Summary Table File: LimSC25.taxa.genus.NoControls.N500.summary_table.tsv

Factor File: 0137_Lim_Metadata_20171019.tsv Output File Root: sickle.genus.tail.t10

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: 10

Samples exclusive to Summary Table:
character(0)

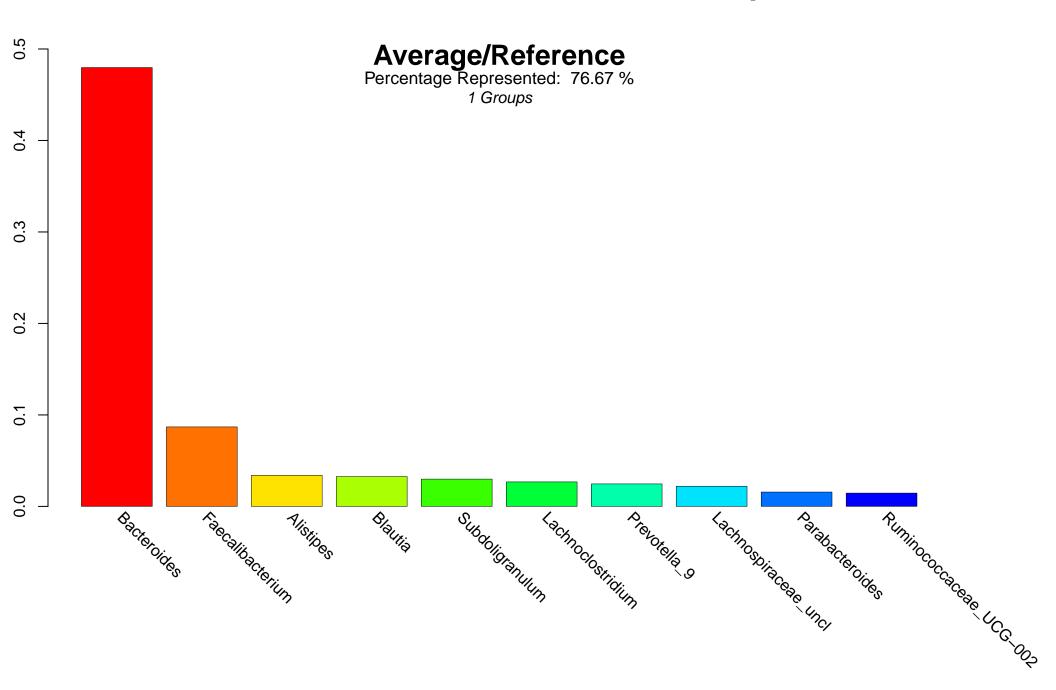
Samples exclusive to Factor Table:

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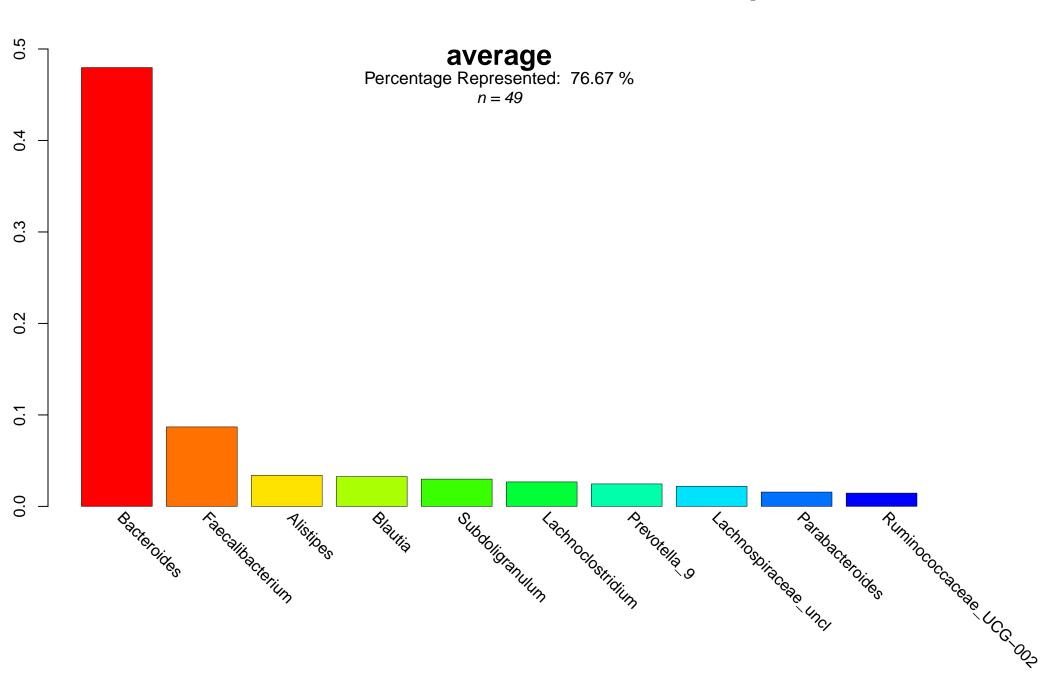
[4] "0000.0386.20170824" "0000.0387.20170825" "0000.0388.20170828"

[7] "0000.0389.Ext.N" "0001.0390.20170913" "0002.0391.20170913"

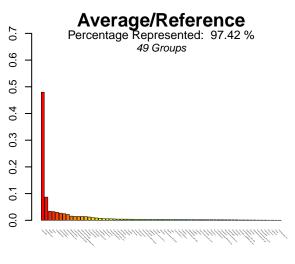
Mean abundances across all samples

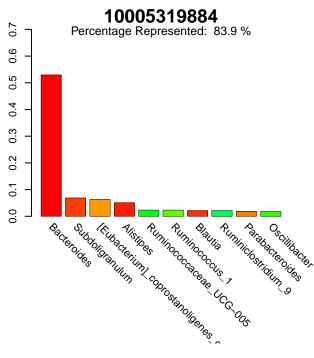


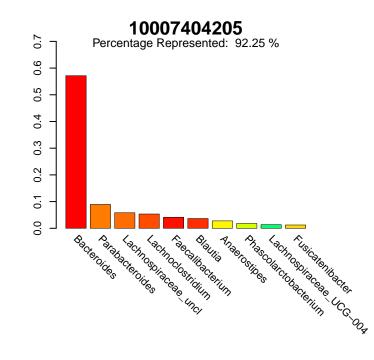
Mean abundances across all samples

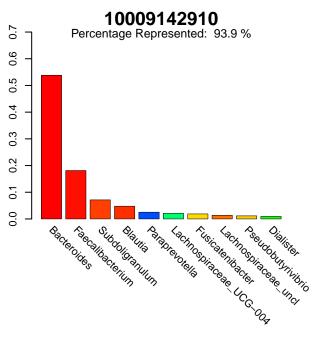


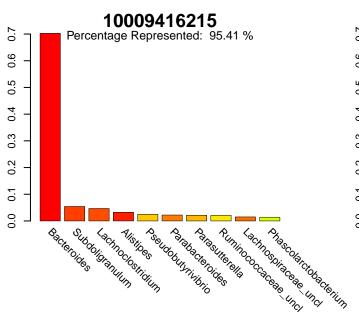
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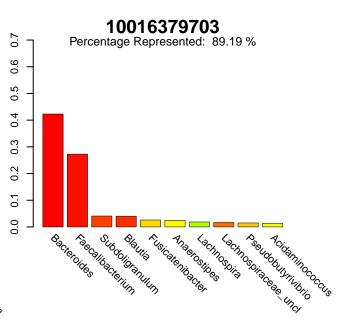




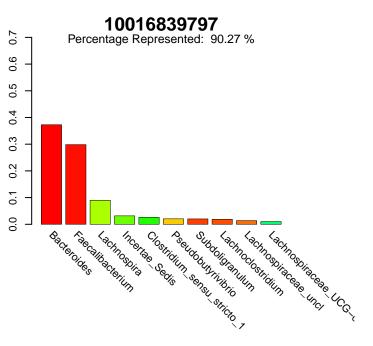




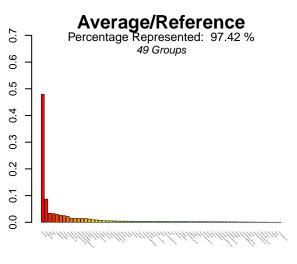


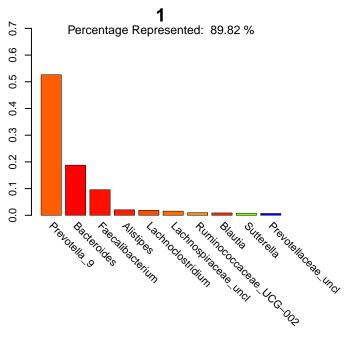


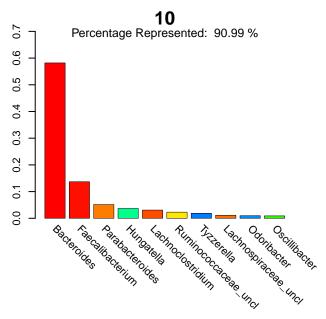
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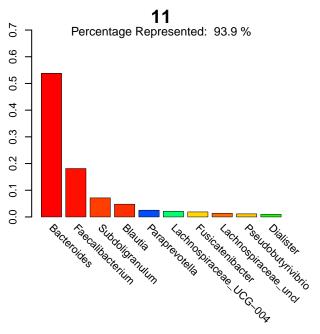


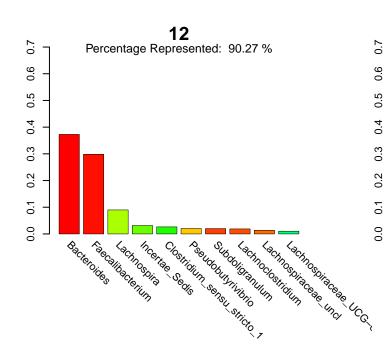
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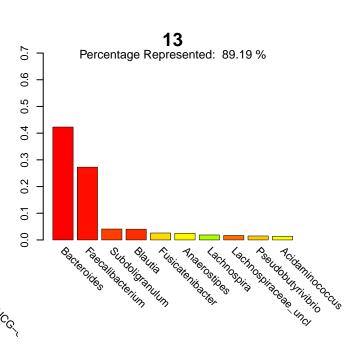




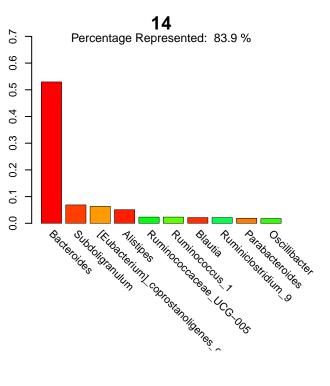




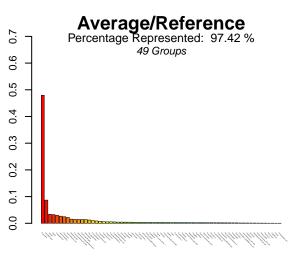


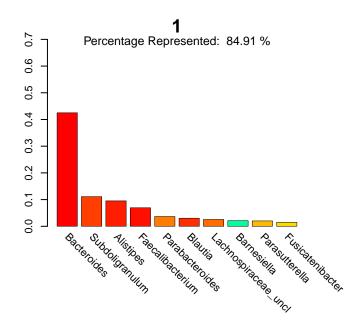


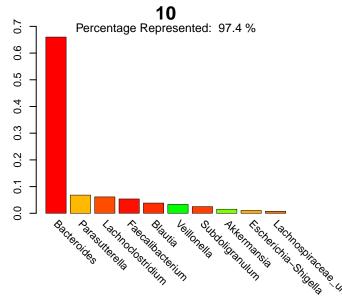
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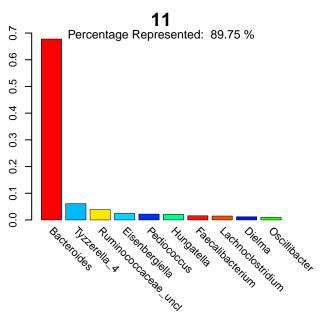


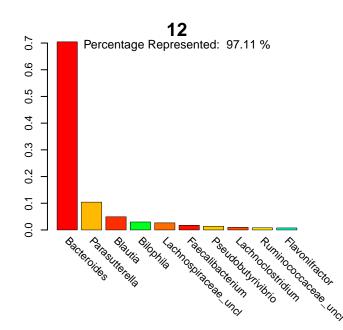
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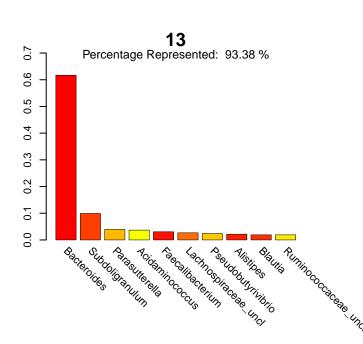




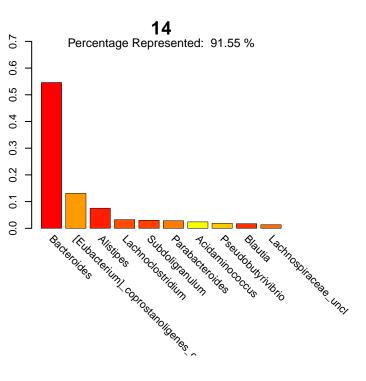




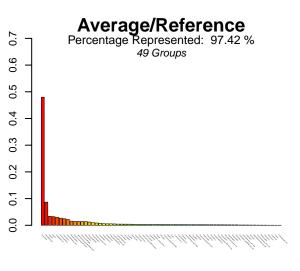


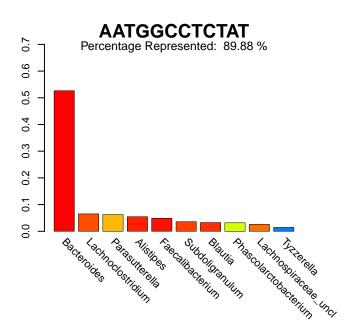


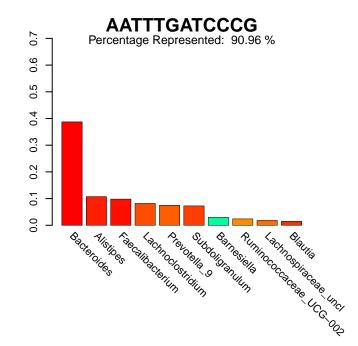
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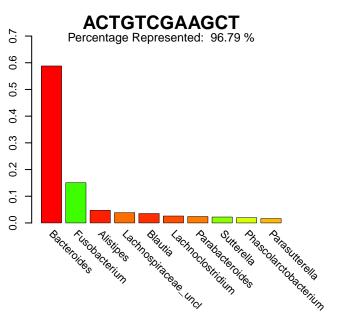


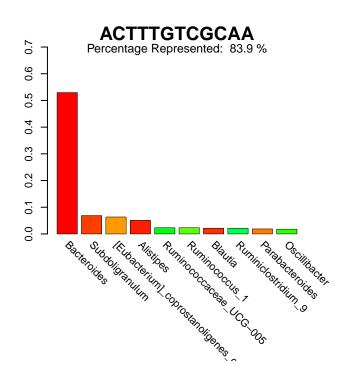
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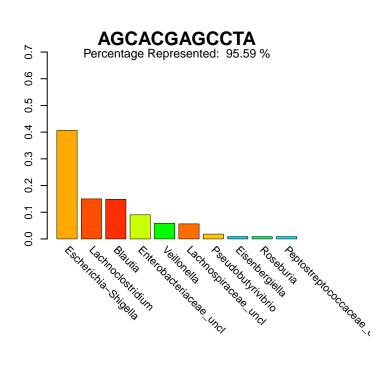




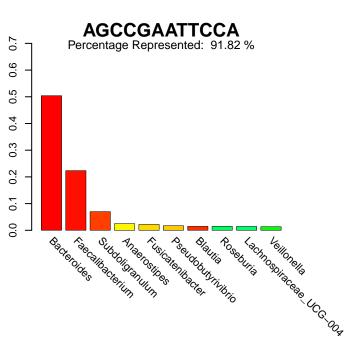




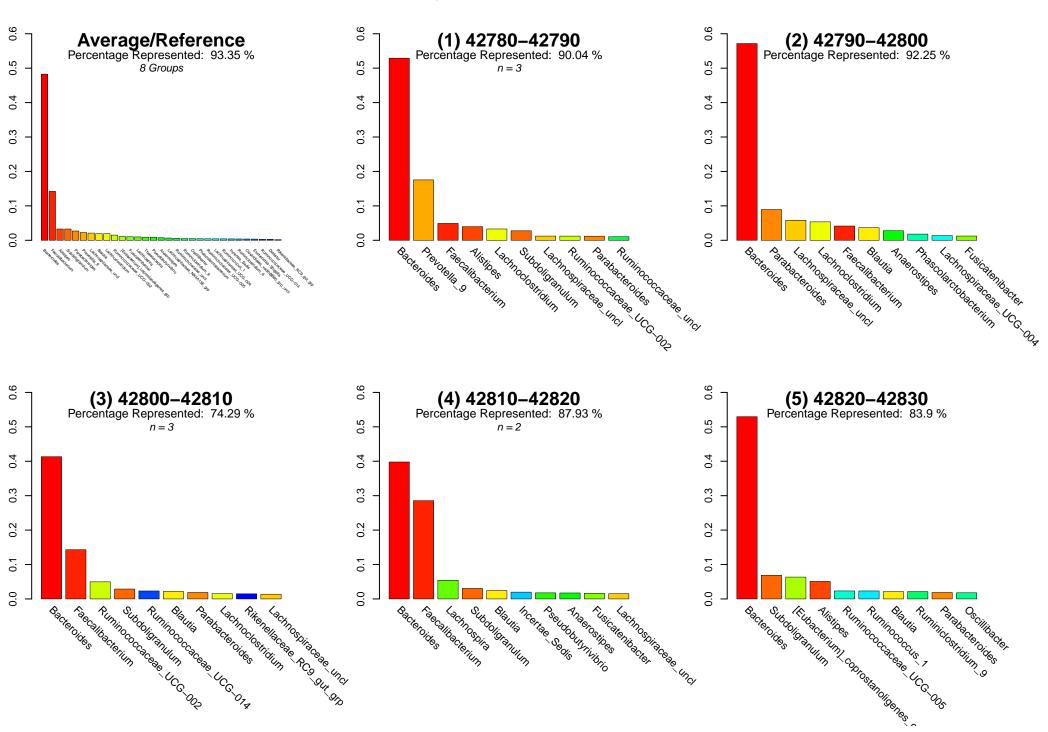




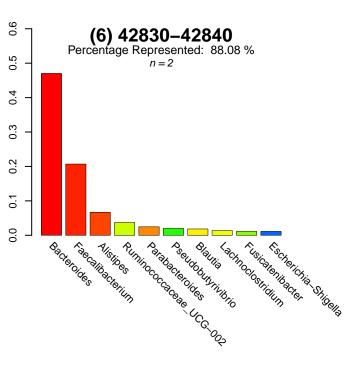
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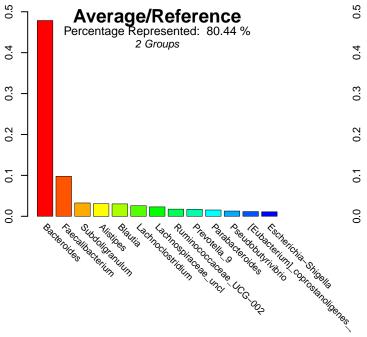
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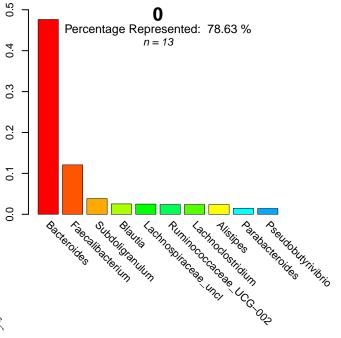


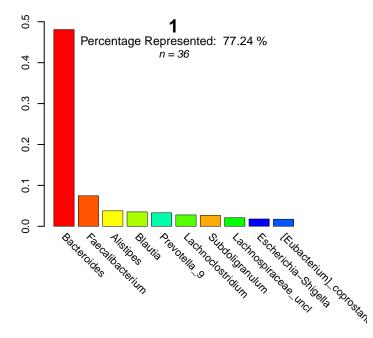
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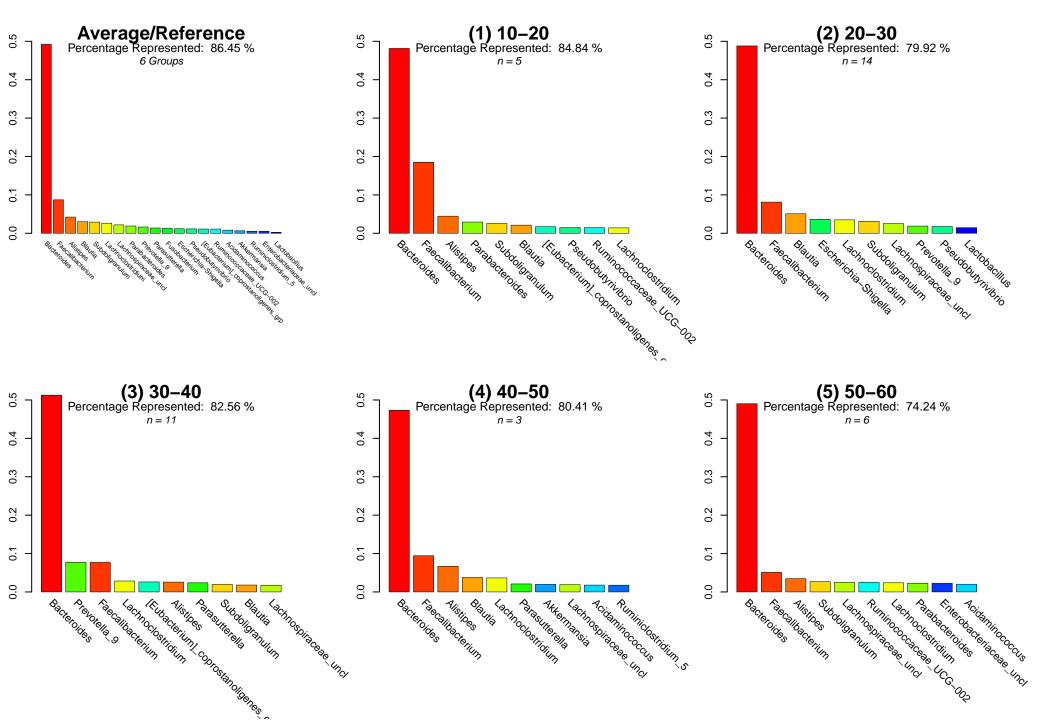
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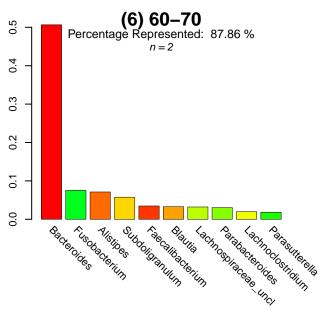




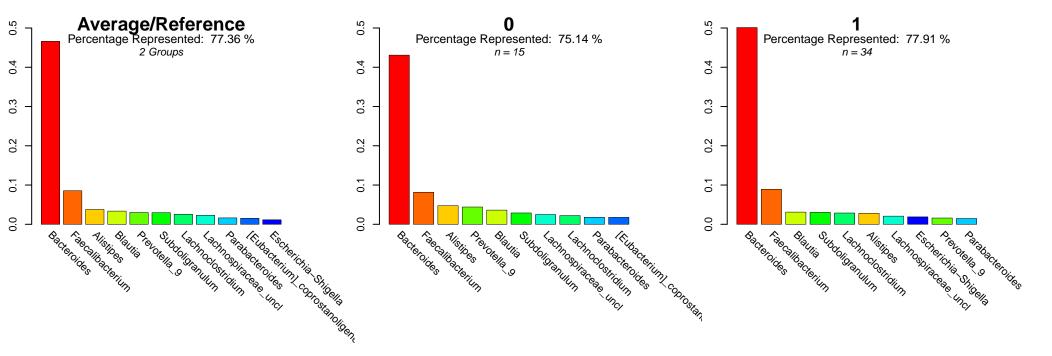
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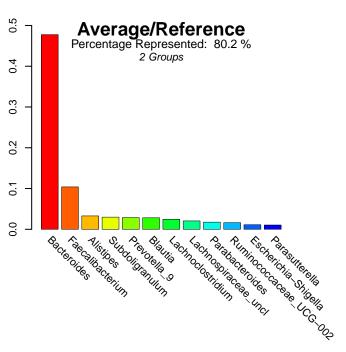
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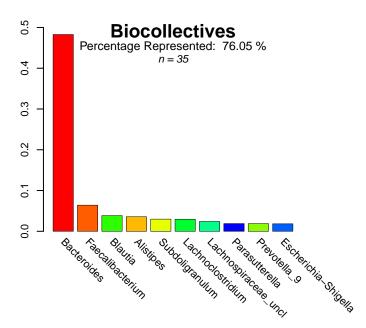


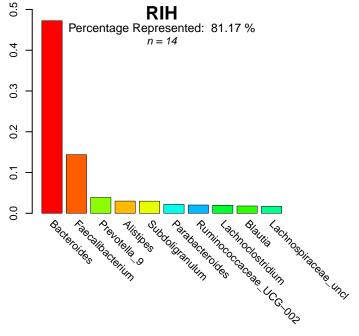
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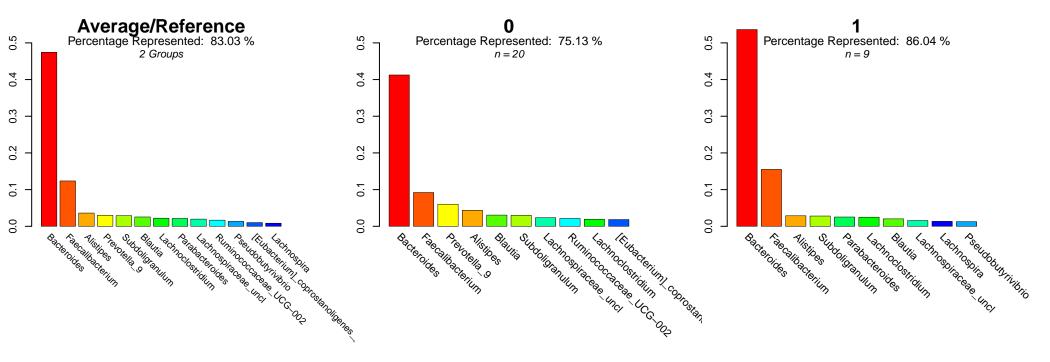
Origin



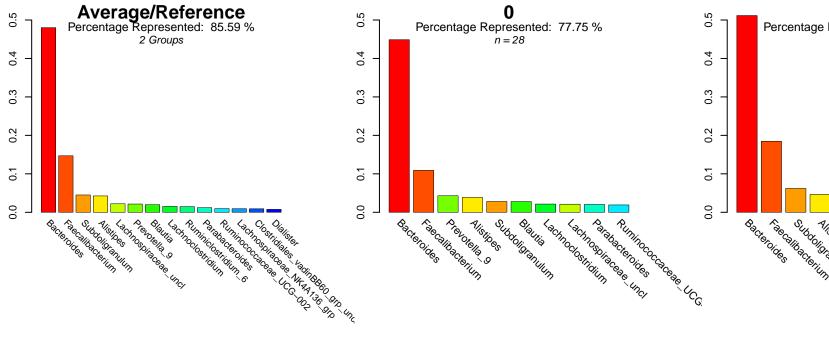


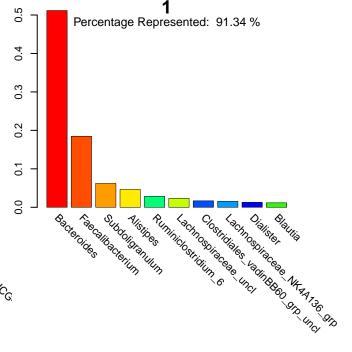


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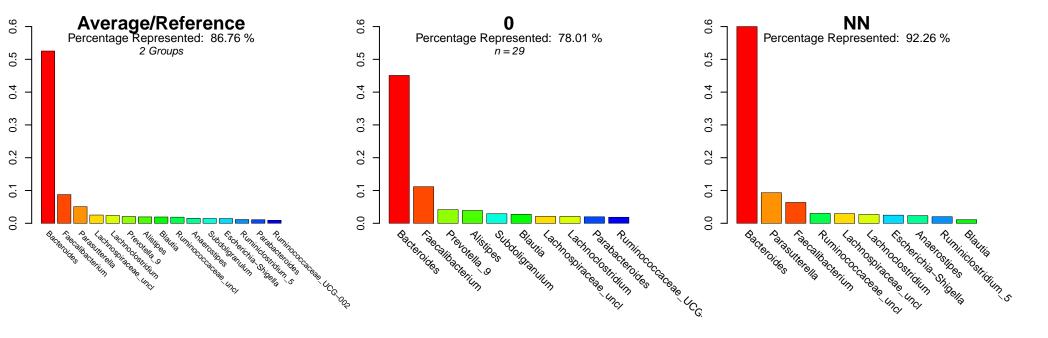


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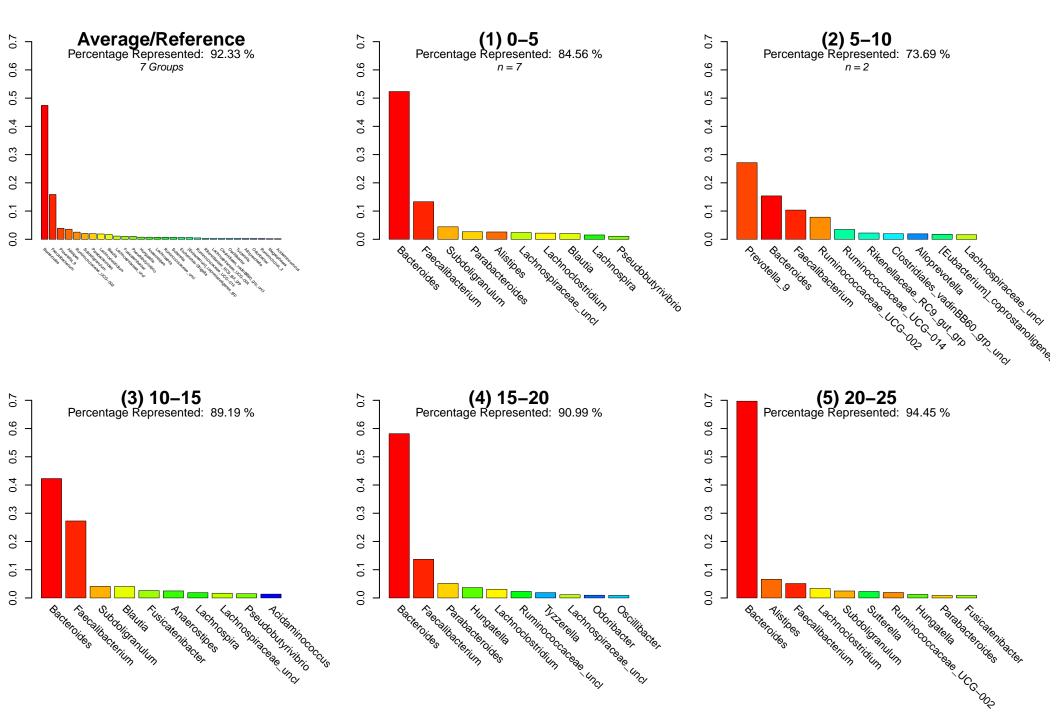




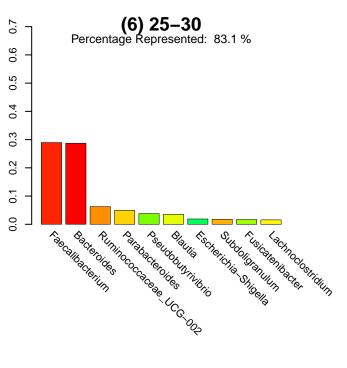
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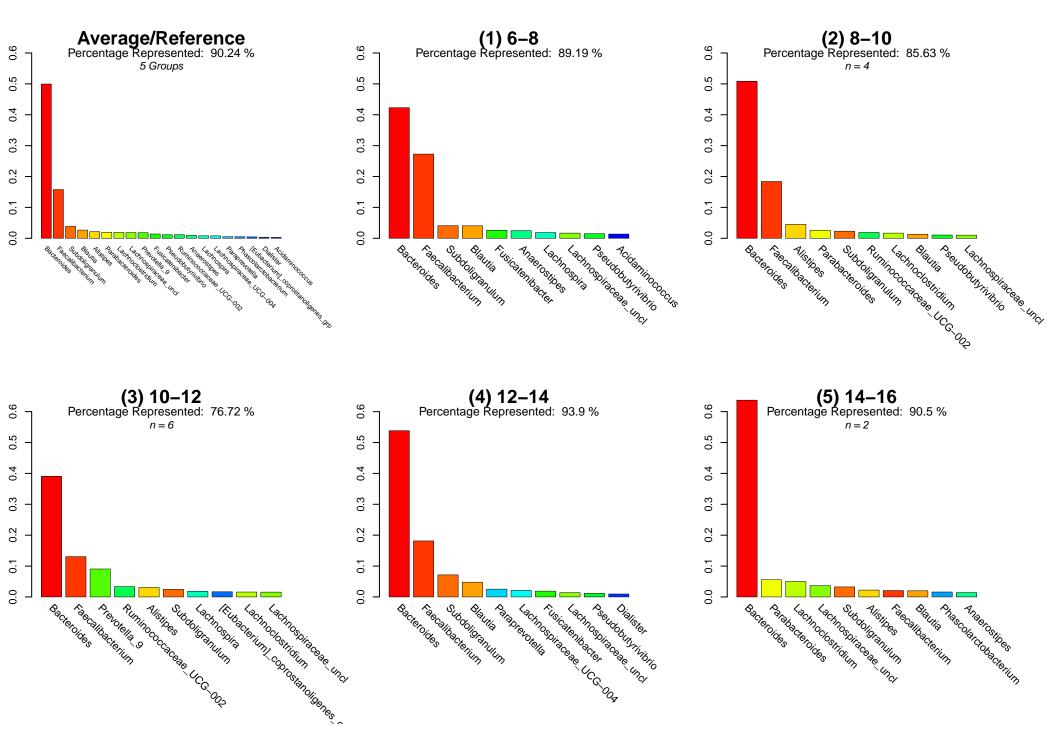
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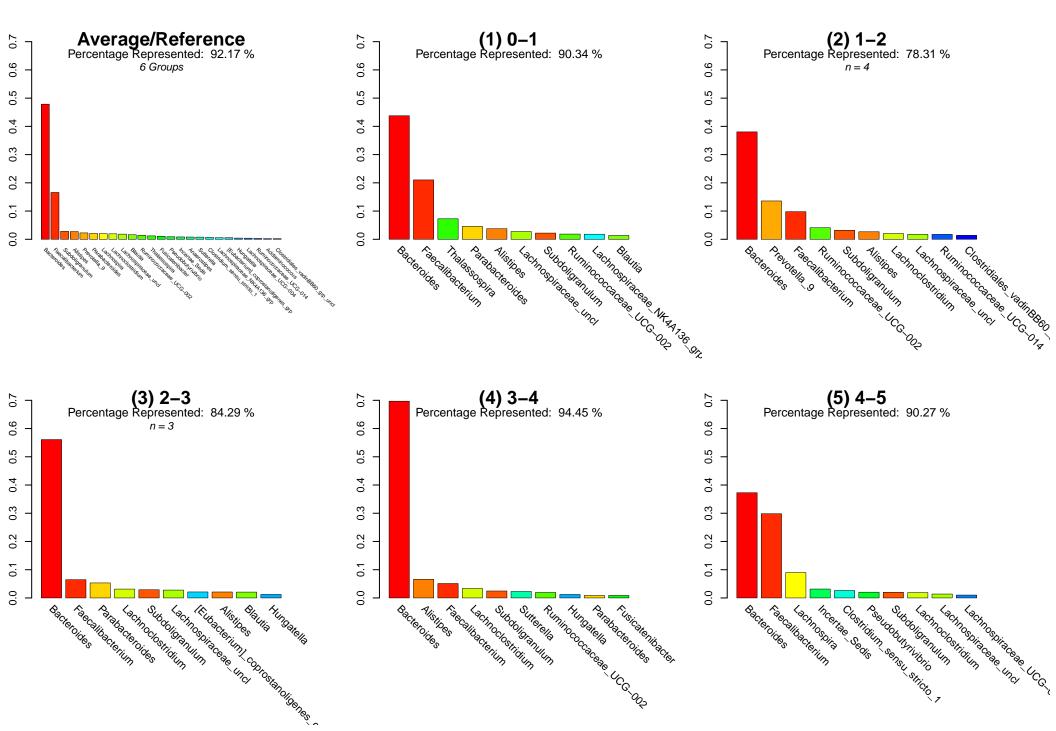
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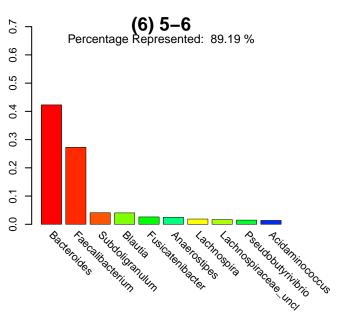
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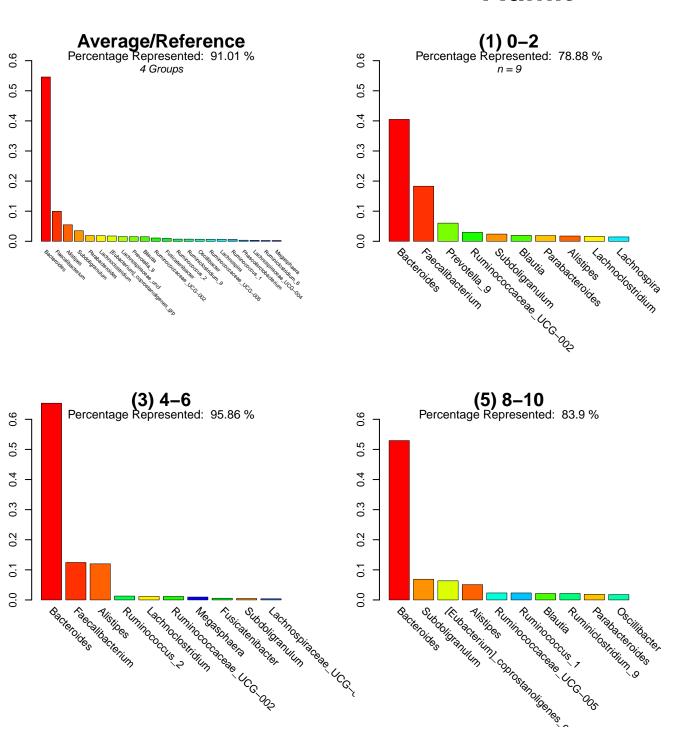
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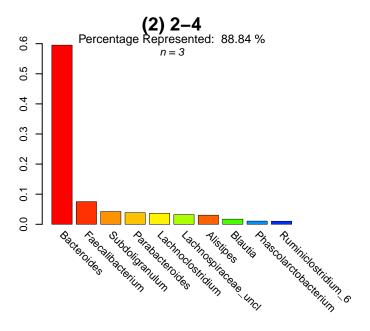


Bilirubin

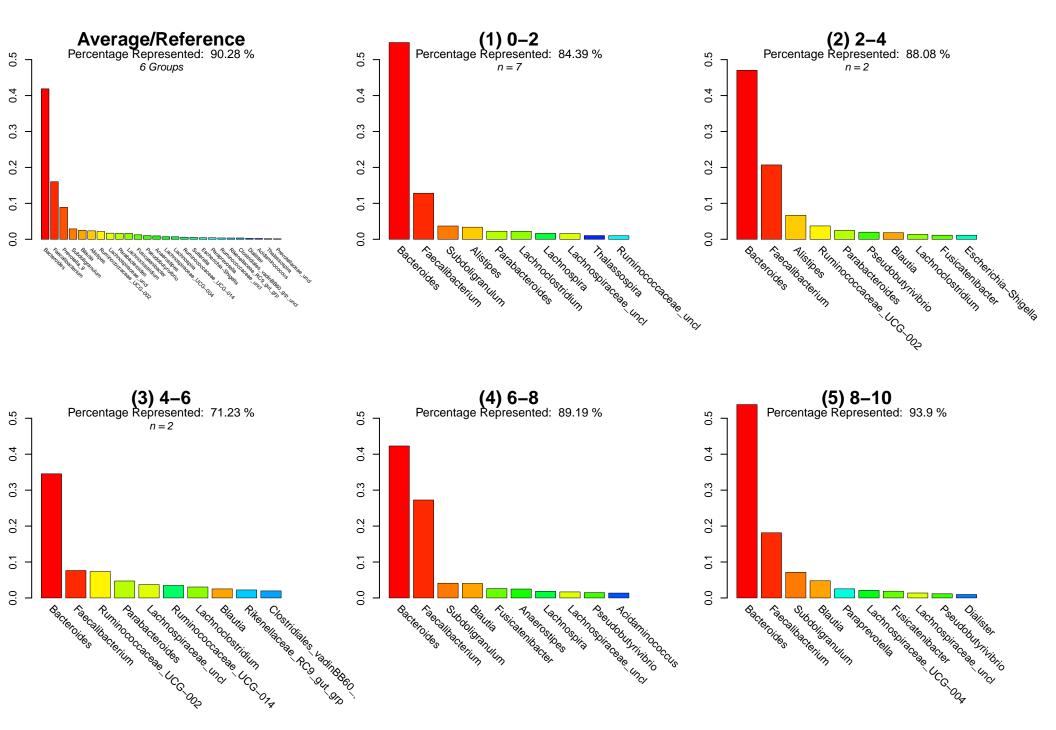


Admit

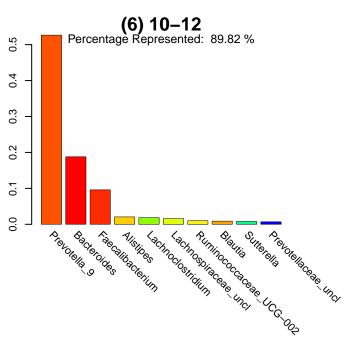




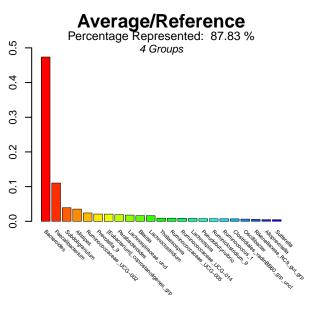
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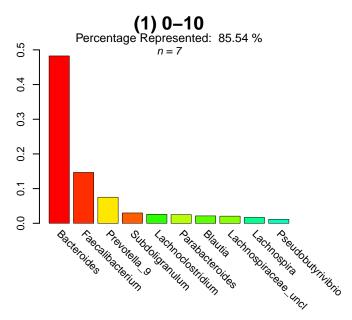


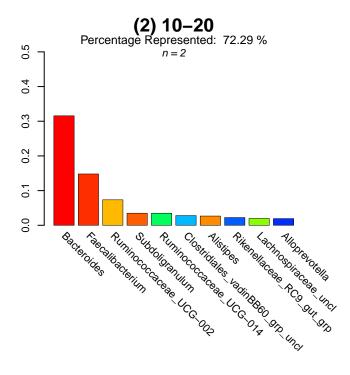
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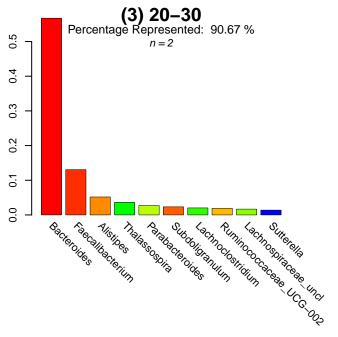


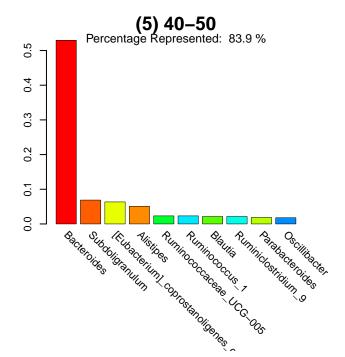
ESR



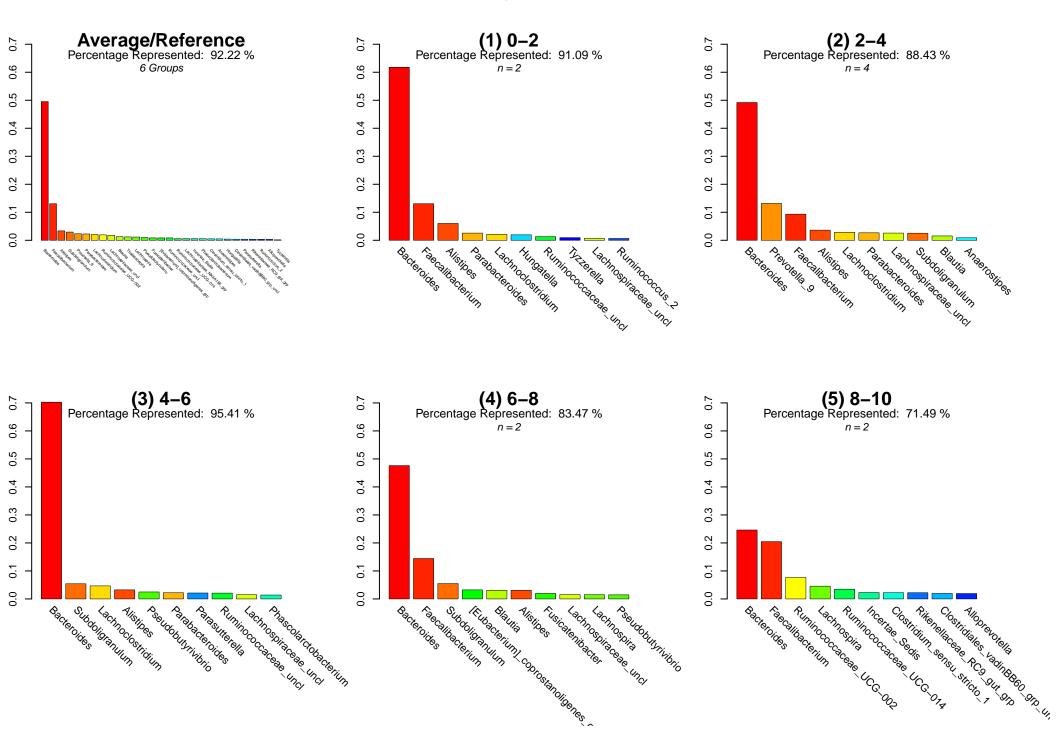




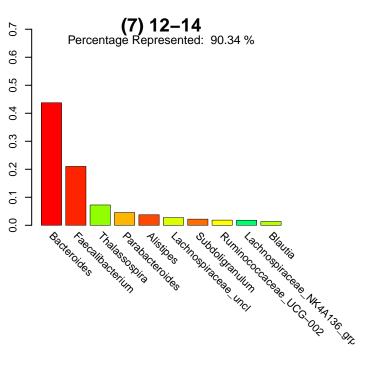




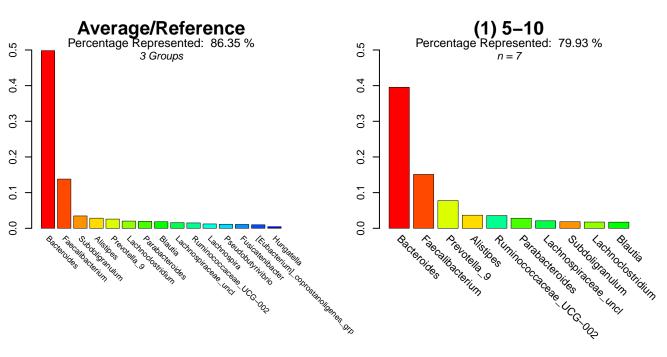
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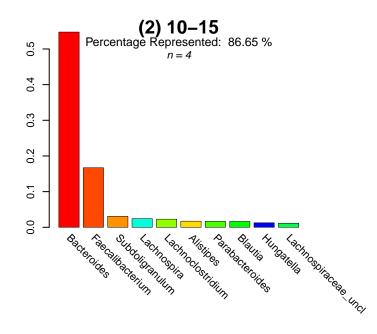


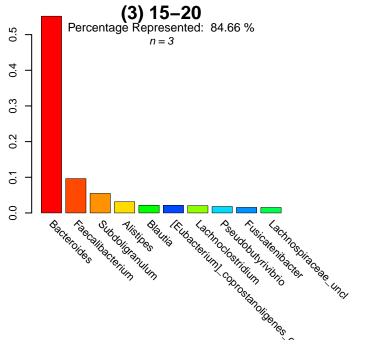
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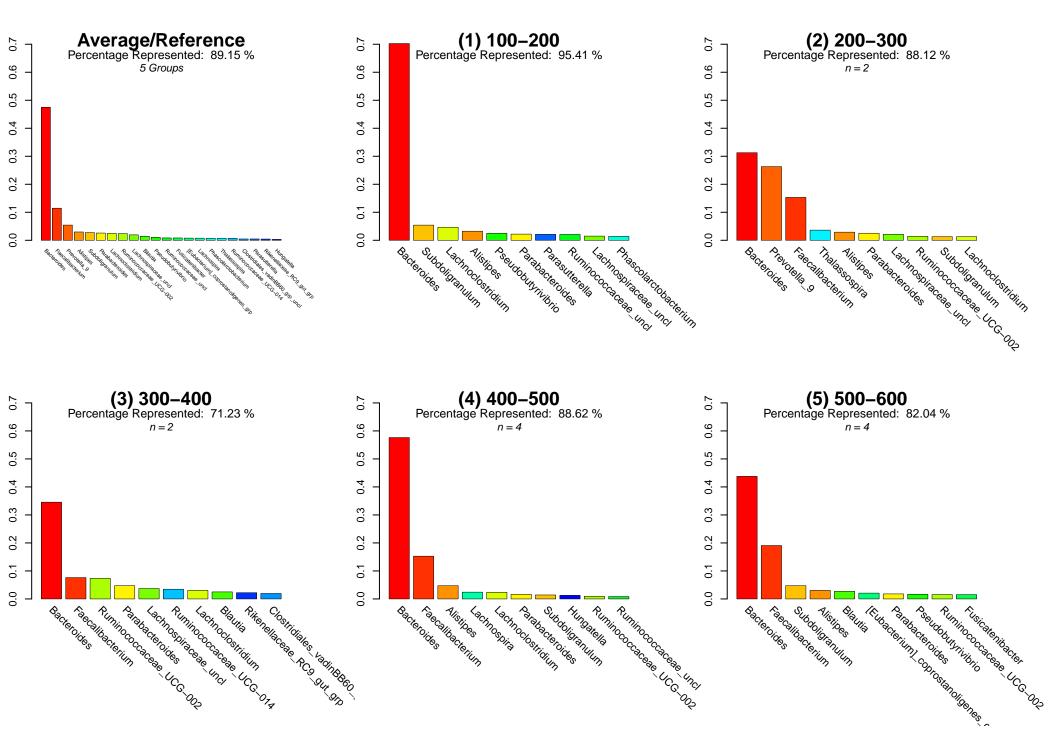
WBC



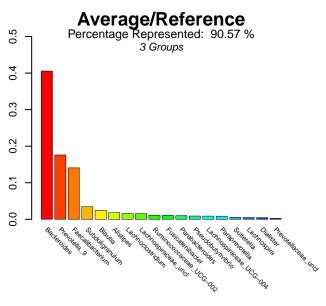


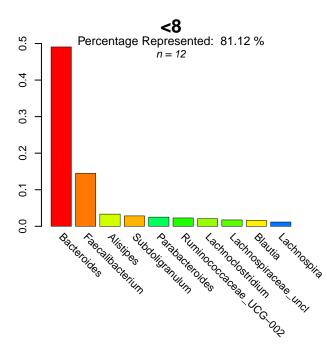


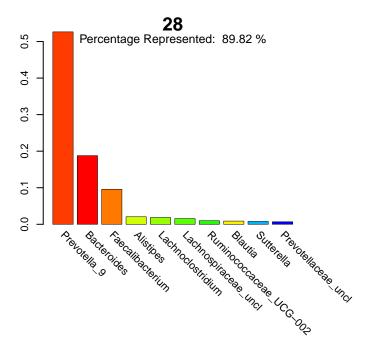
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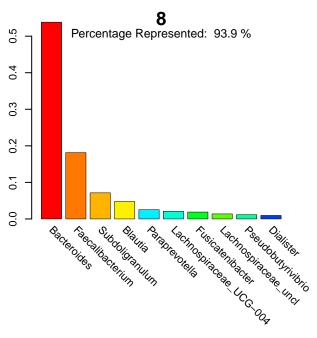


Hapto

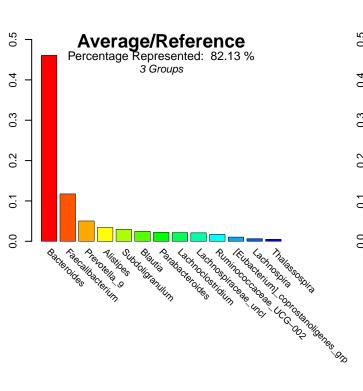


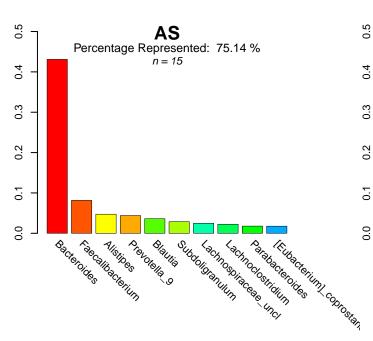


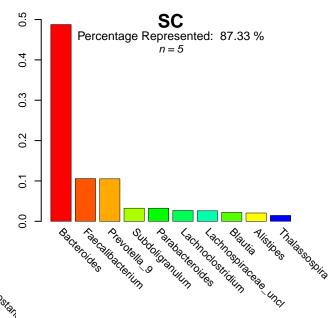


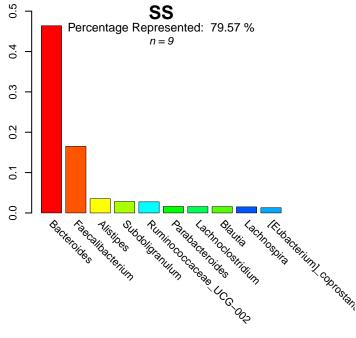


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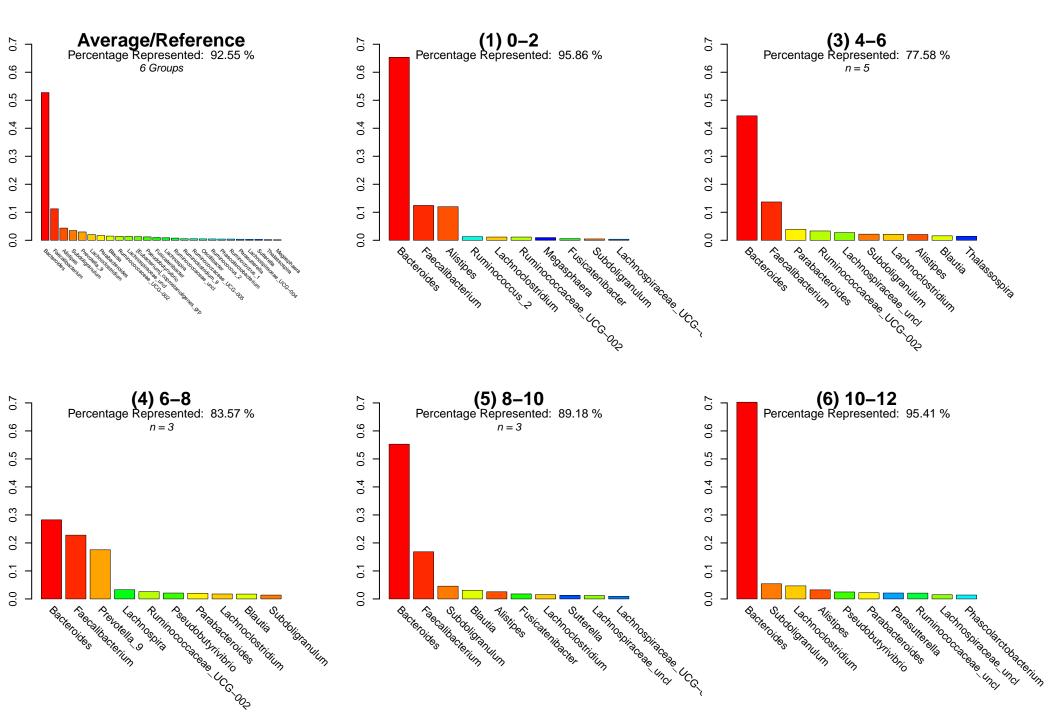




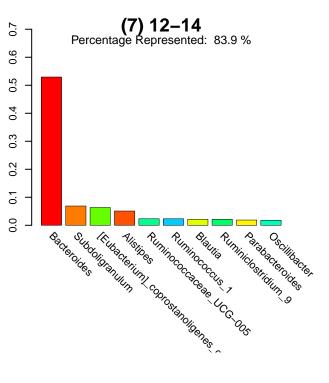




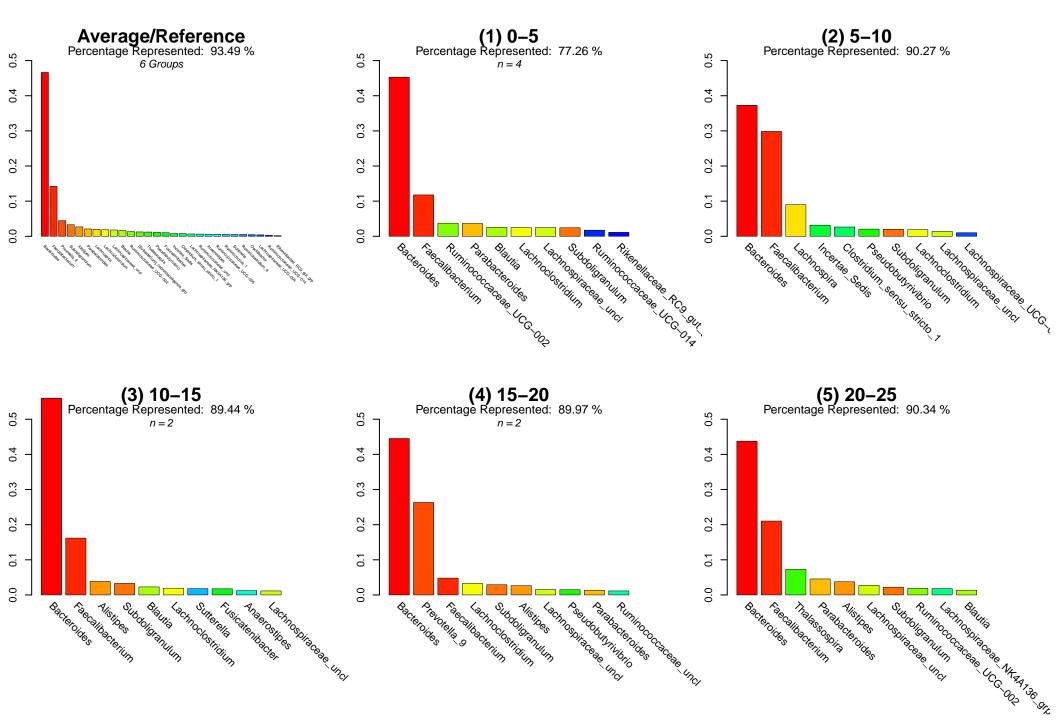
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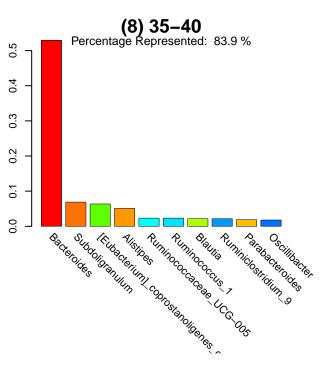
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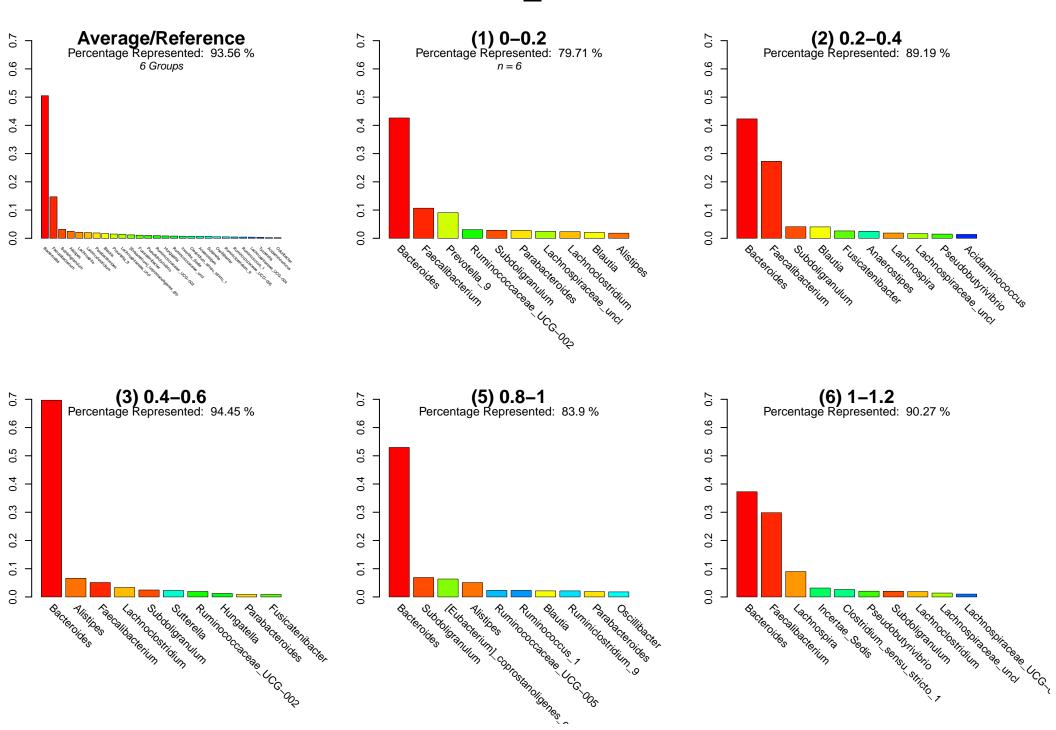
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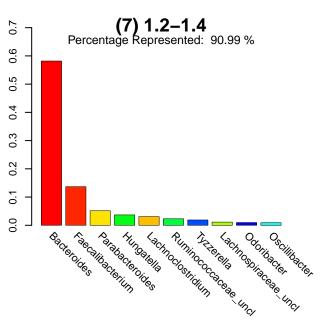
SCChi_CD16



CD62lo_CD11b



CD62lo_CD11b



CD62lo_CD184

