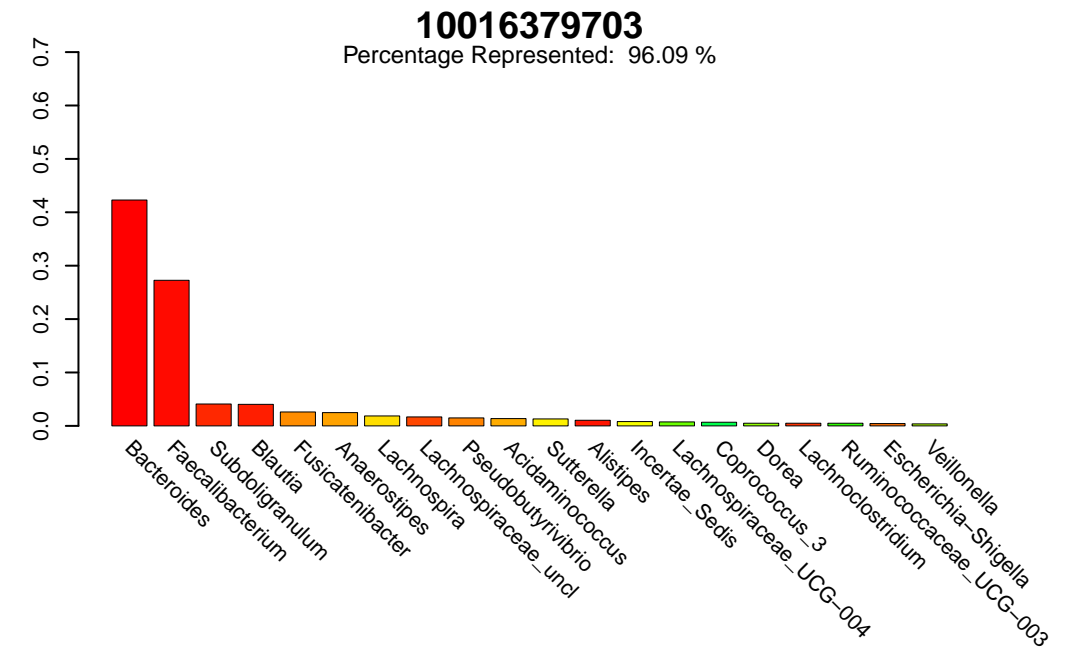
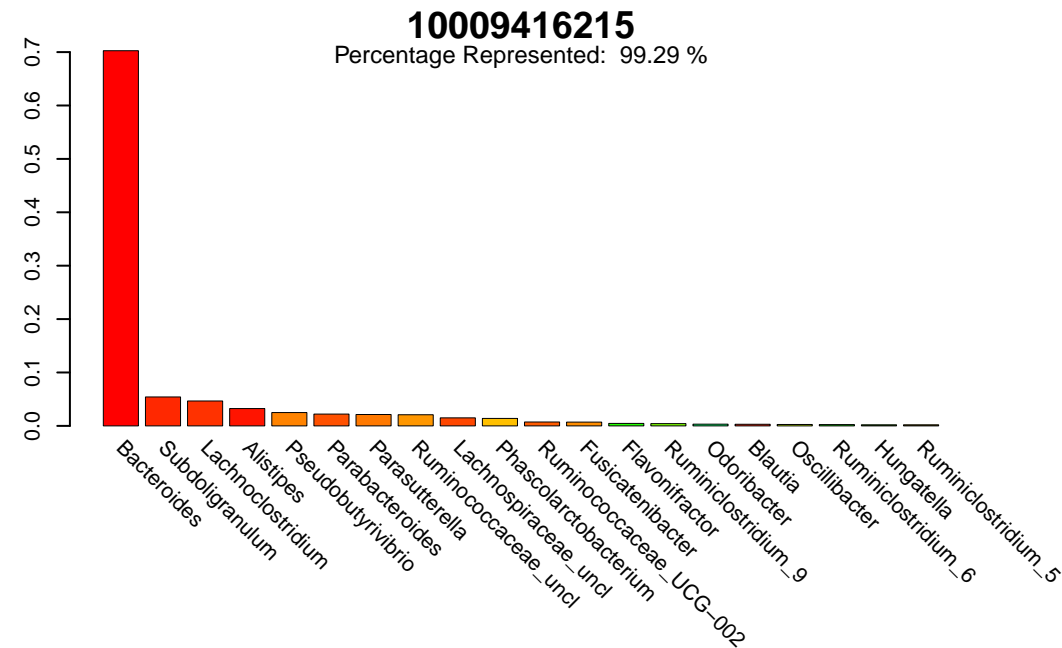
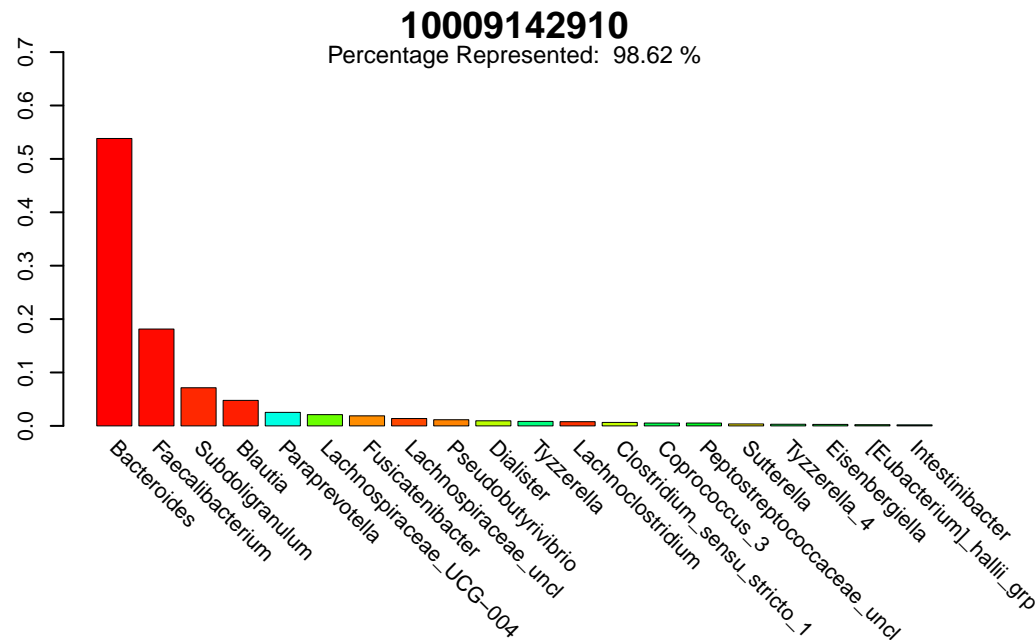
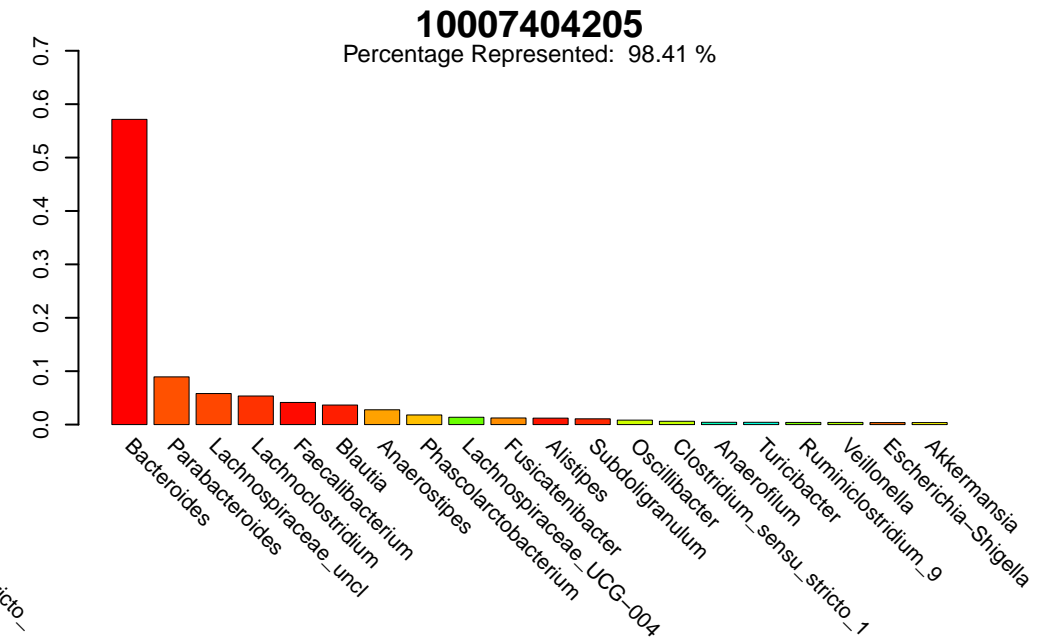
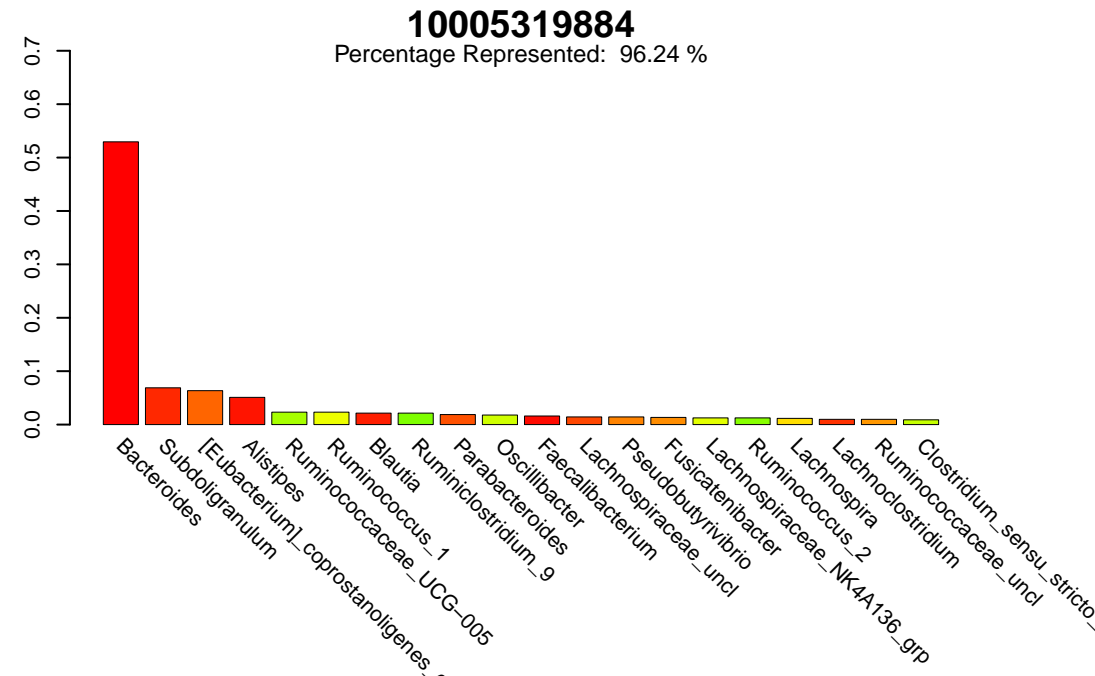
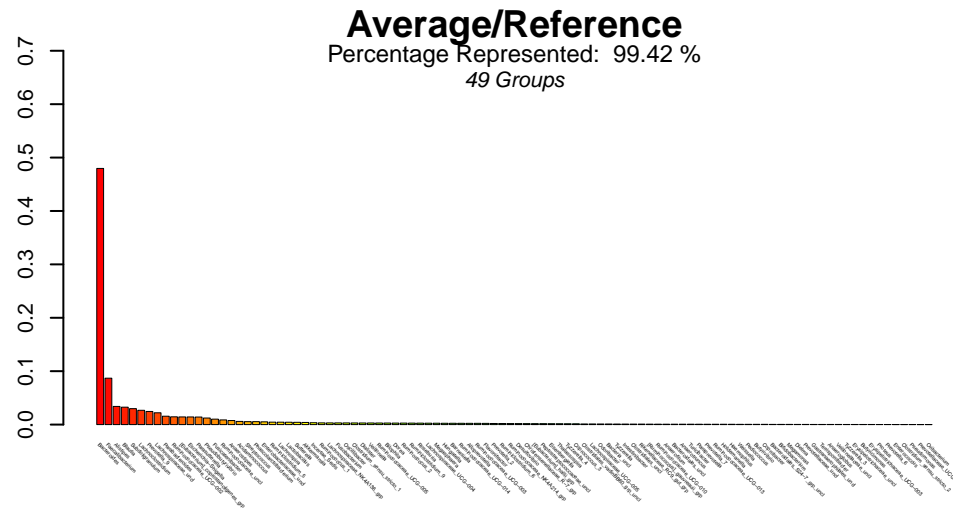
1 Groups



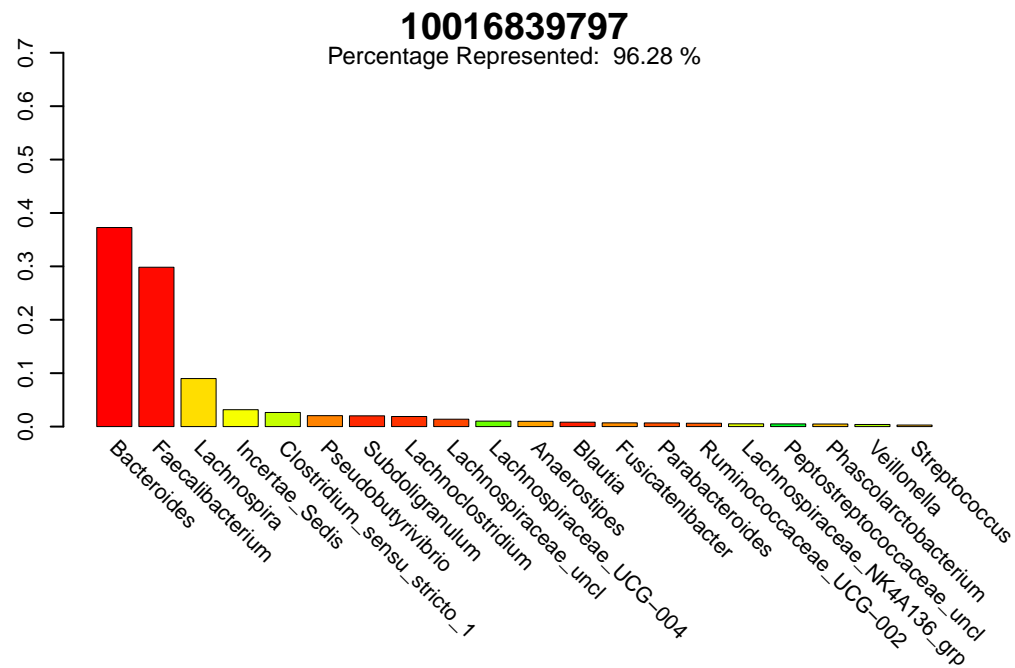
# Mean abundances across all samples



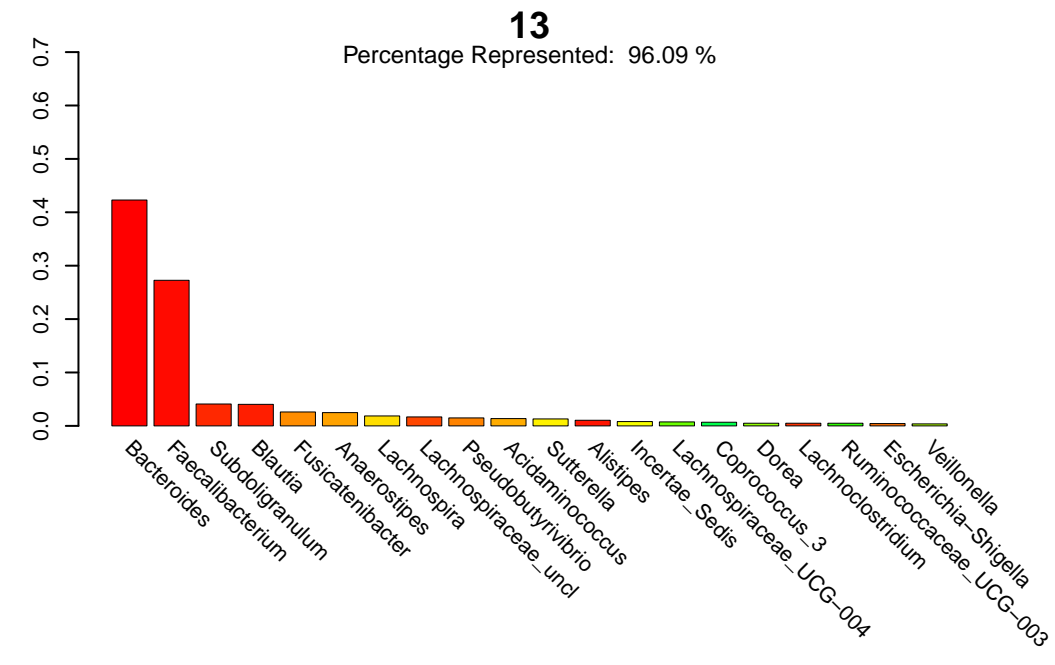
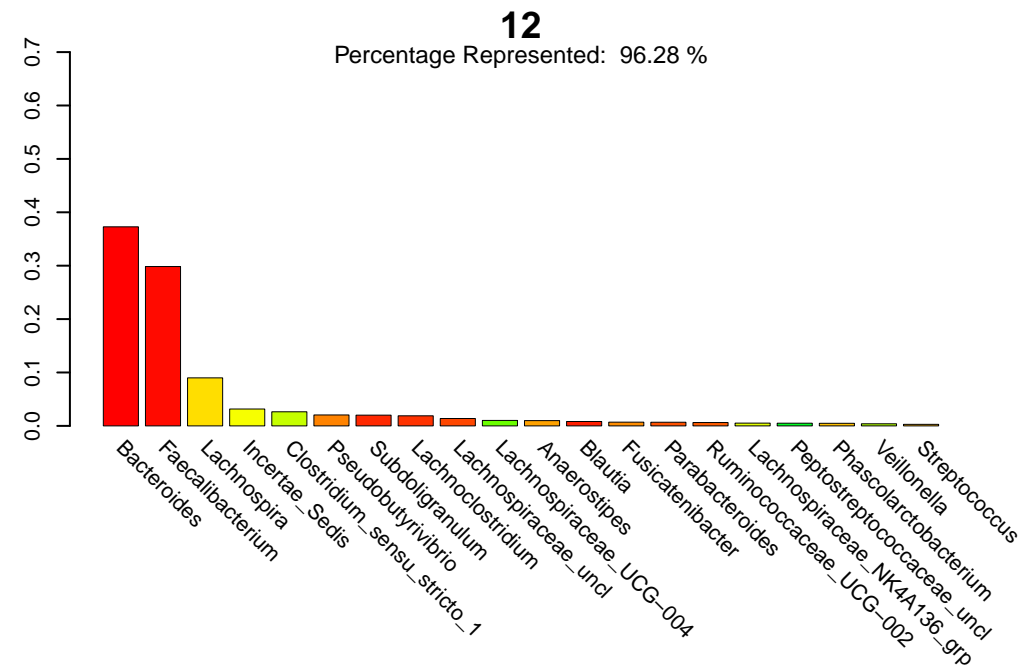
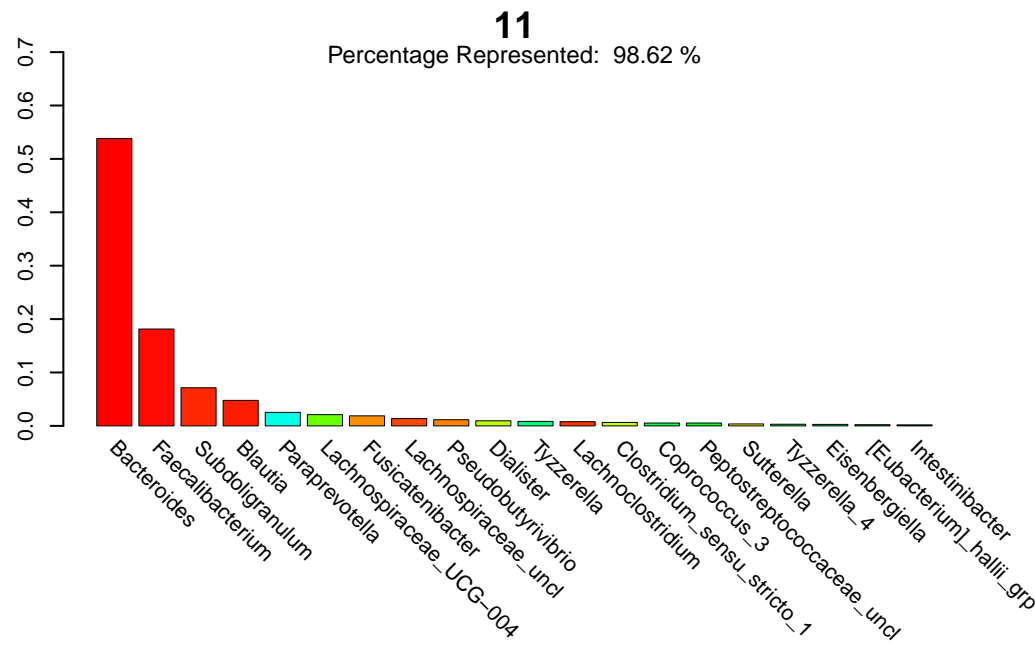
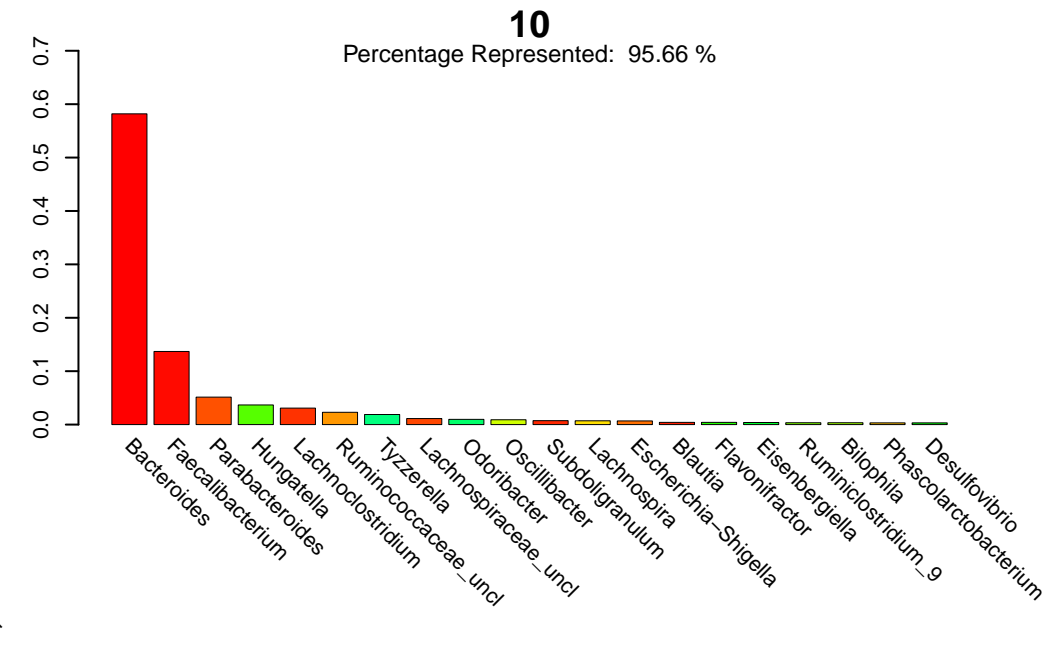
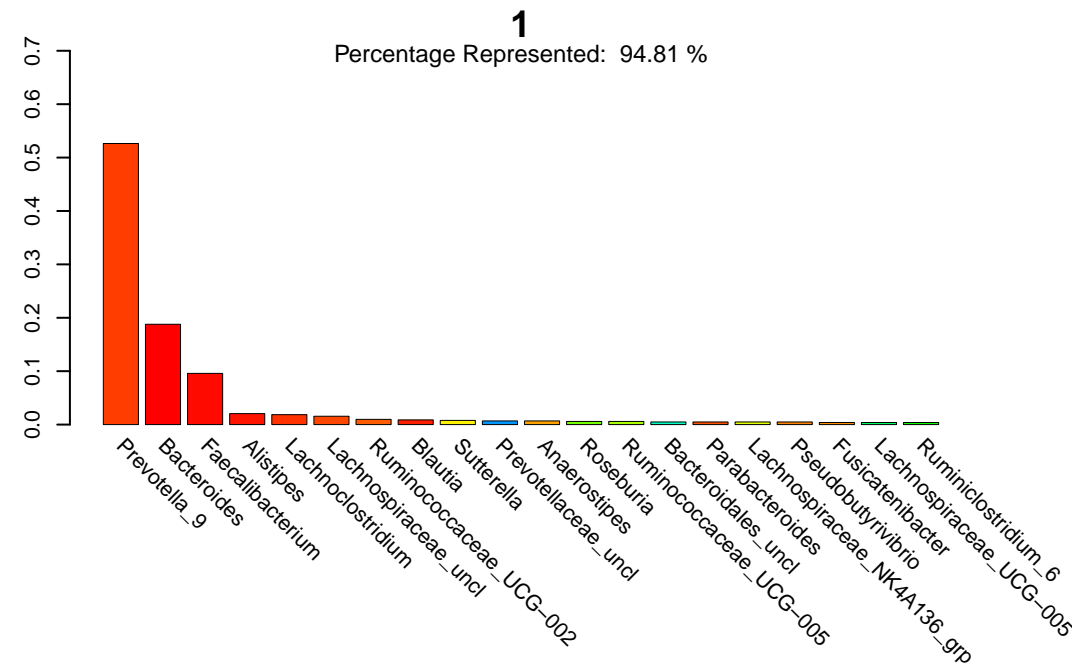
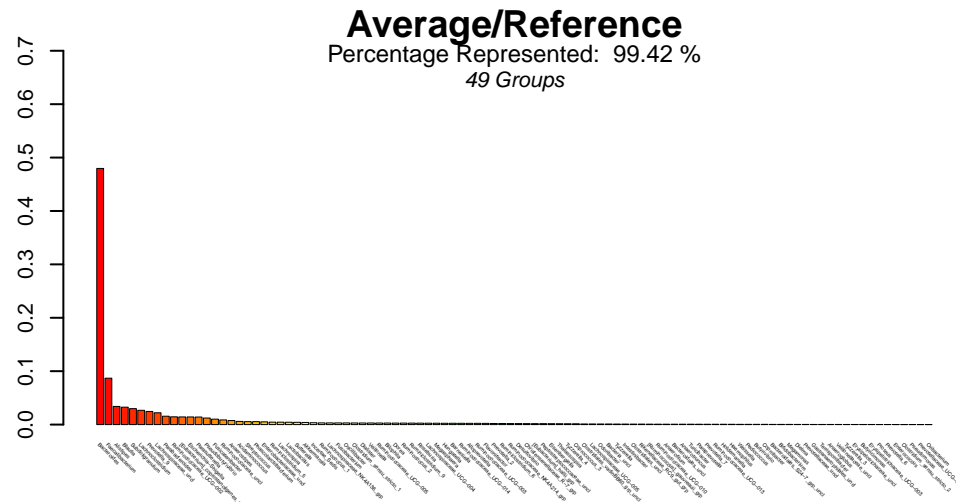
# Identifier



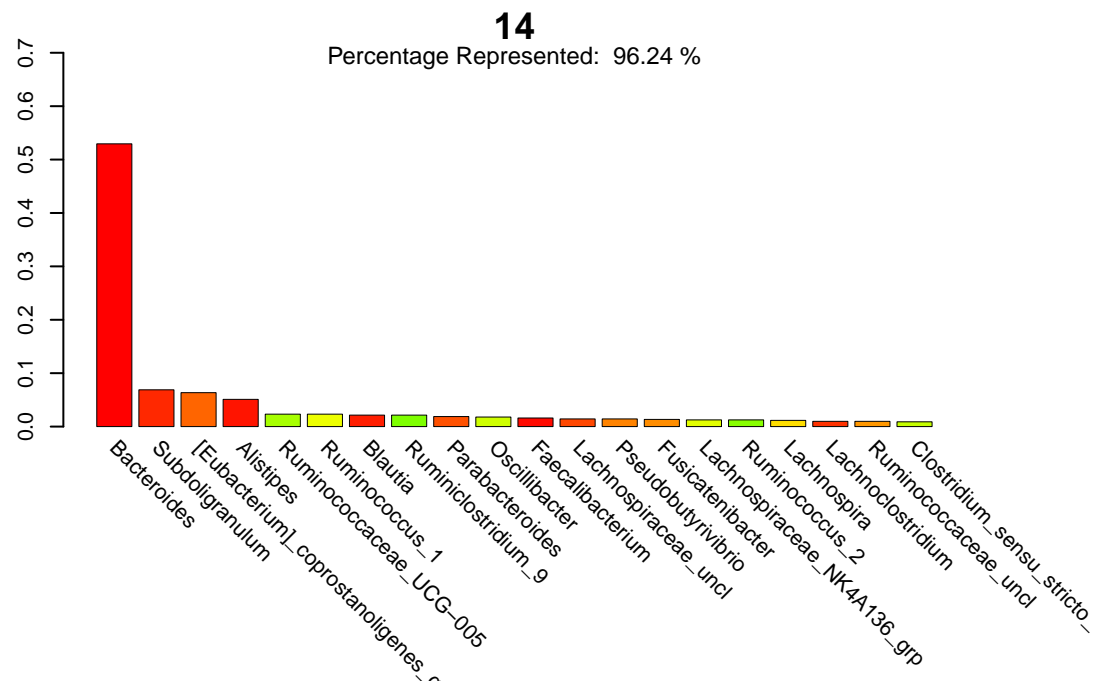
# Identifier



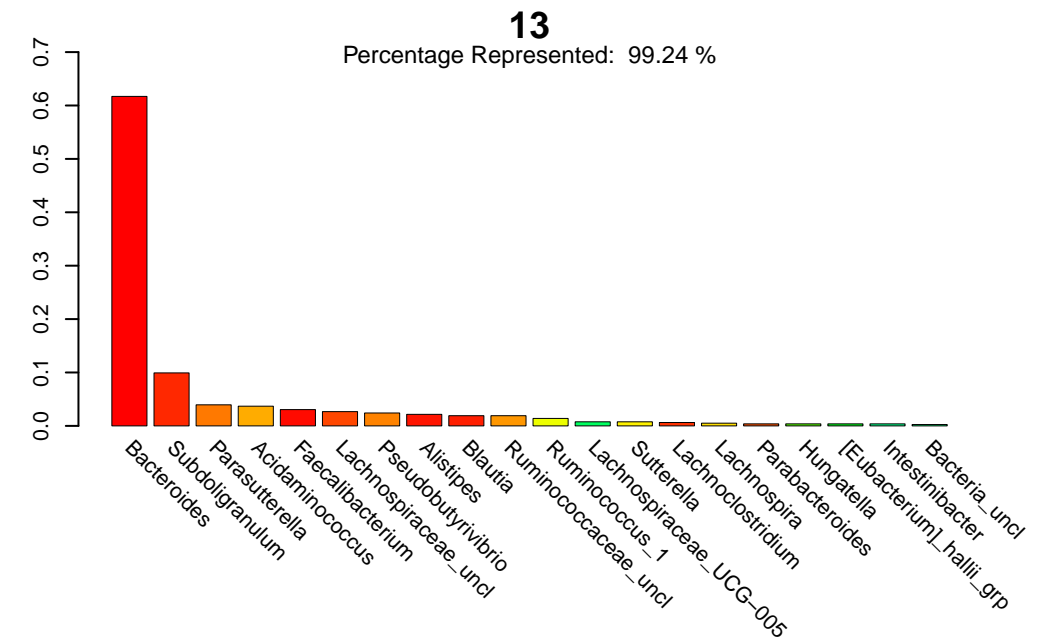
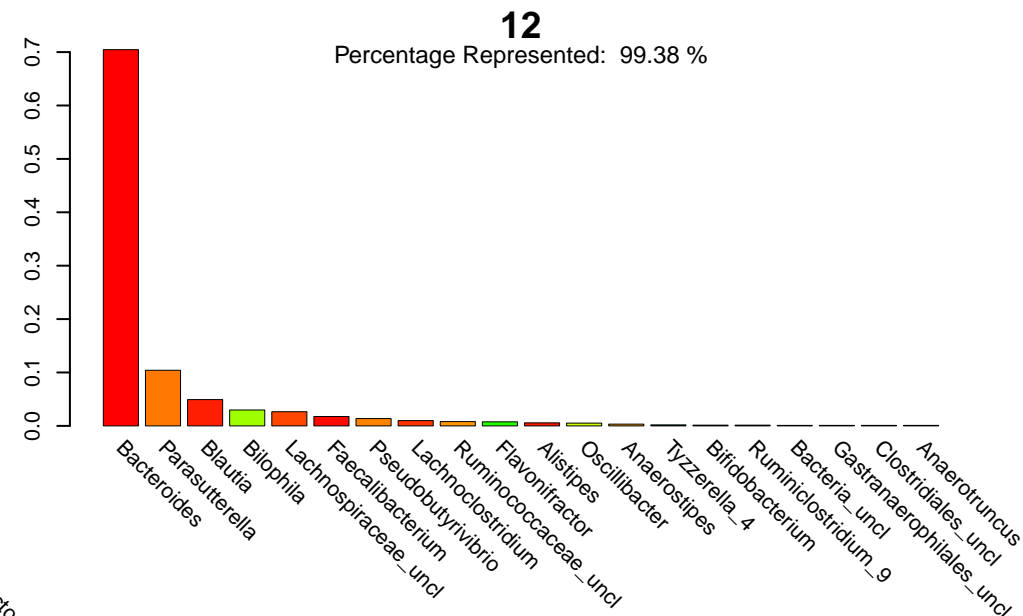
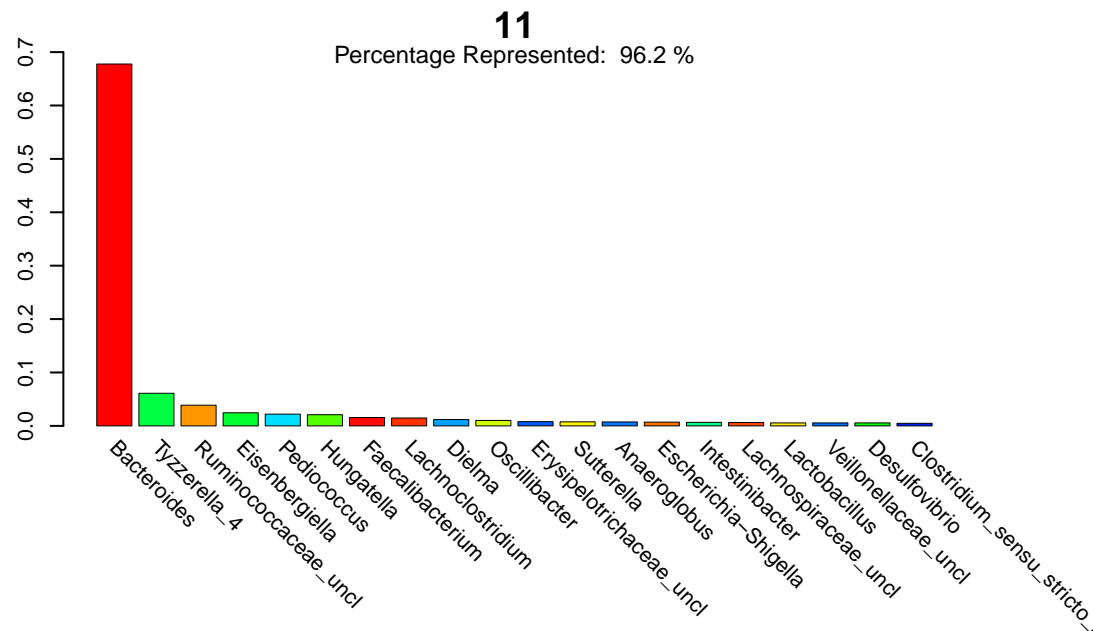
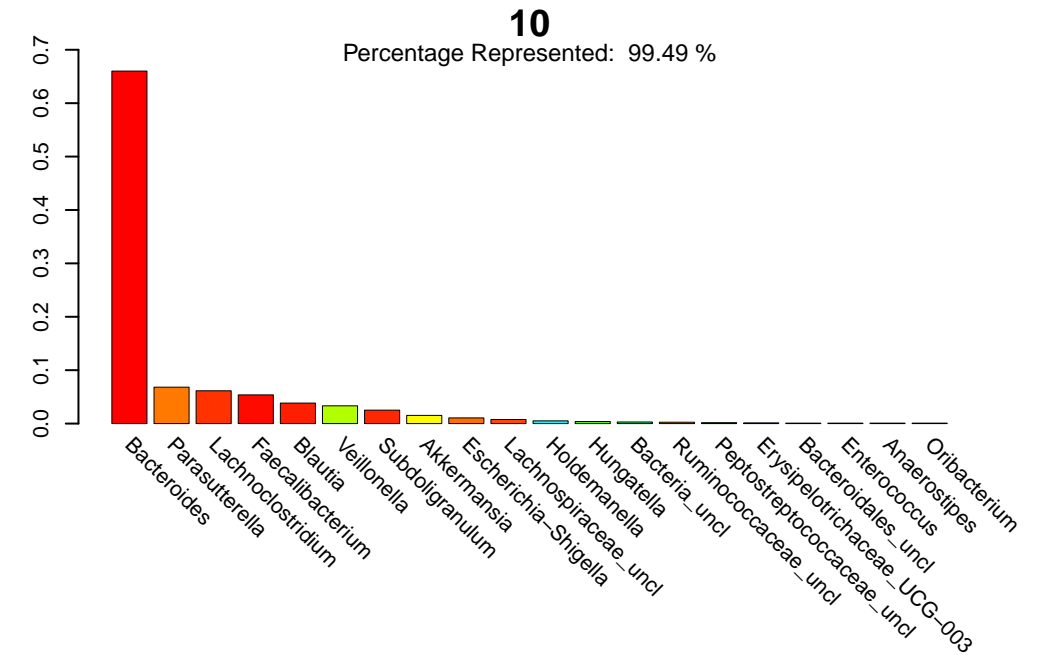
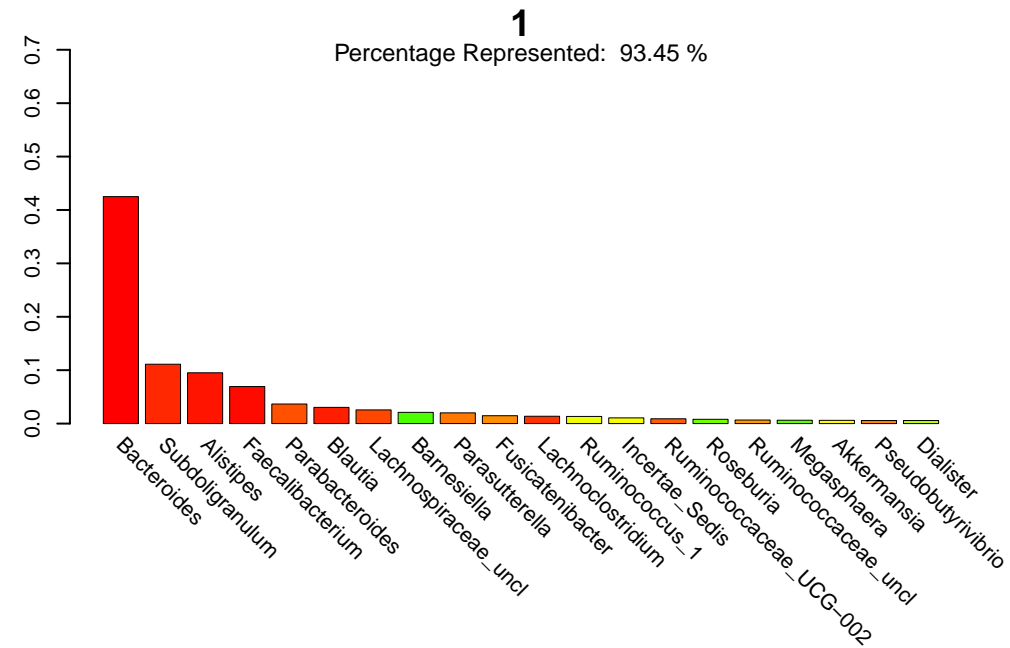
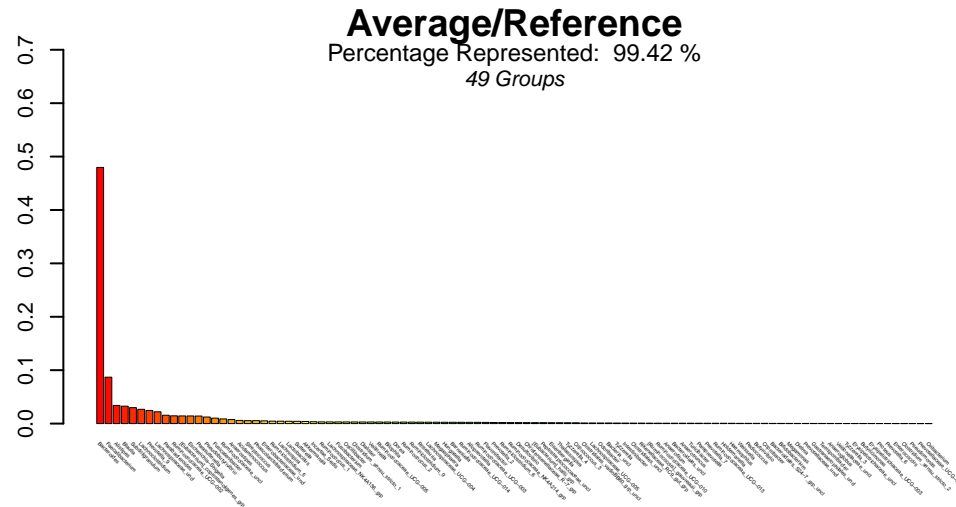
# RIH\_UPN



RIH\_UPN

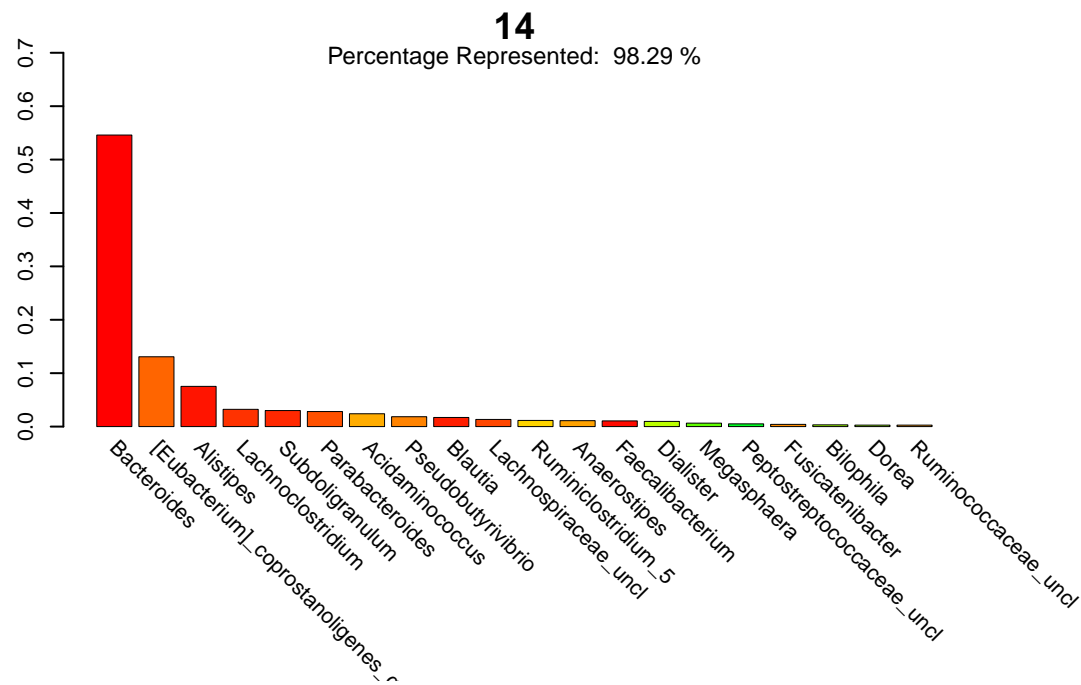


# PCR\_Temporary\_Number

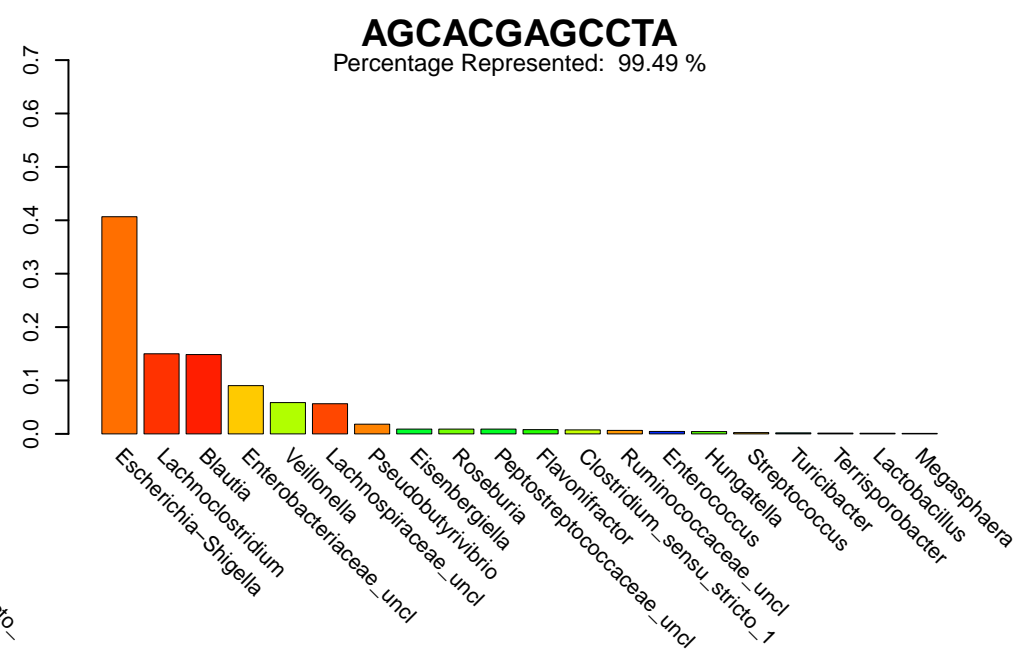
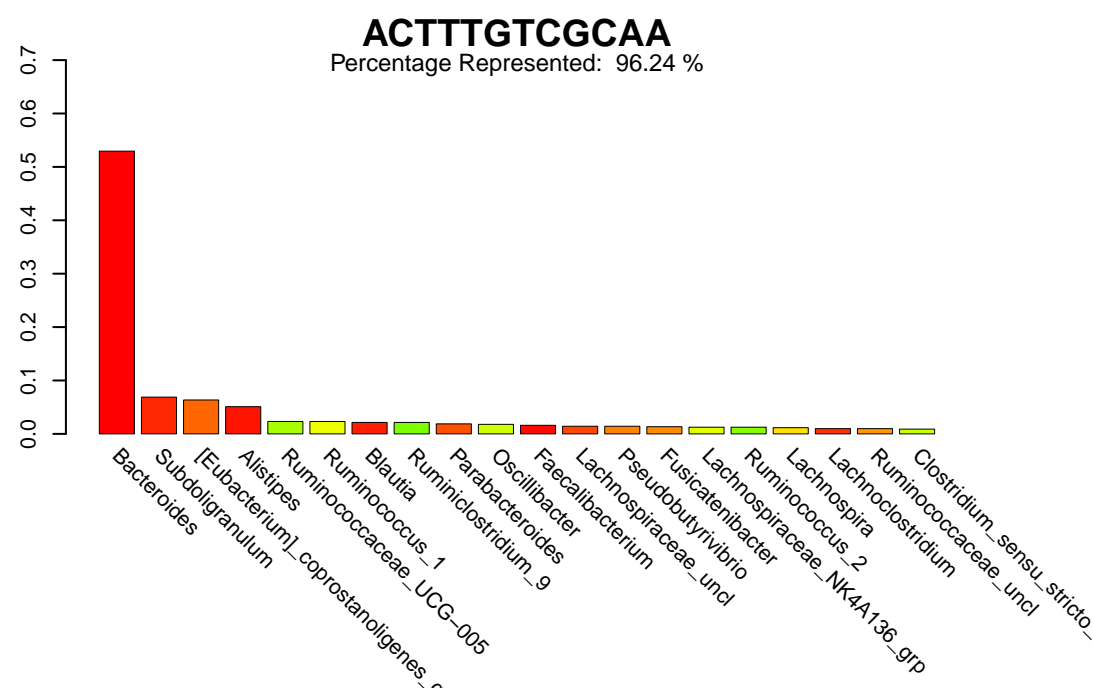
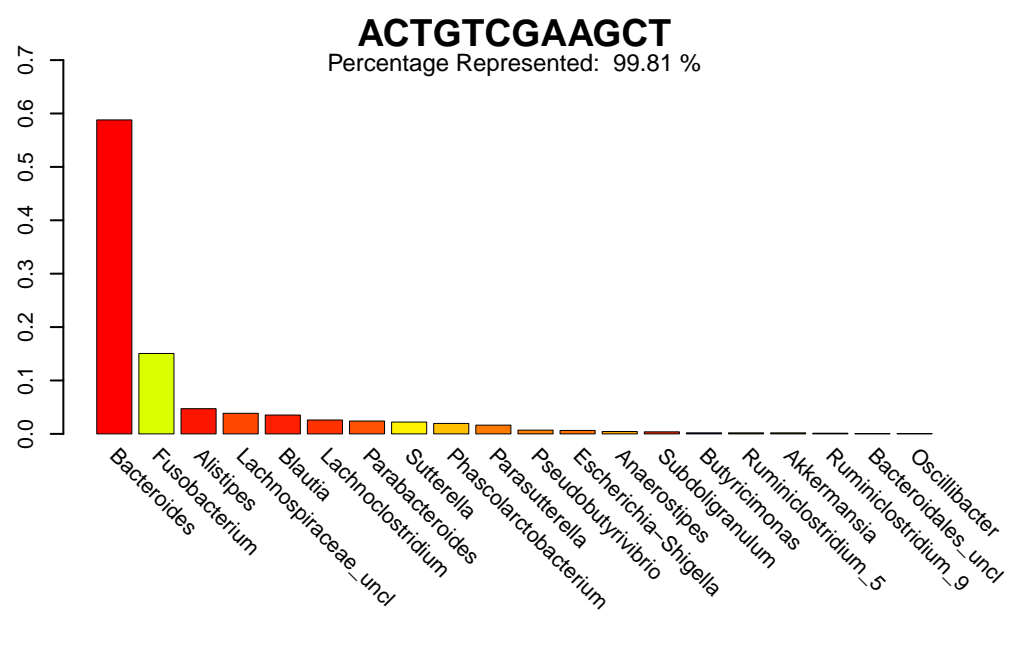
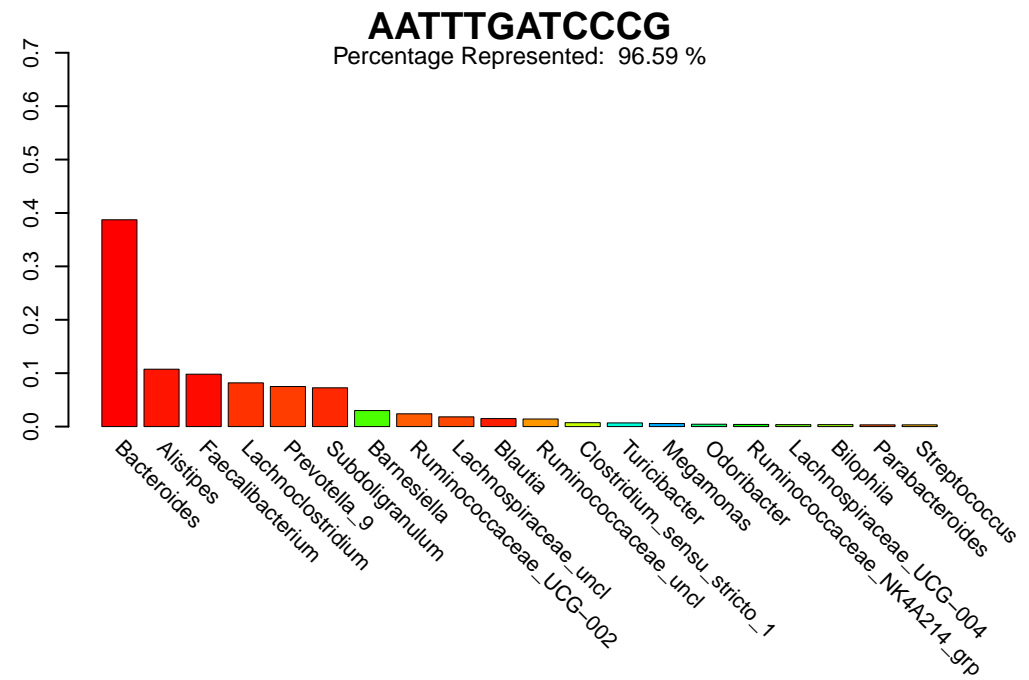
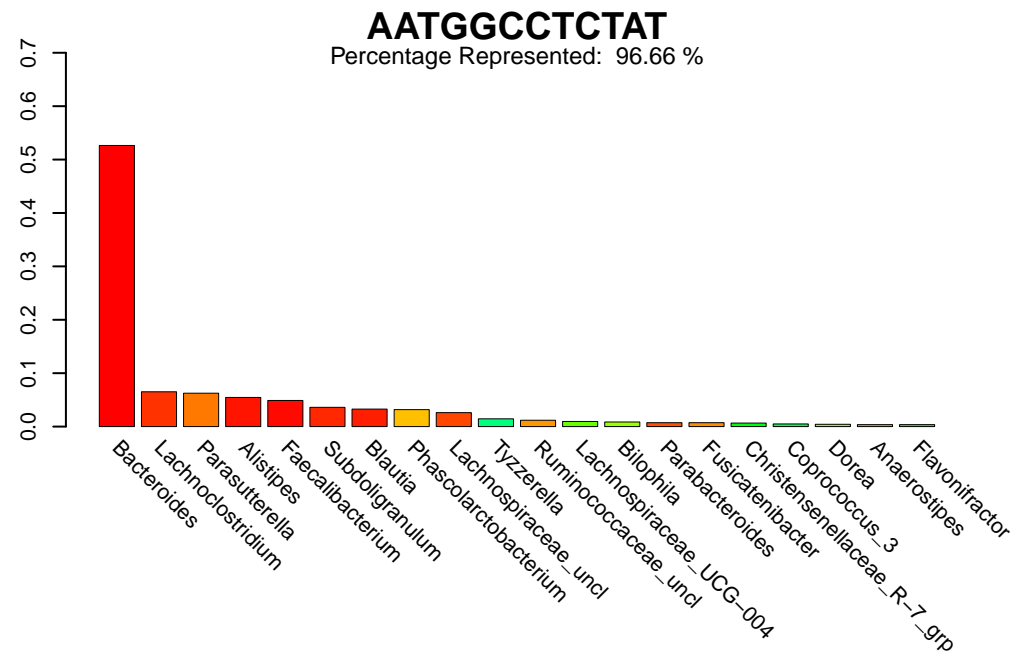
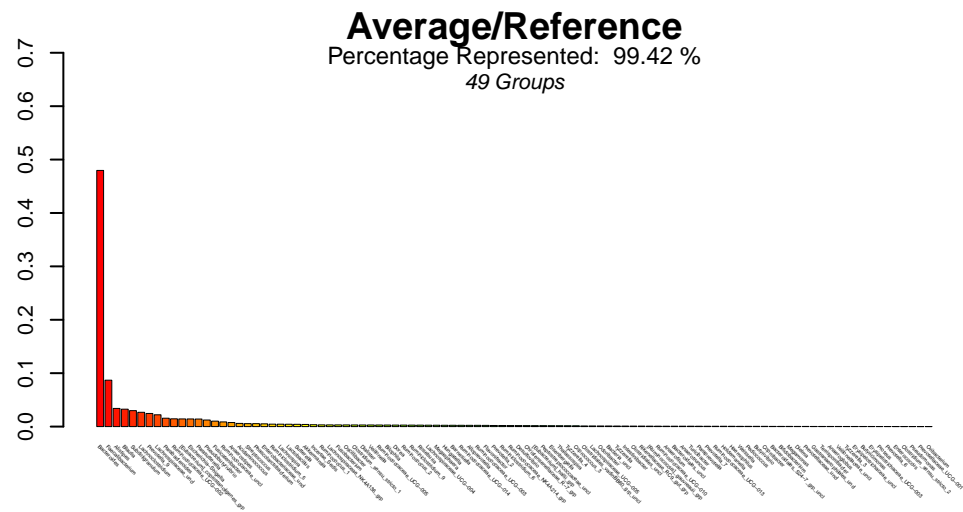




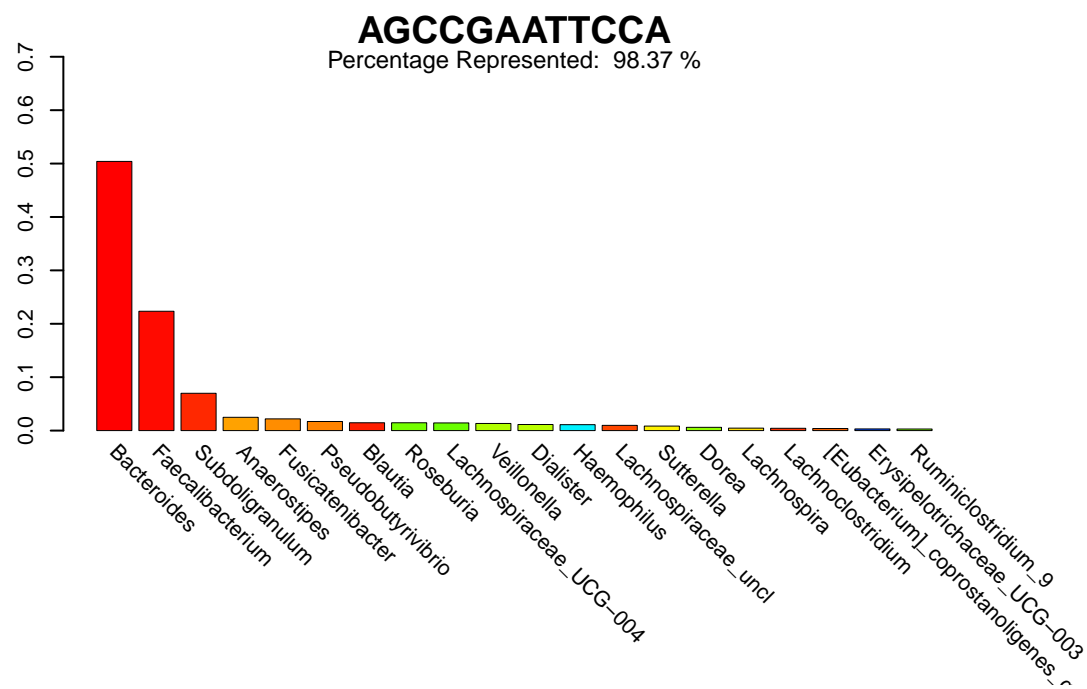
# PCR\_Temporary\_Number



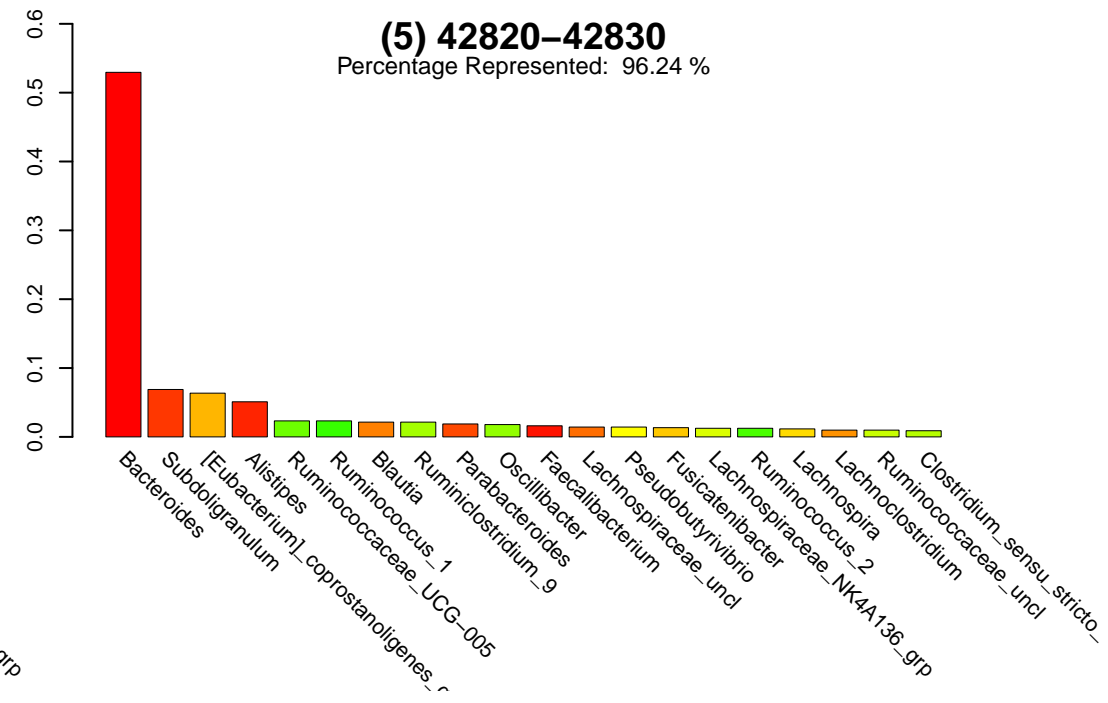
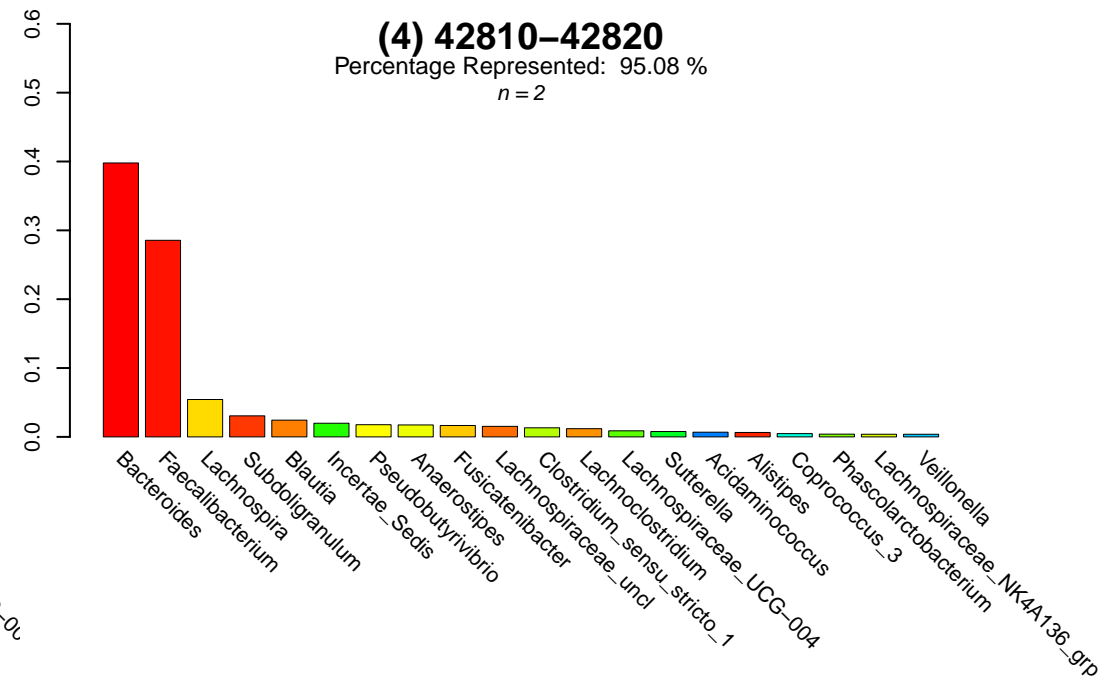
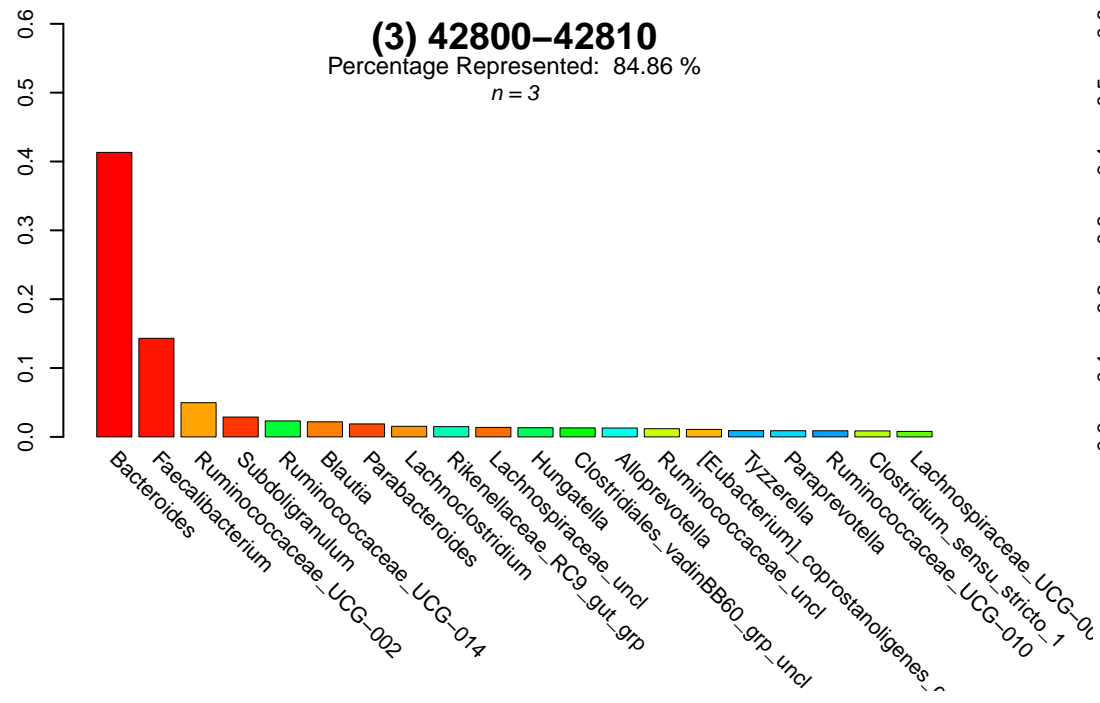
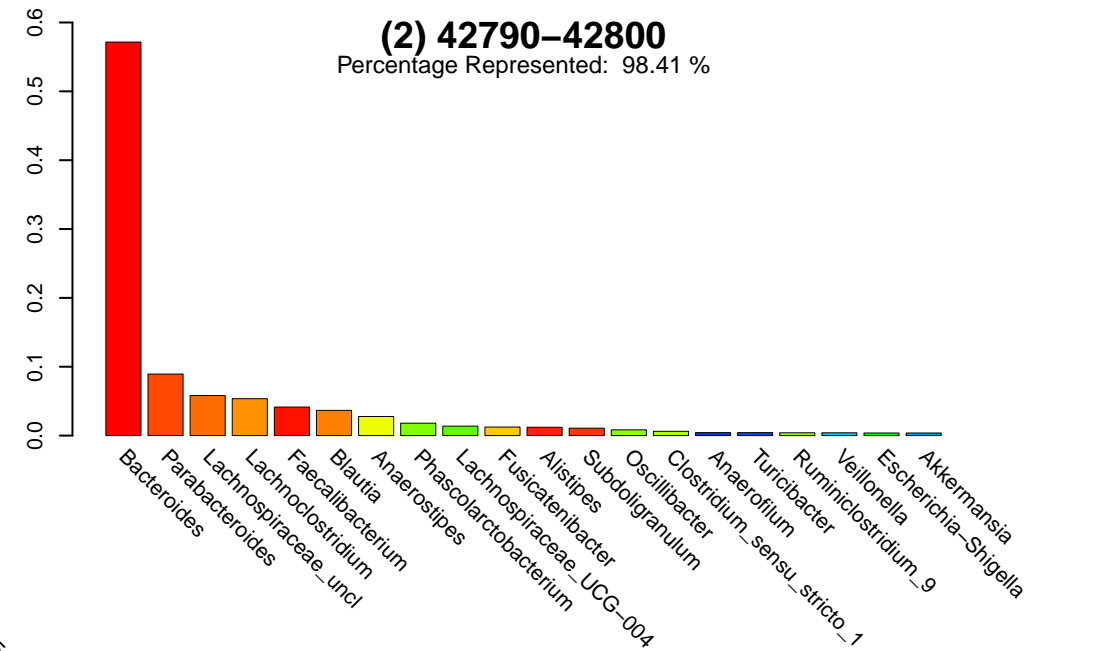
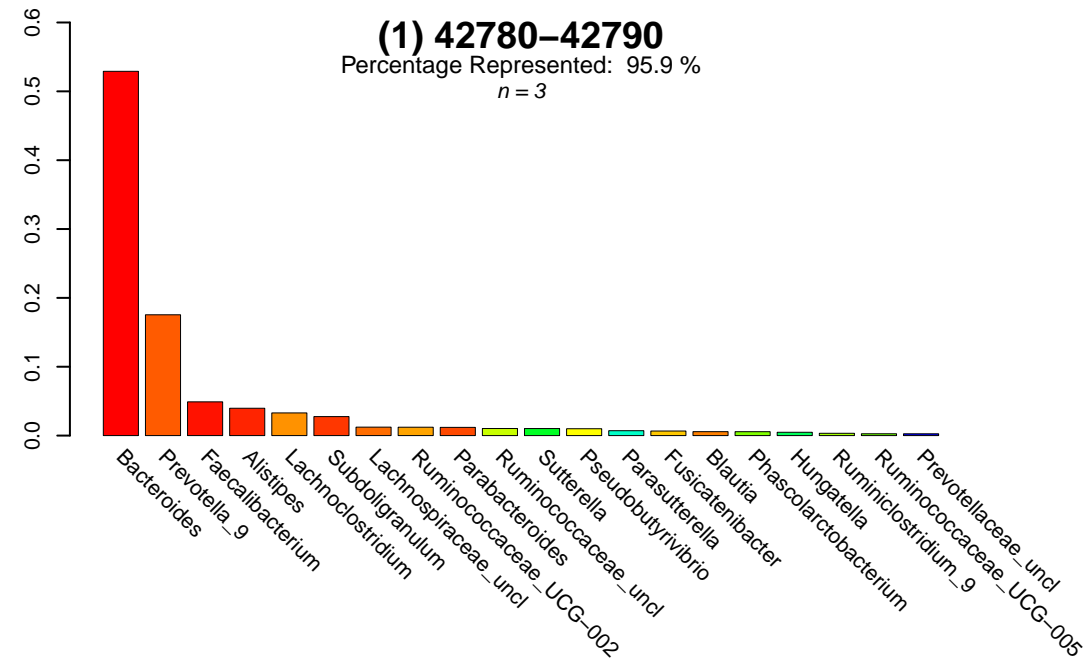
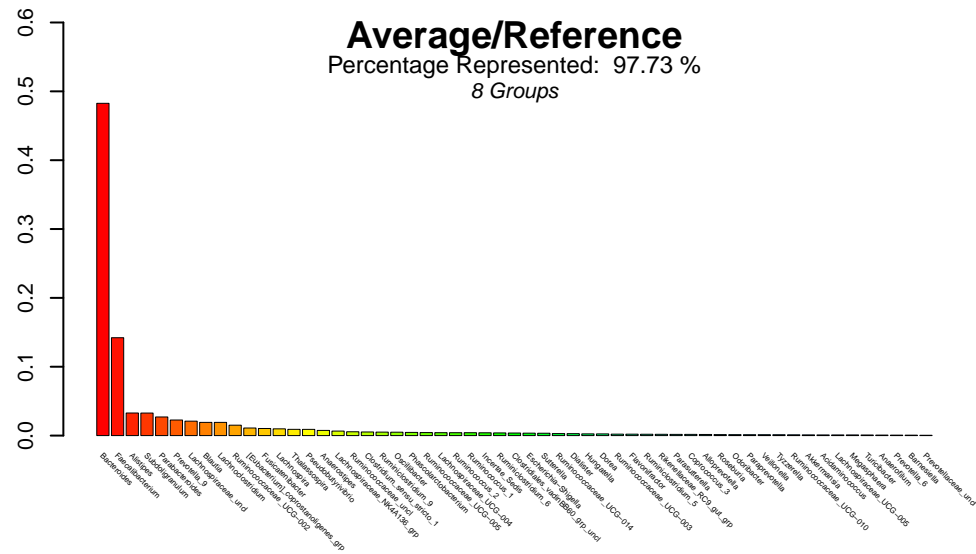
# Barcode.RC



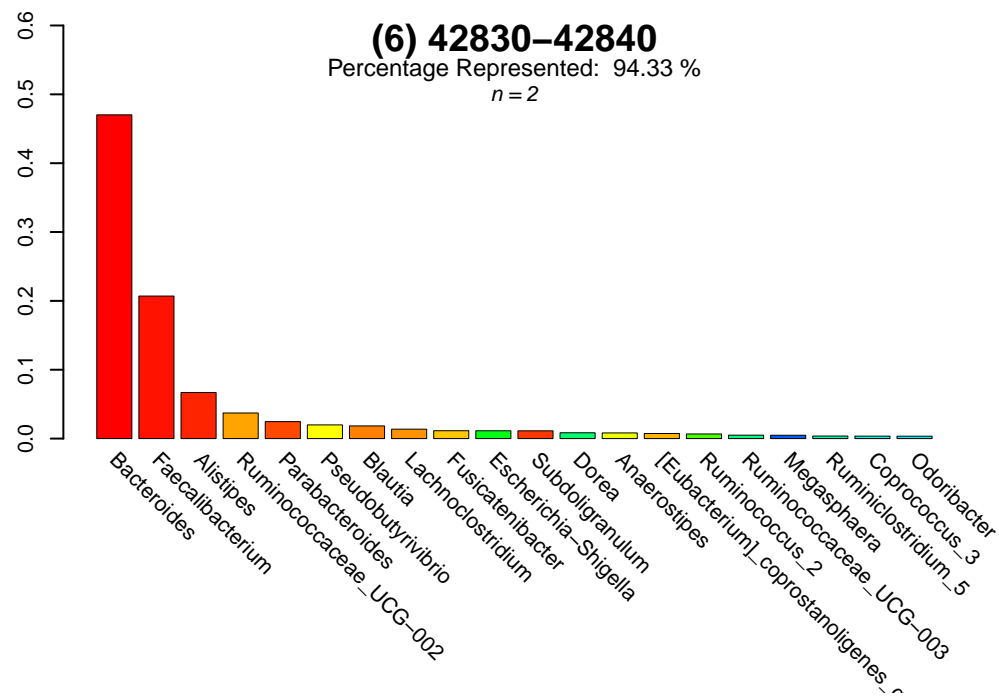
# Barcode.RC



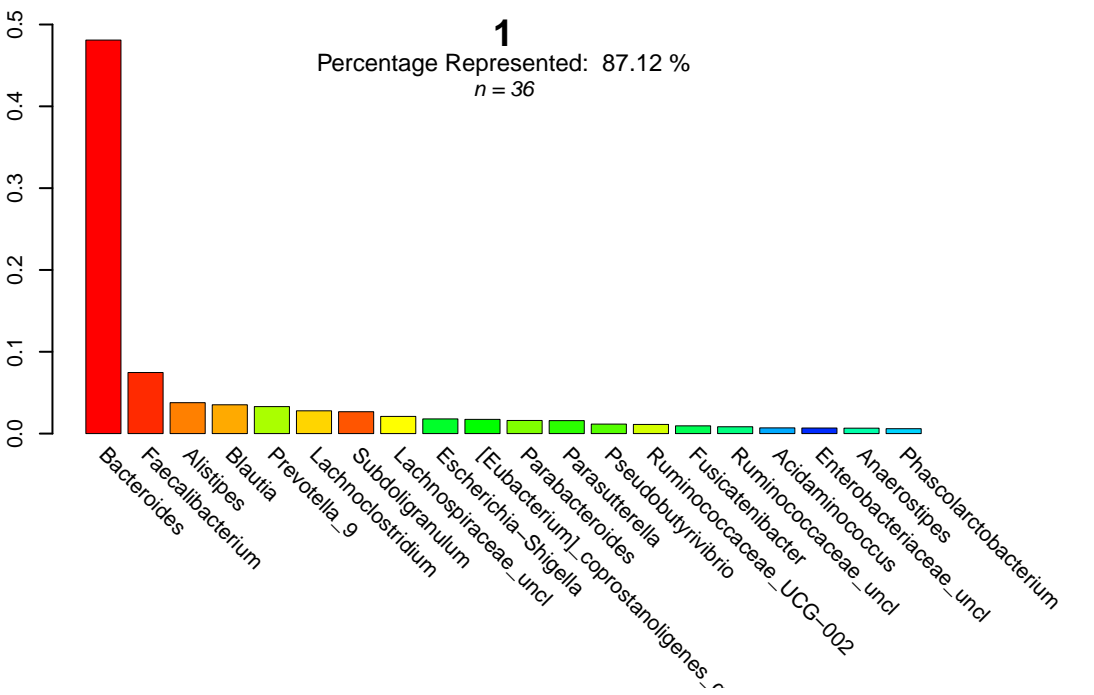
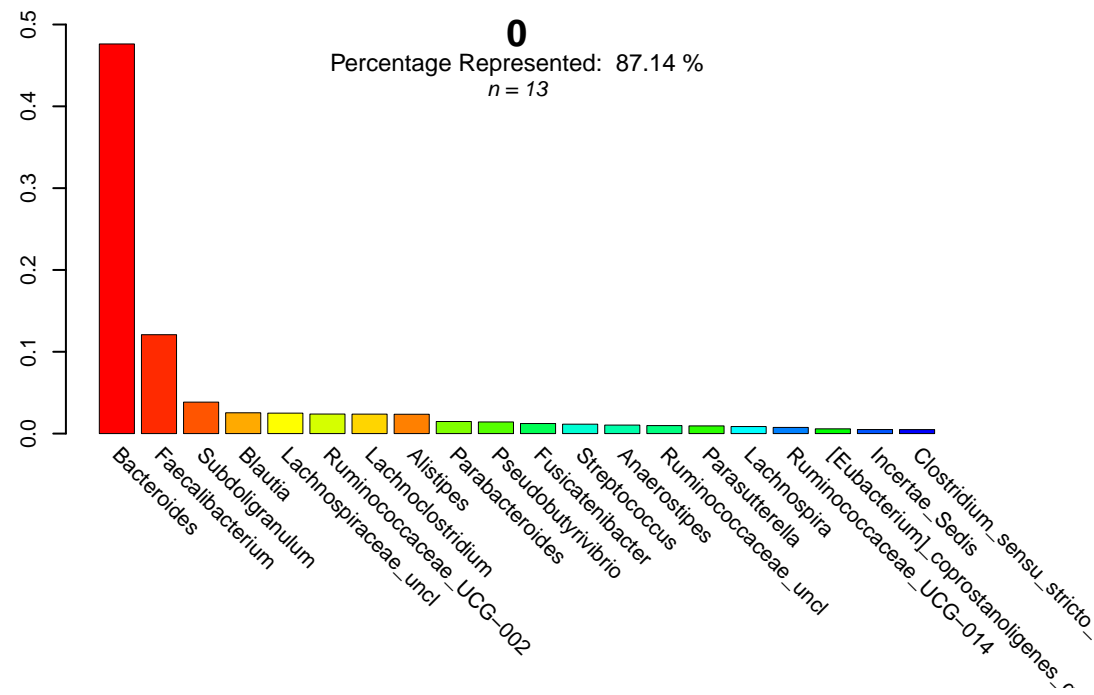
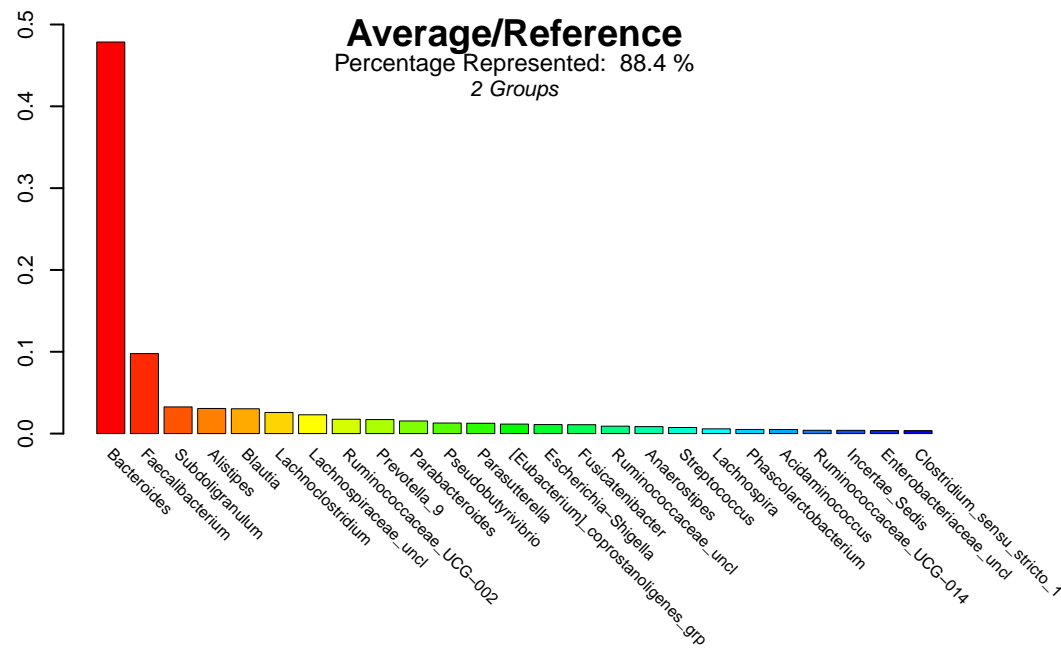
## Consent.Date



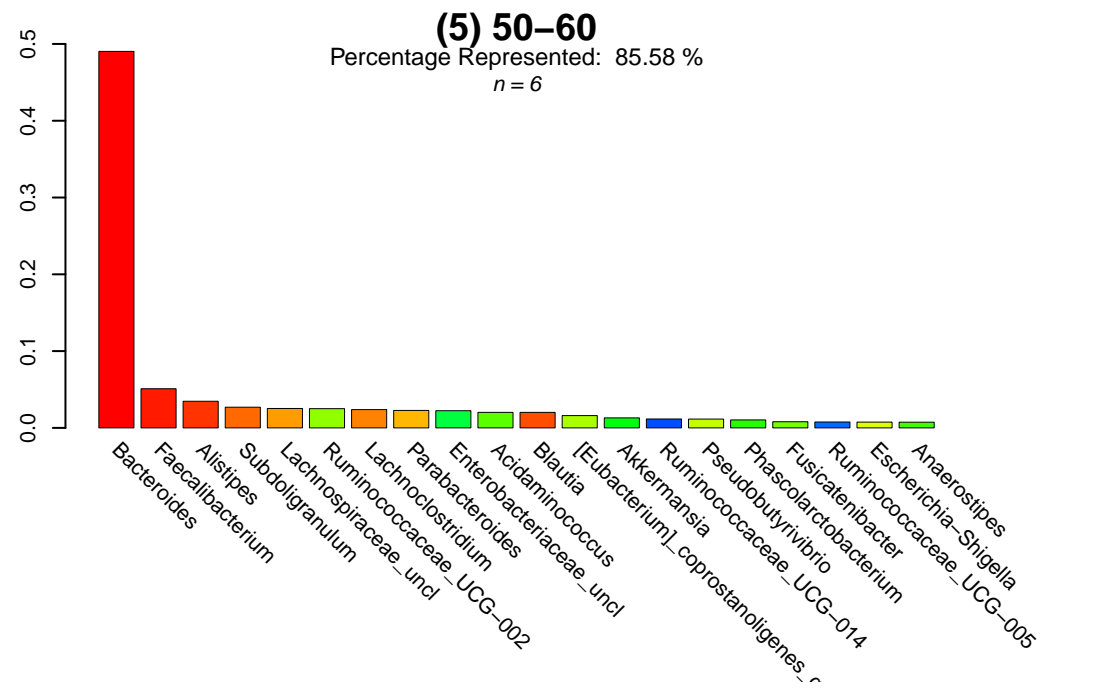
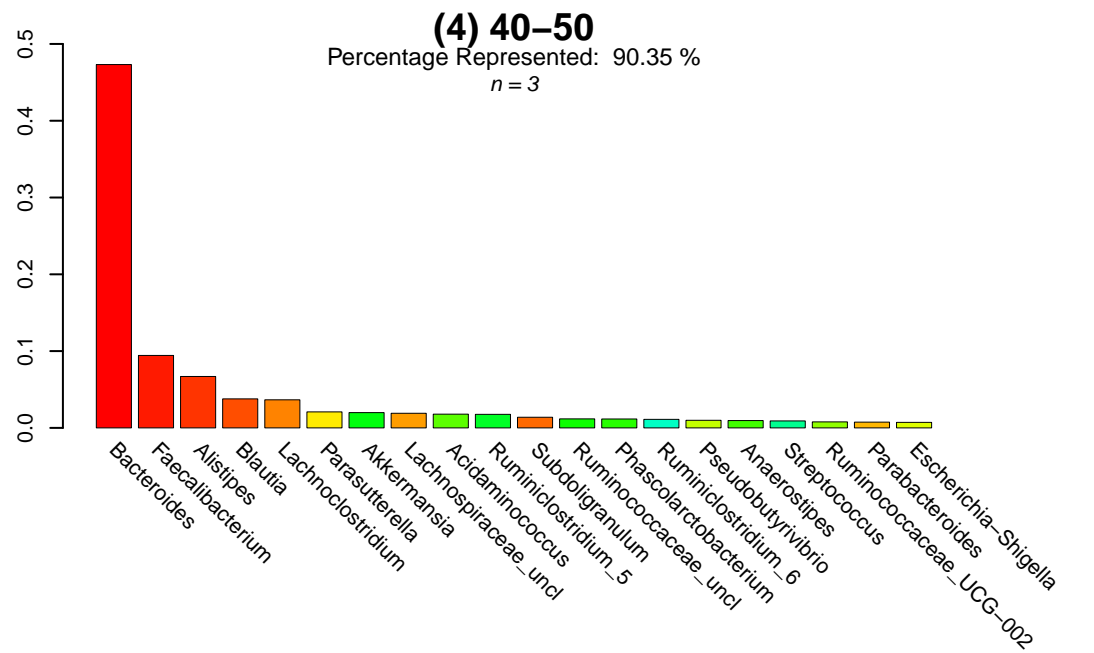
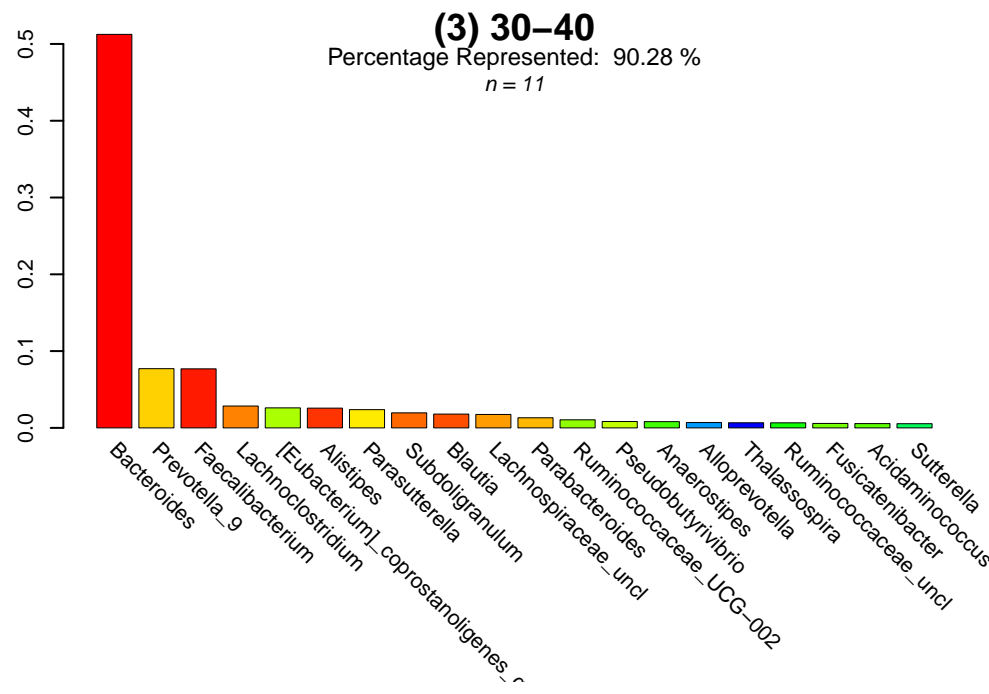
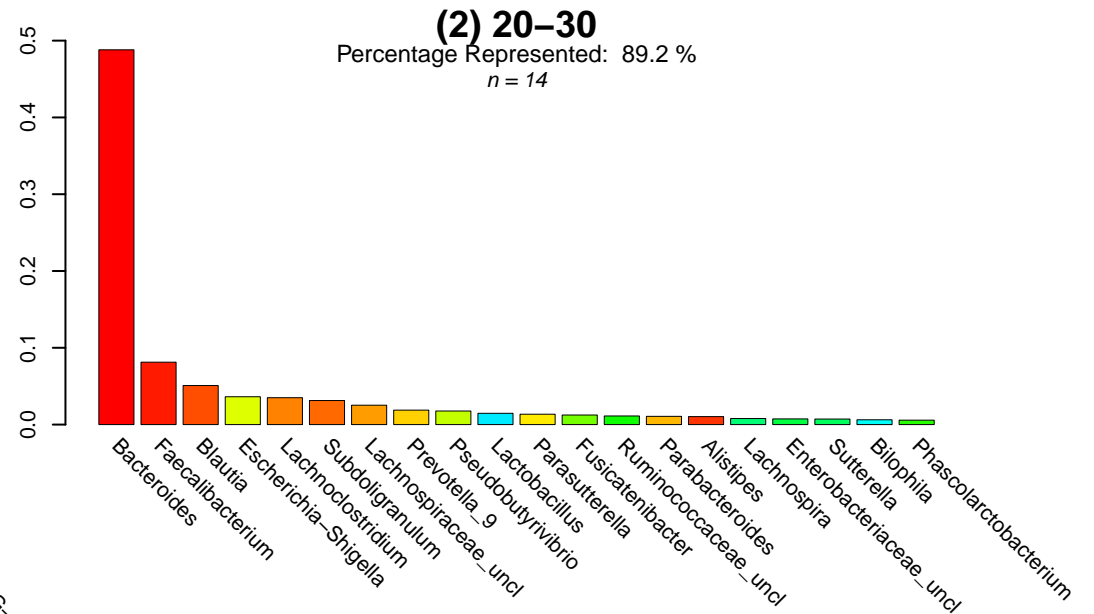
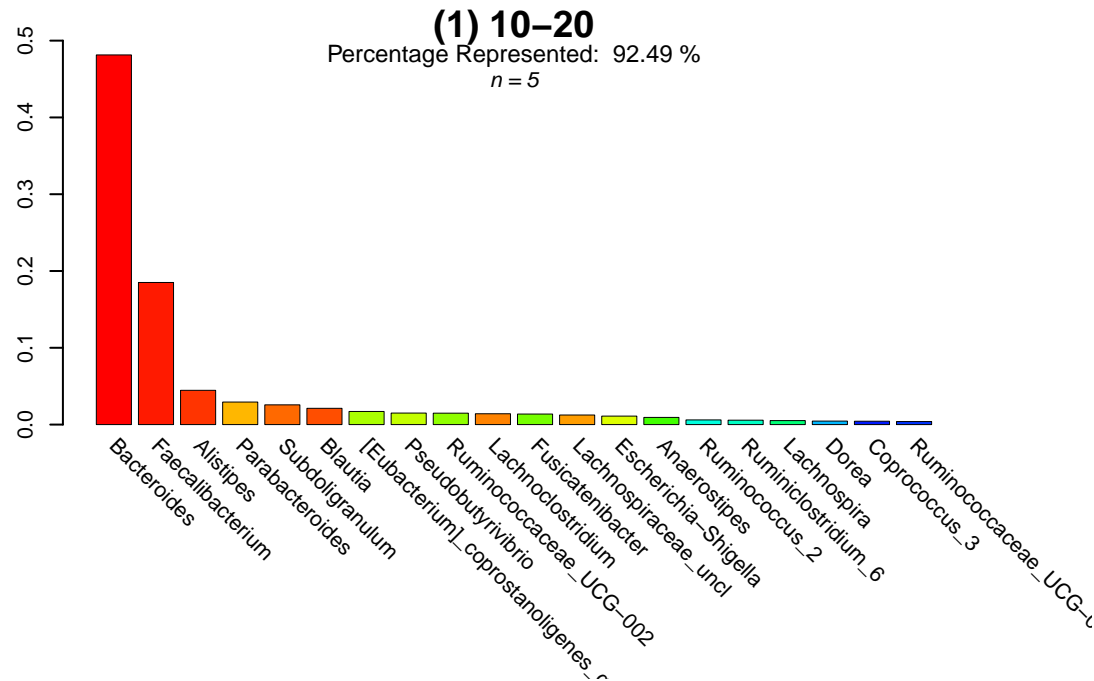
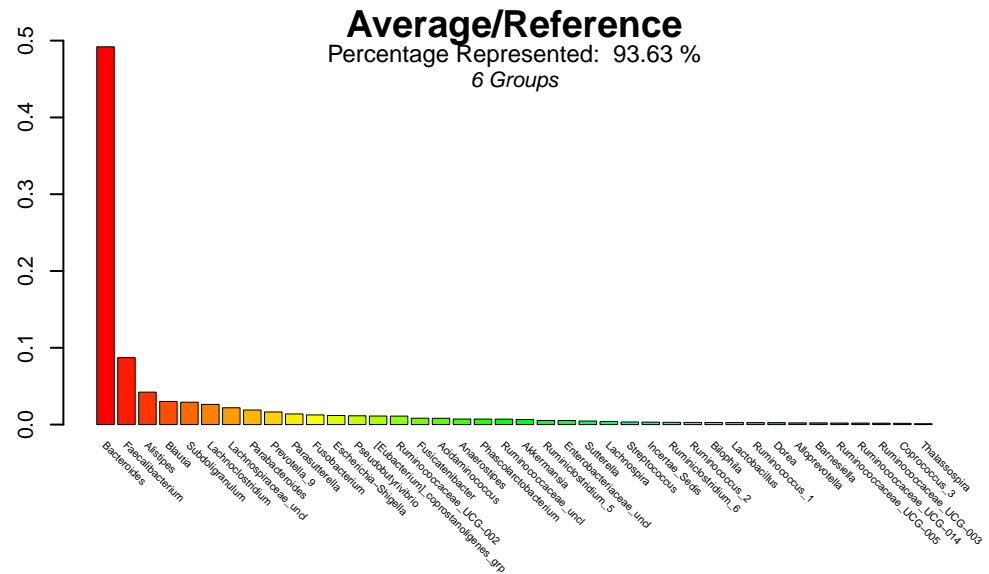
# Consent.Date



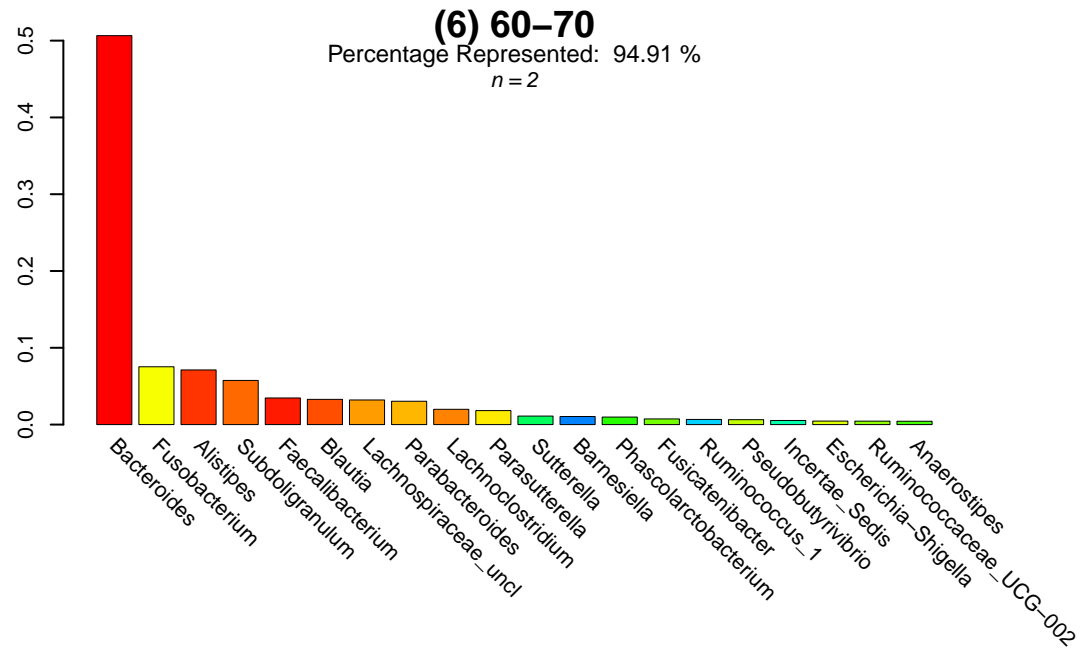
isFemale



# Age

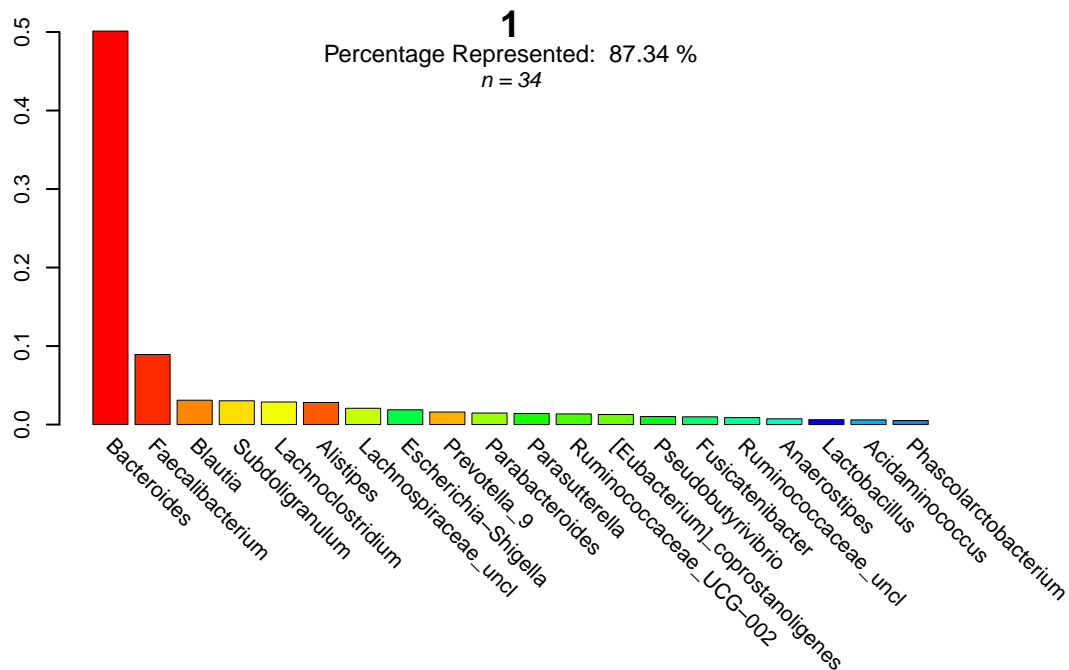
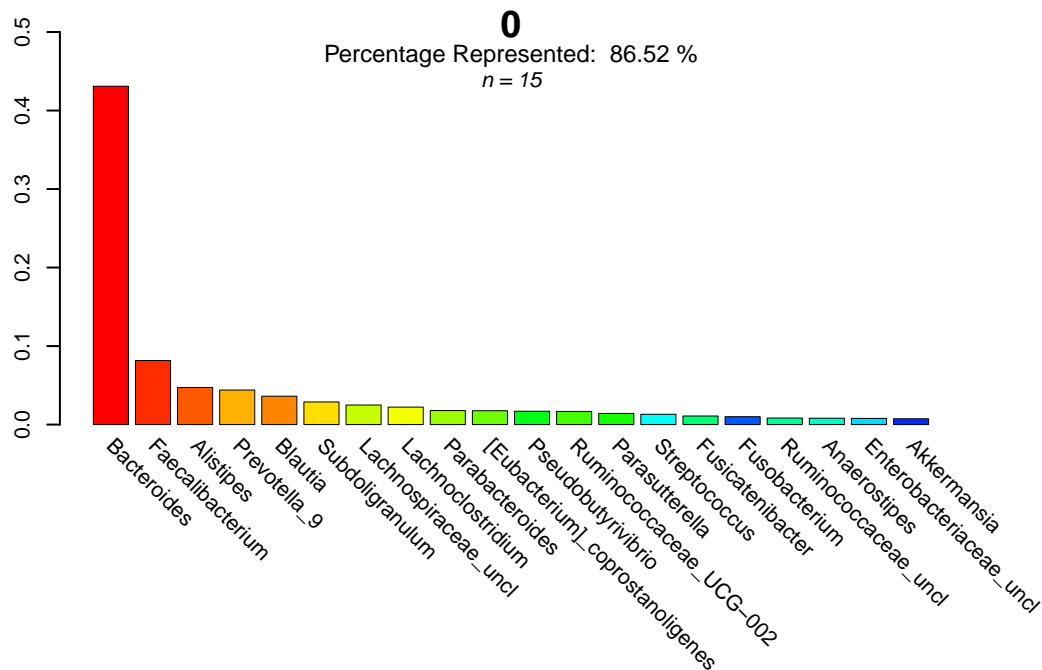
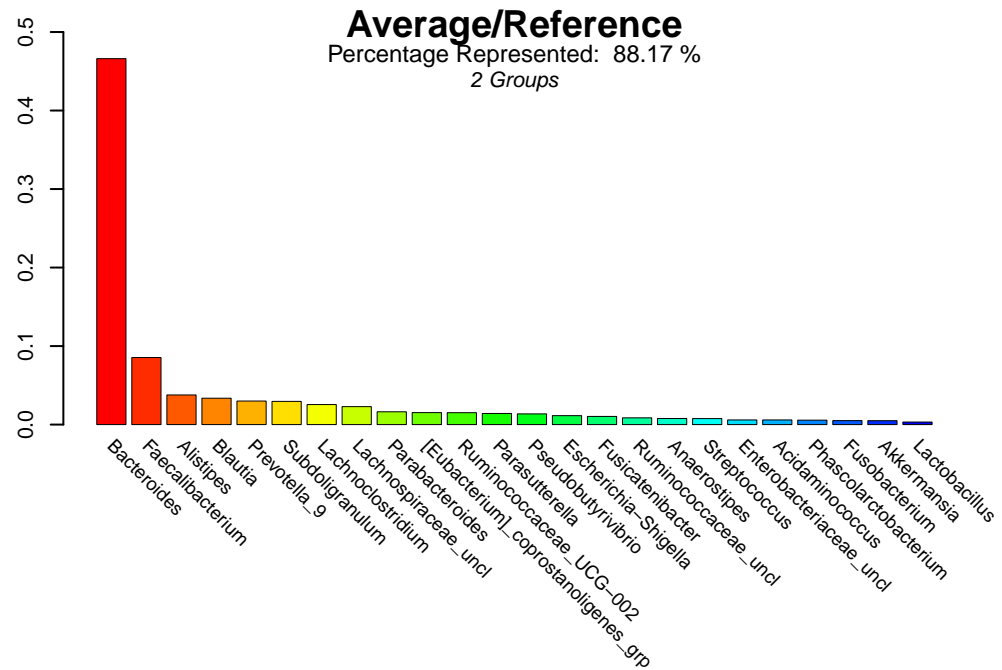


# Age

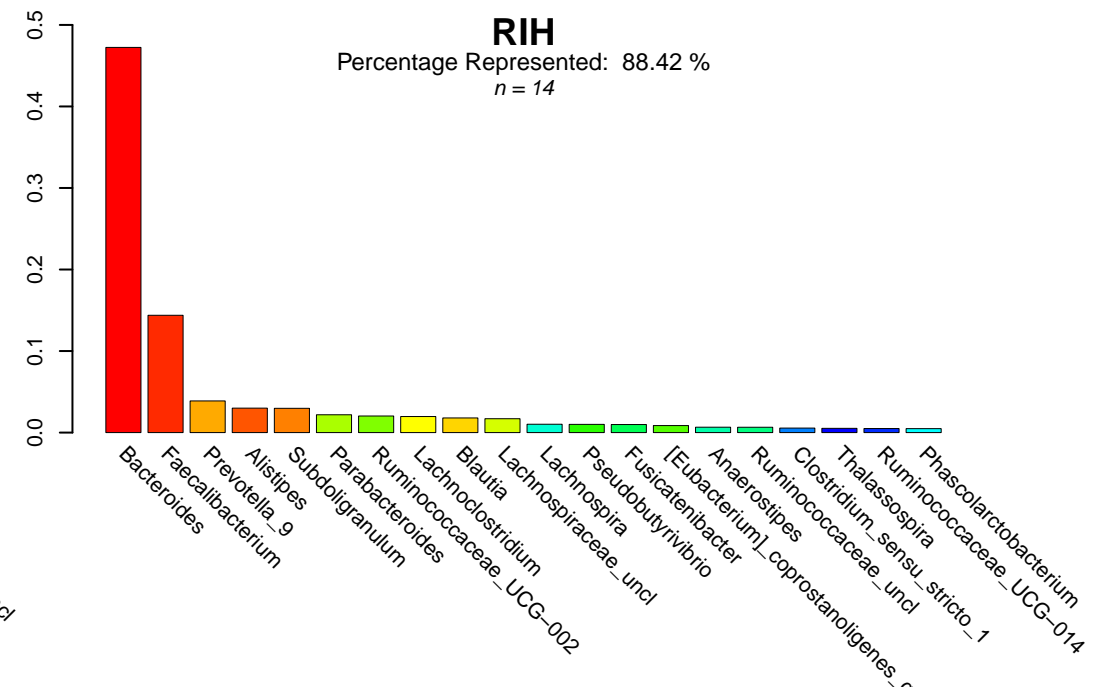
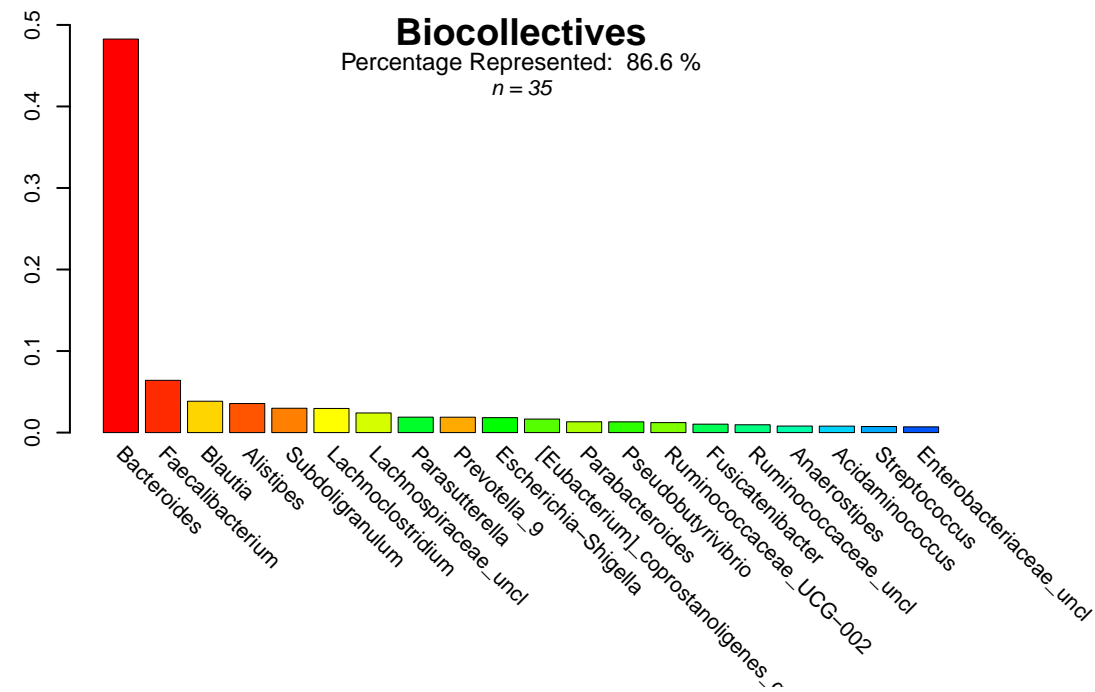
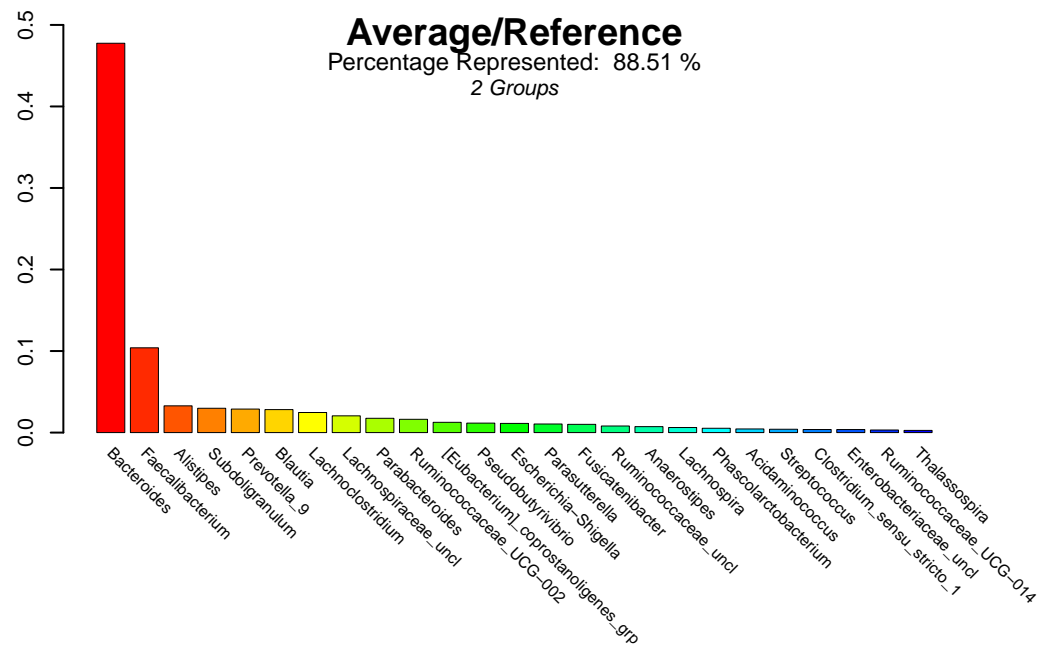




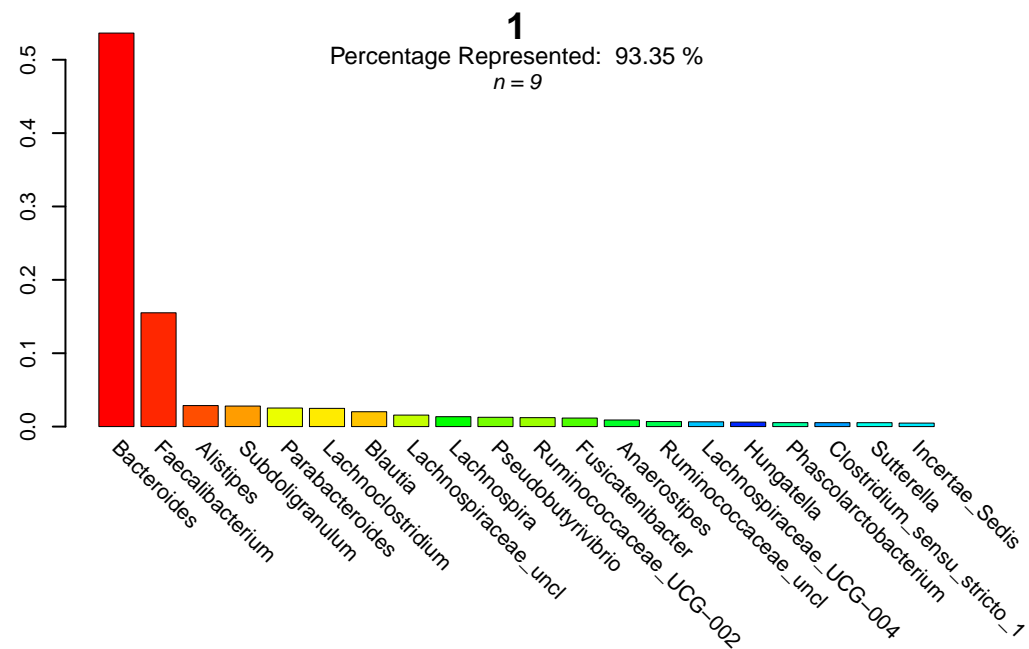
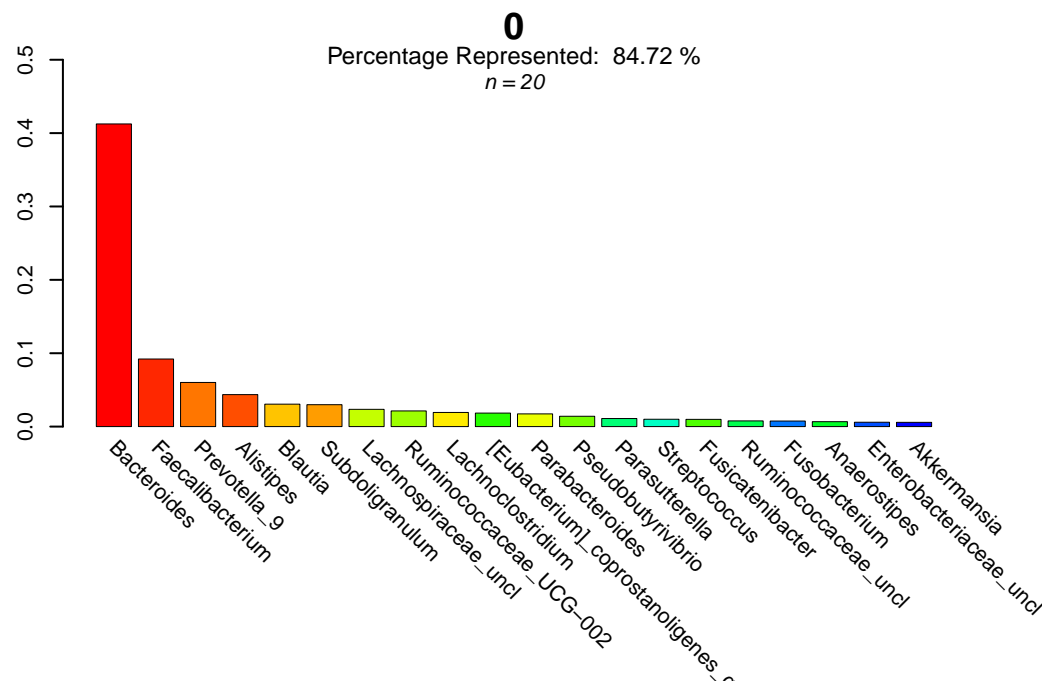
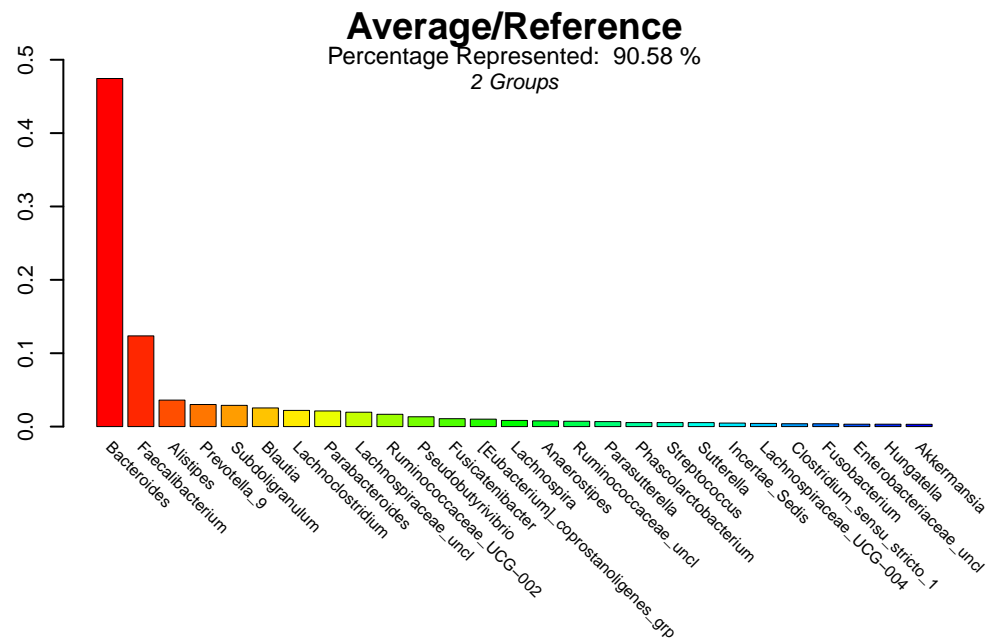
# isCase



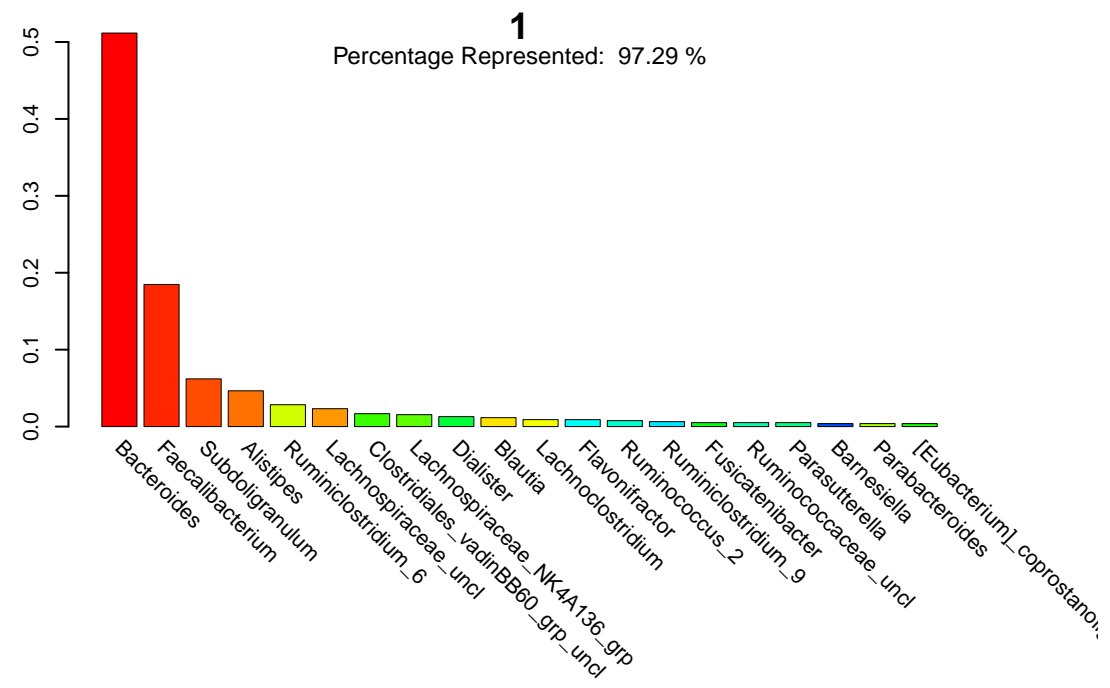
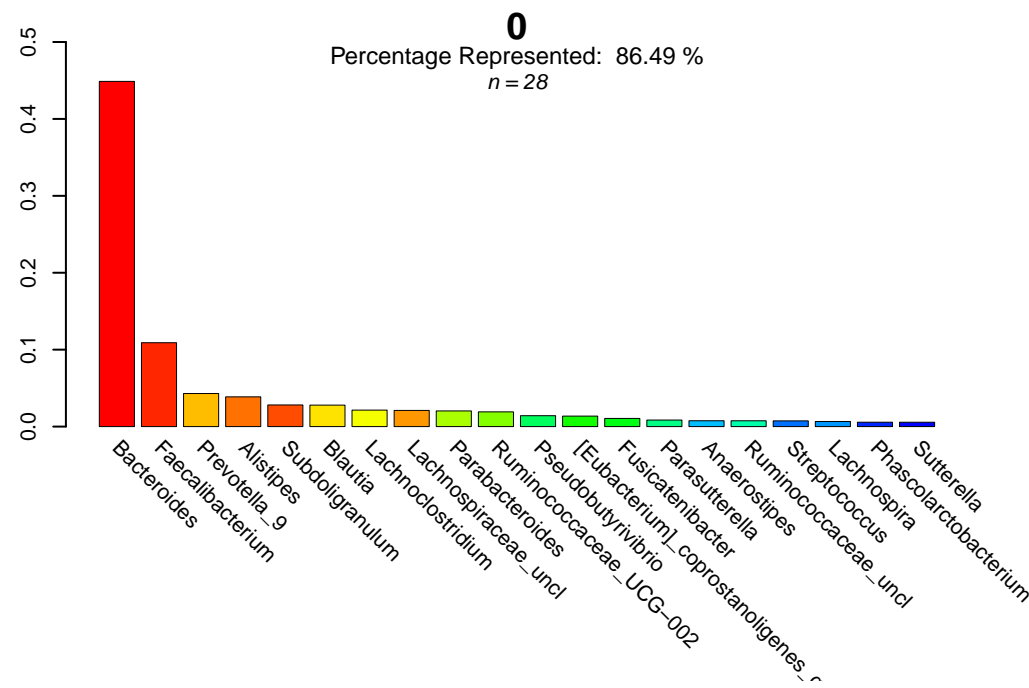
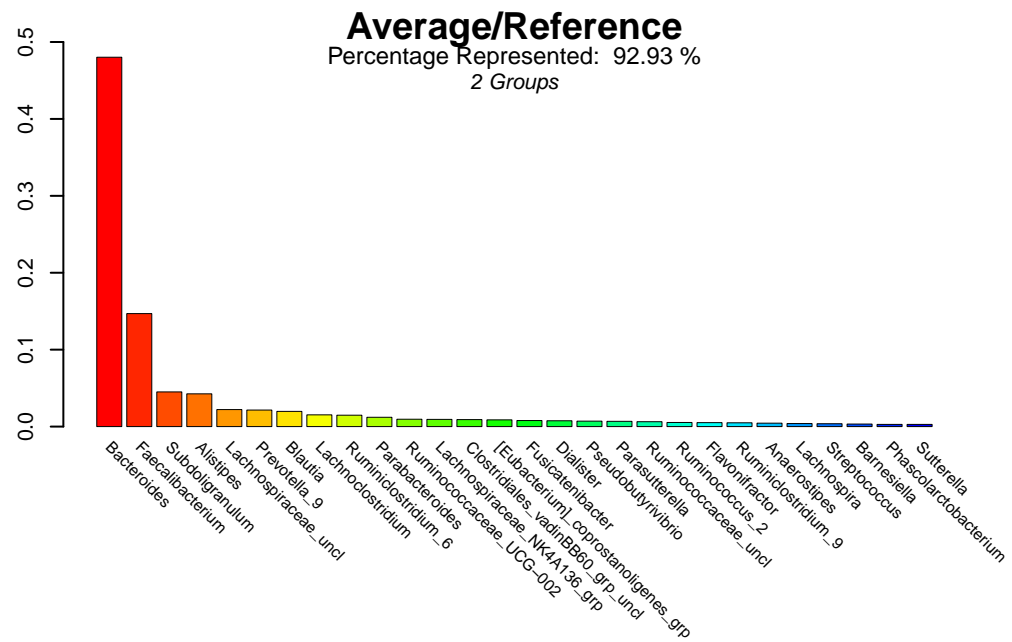
# Origin



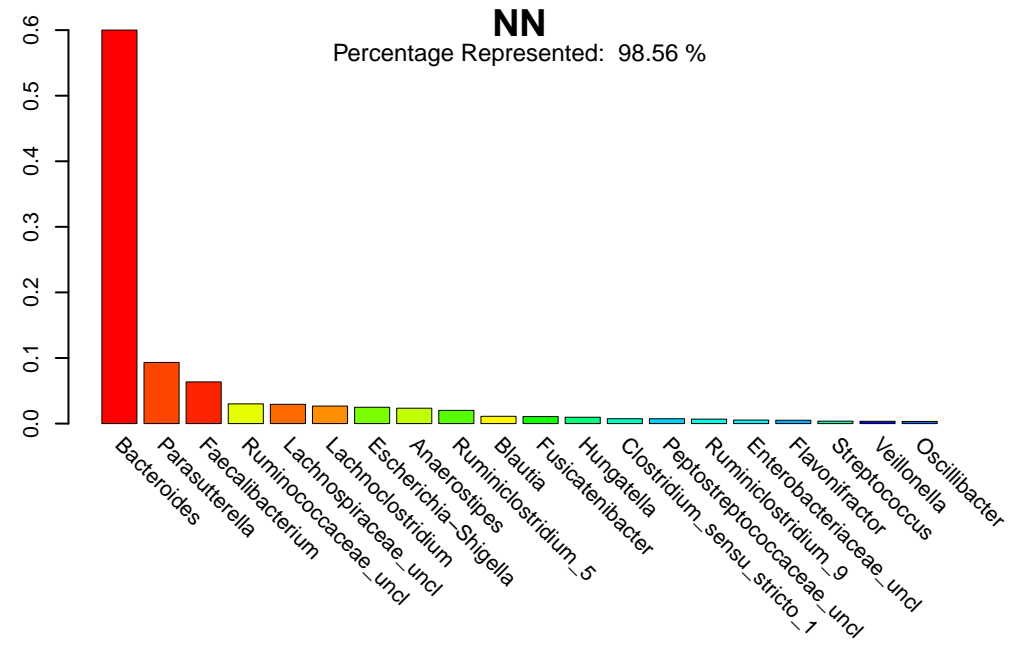
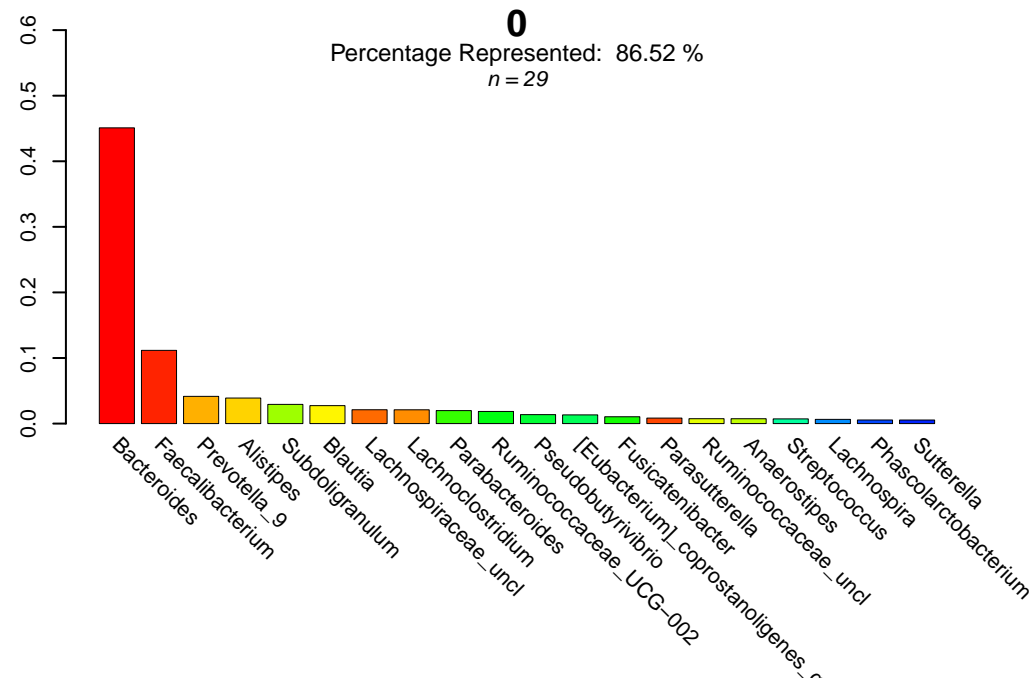
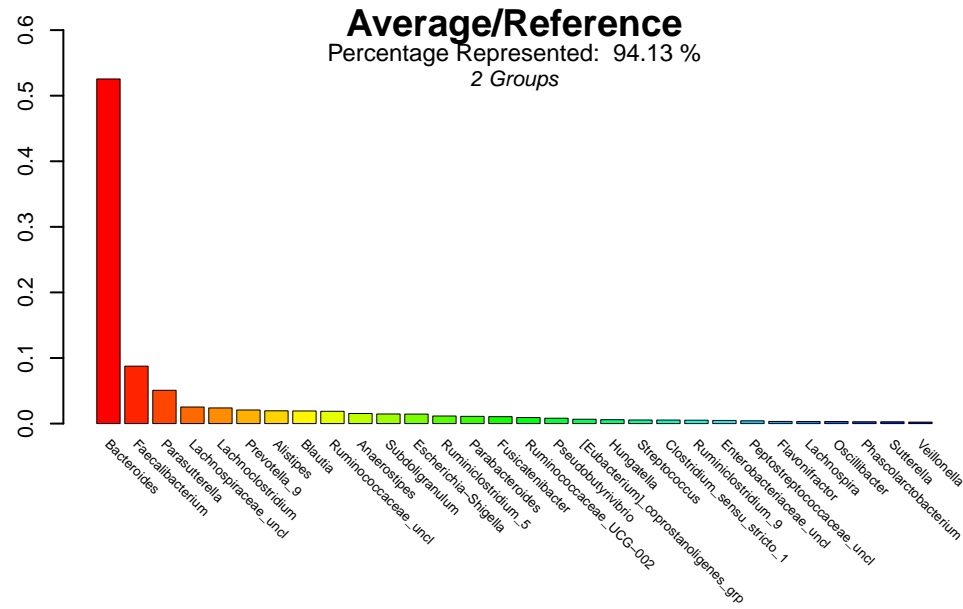
# hadHydrea



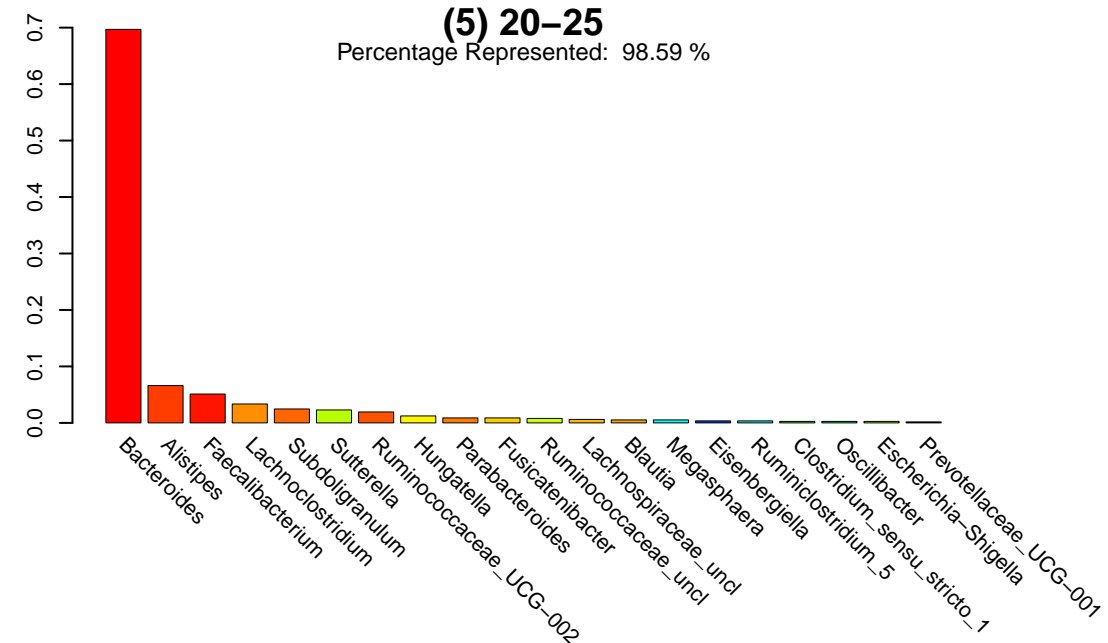
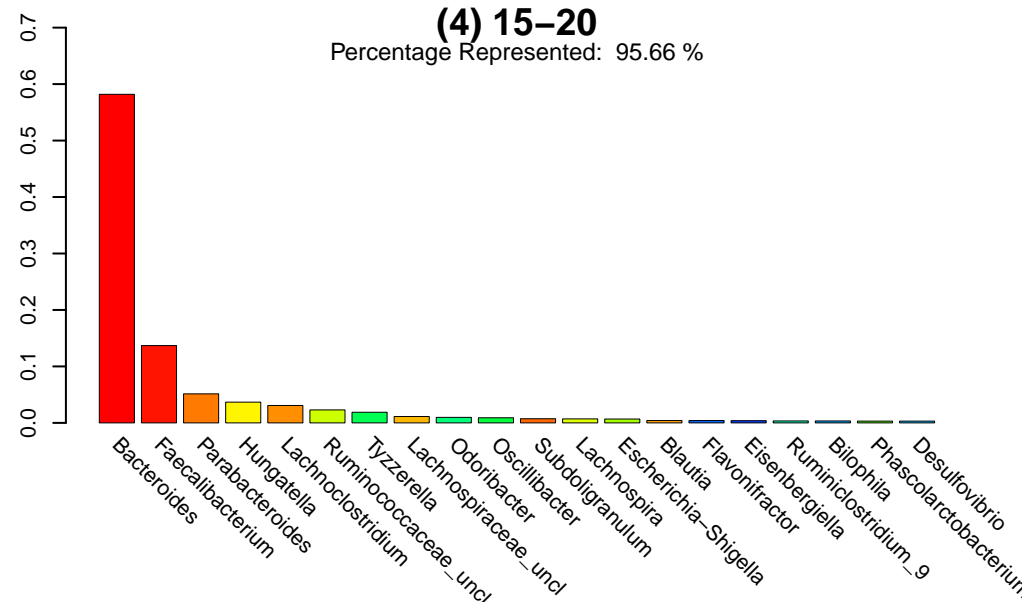
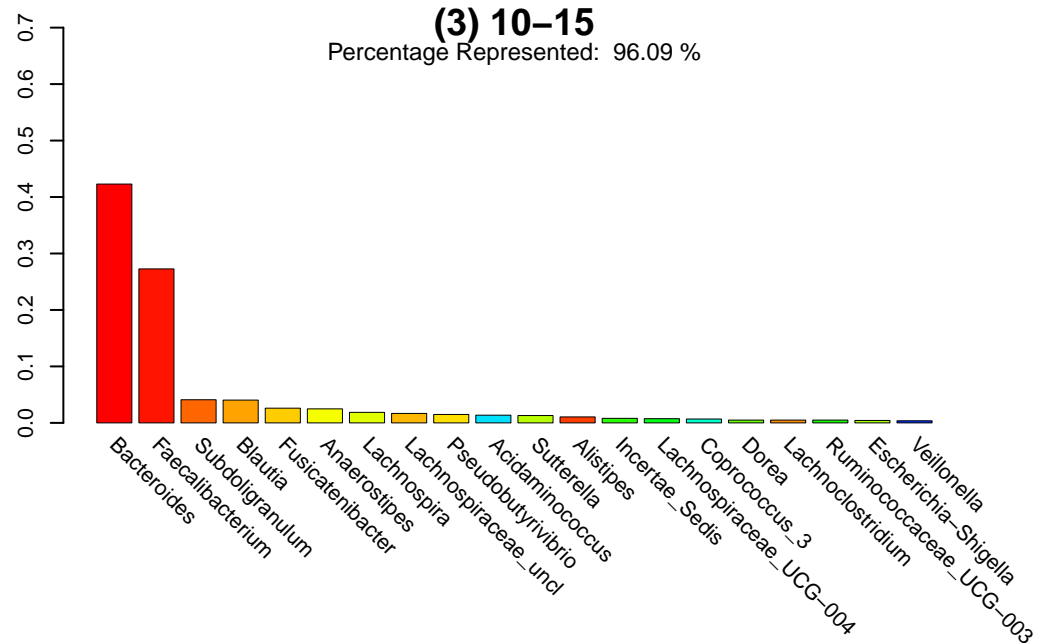
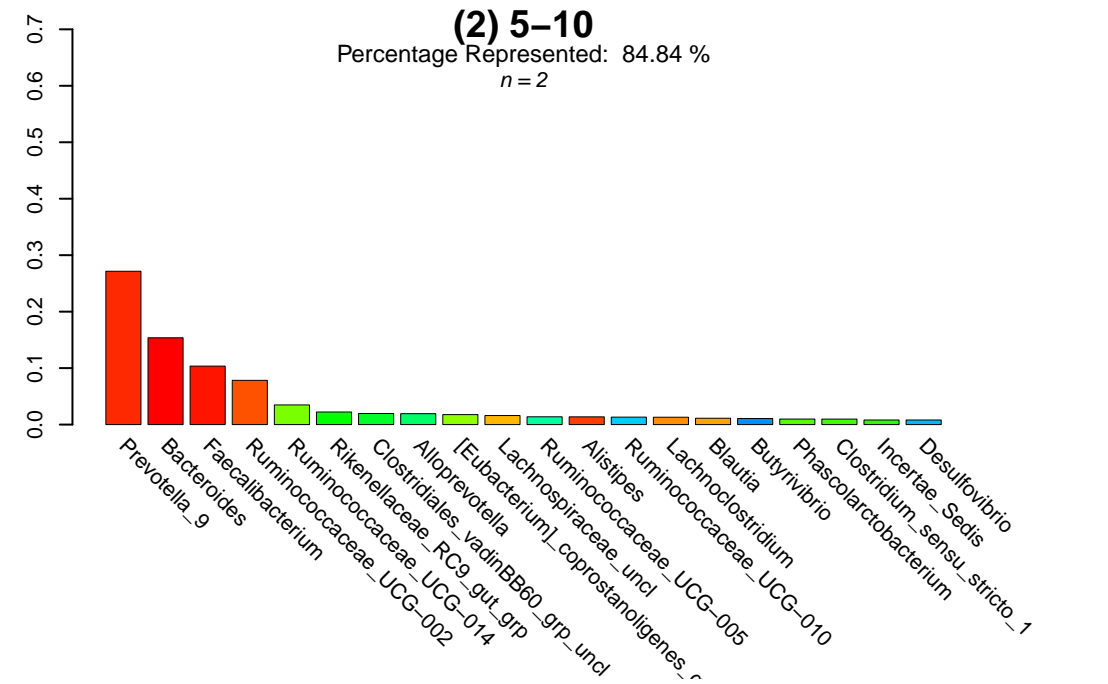
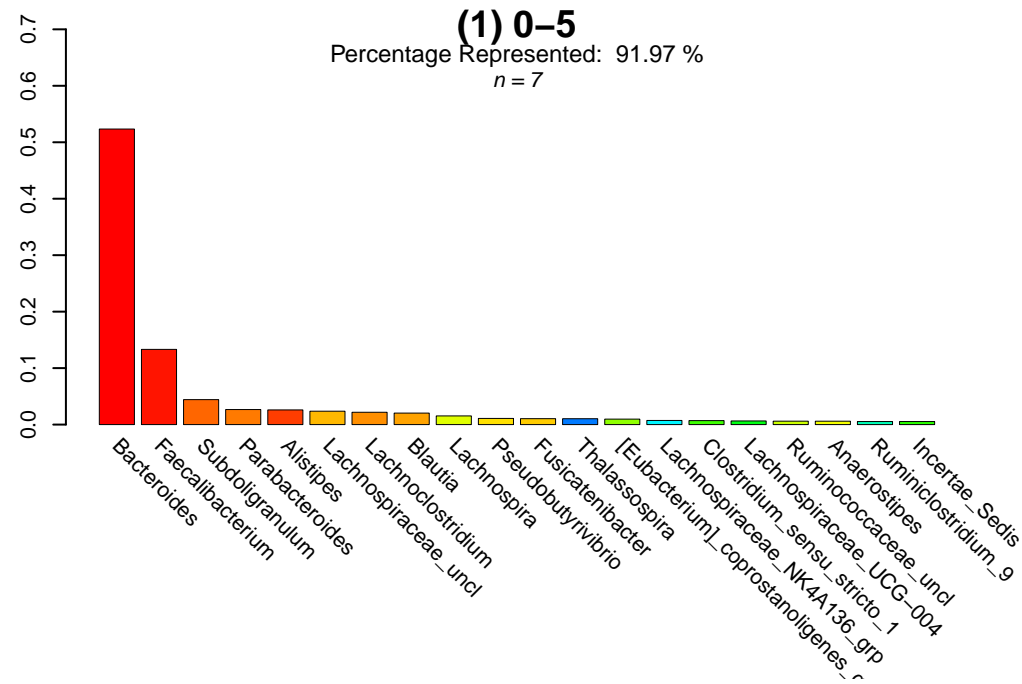
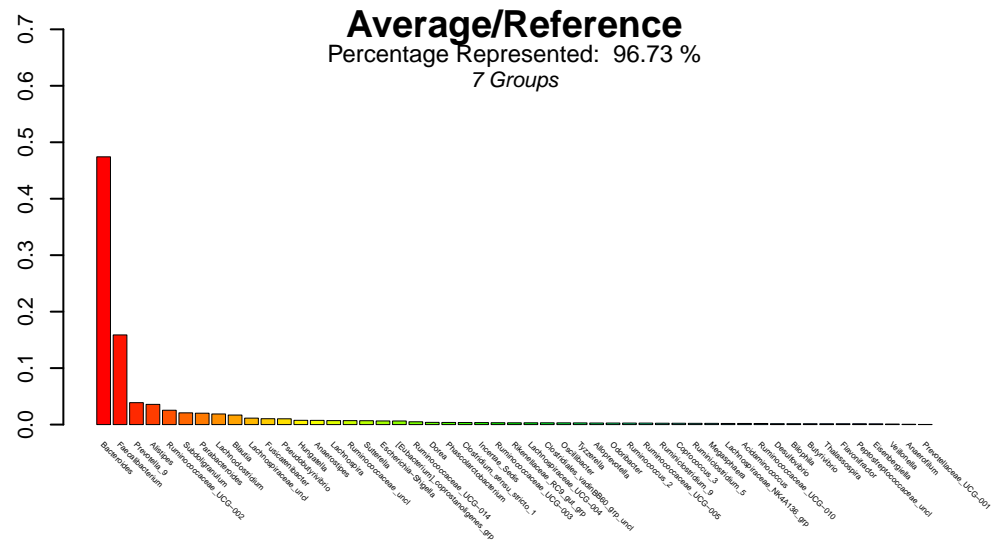
hadExch



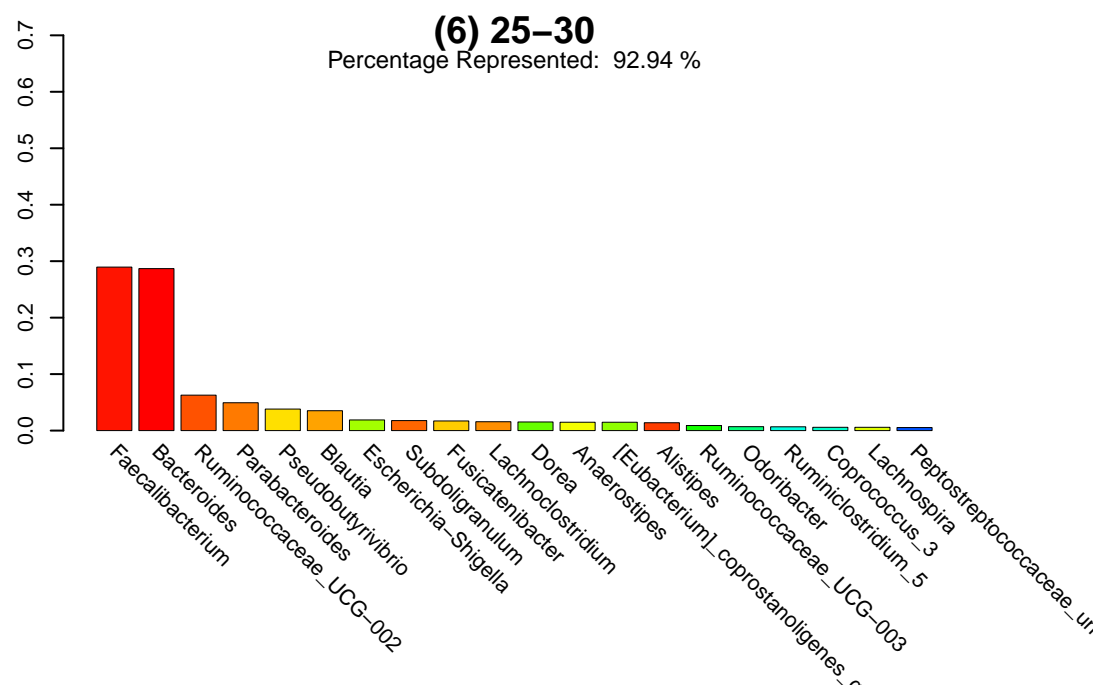
onPCN



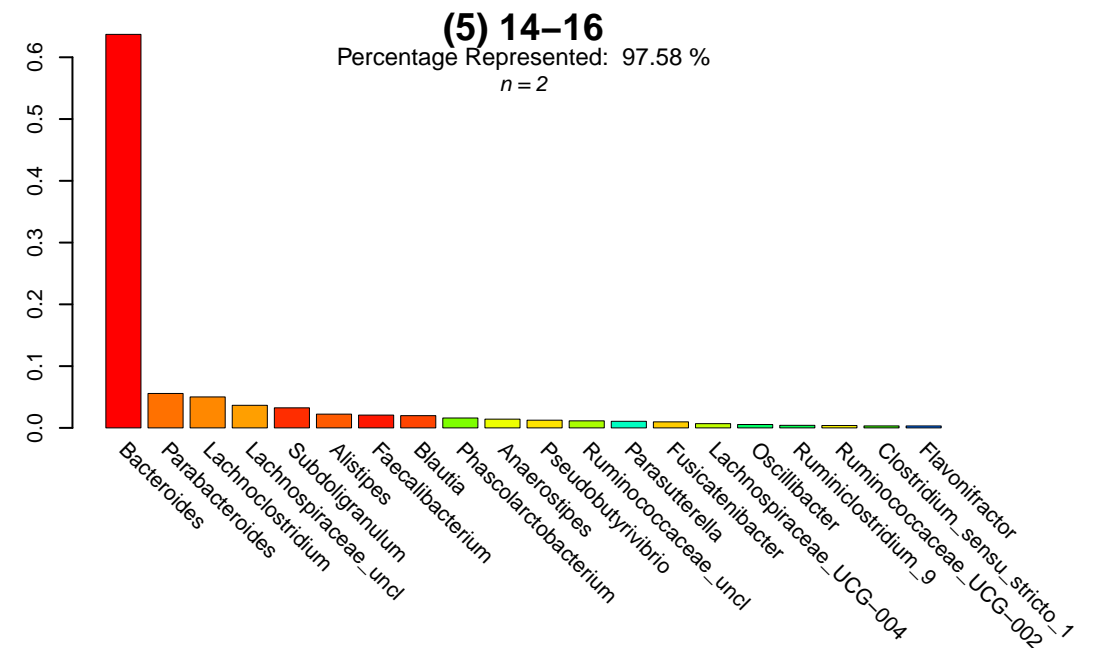
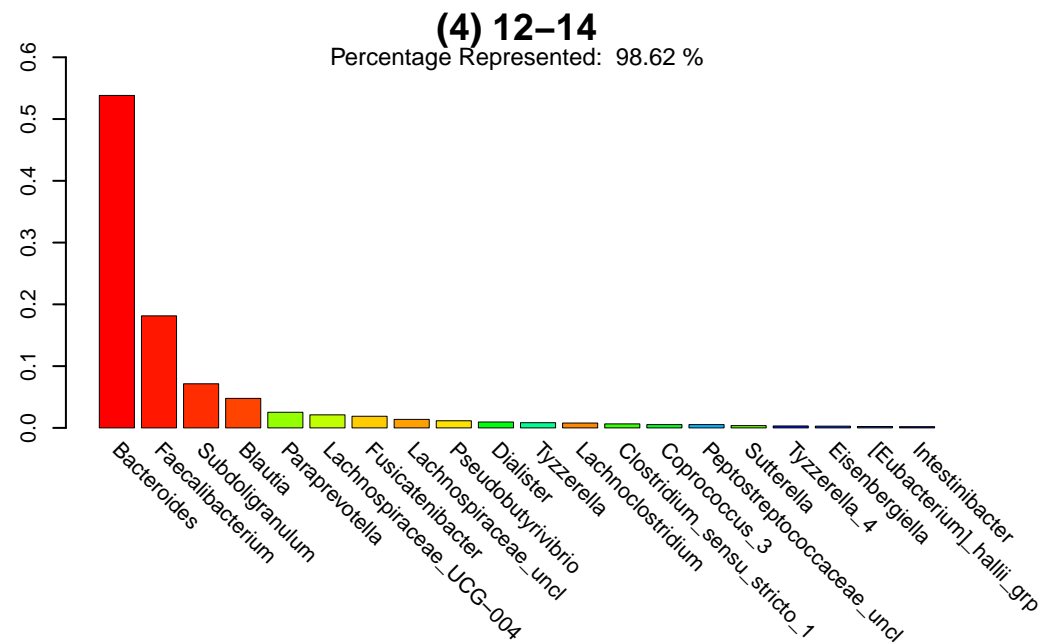
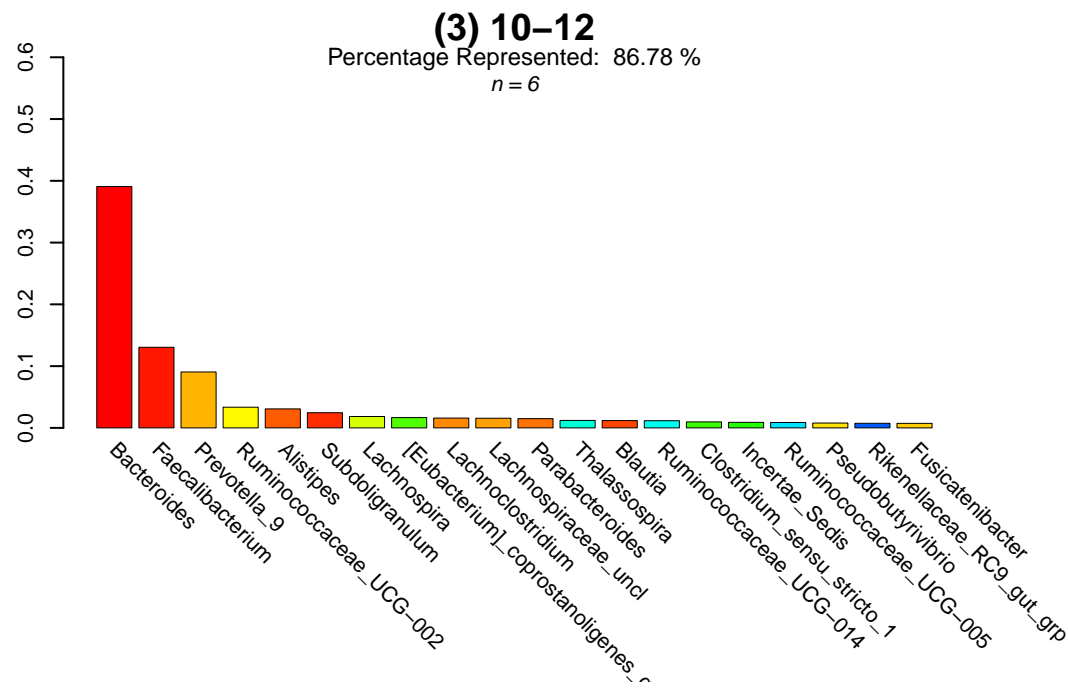
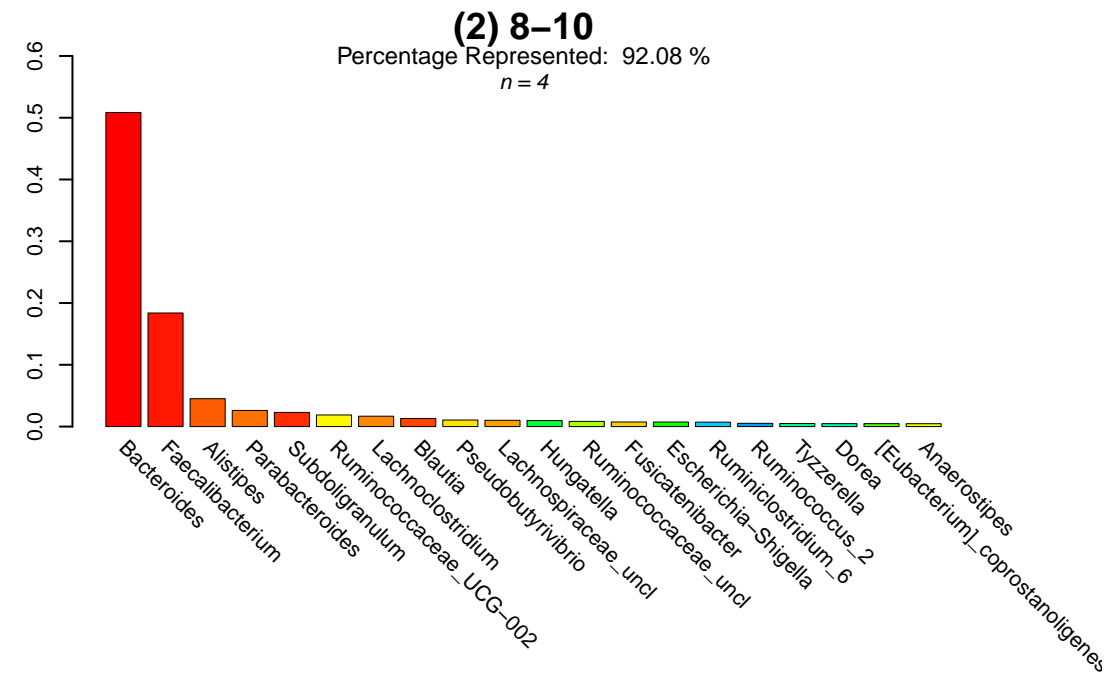
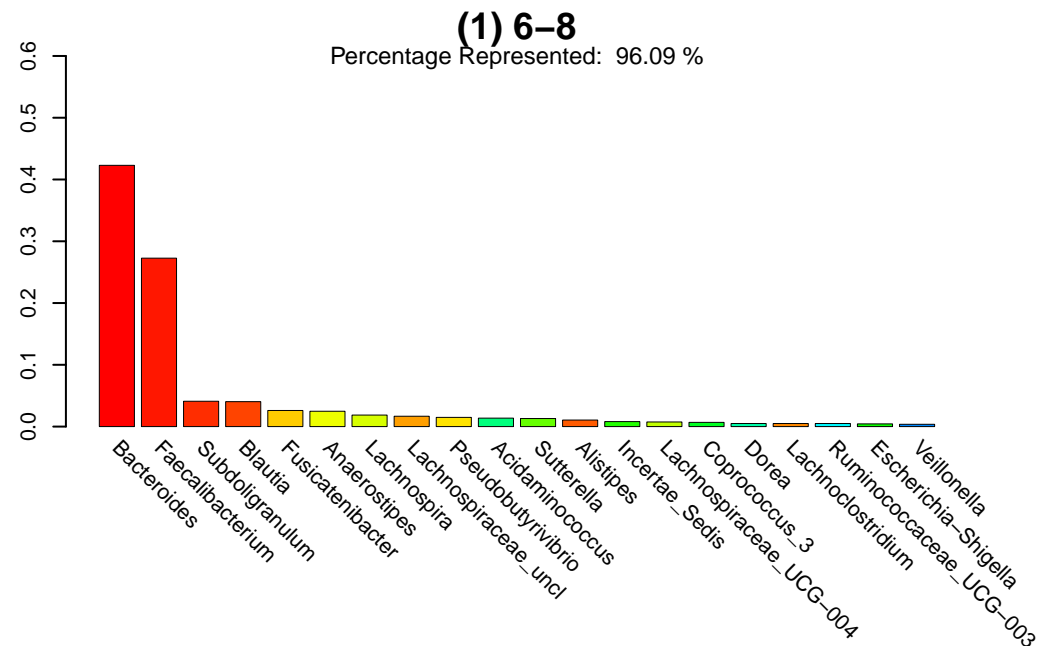
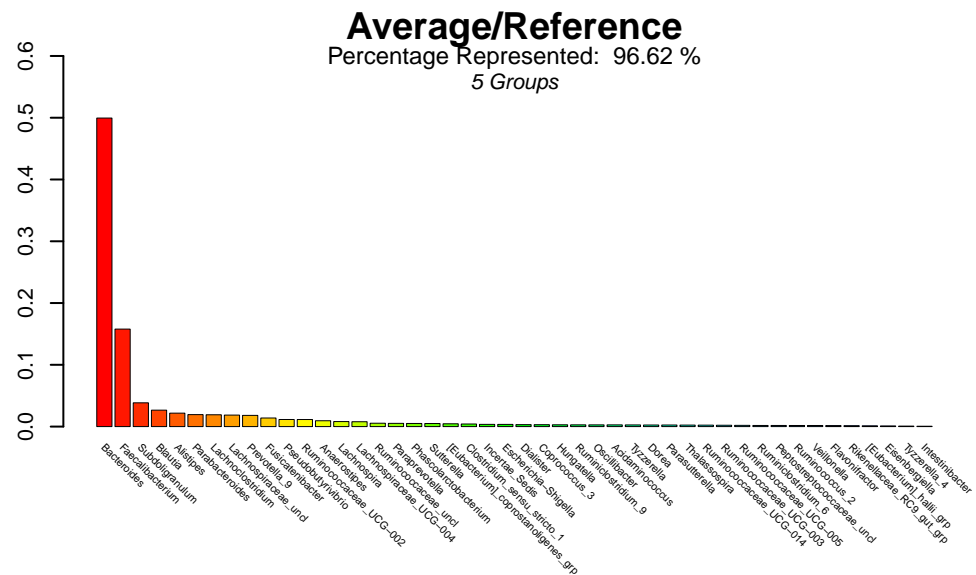
# HbF



# HbF

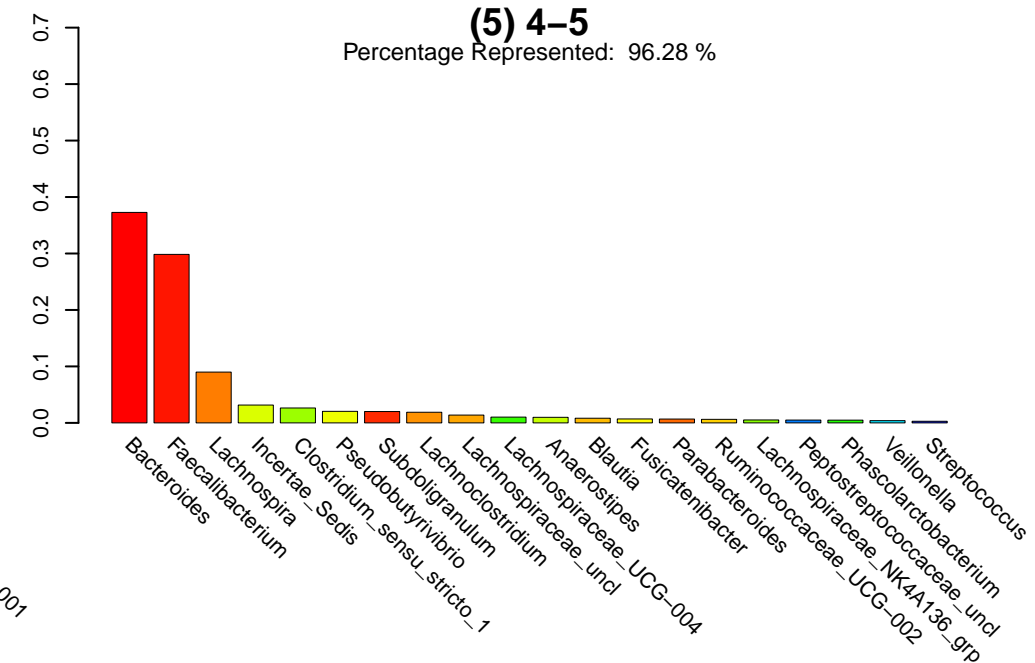
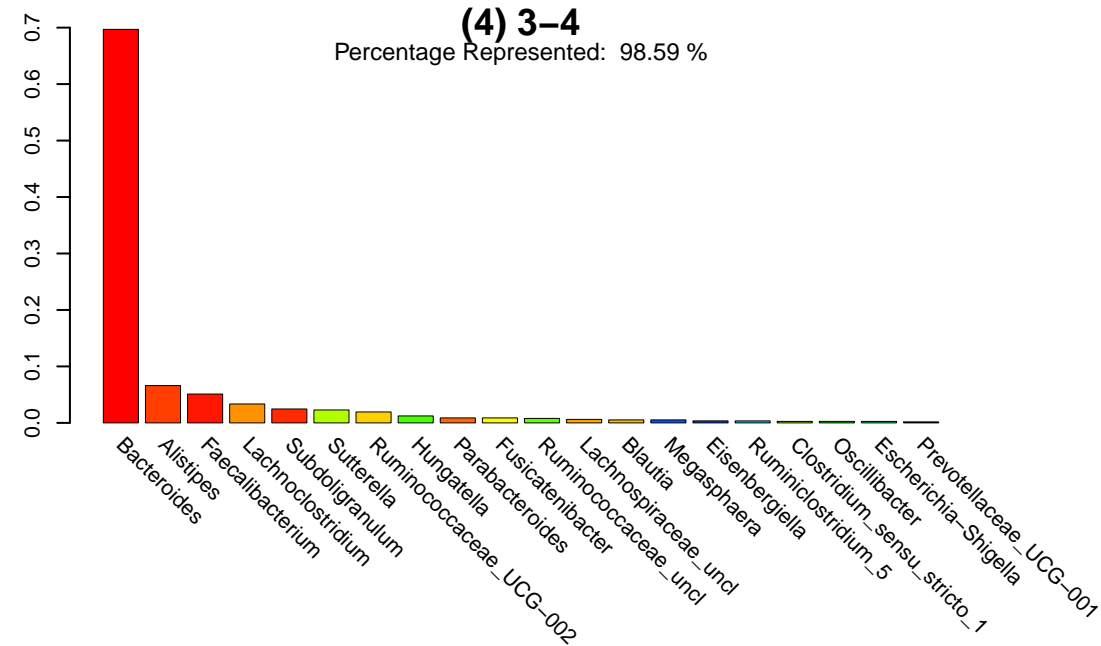
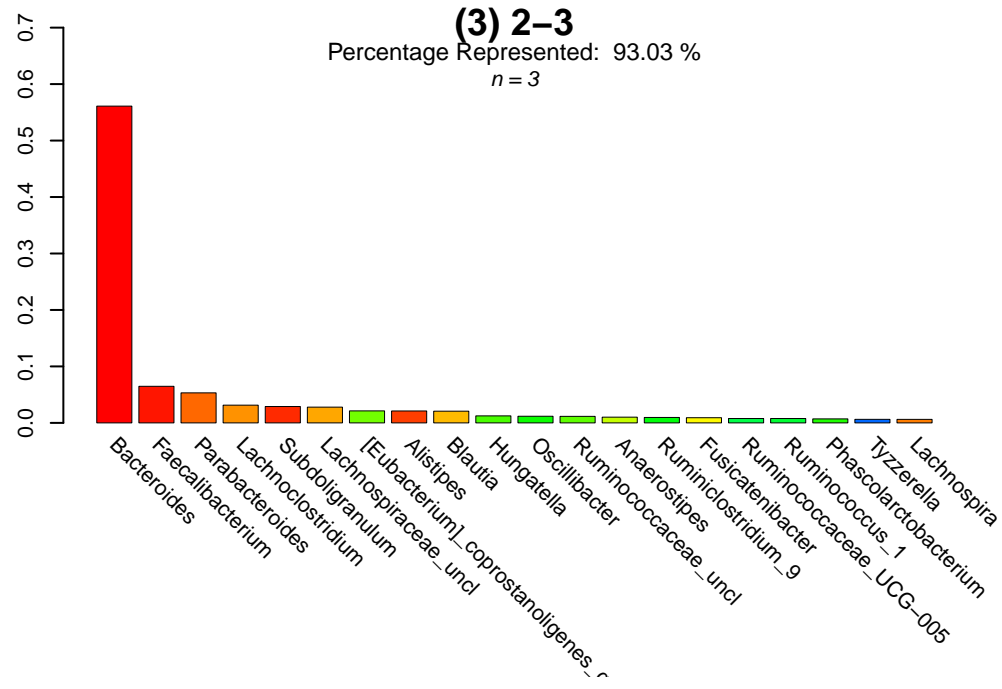
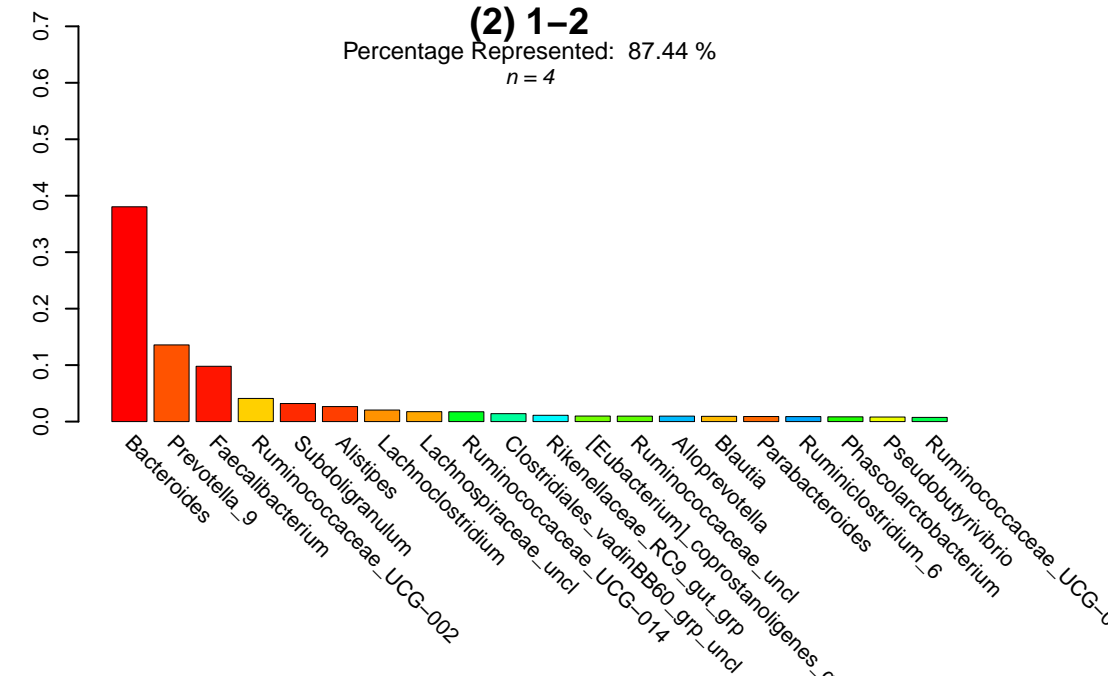
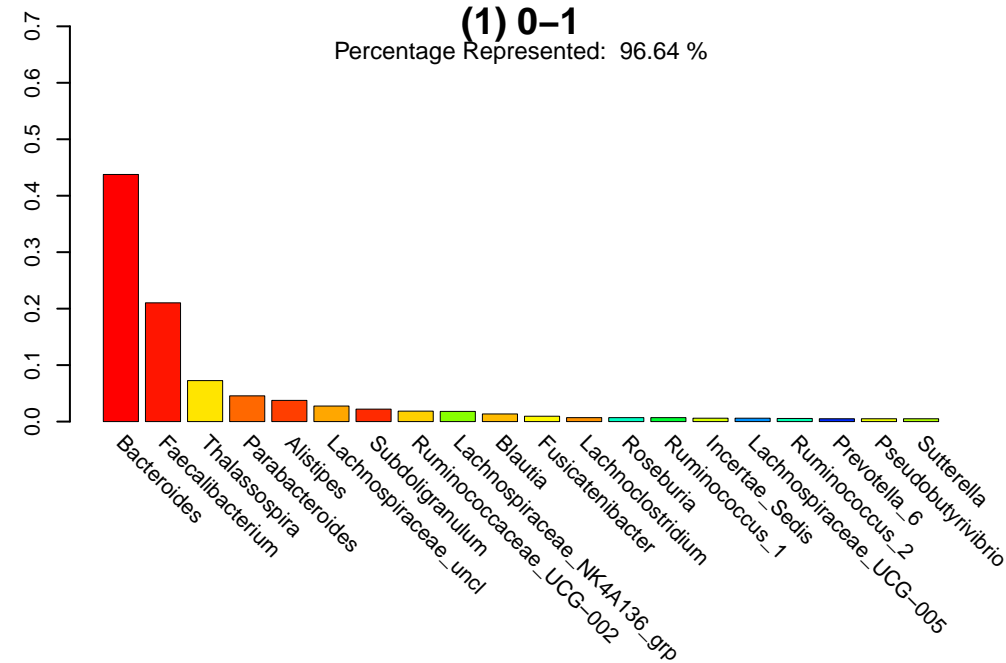
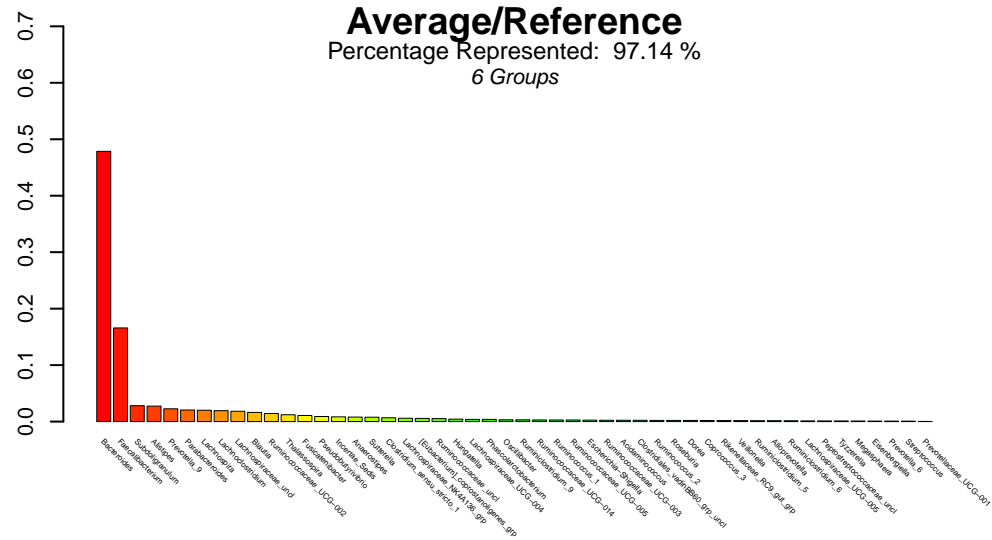


# Hb

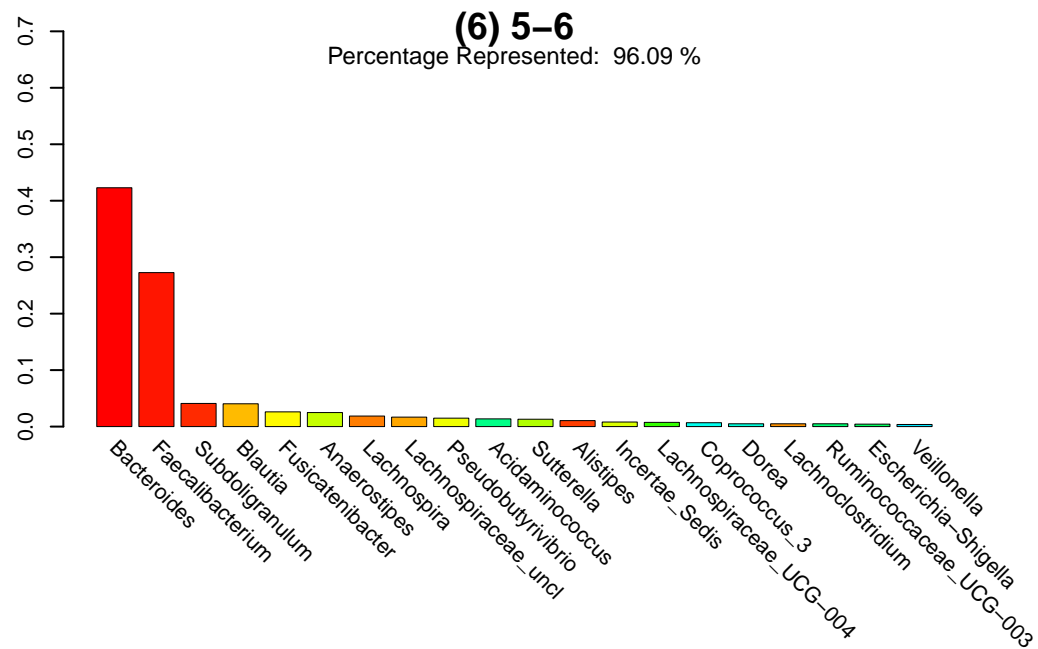




# Bilirubin



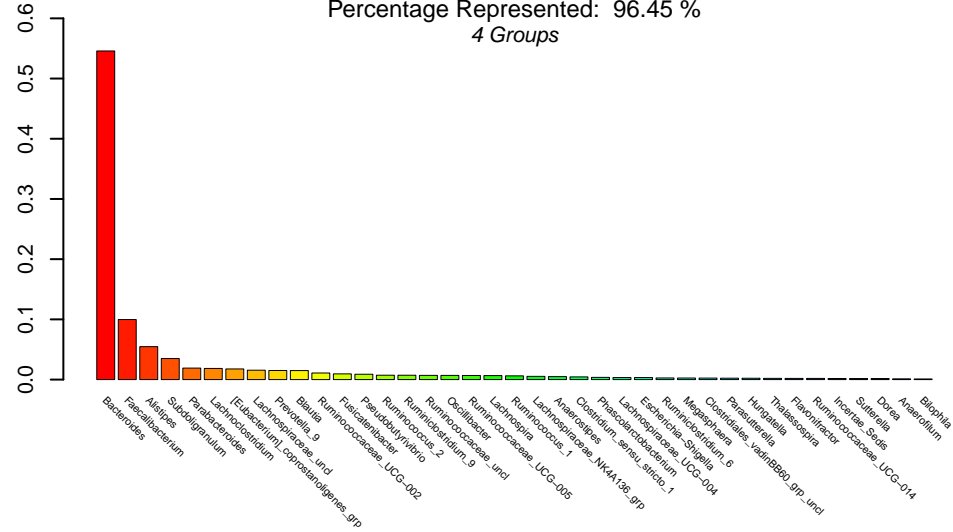
# Bilirubin



# Admit

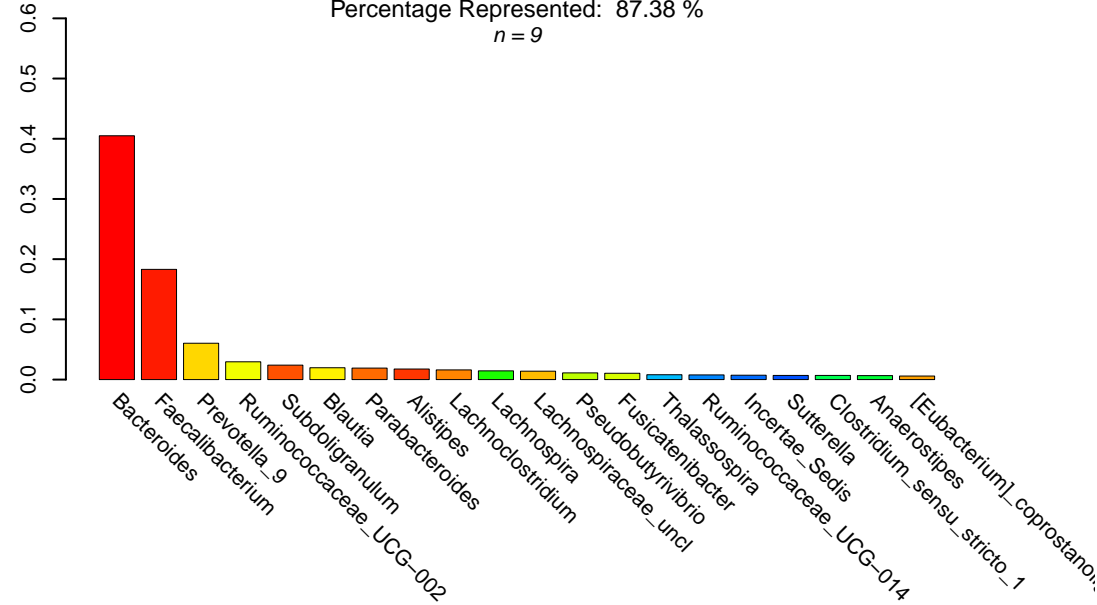
## Average/Reference

Percentage Represented: 96.45 %  
4 Groups



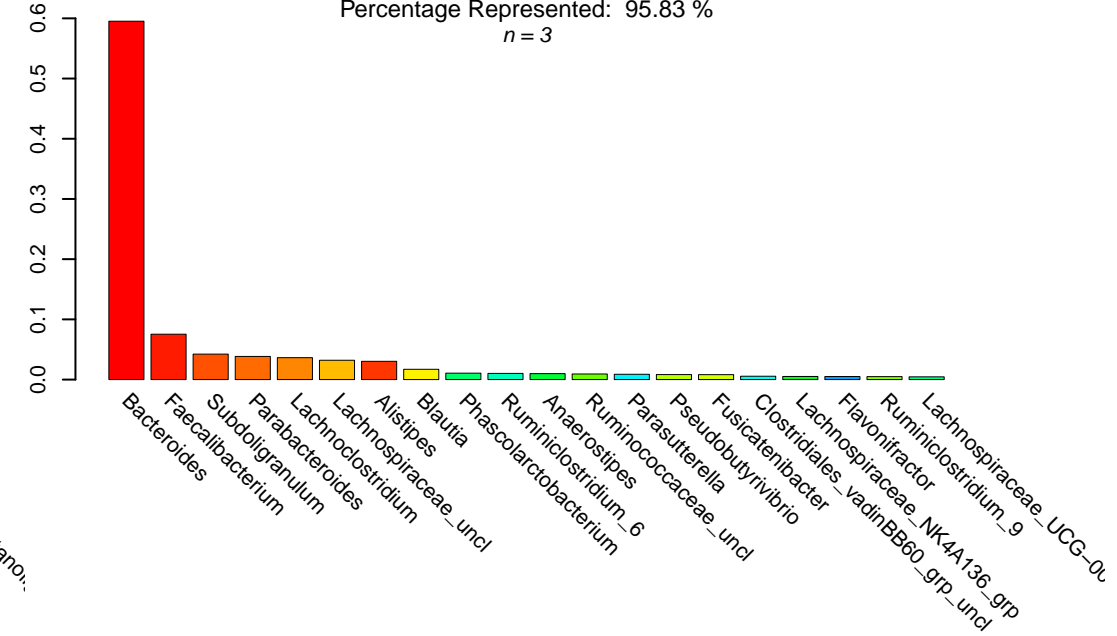
## (1) 0-2

Percentage Represented: 87.38 %  
n = 9



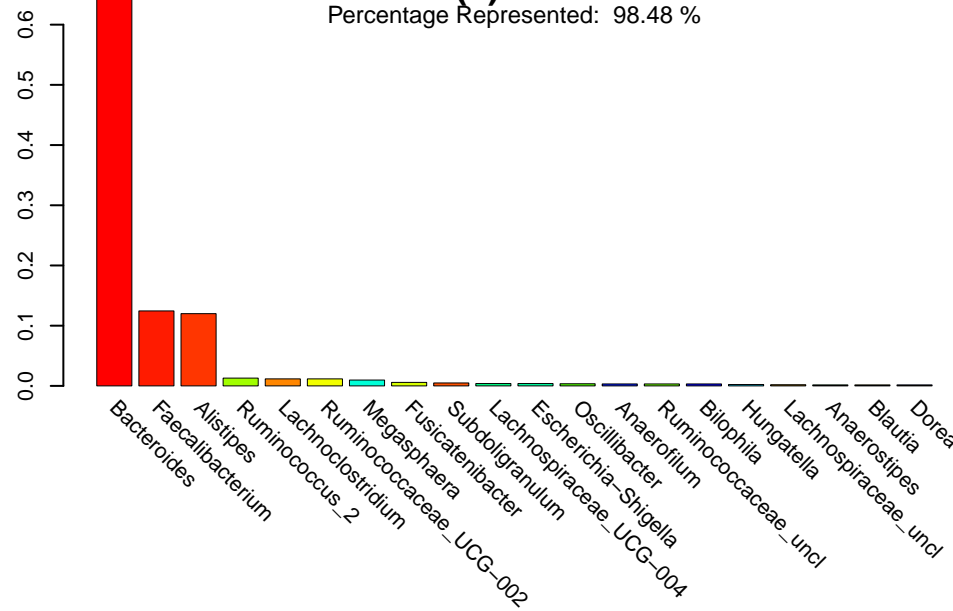
## (2) 2-4

Percentage Represented: 95.83 %  
n = 3



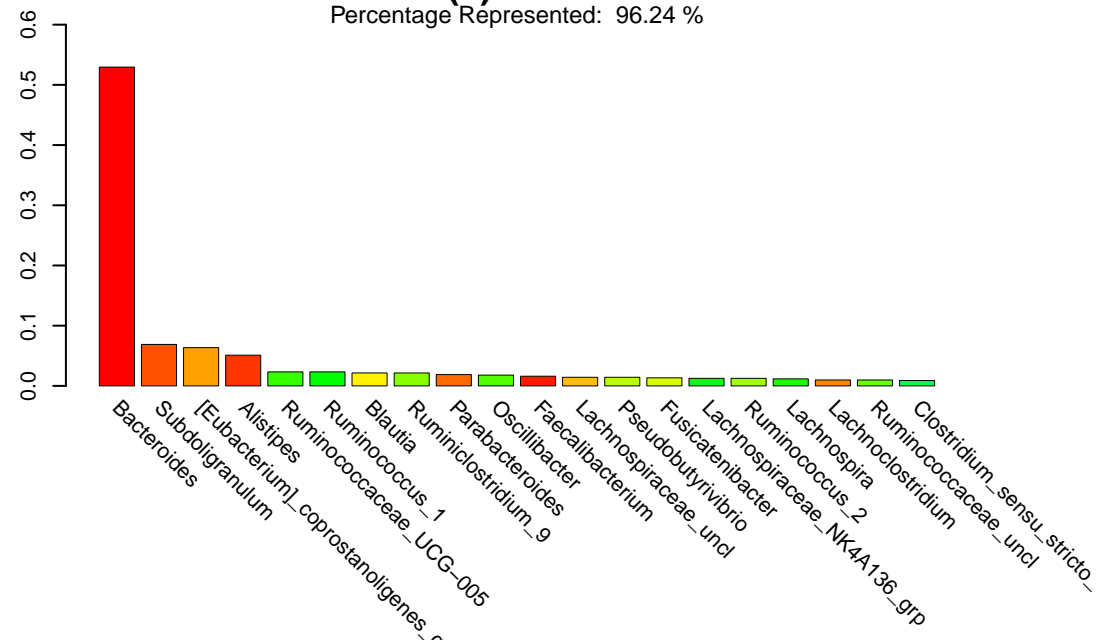
## (3) 4-6

Percentage Represented: 98.48 %

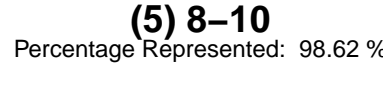
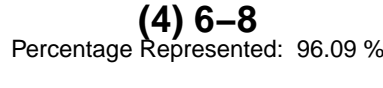
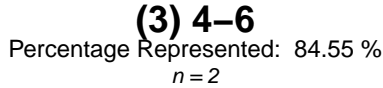
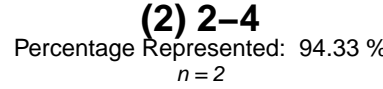
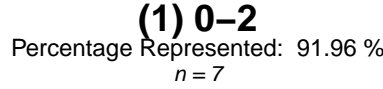


## (5) 8-10

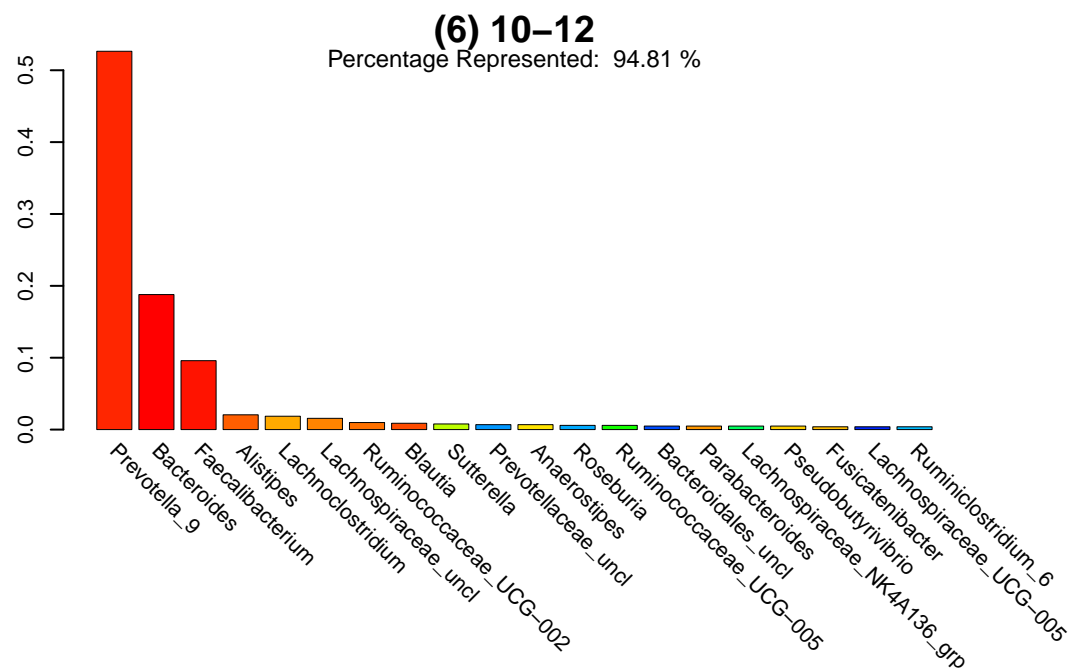
Percentage Represented: 96.24 %



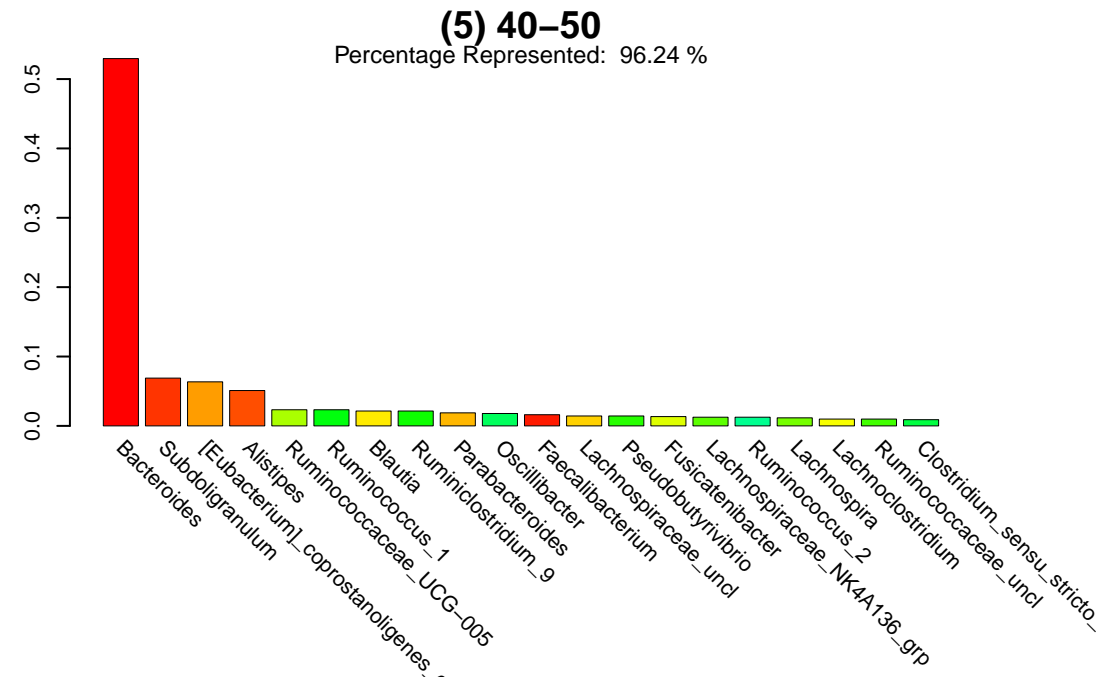
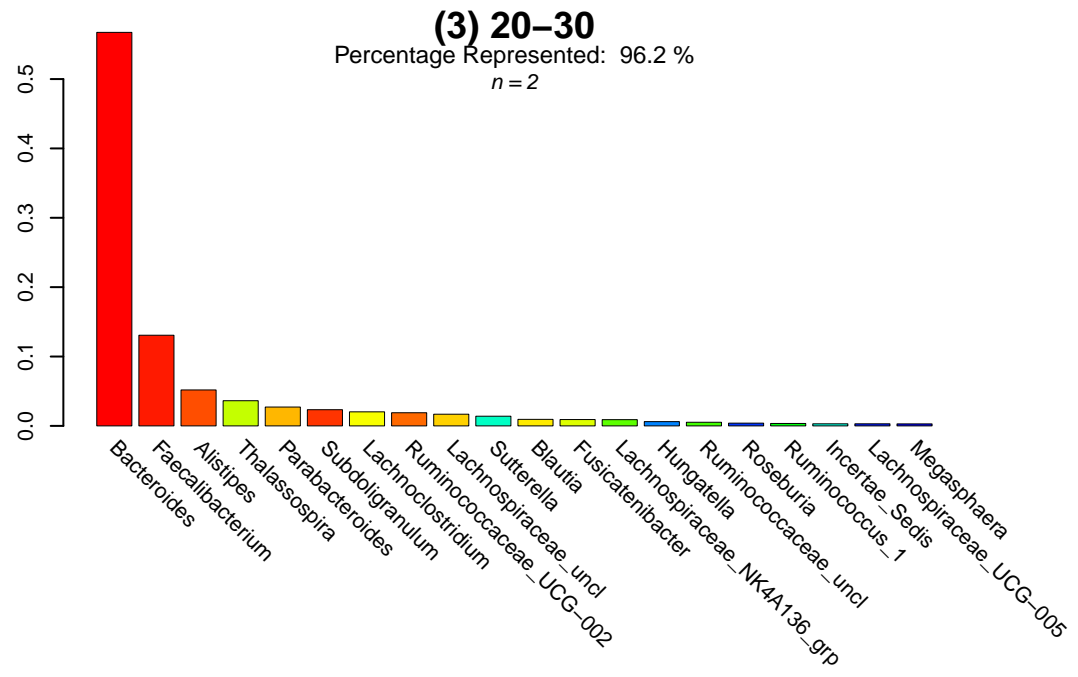
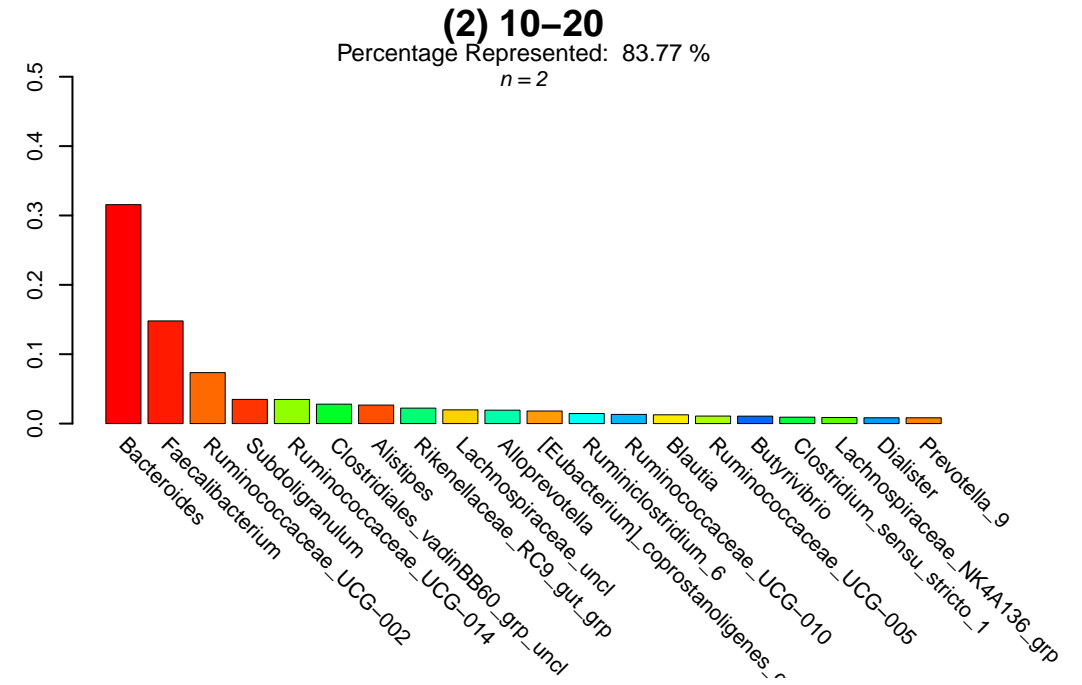
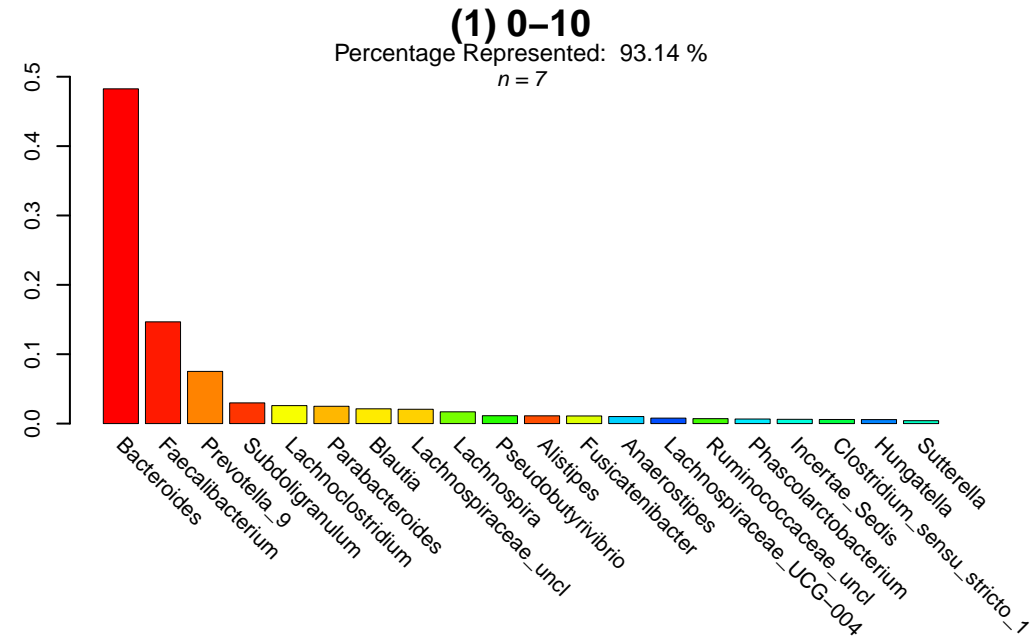
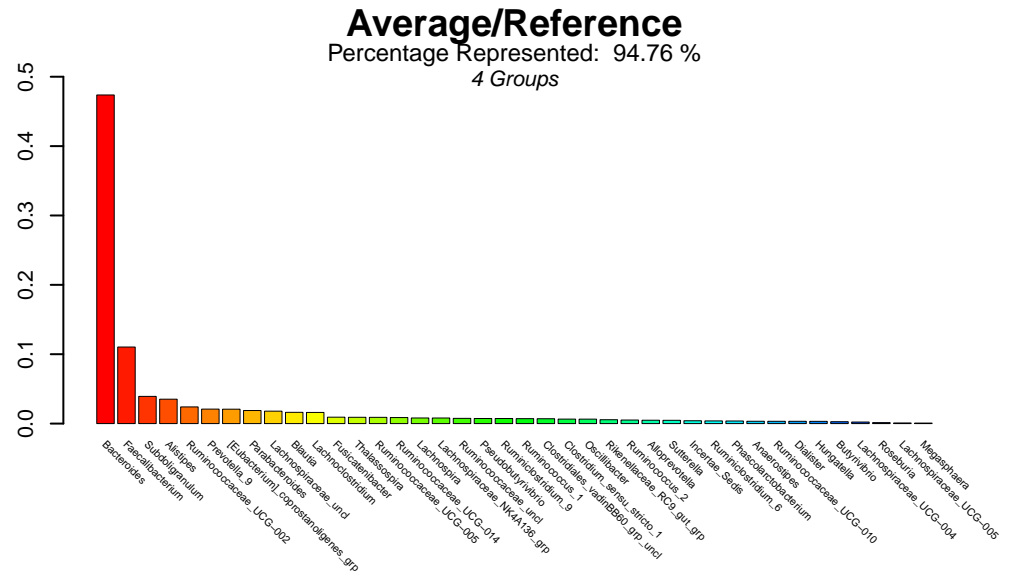
# LastVOC



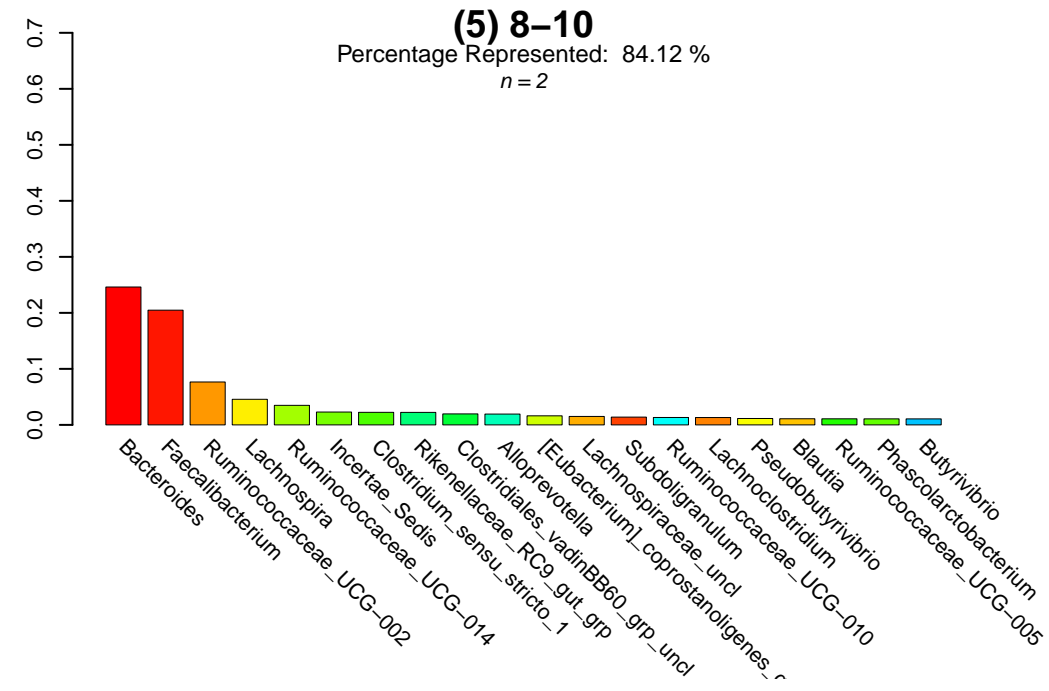
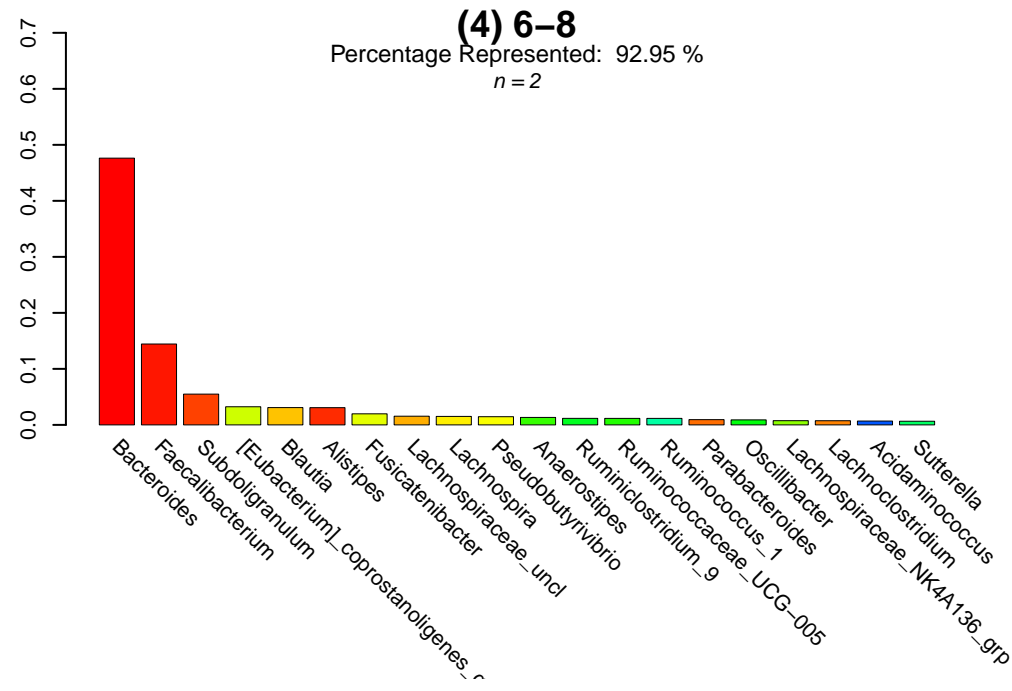
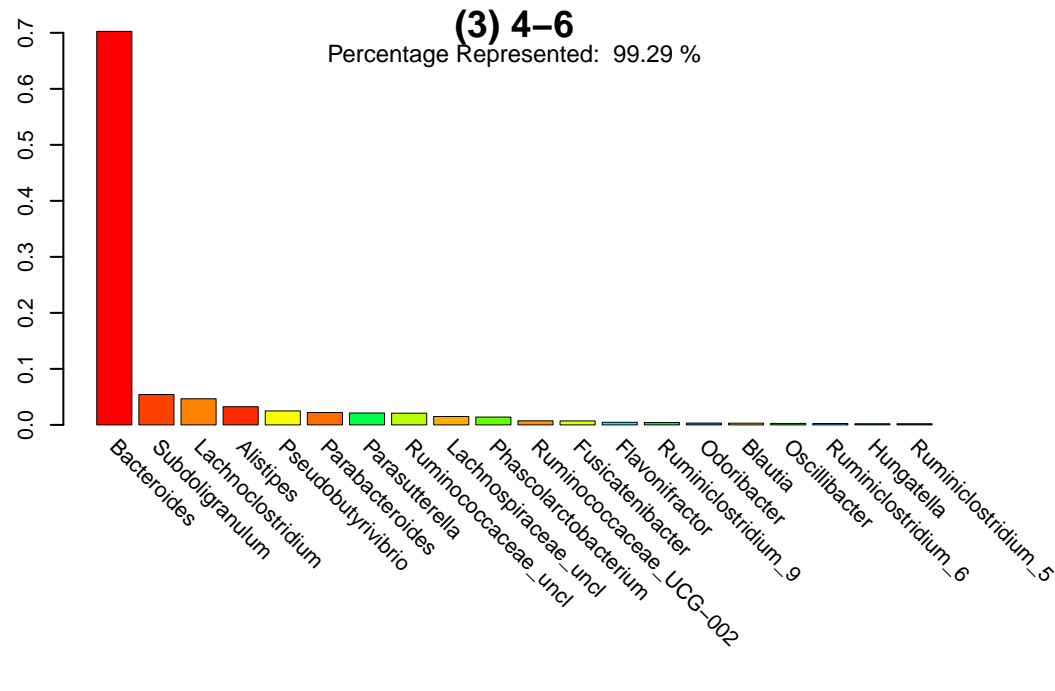
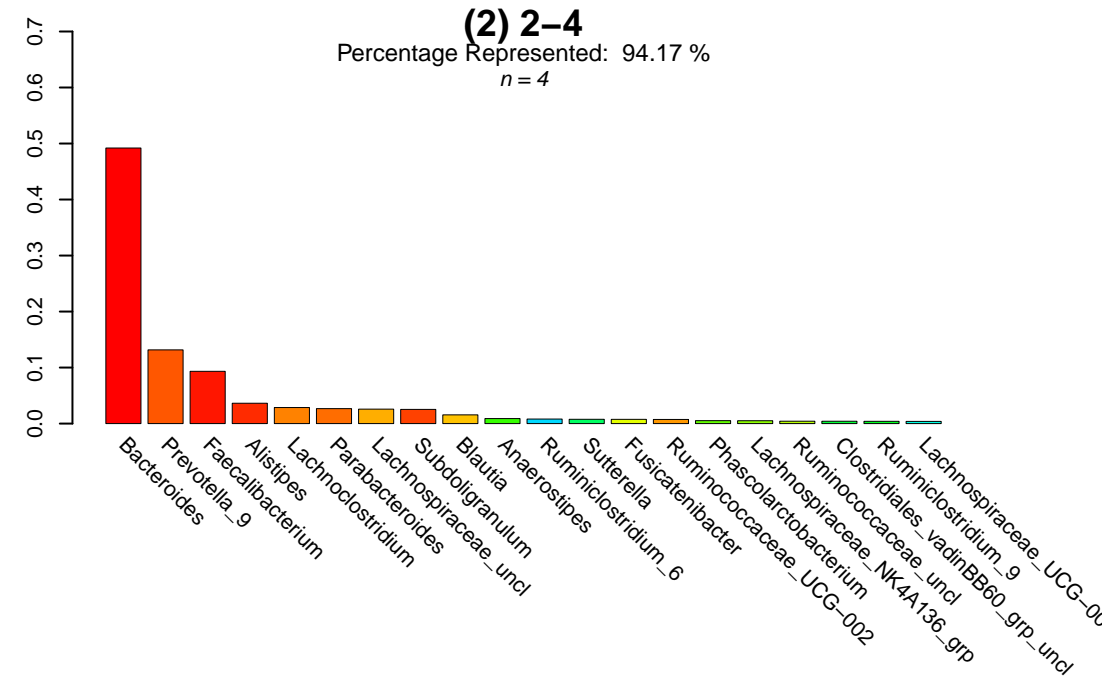
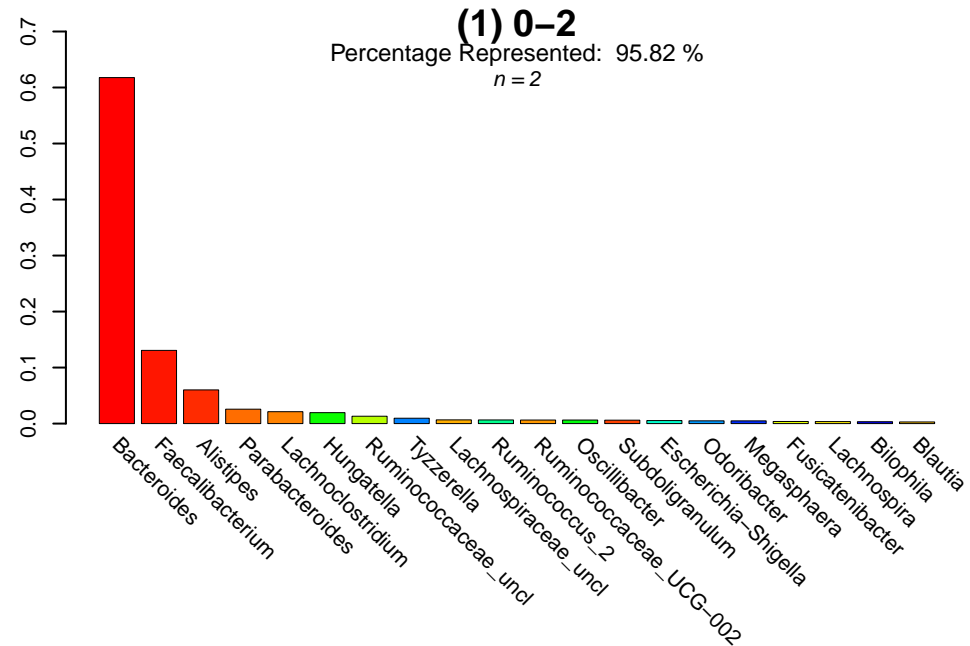
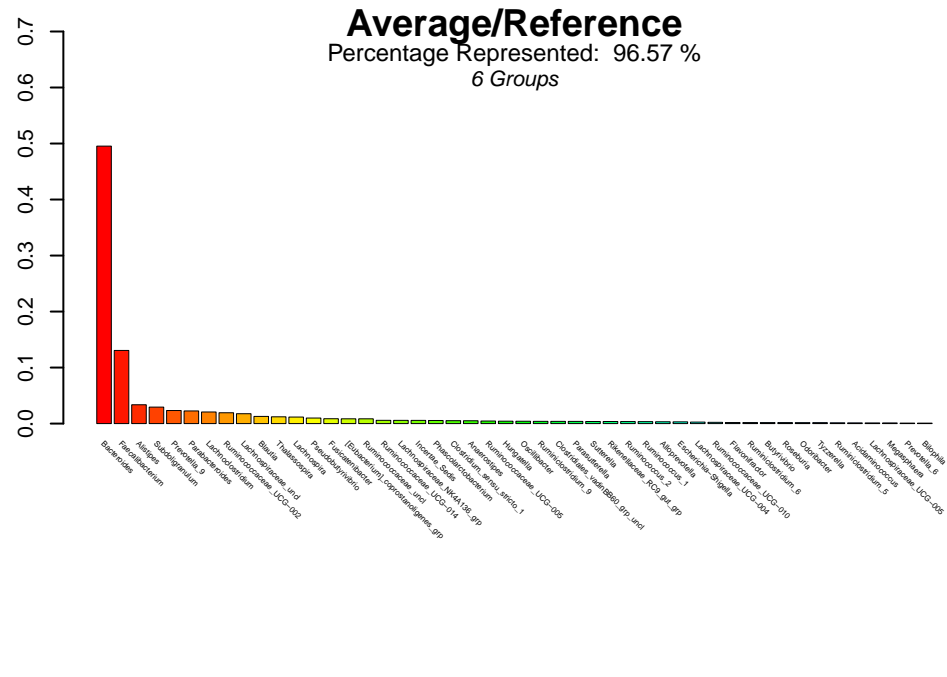
# LastVOC



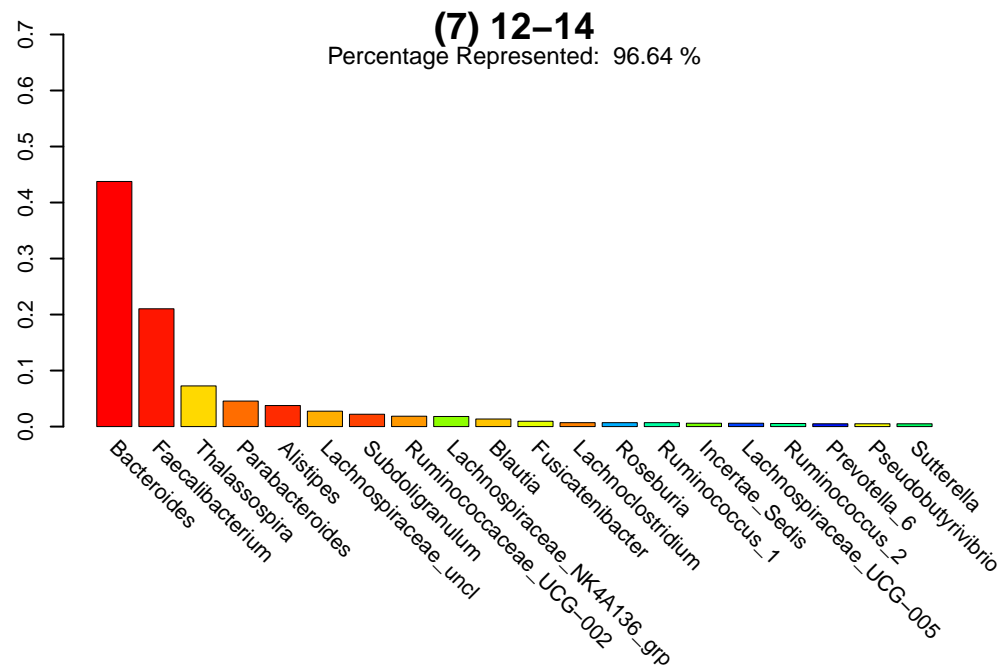
# ESR



# CRP

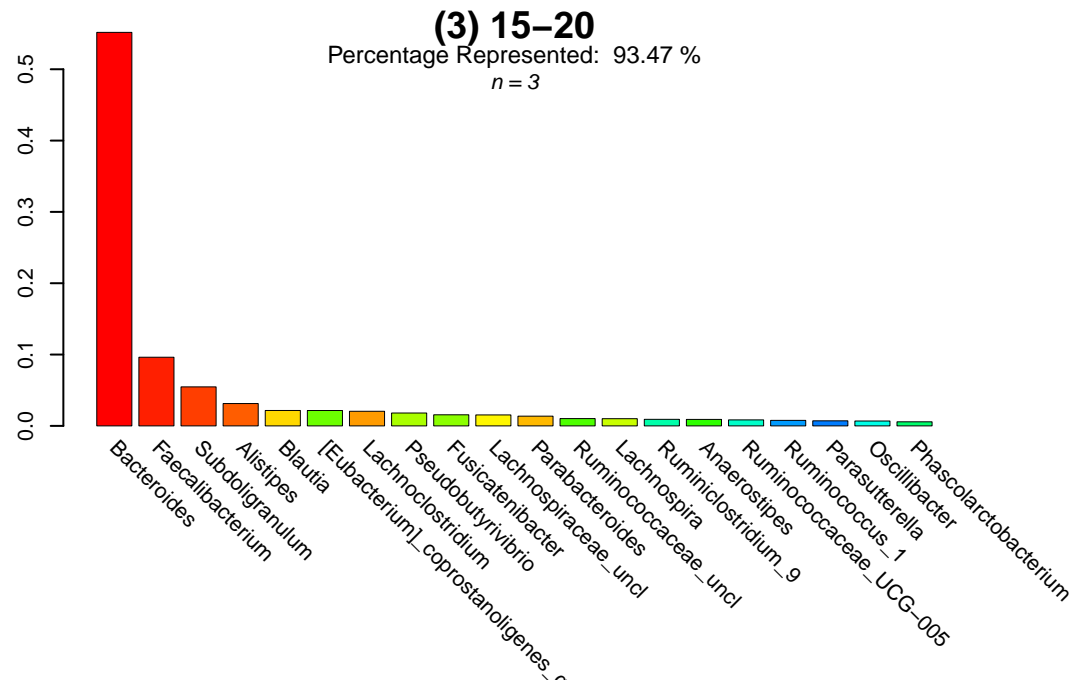
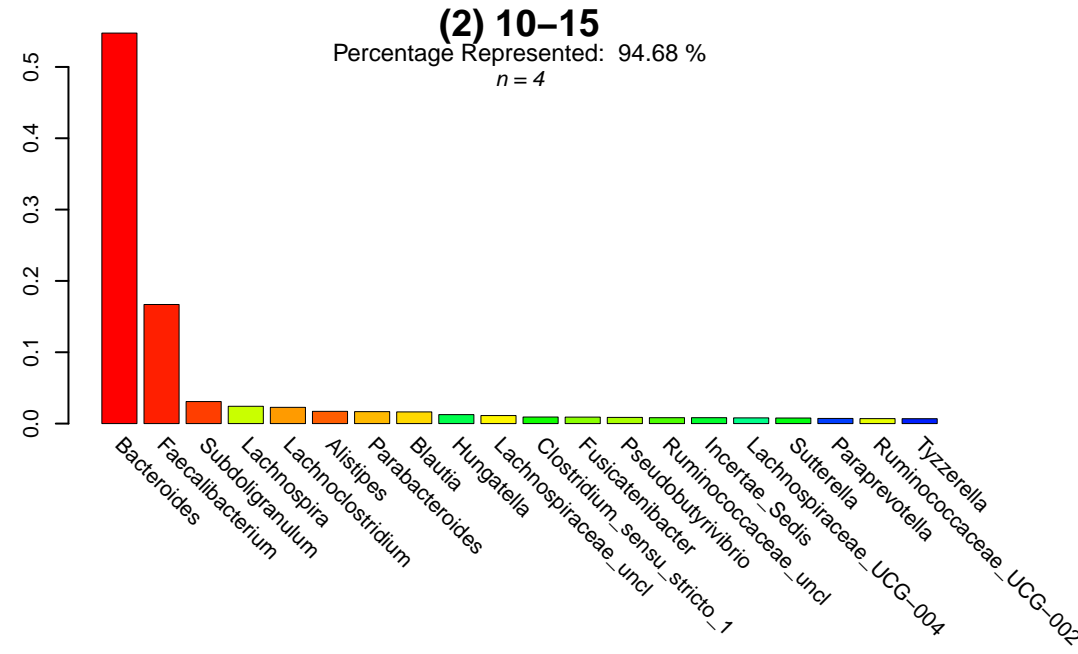
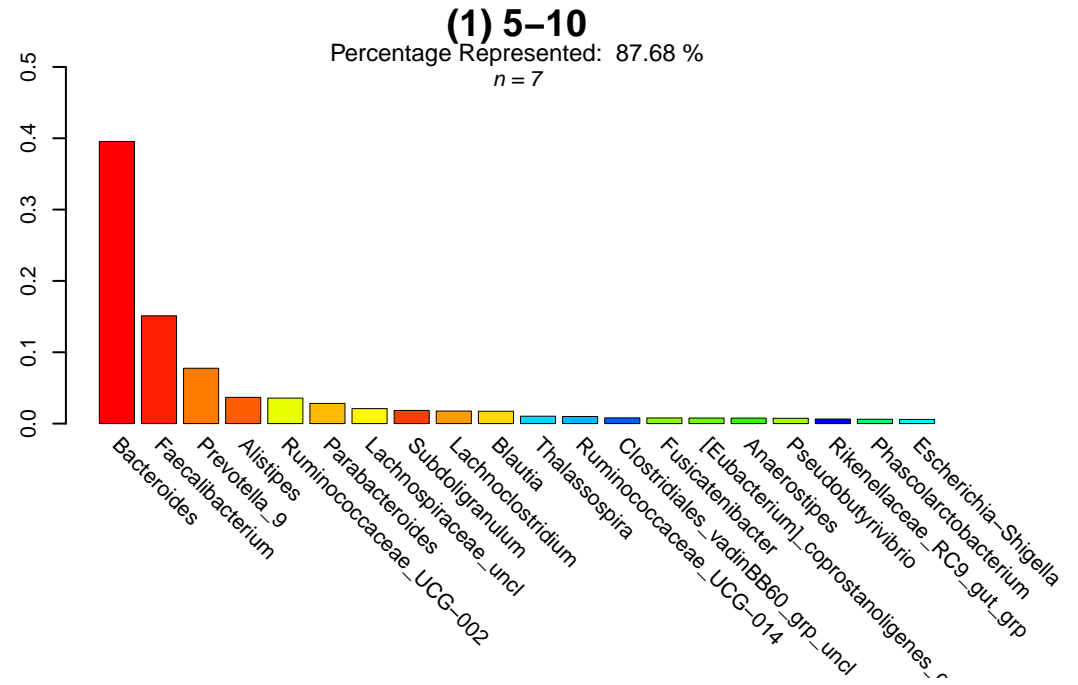
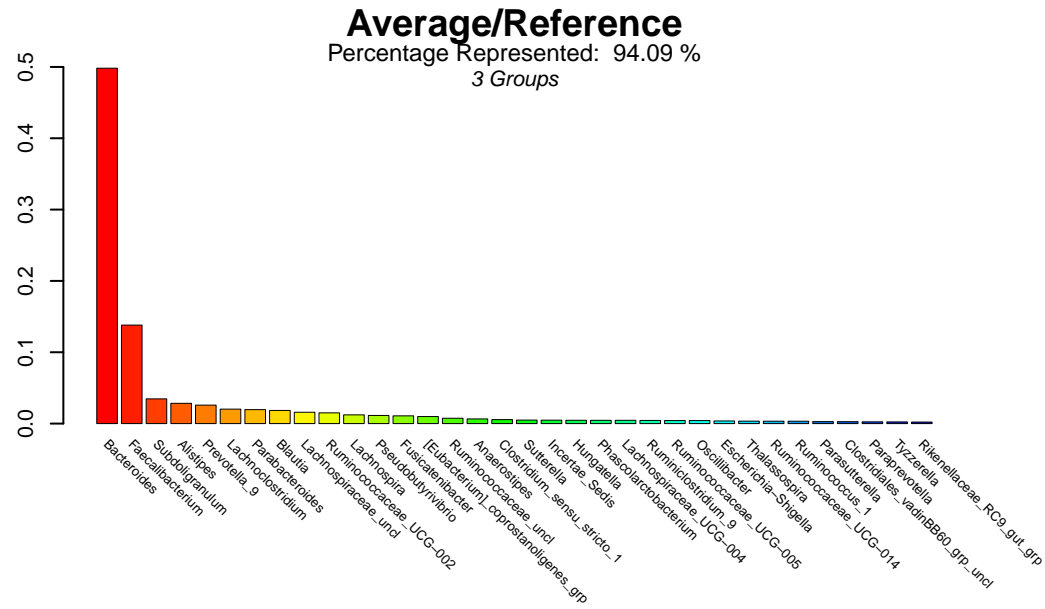


# CRP

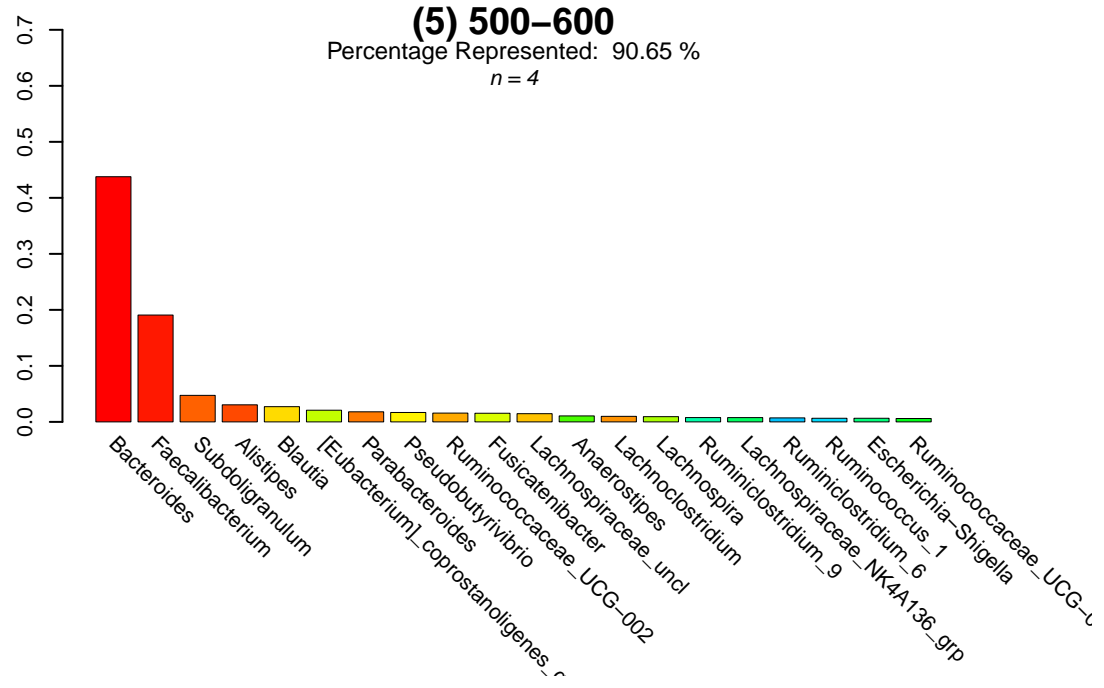
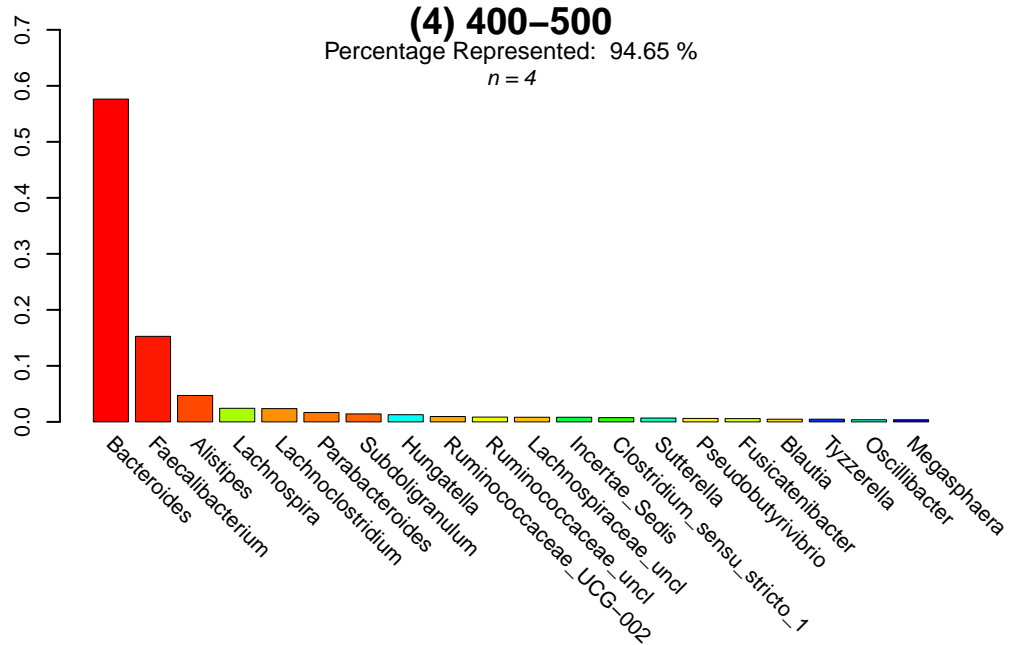
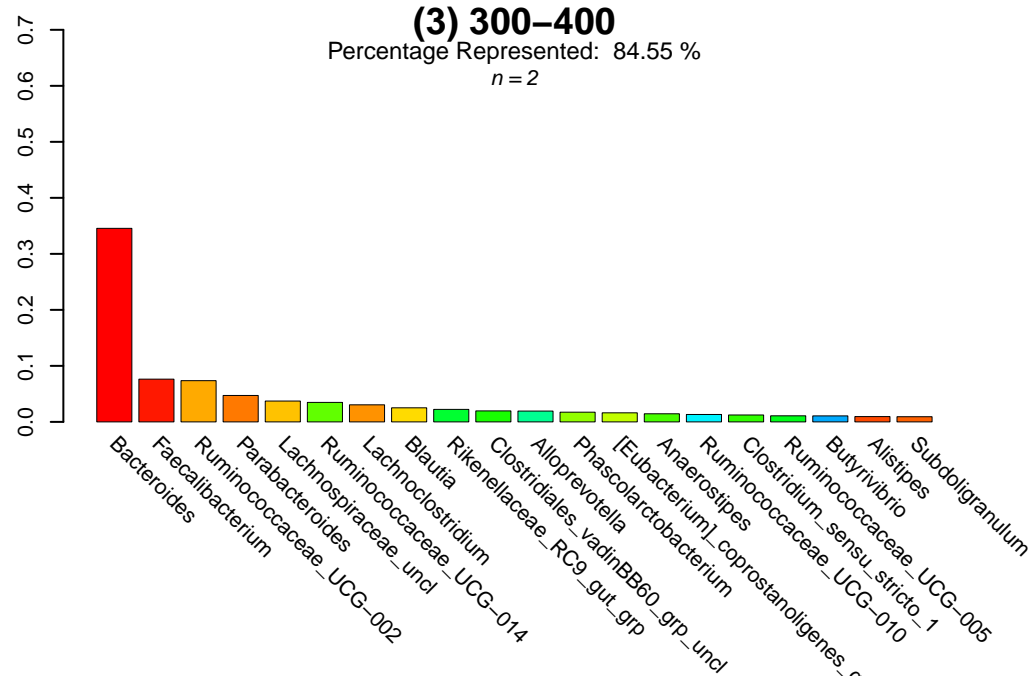
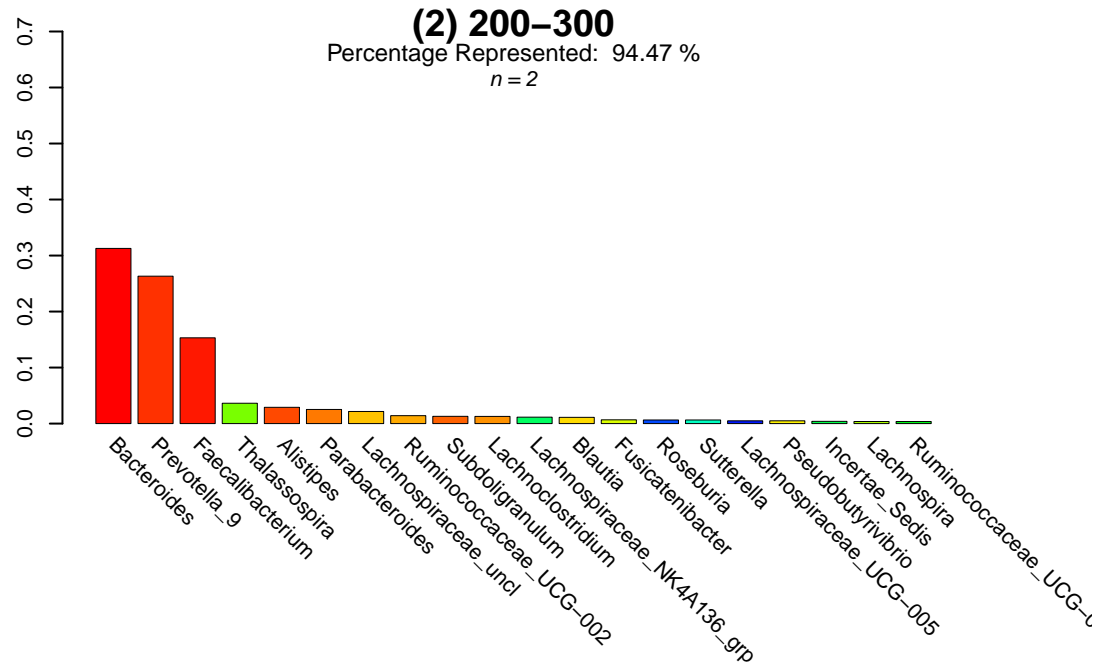
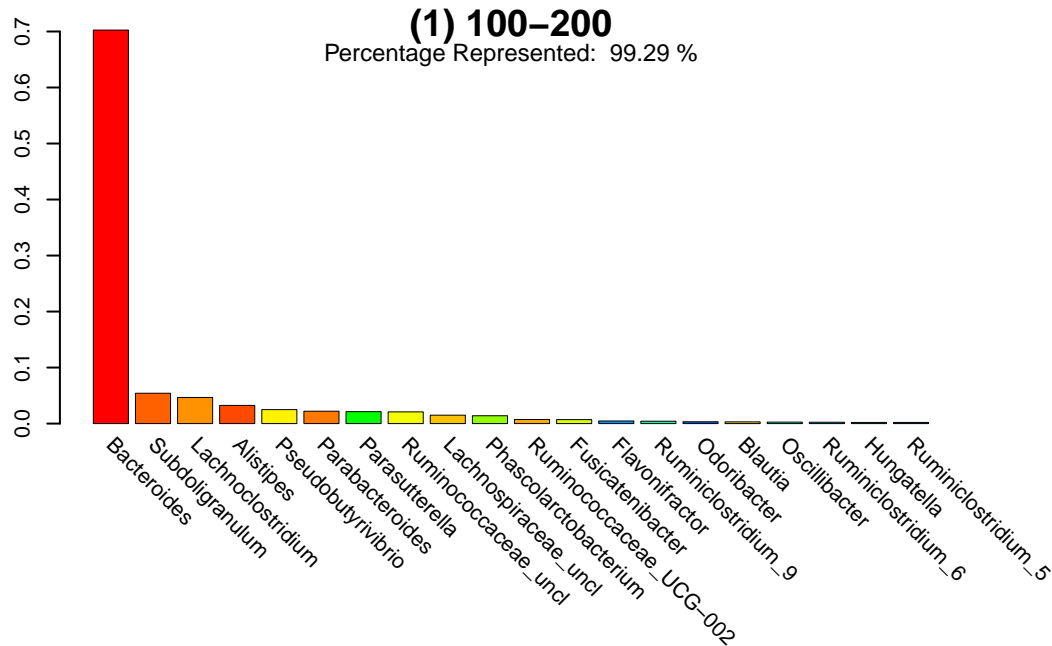
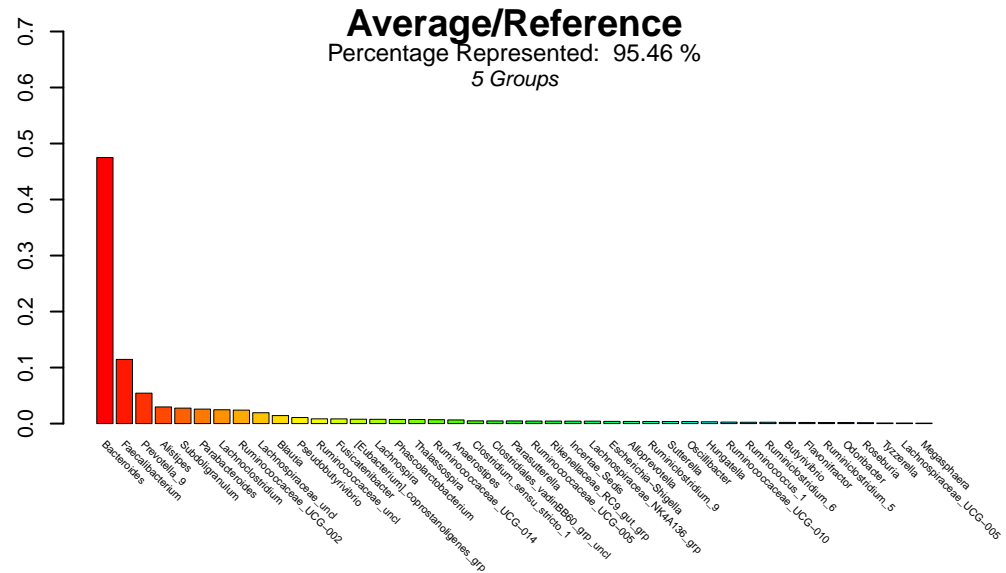




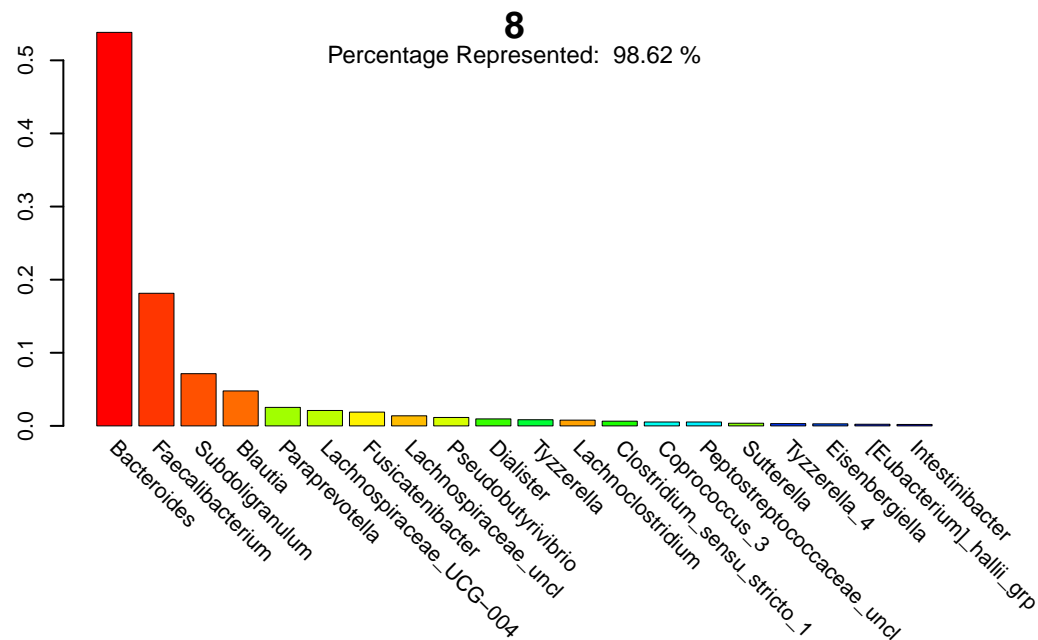
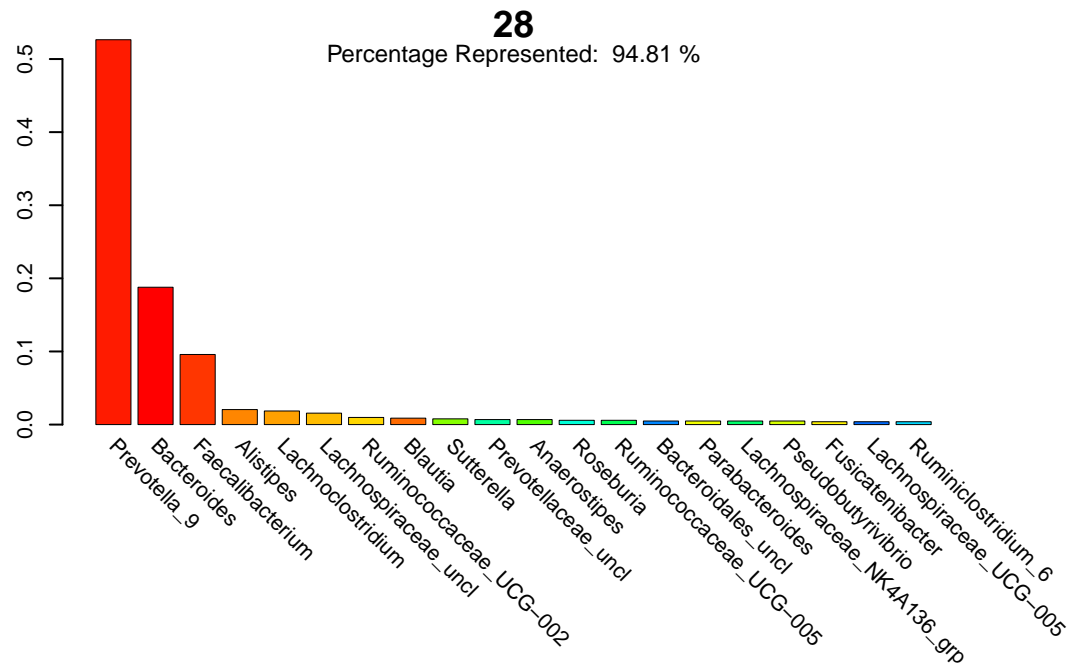
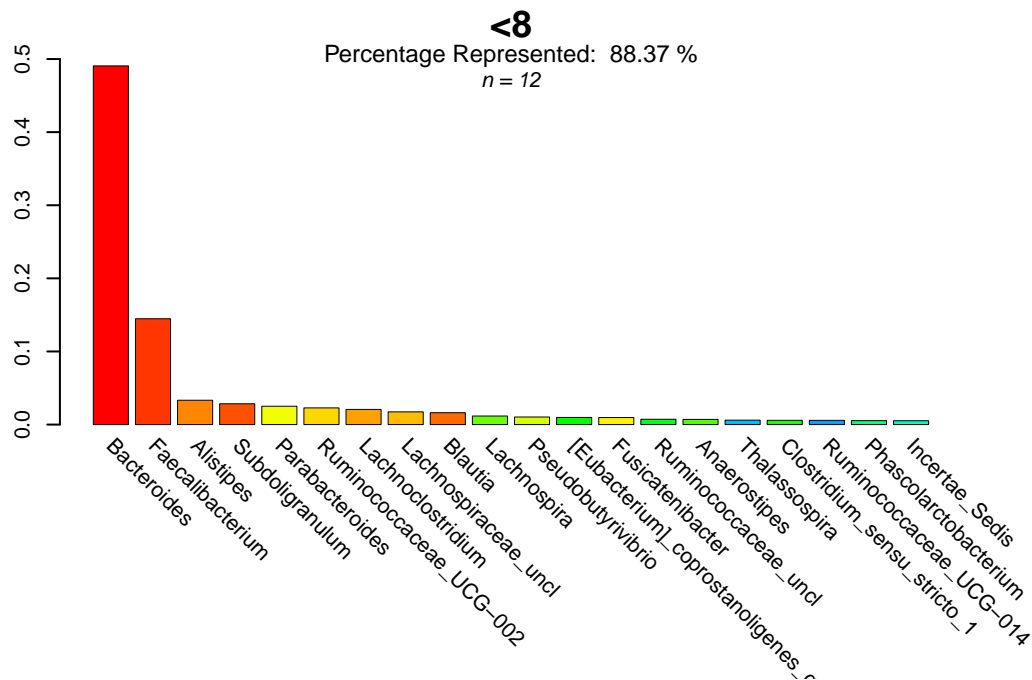
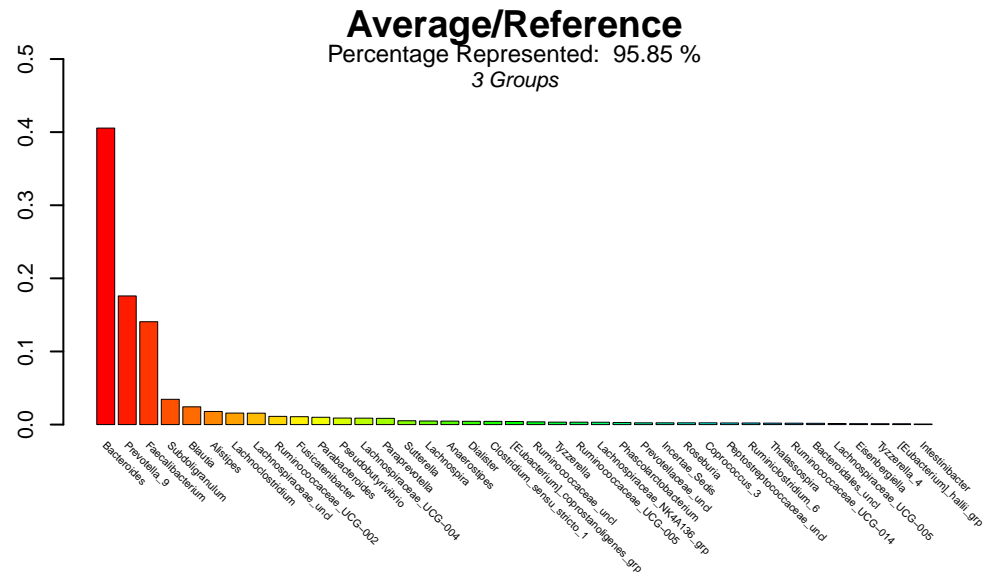
# WBC



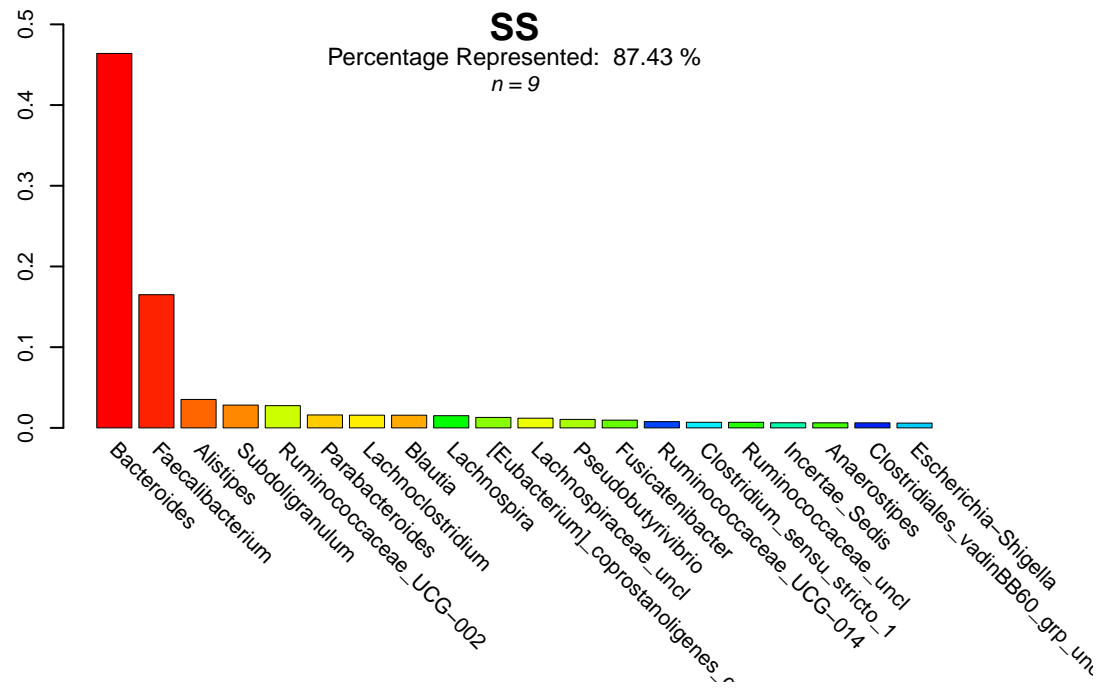
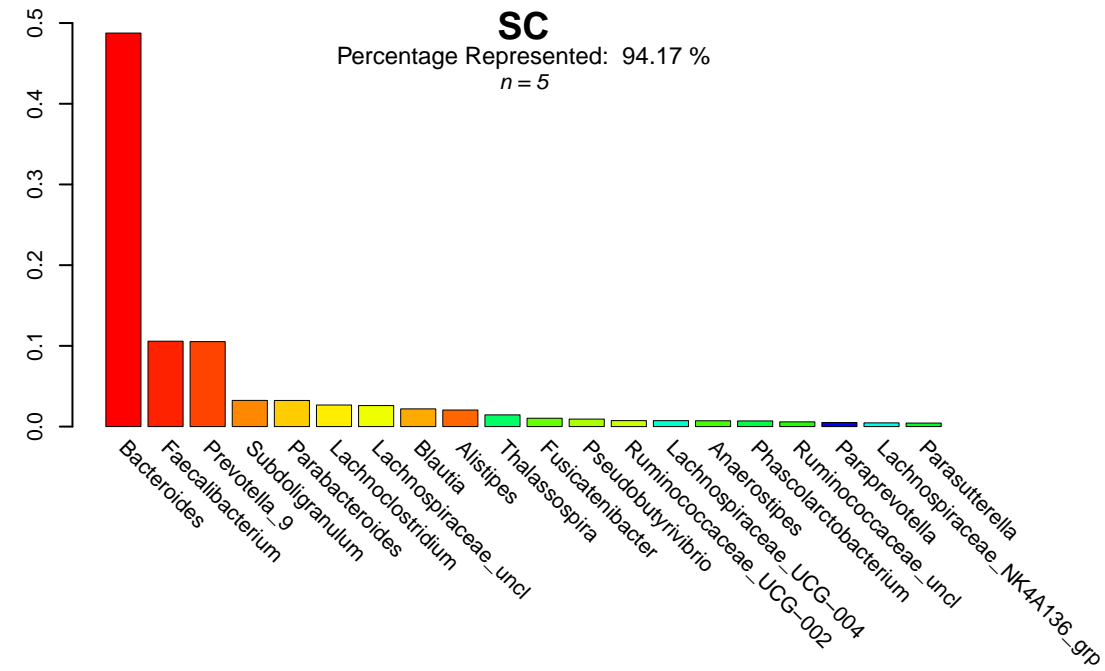
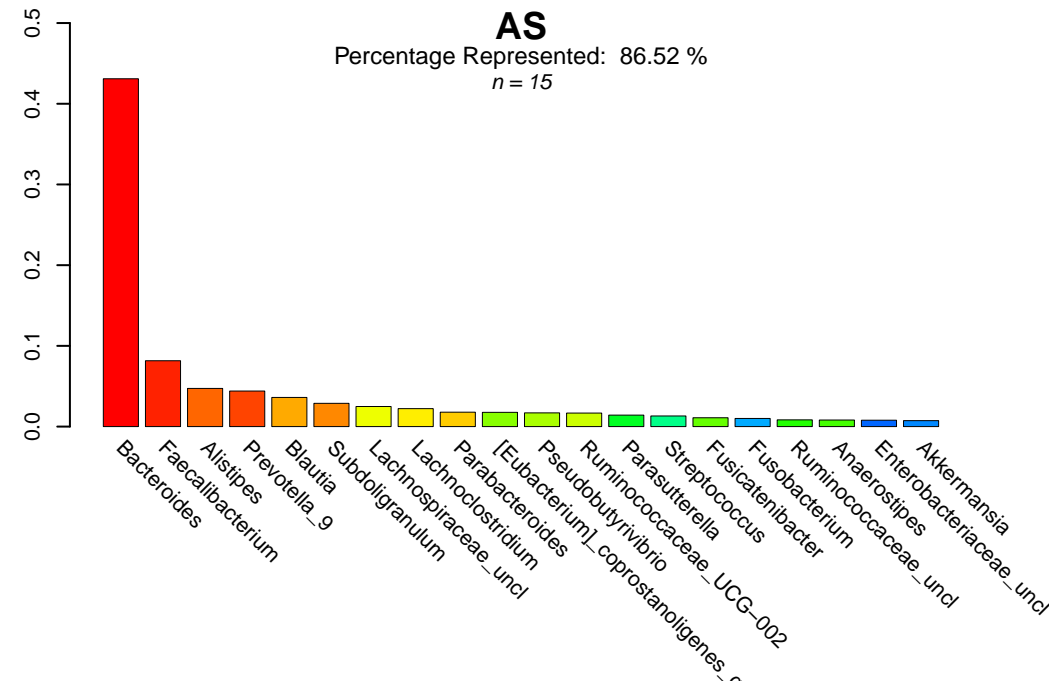
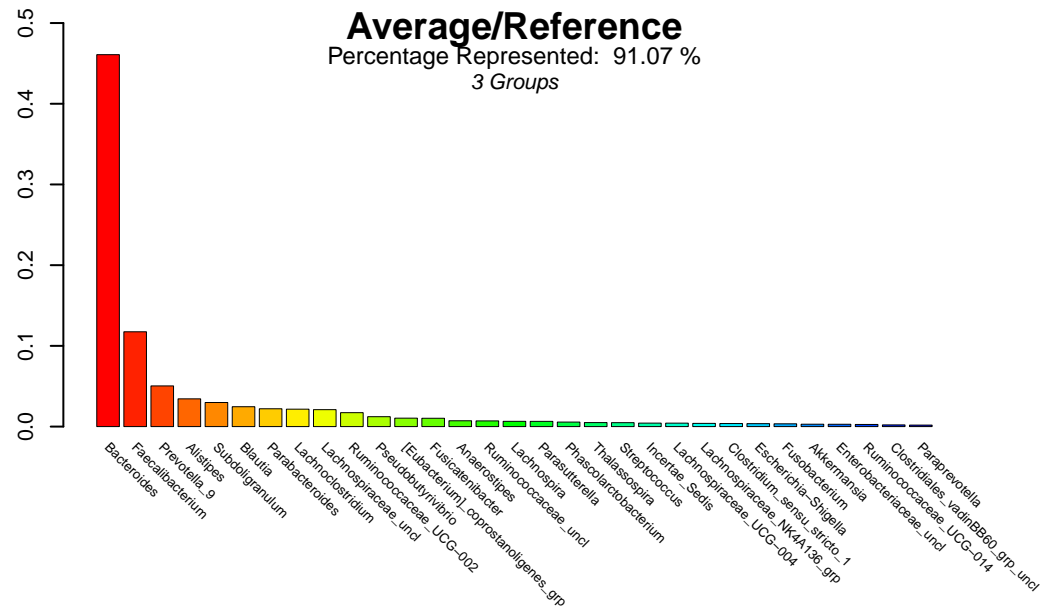
# LDH



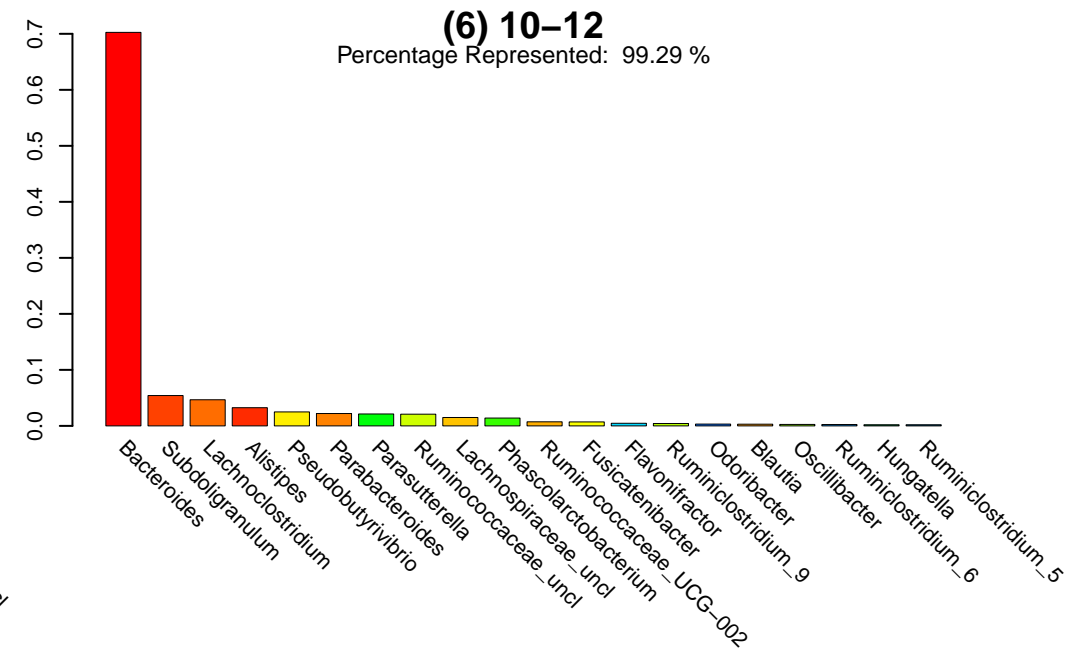
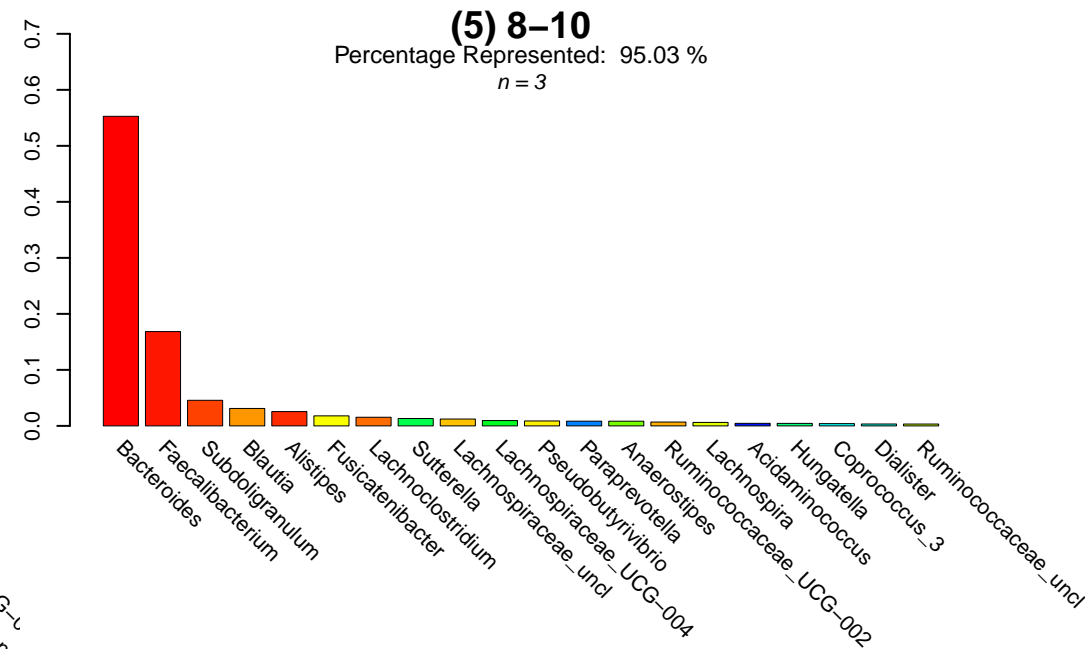
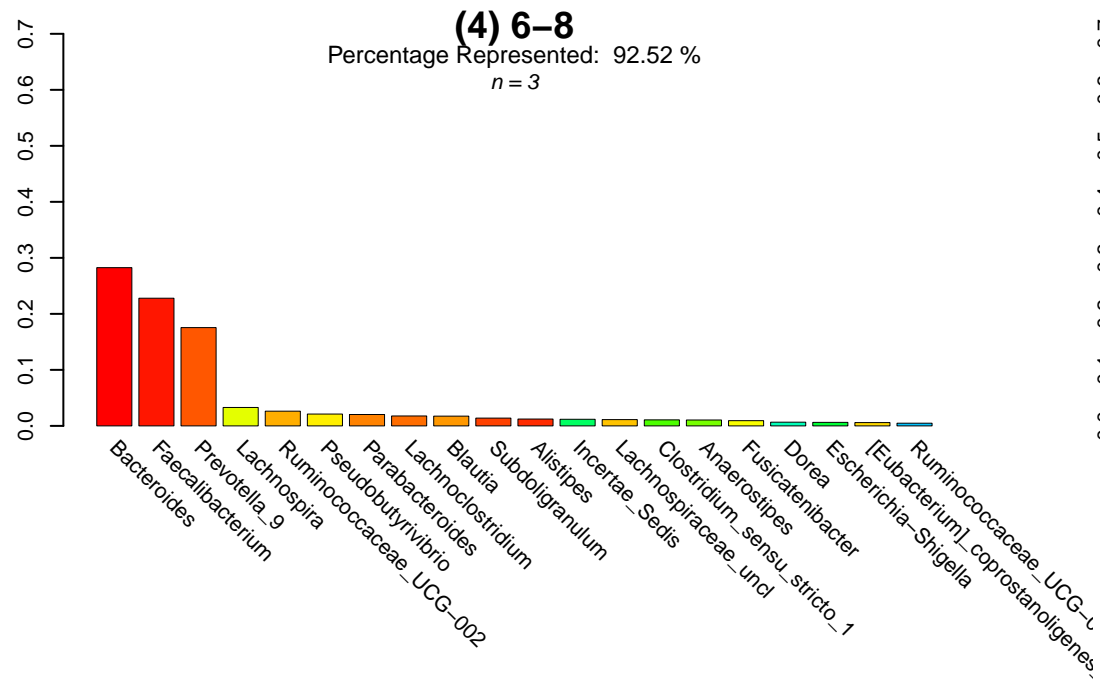
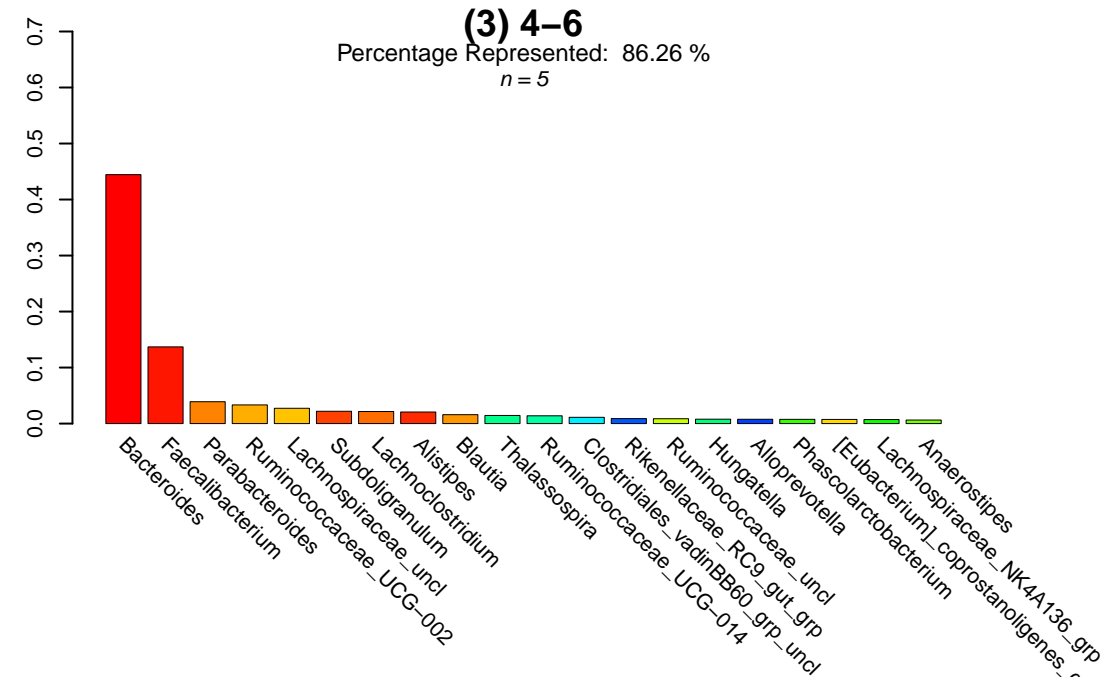
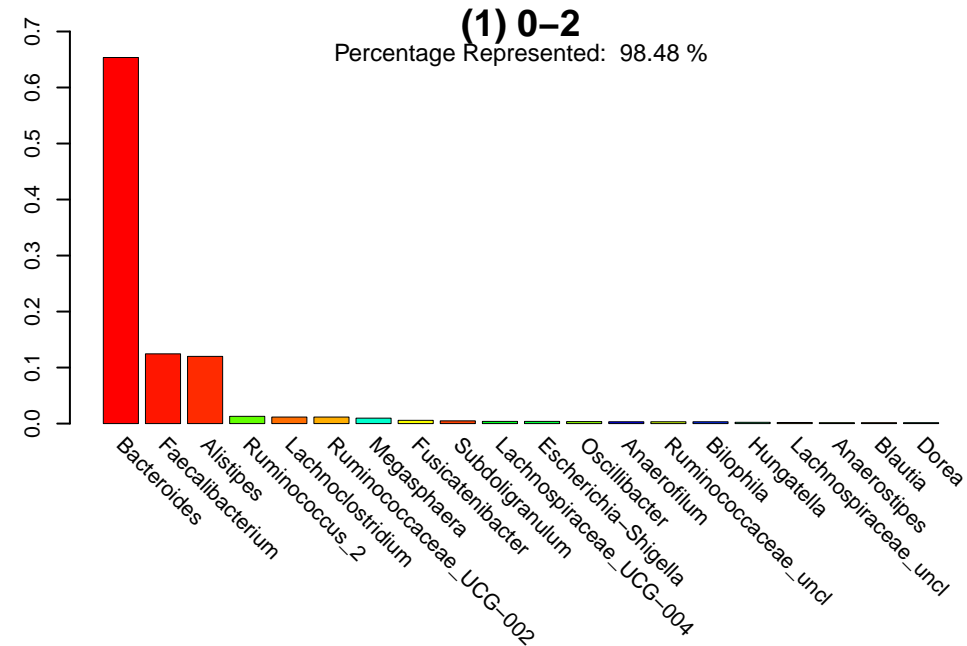
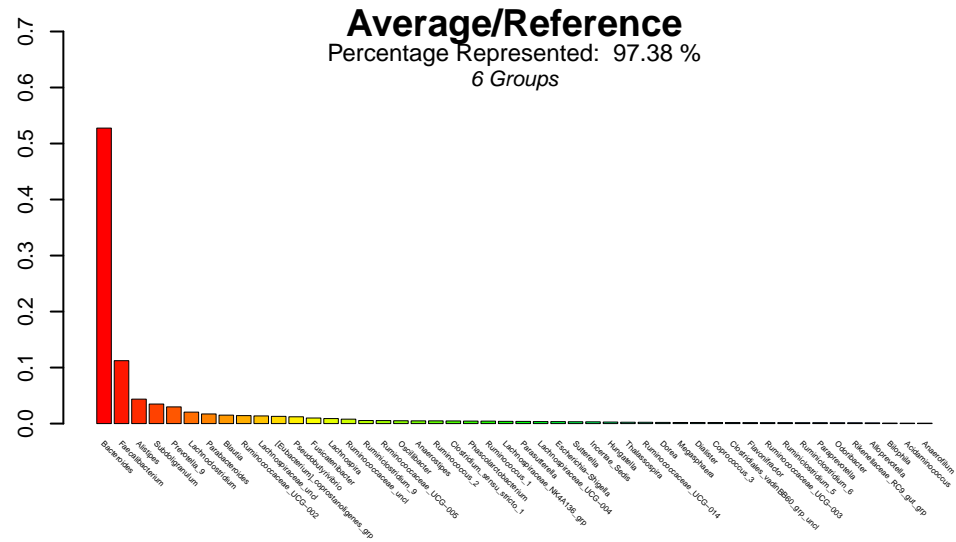
Hapto



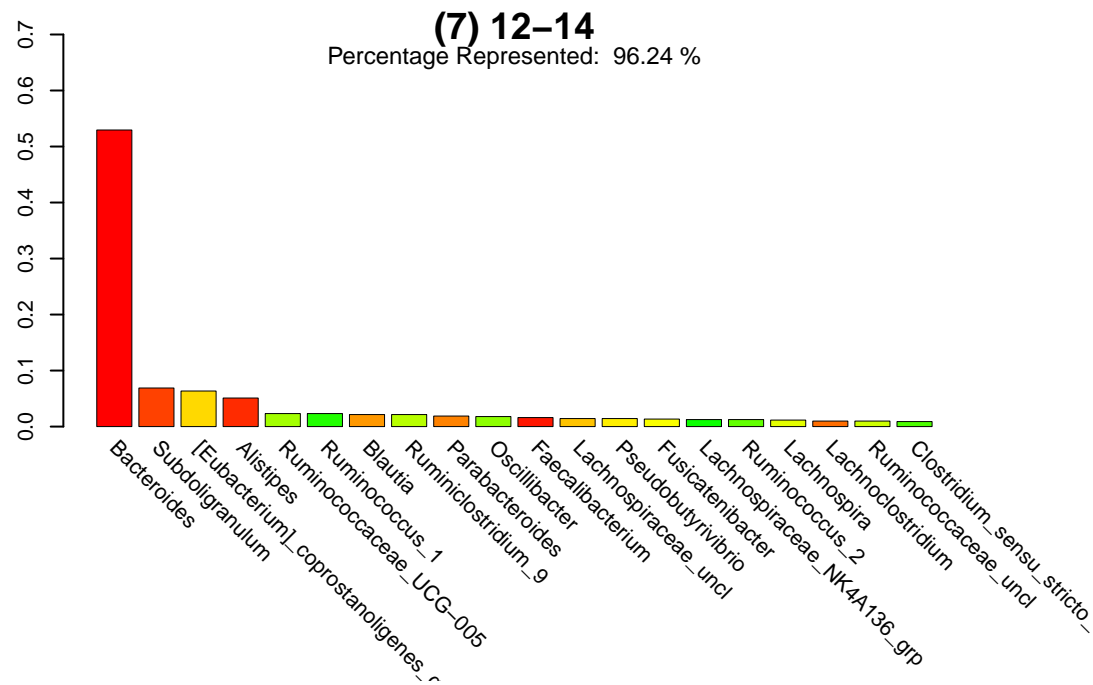
# Type



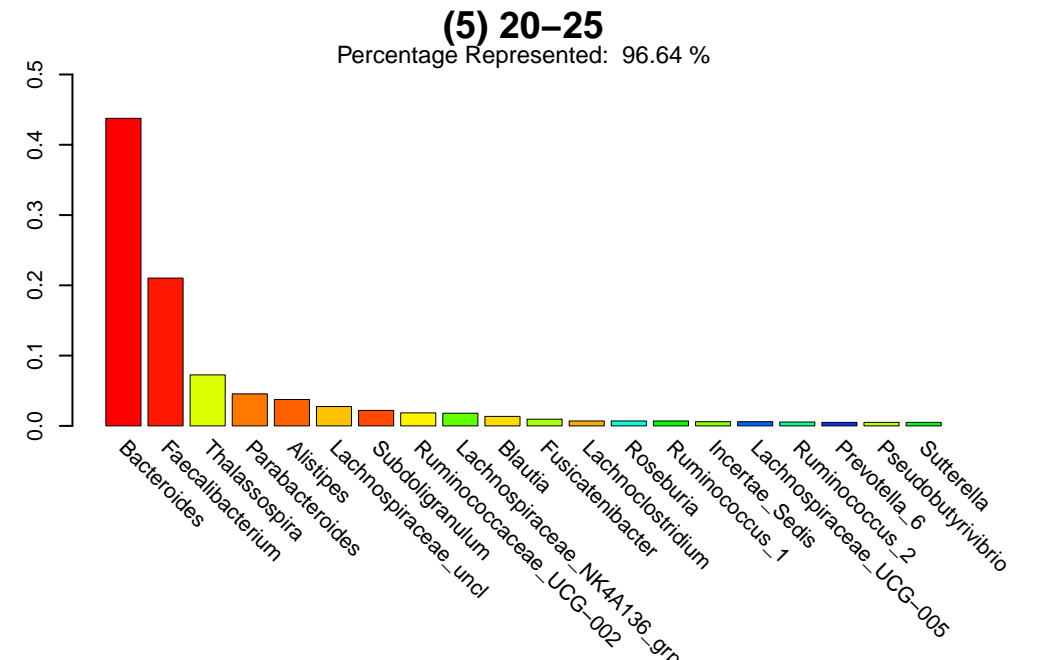
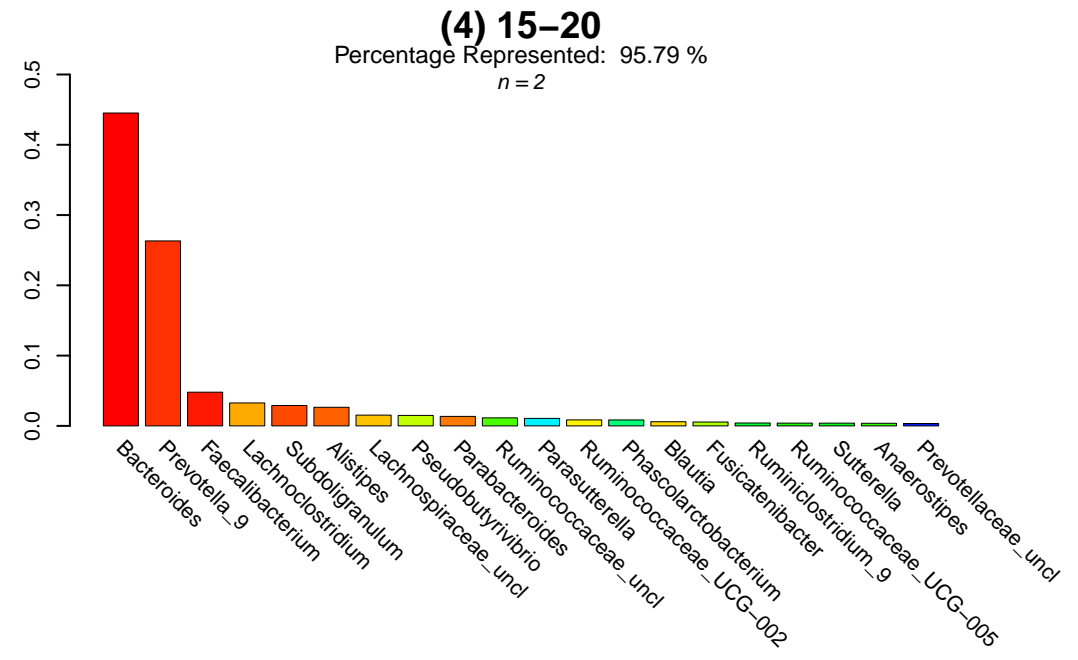
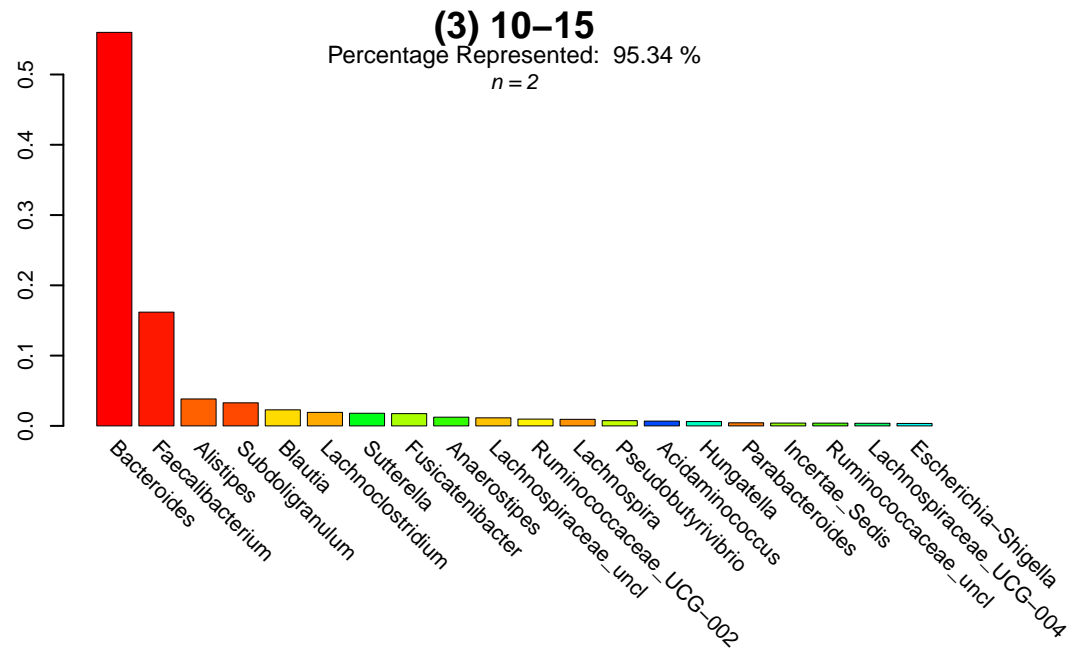
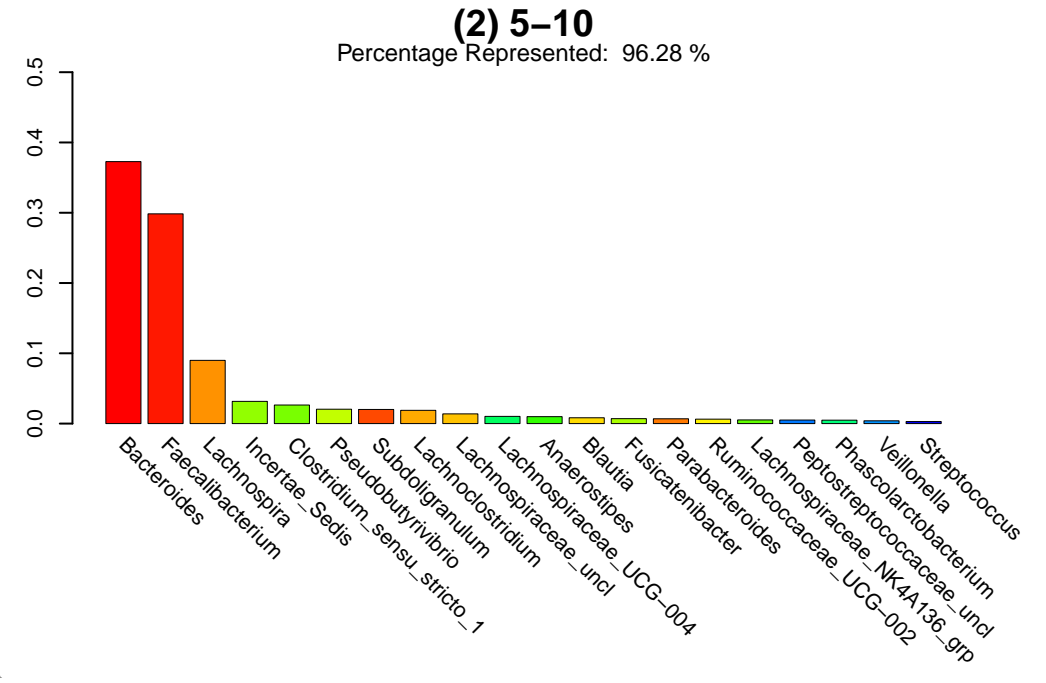
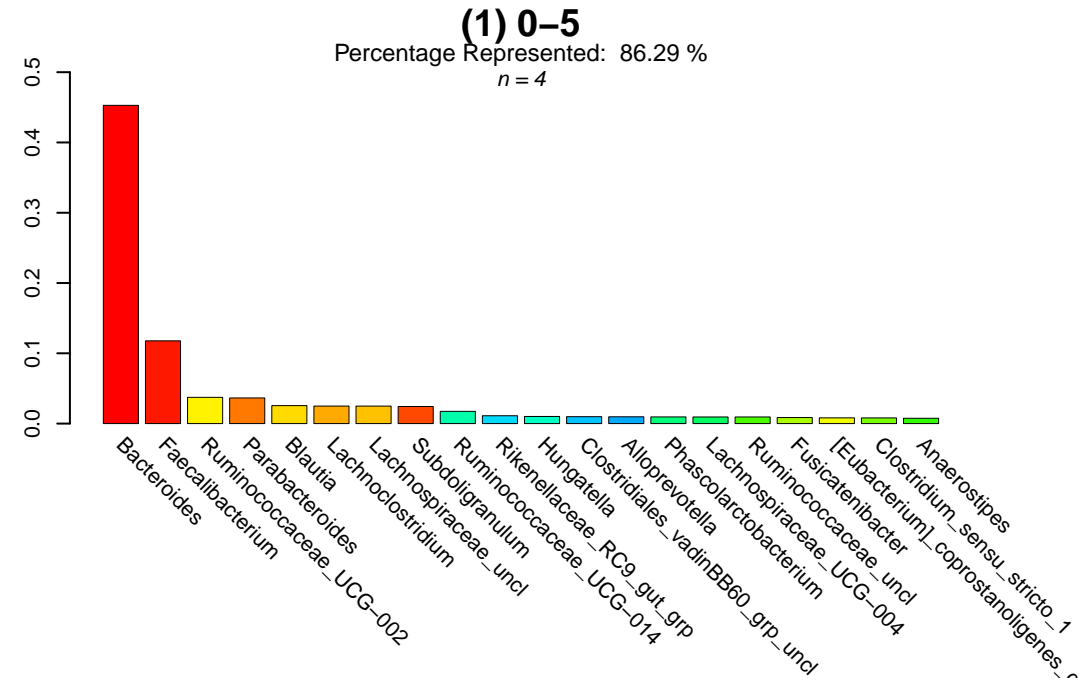
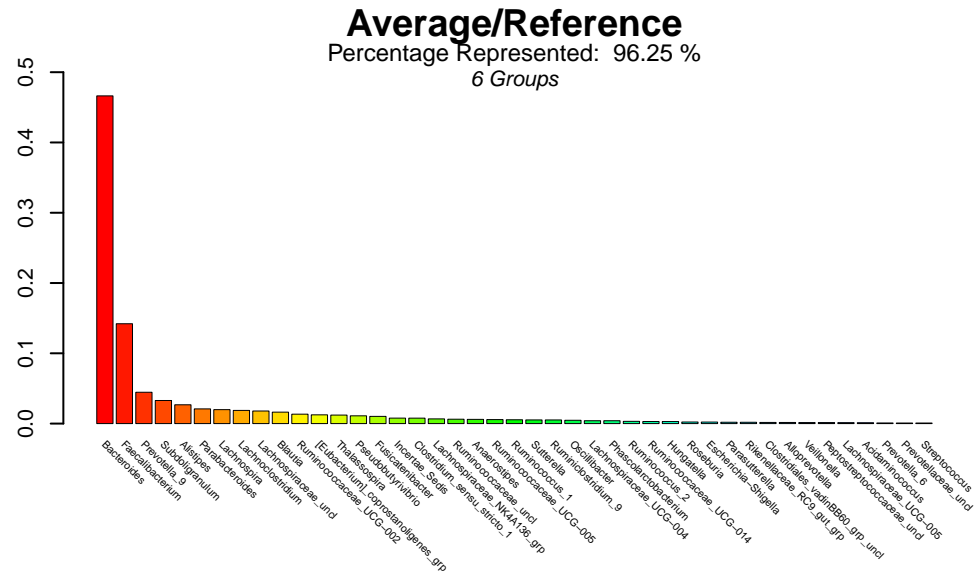
# Neut



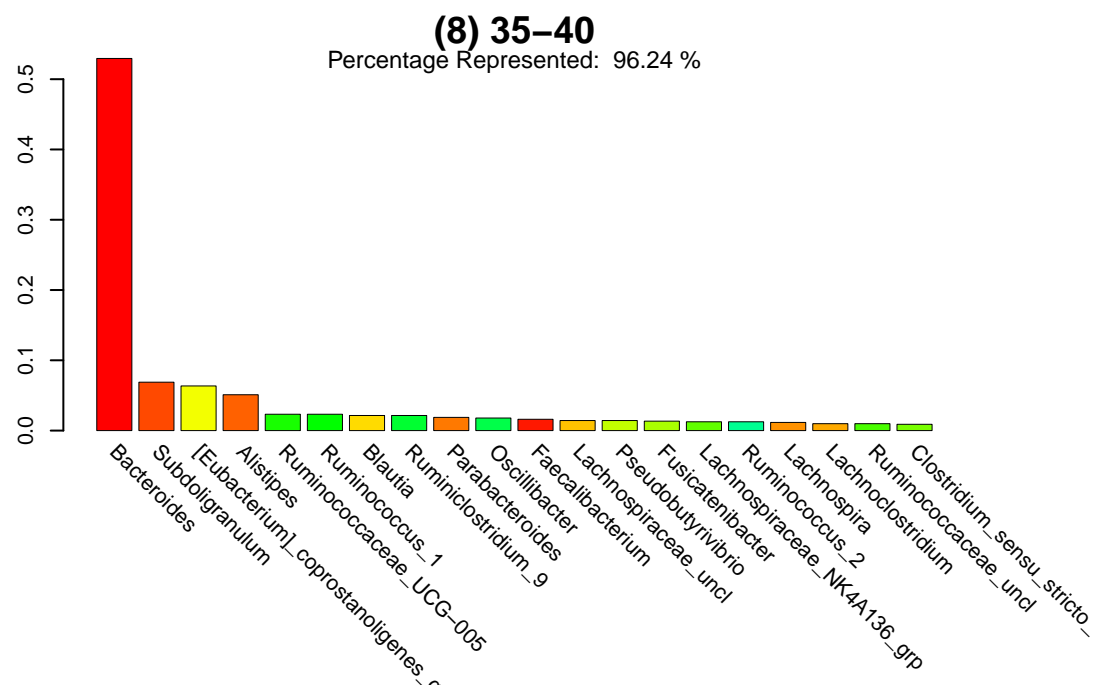
Neut



# SCChi\_CD16

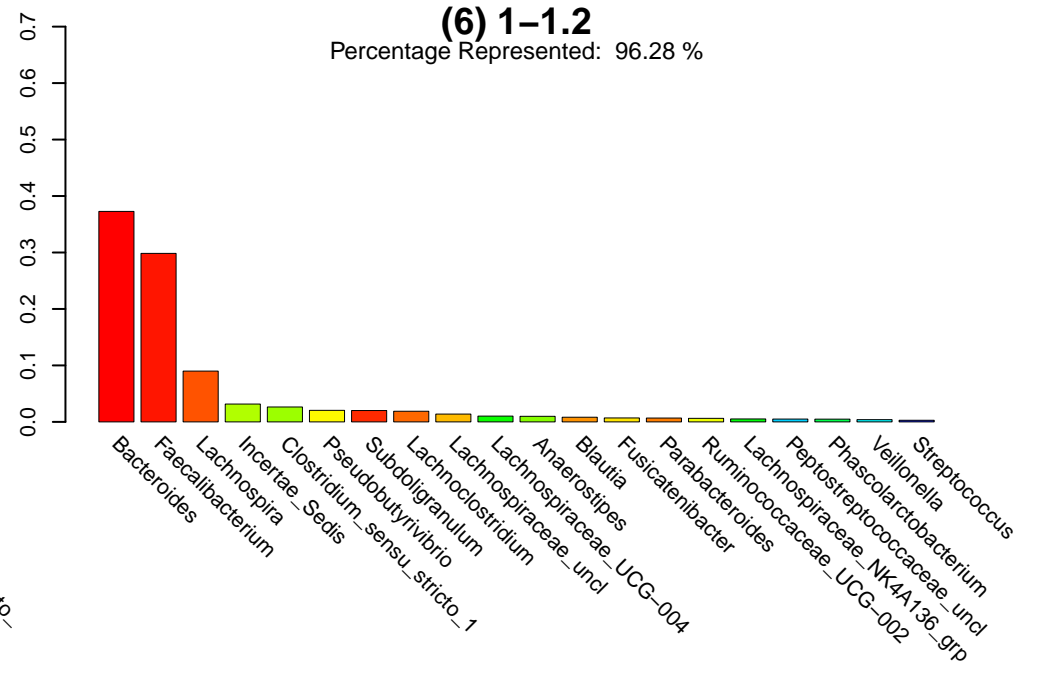
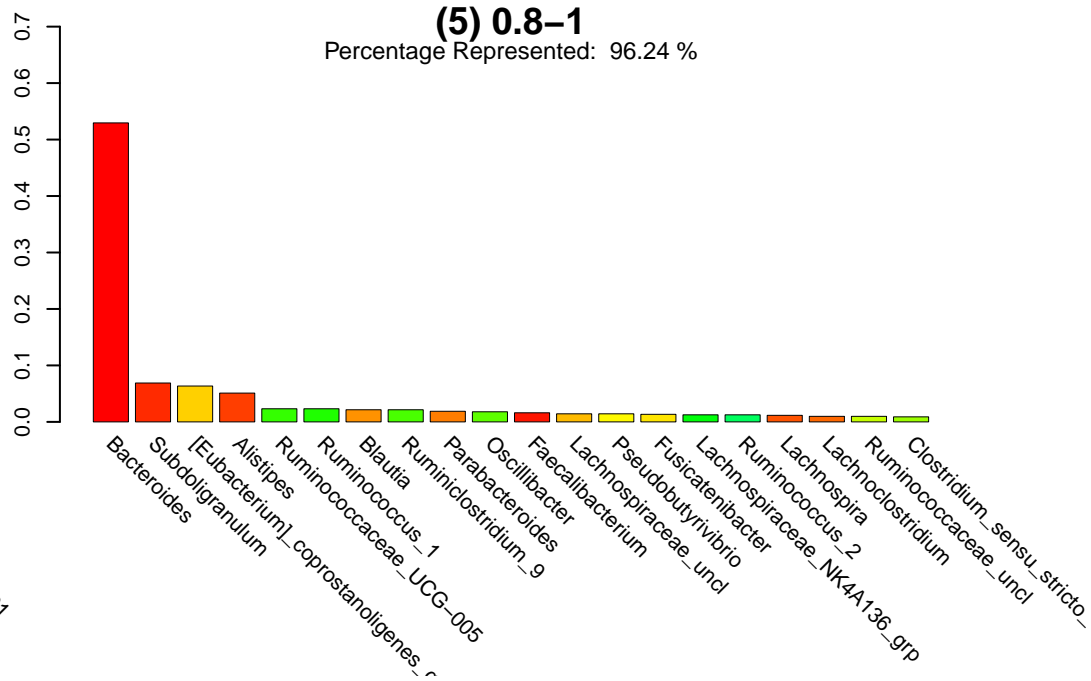
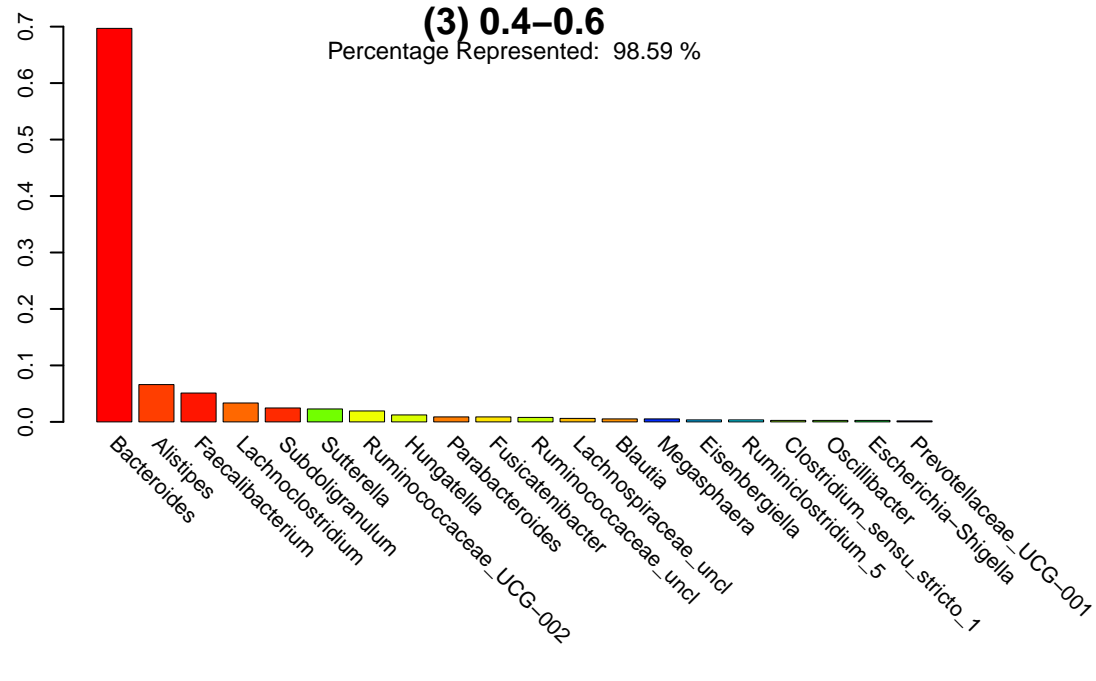
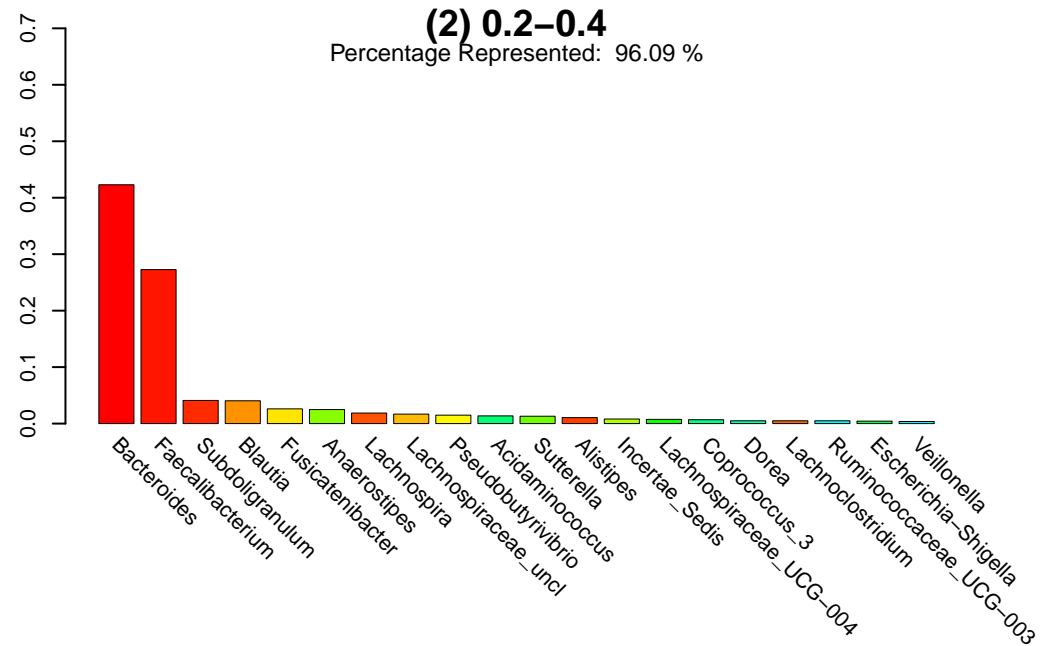
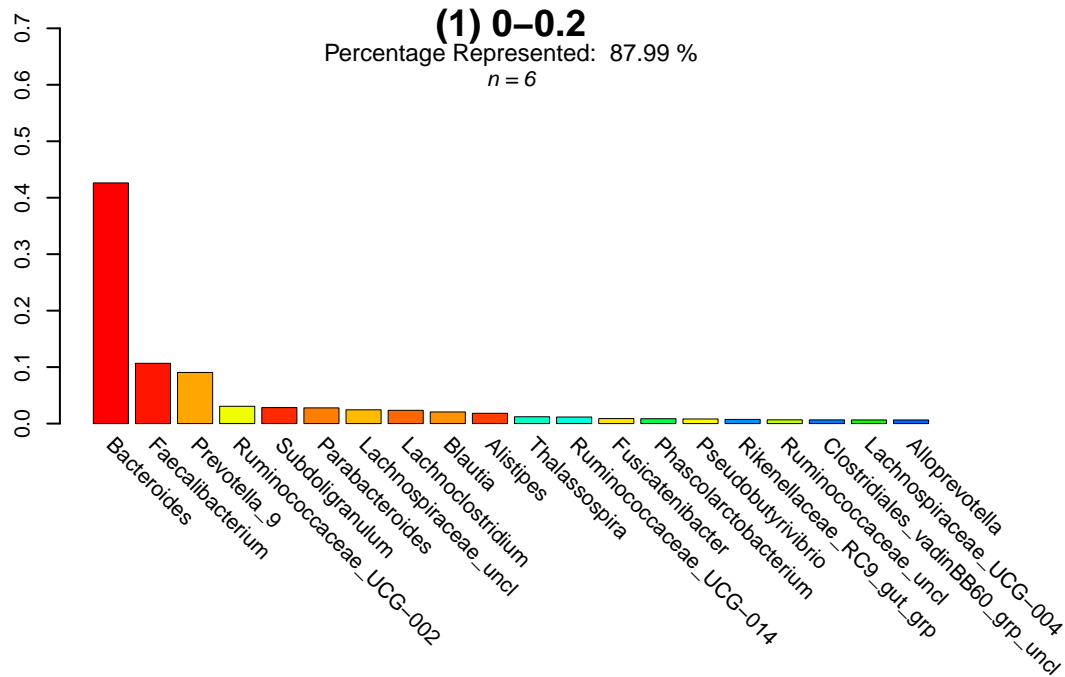
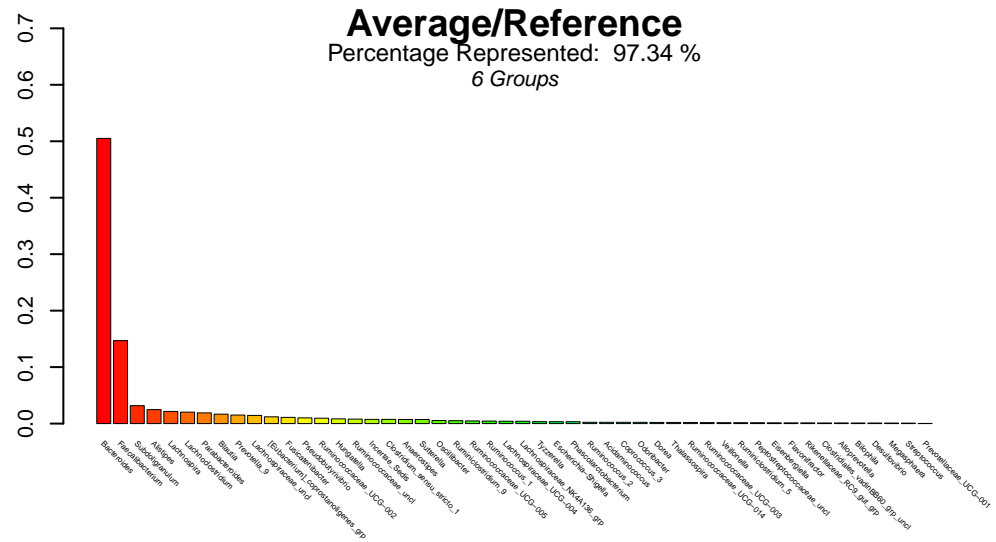


# SCCChi\_CD16

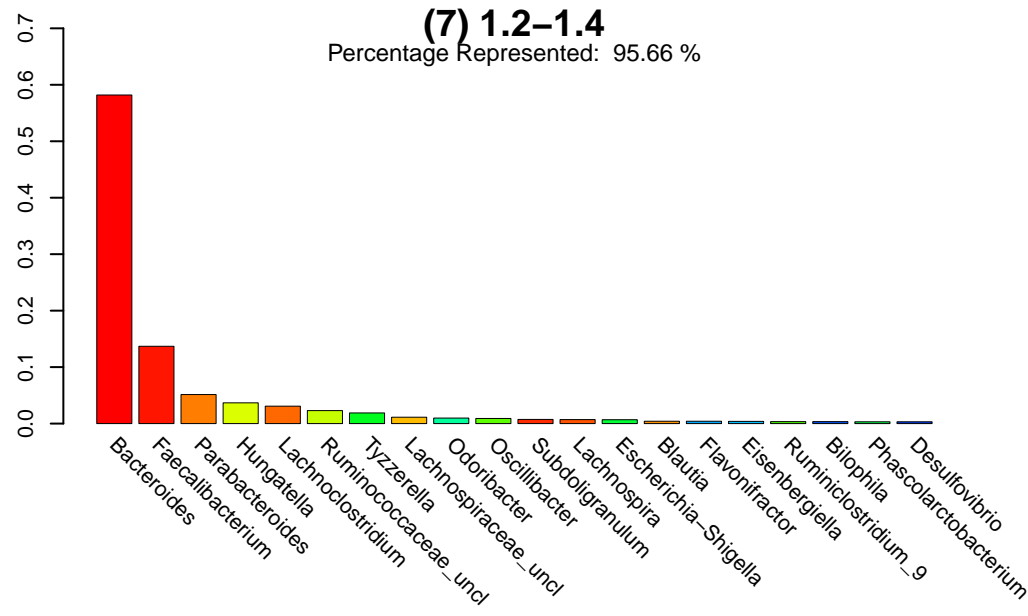




# CD62lo\_CD11b



# CD62lo\_CD11b



CD62lo\_CD184

