

Summary Table File: LimSC25.taxa.genus.NoControls.N500.summary_table.tsv
Factor File: 0137_Lim_Metadata_20171019.tsv
Output File Root: sickle.genus.tail.t4

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: 4
Number of Unique Categories across top categories extracted: 36

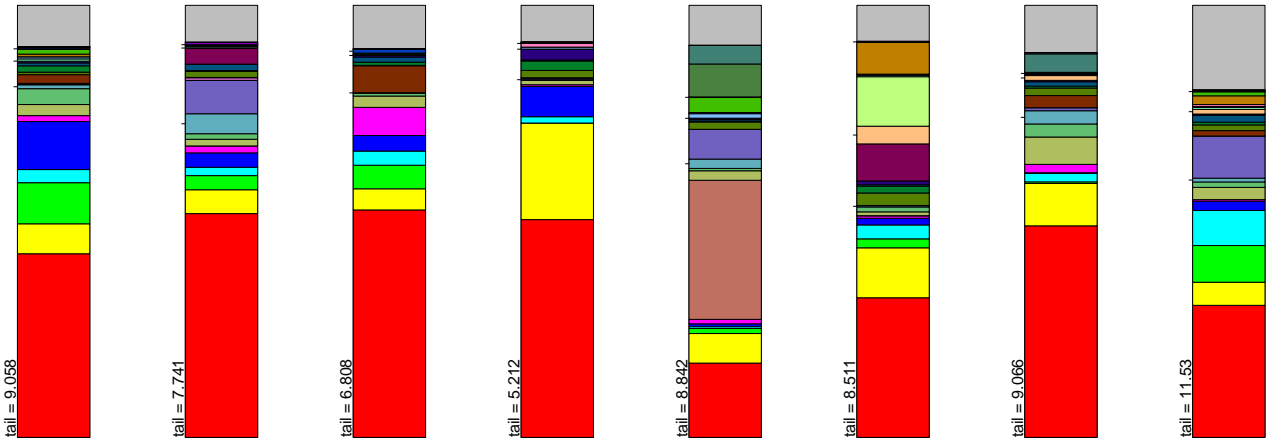
Samples exclusive to Summary Table:
character(0)

Samples exclusive to Factor Table:

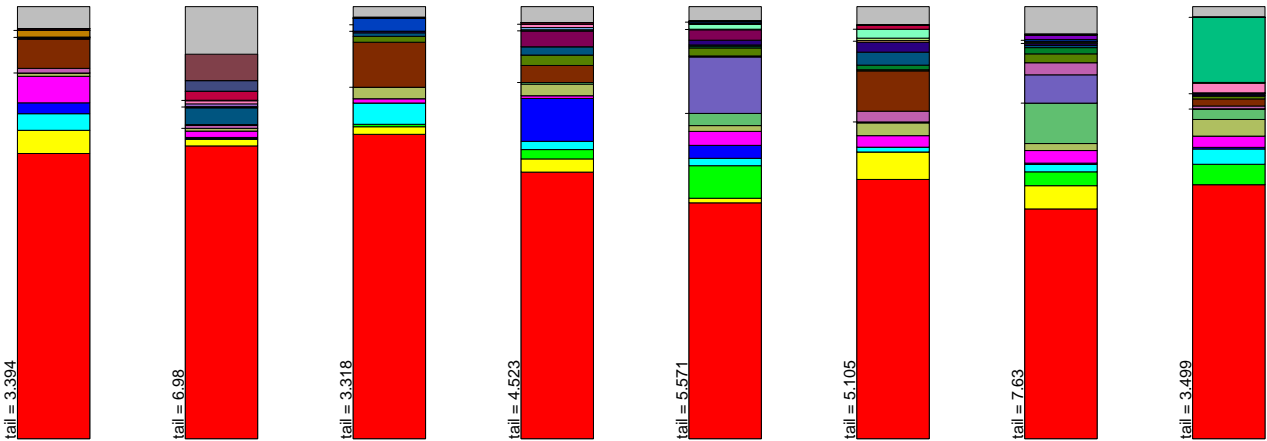
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By Sample ID

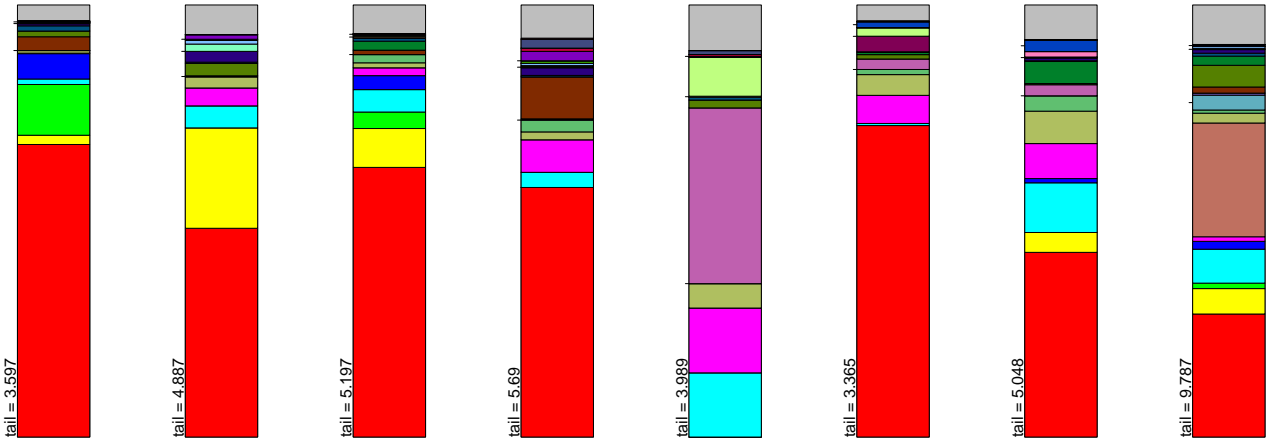
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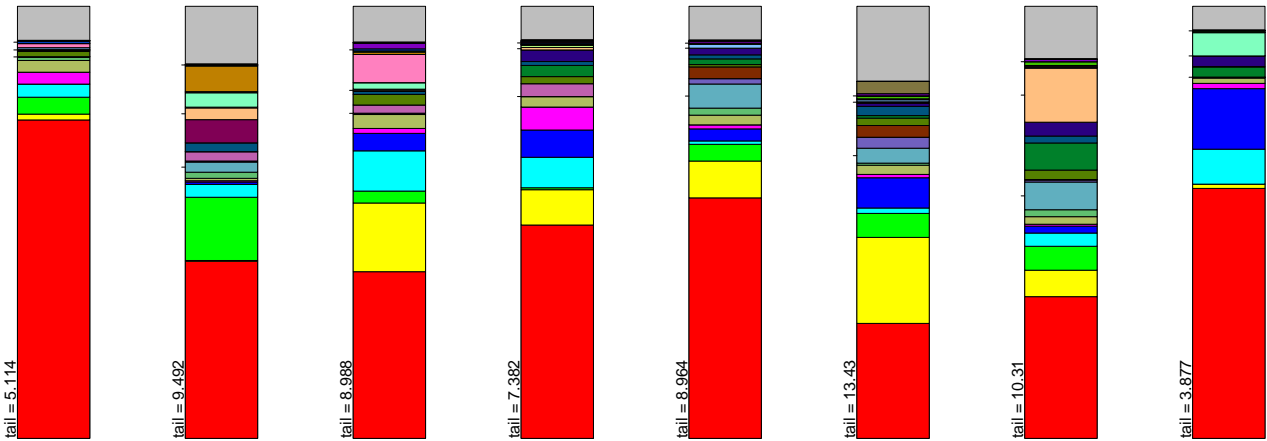
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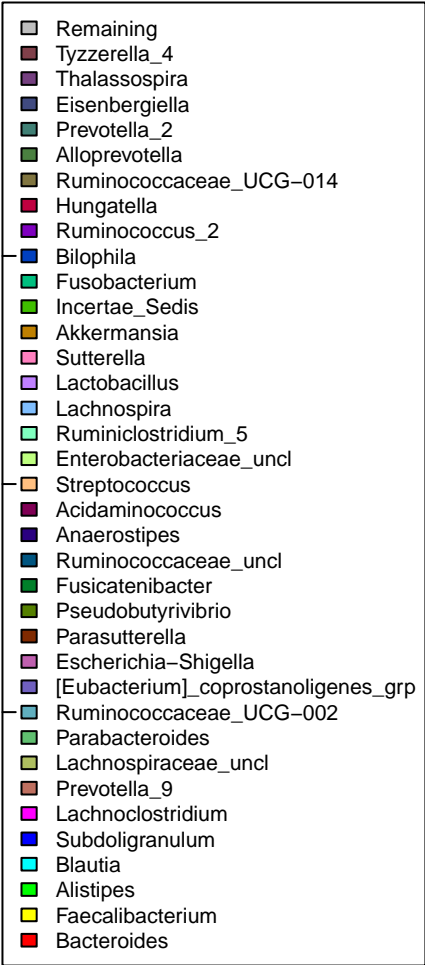
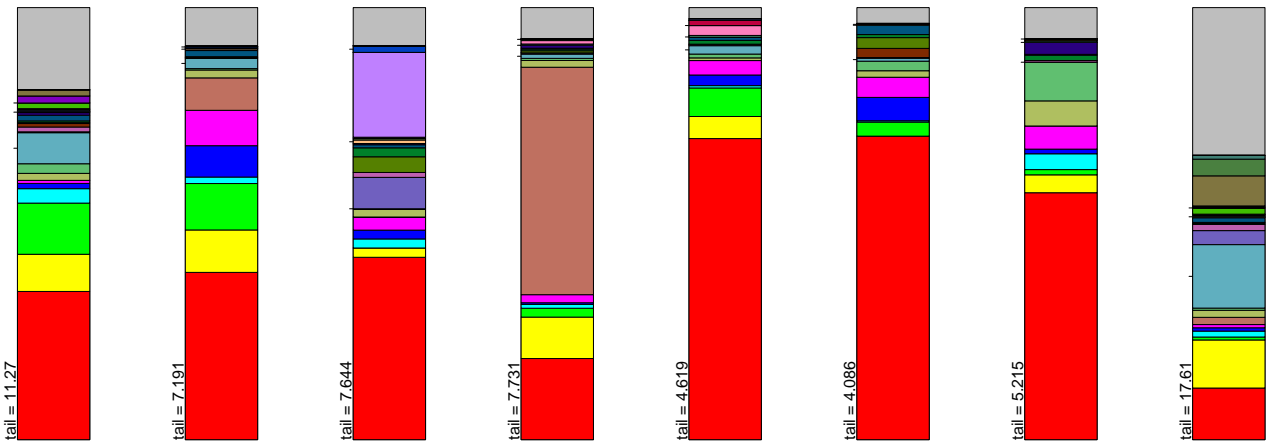
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0137.BC000343.01.ST 0137.BC000296.01.ST 0137.BC000297.01.ST 0137.BC000362.01.ST 0137.BC001412.01.ST 0137.BC001413.01.ST 0137.BC001415.01.ST 0137.BC001418.01.ST

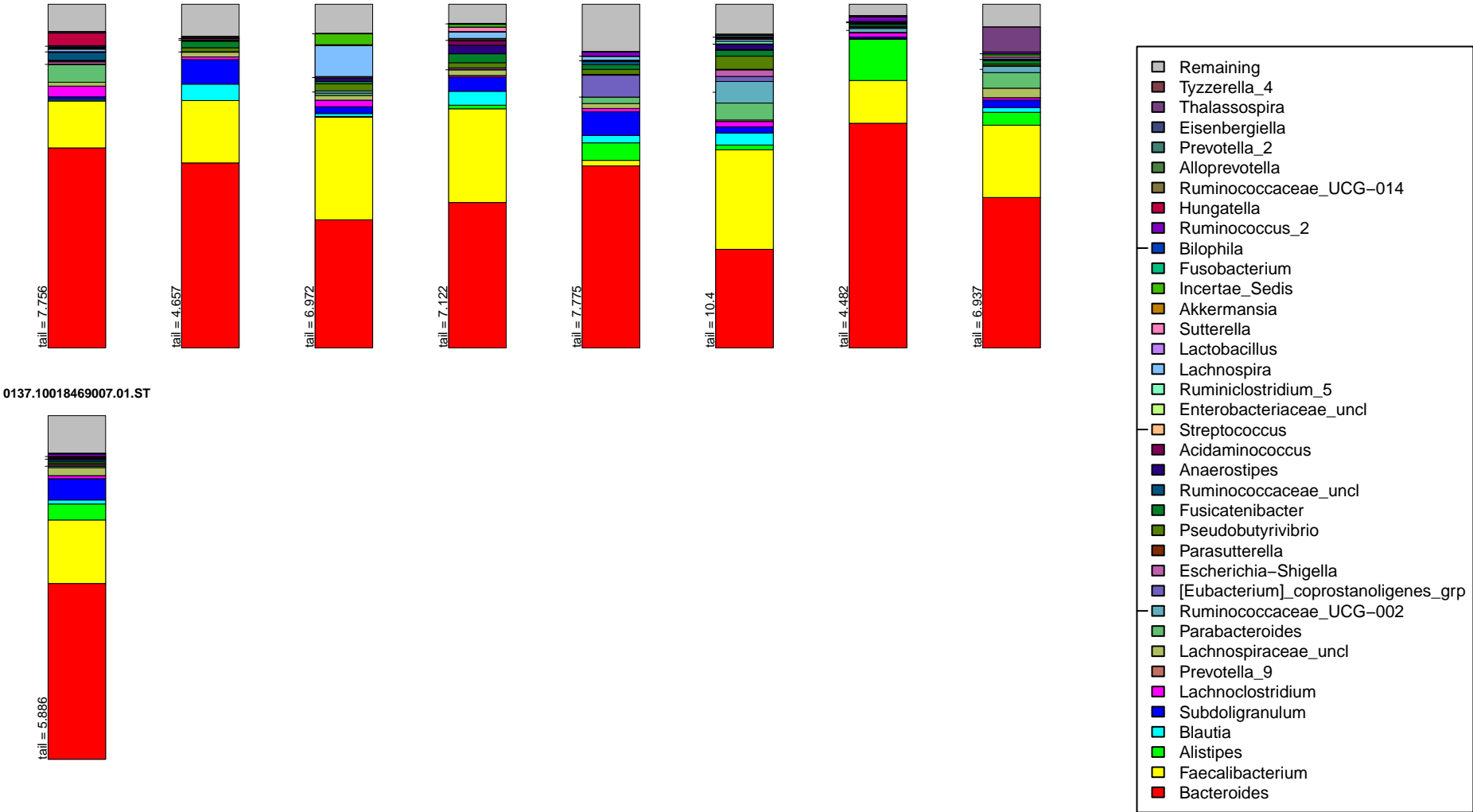


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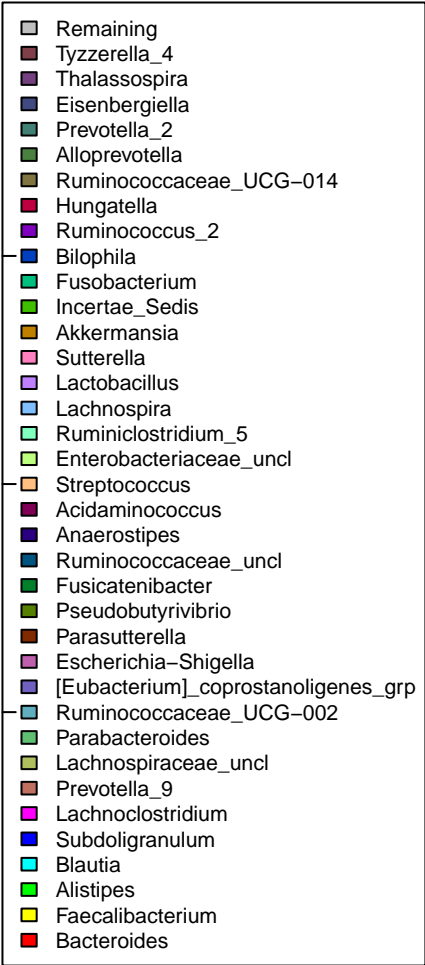
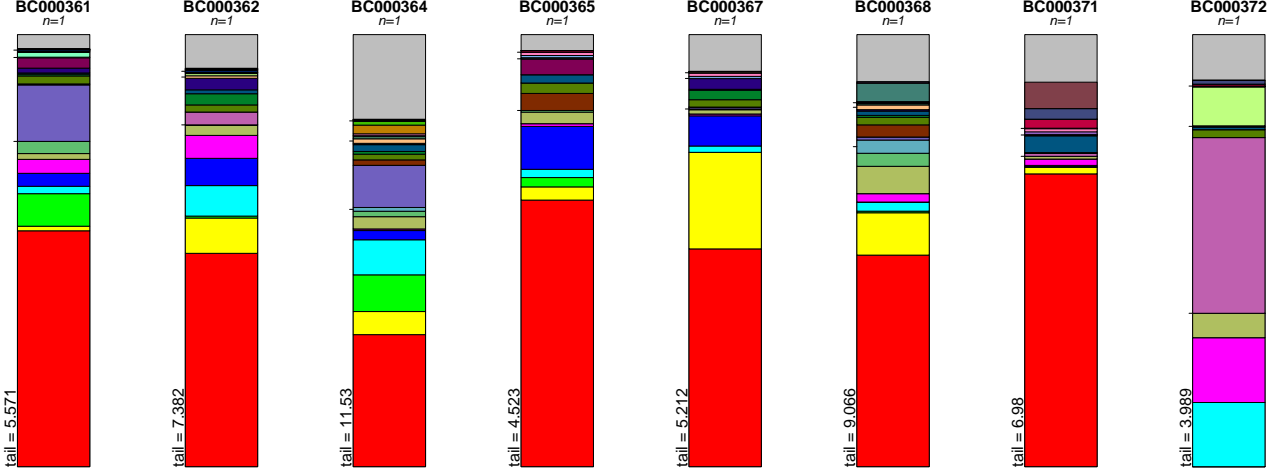
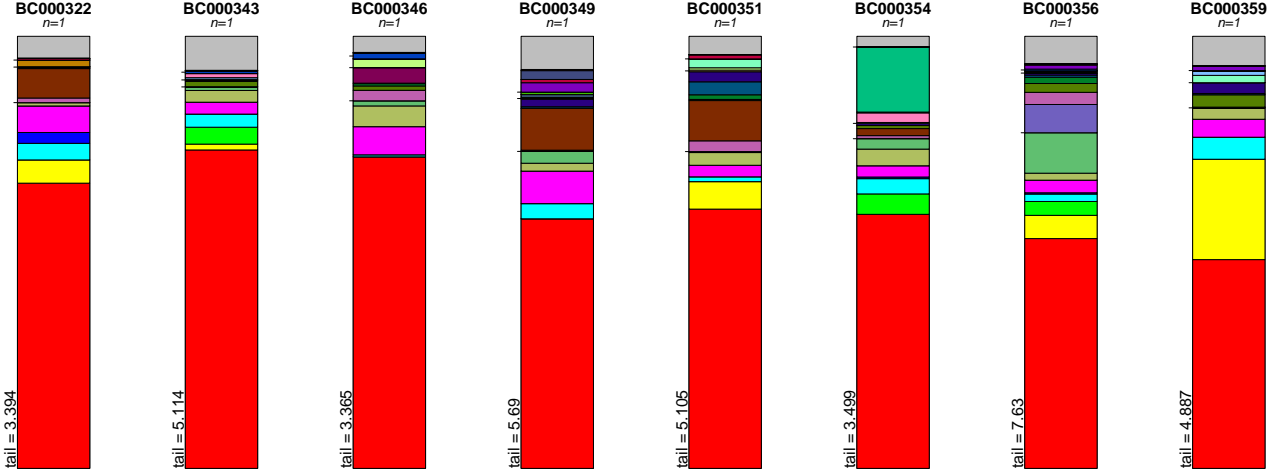
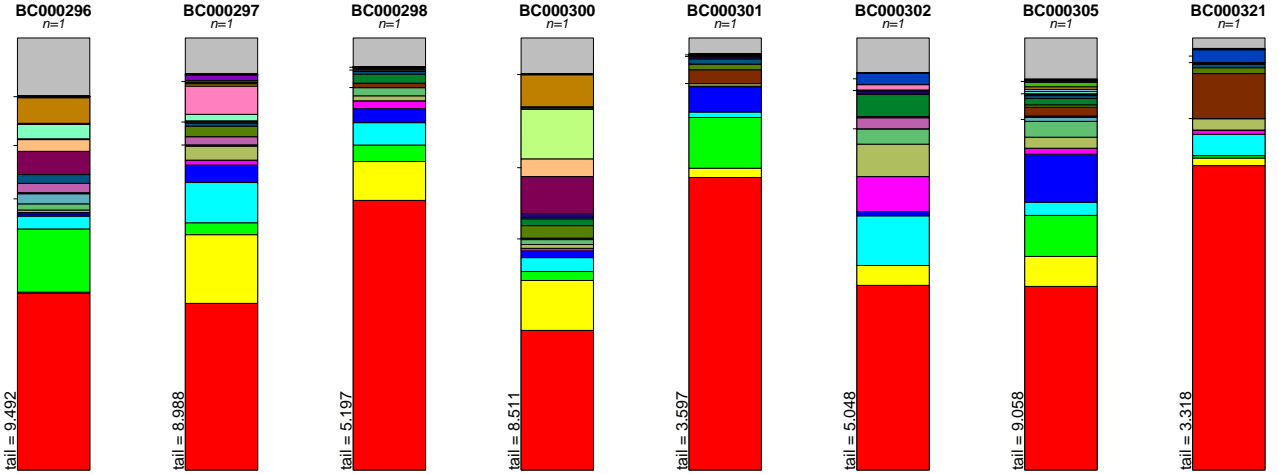
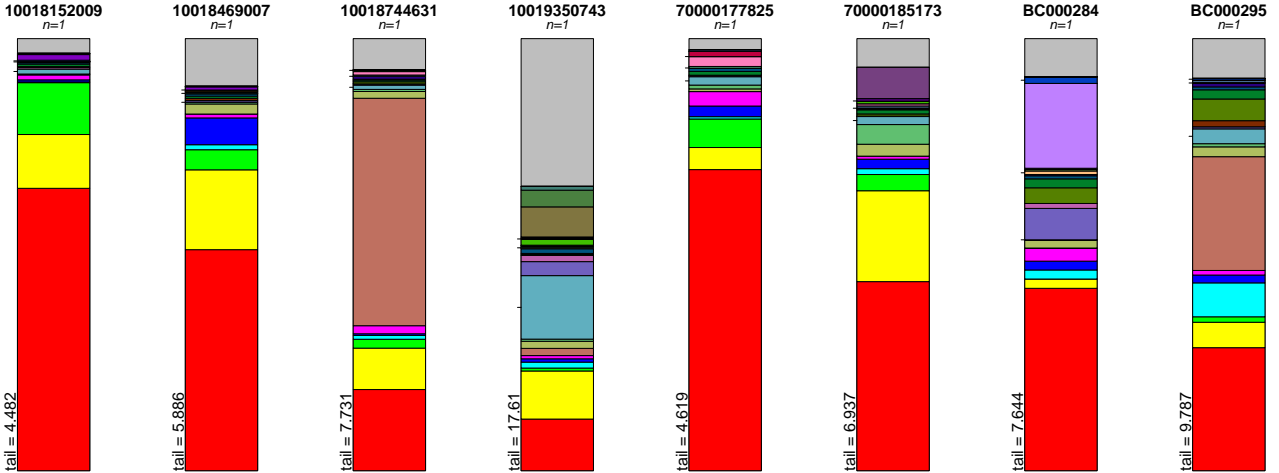
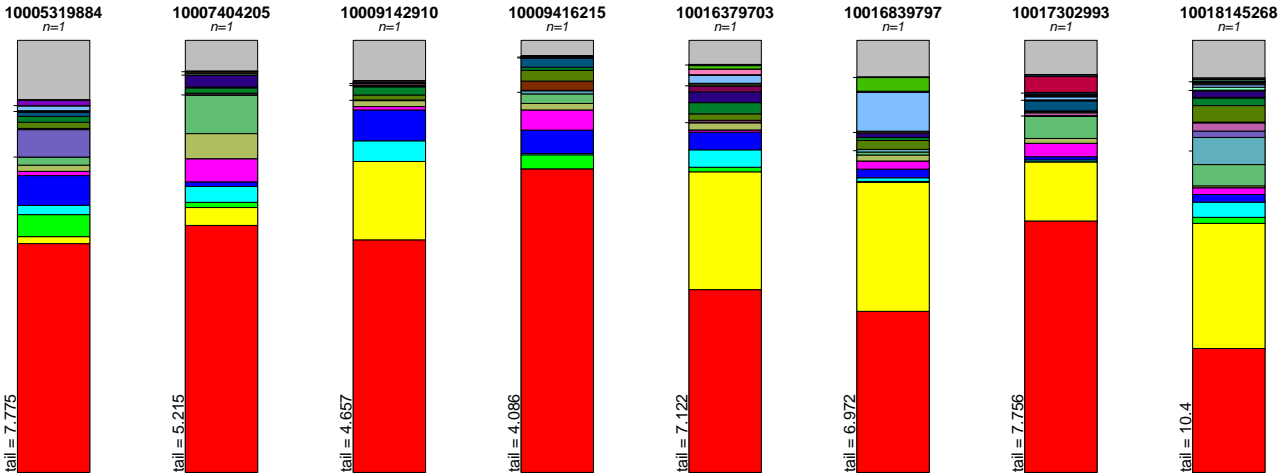


By Sample ID

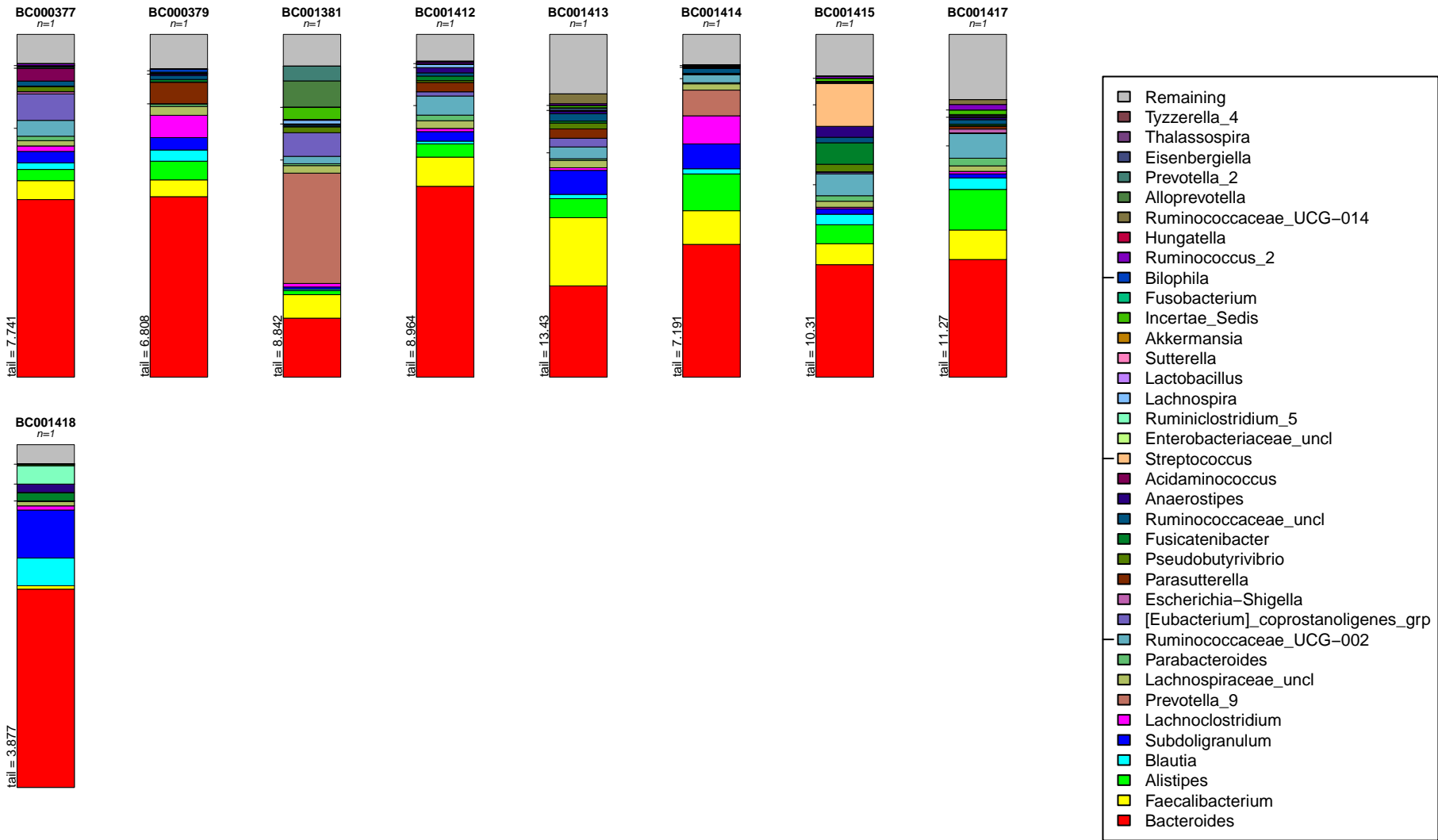
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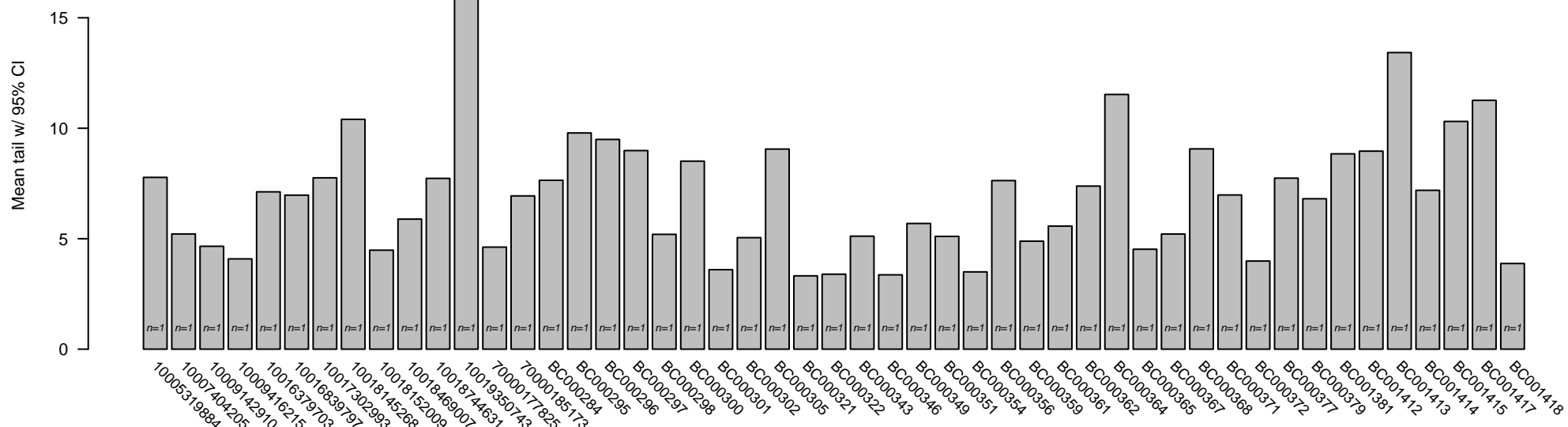
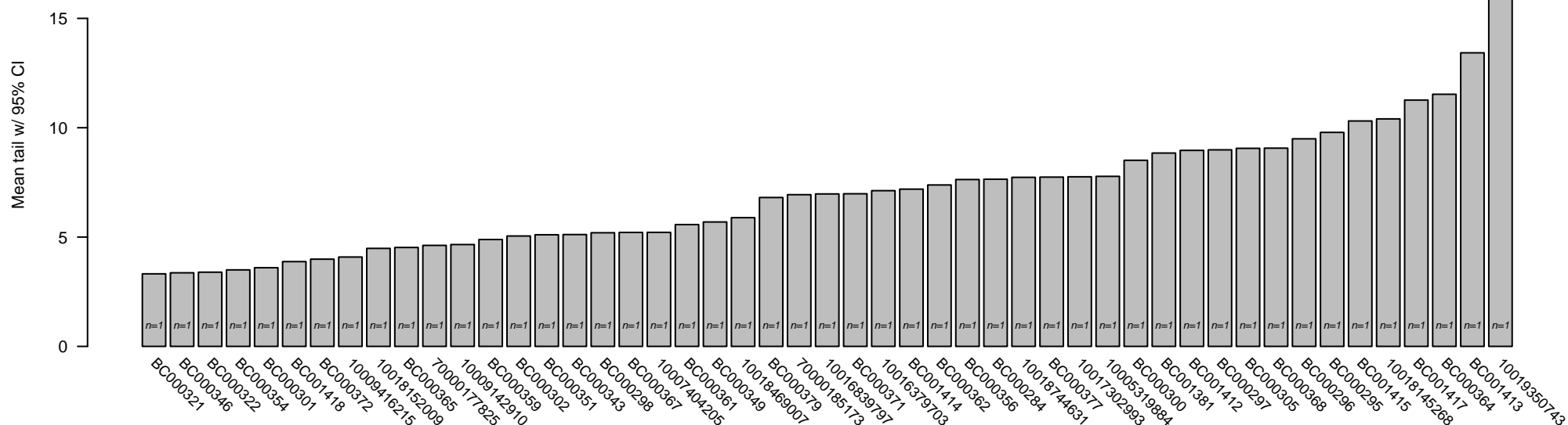
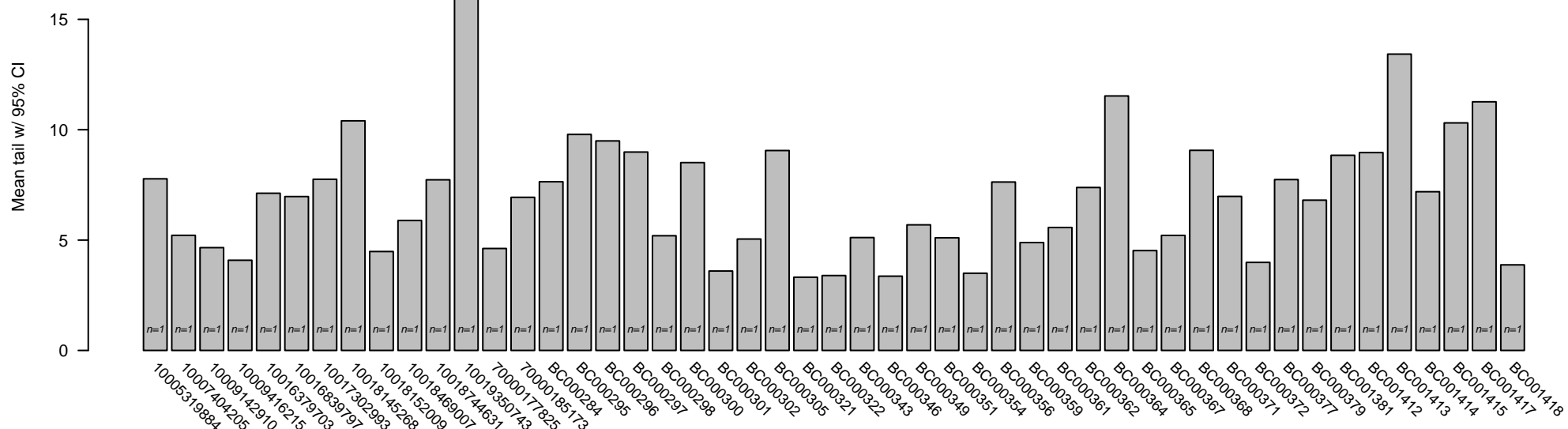
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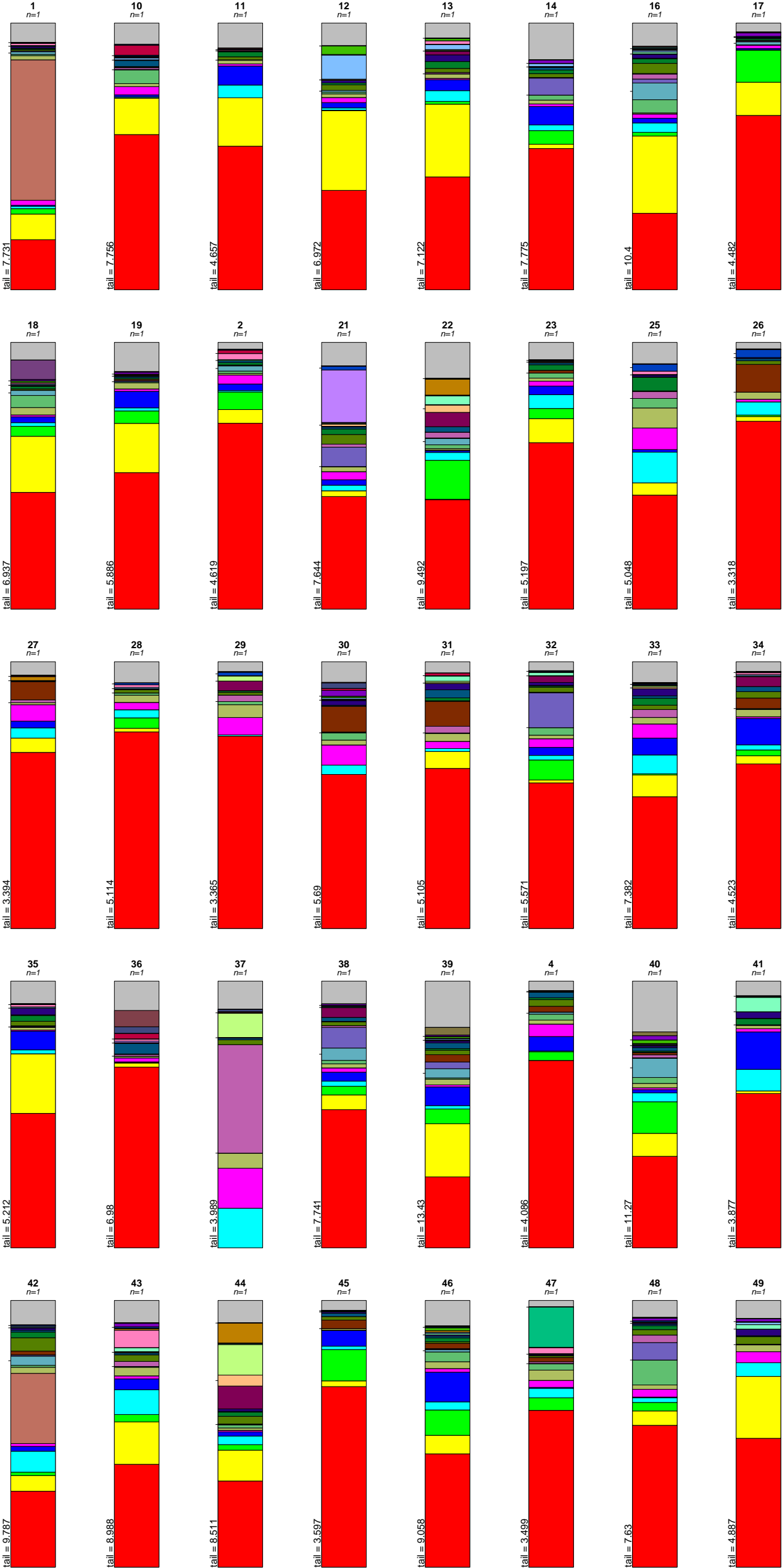
Identifier



Identifier: Ordered by Category

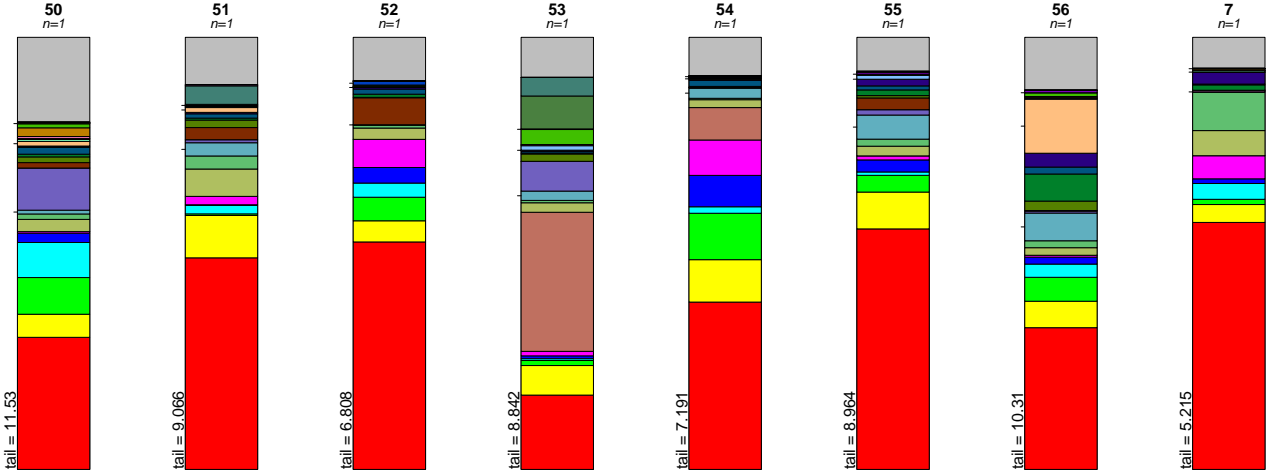
**Identifier: Ordered by Diversity****Identifier: Ordered by Sample Size**

RIH_UPN



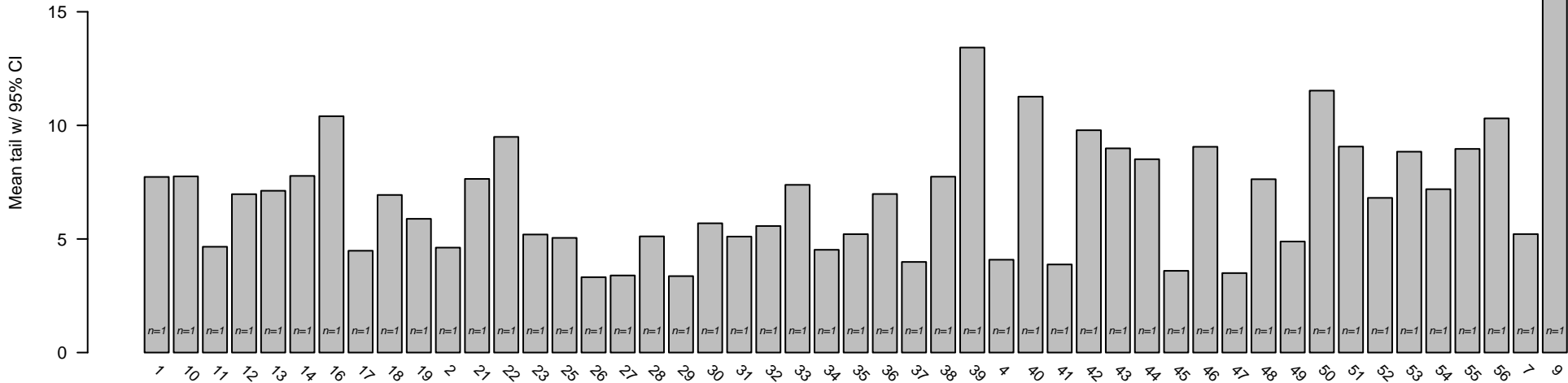
- Remaining
- Tyzzrella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

RIH_UPN

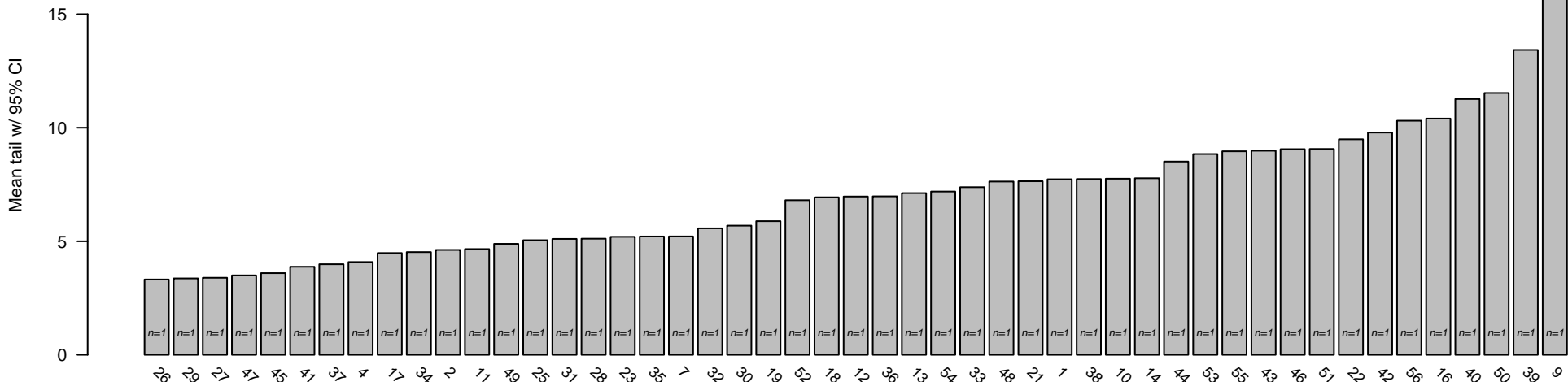


- Remaining
- Tyzzarella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

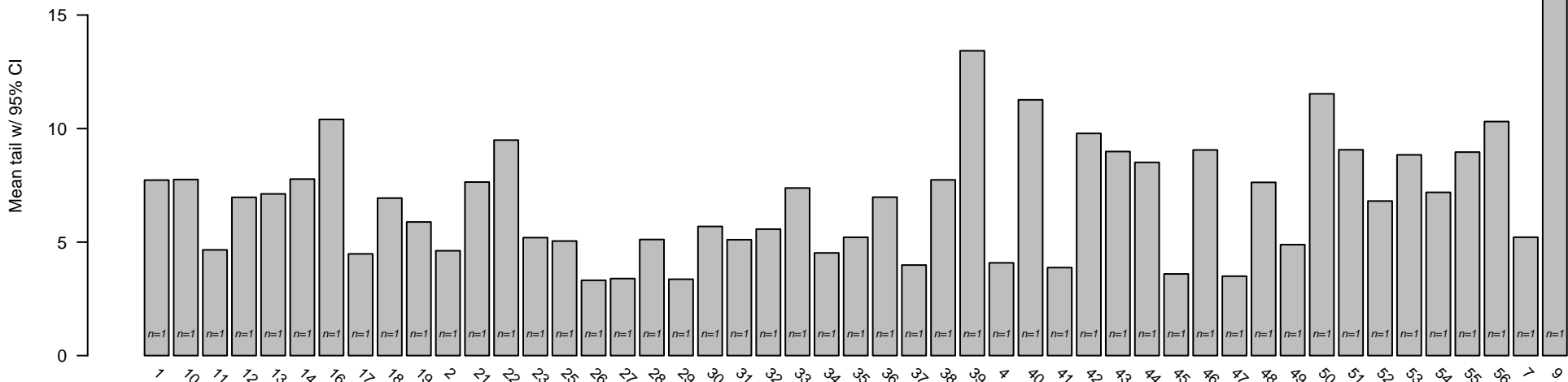
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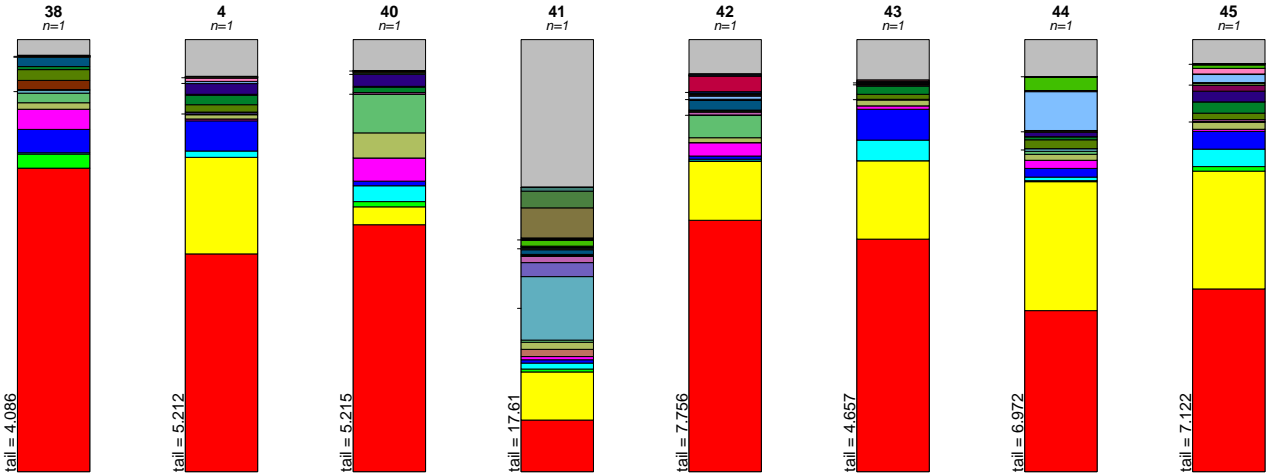
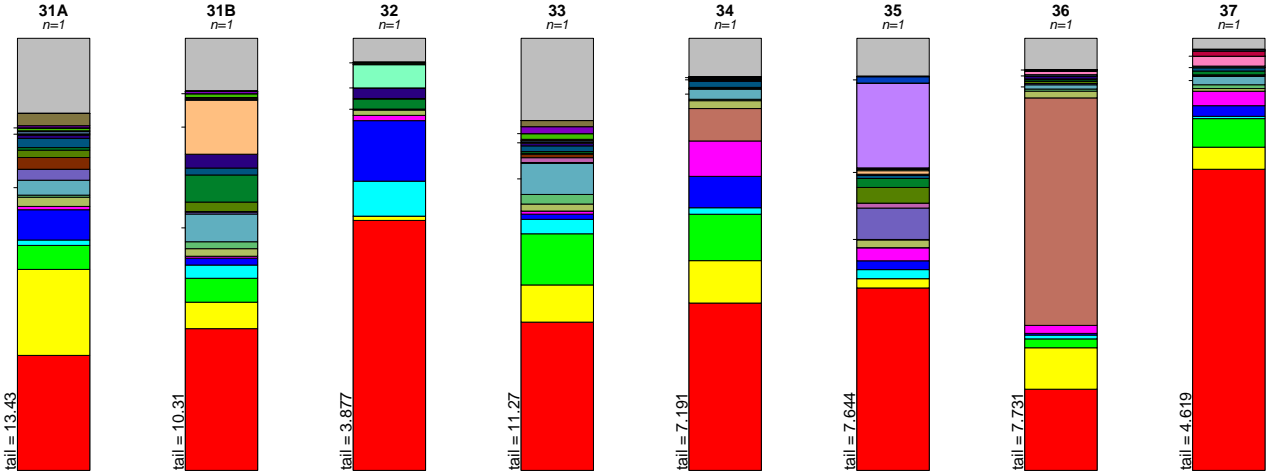
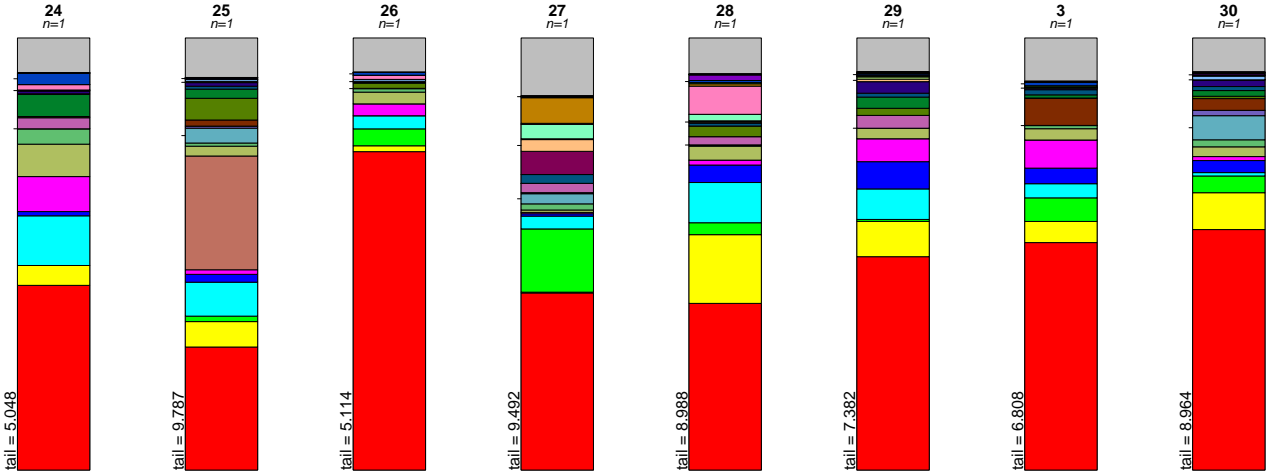
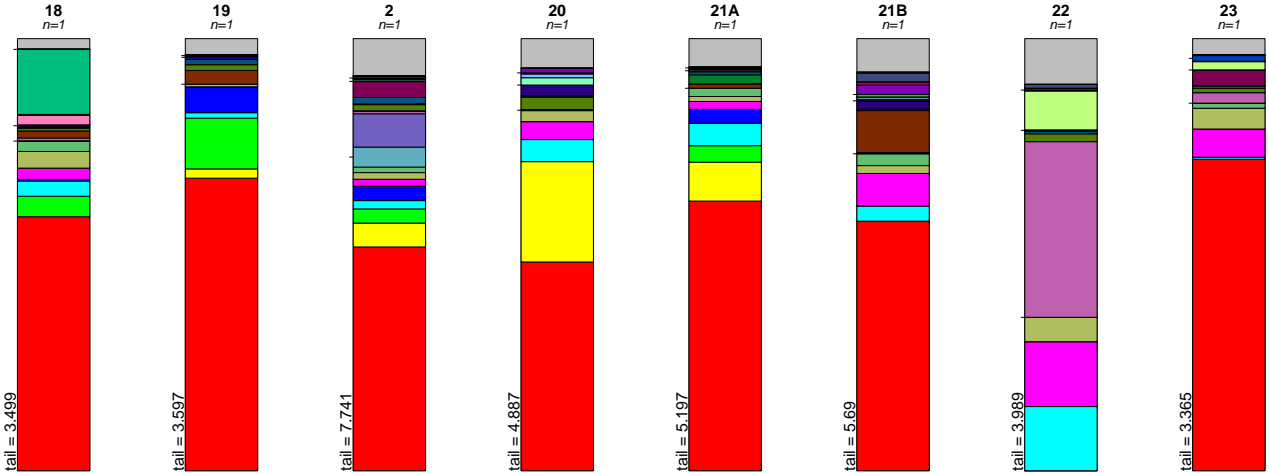
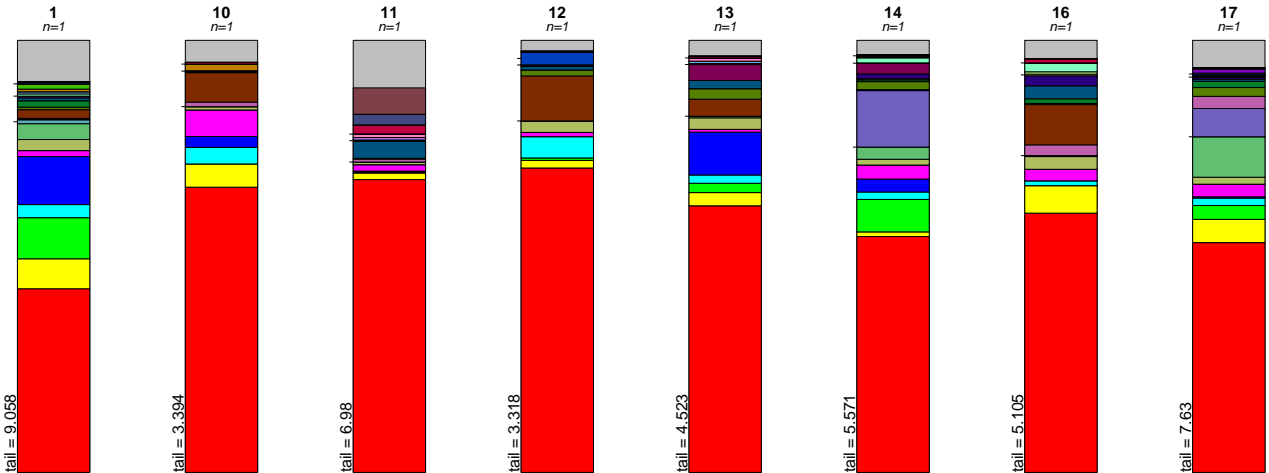
RIH_UPN: Ordered by Diversity



RIH_UPN: Ordered by Sample Size

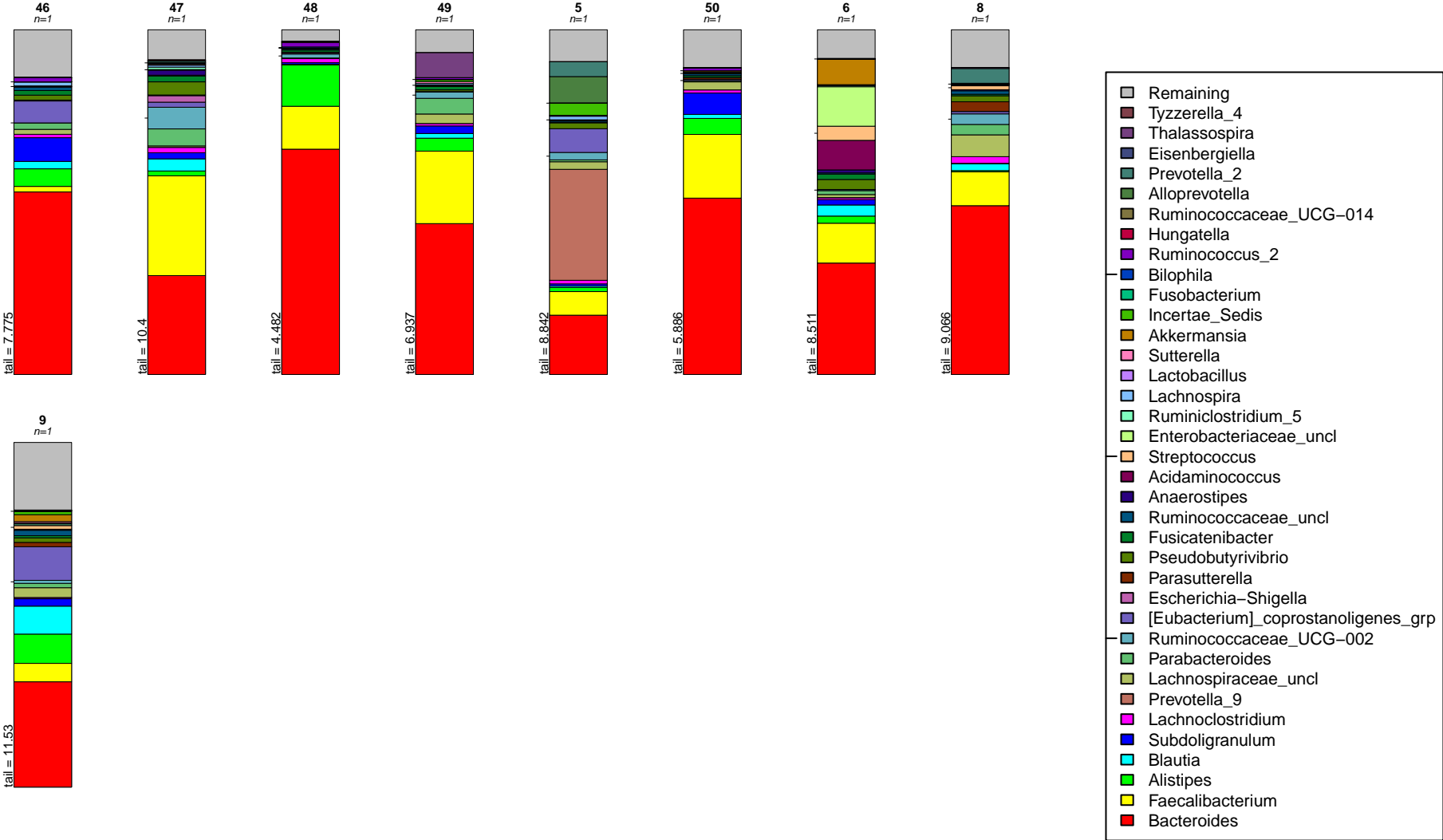


PCR_Temporary_Number

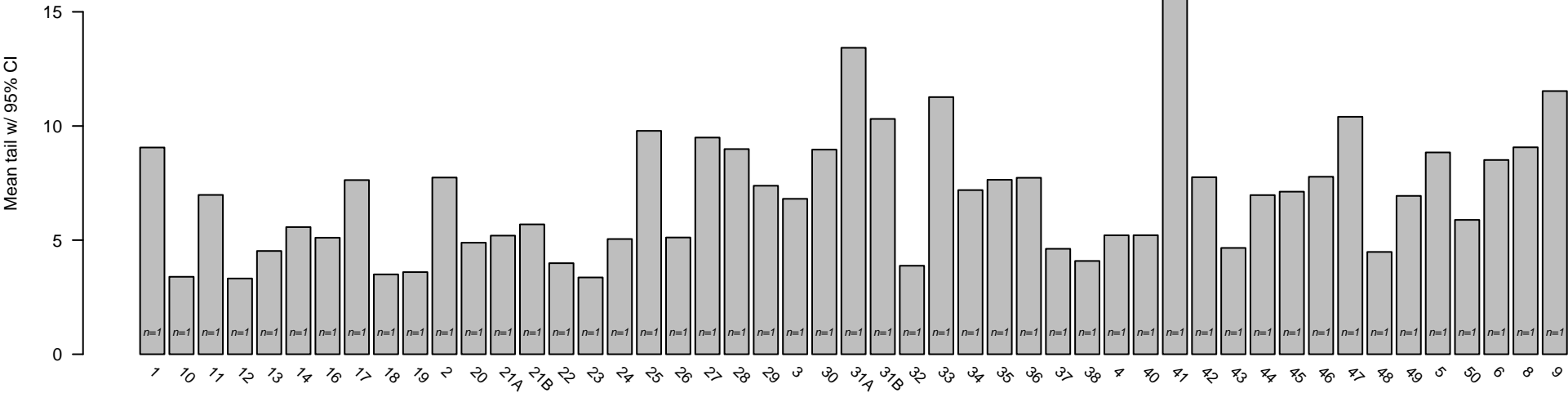


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

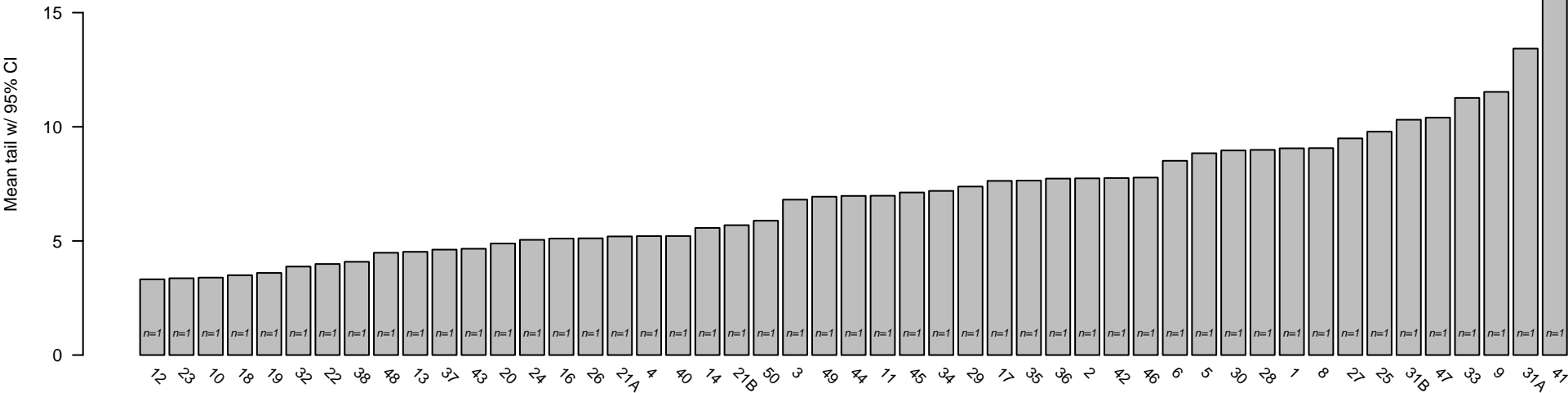
PCR_Temporary_Number



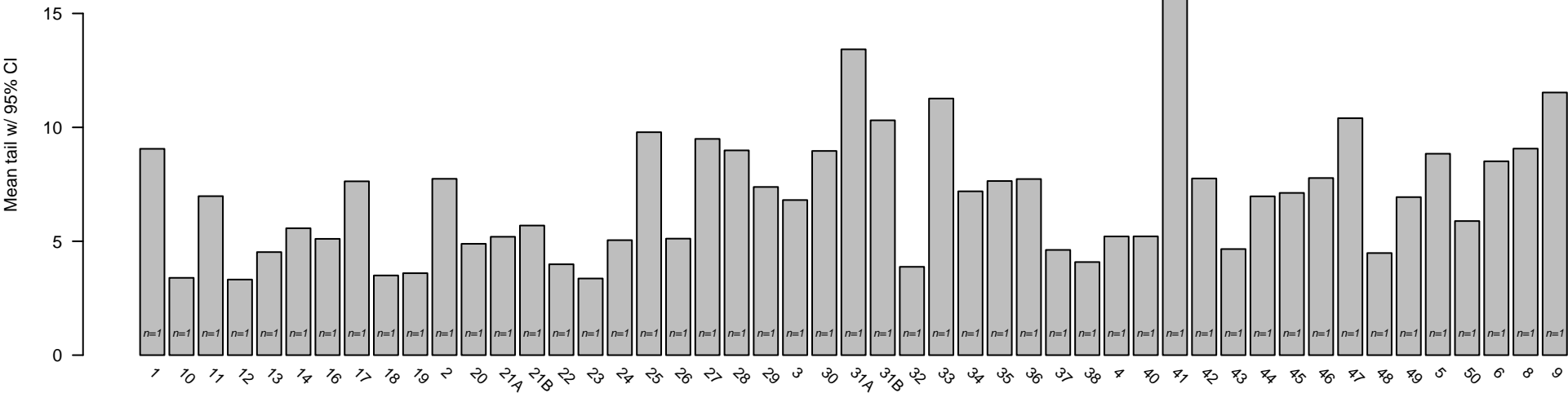
PCR_Temporary_Number: Ordered by Category



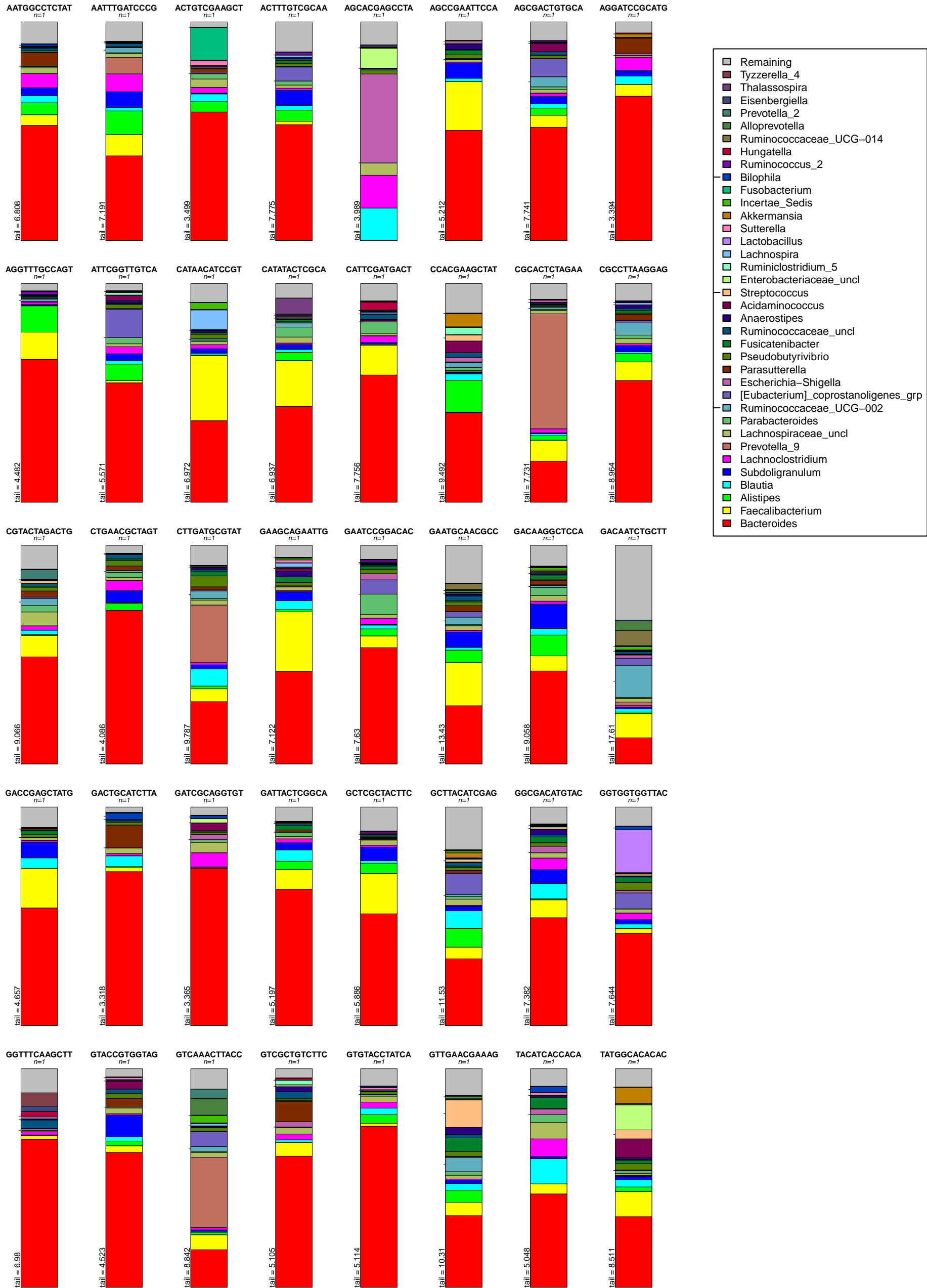
PCR_Temporary_Number: Ordered by Diversity



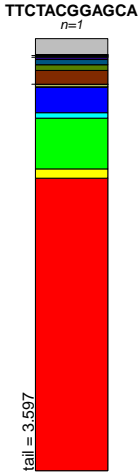
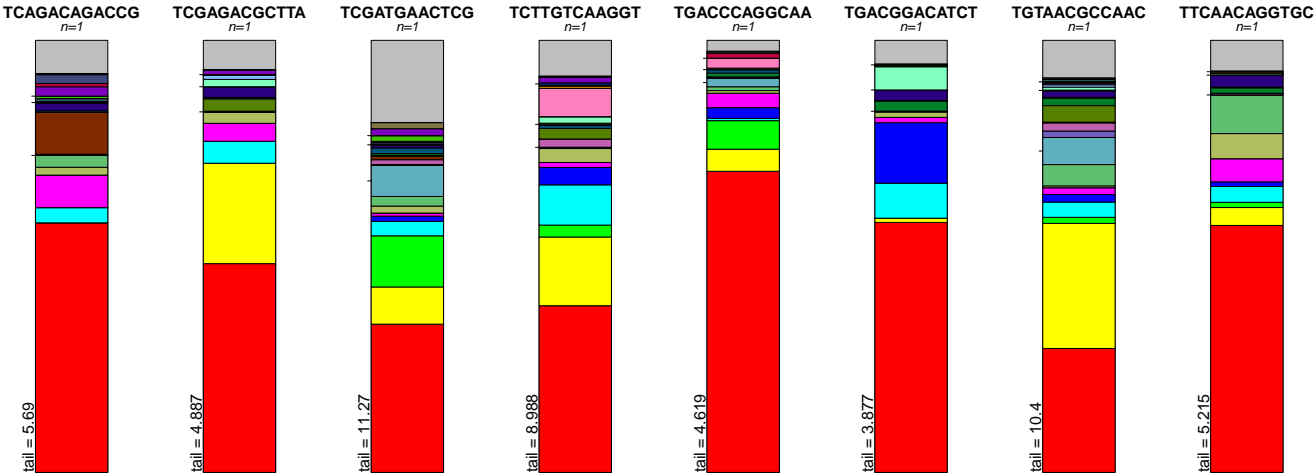
PCR_Temporary_Number: Ordered by Sample Size



Barcode.RC

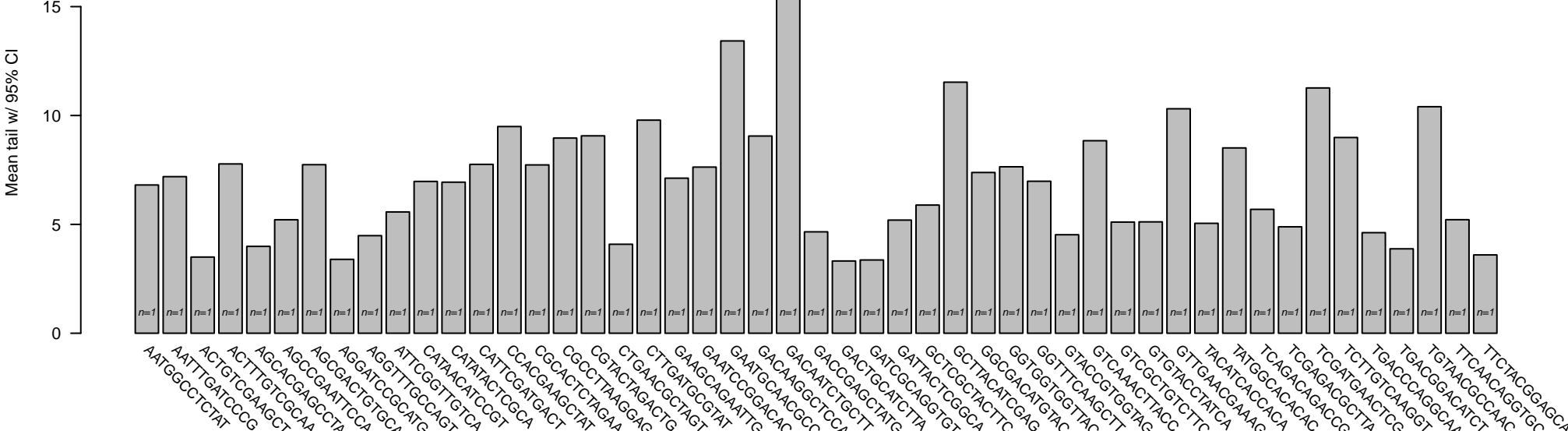


Barcode.RC

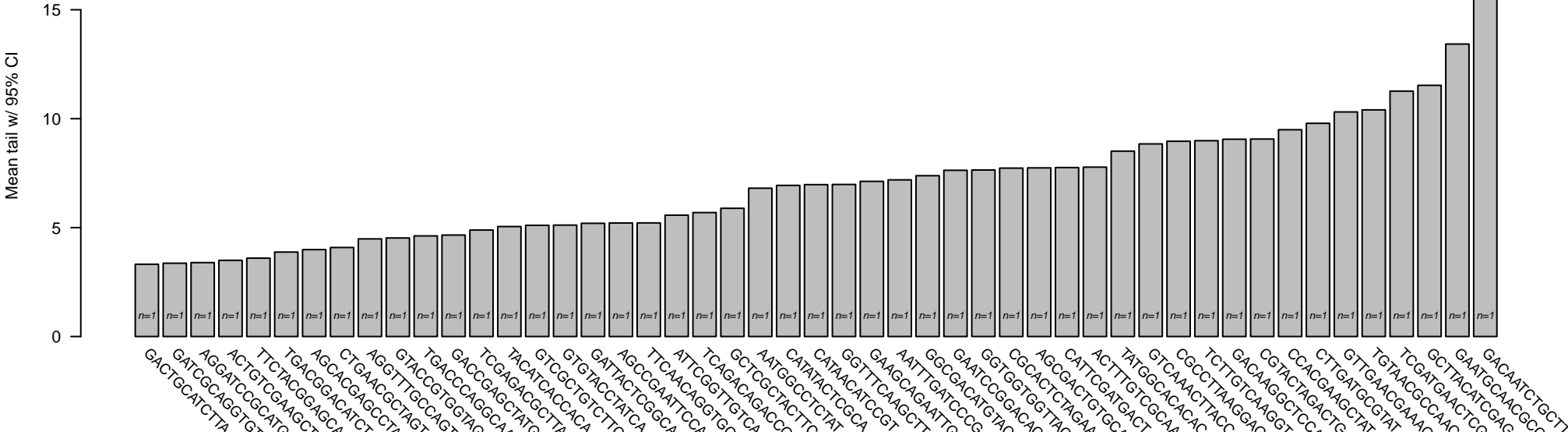


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

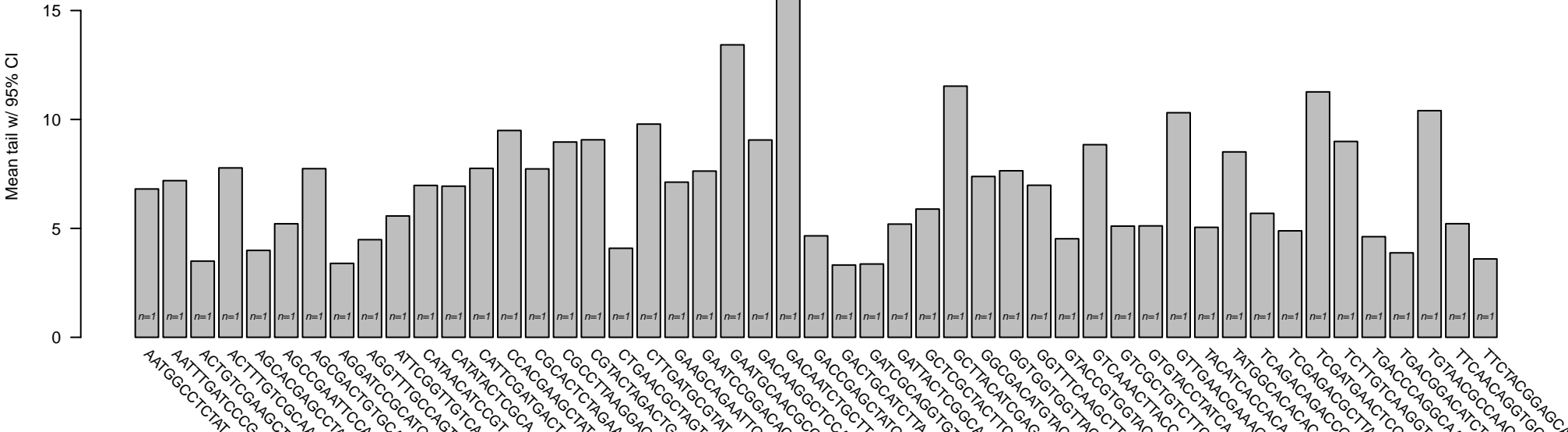
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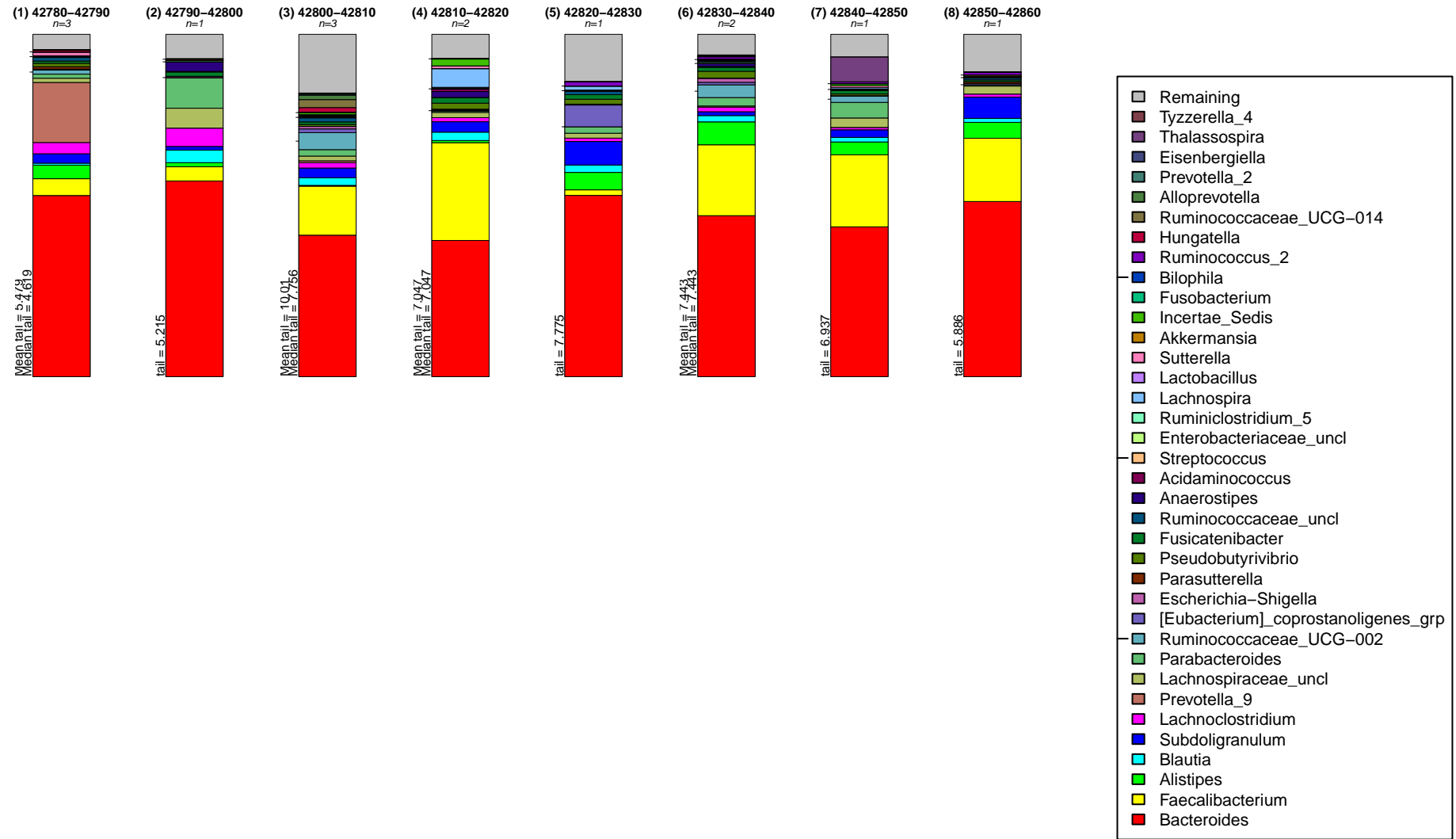
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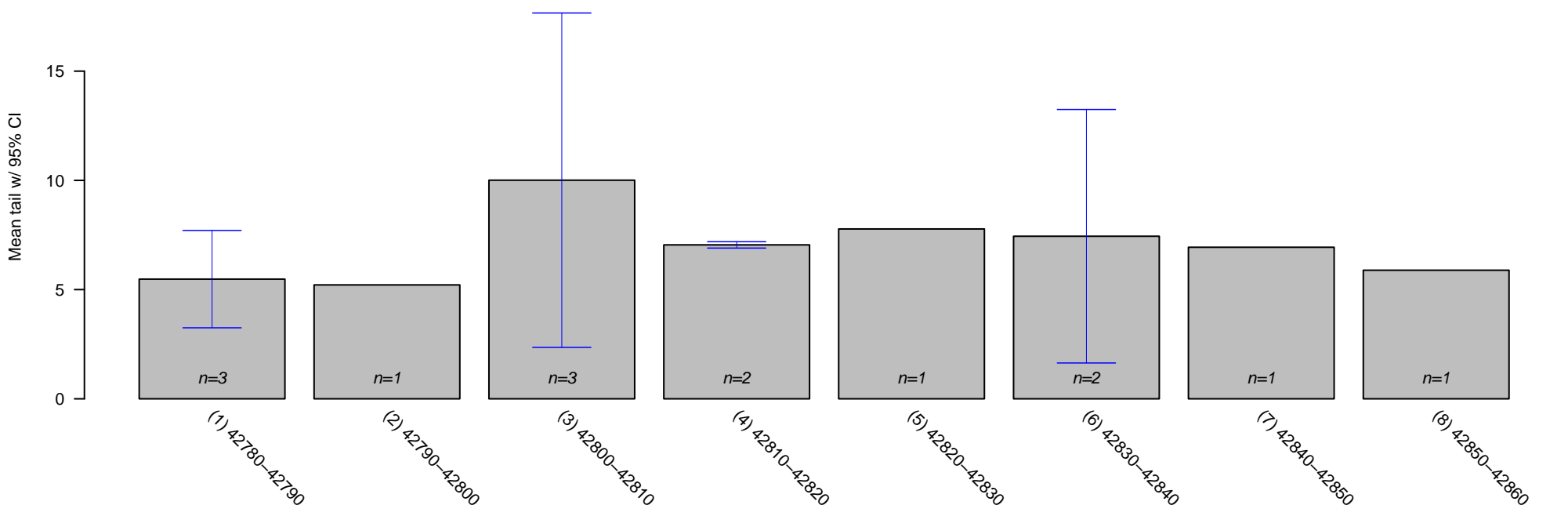
Barcode-RC: Ordered by Sample Size



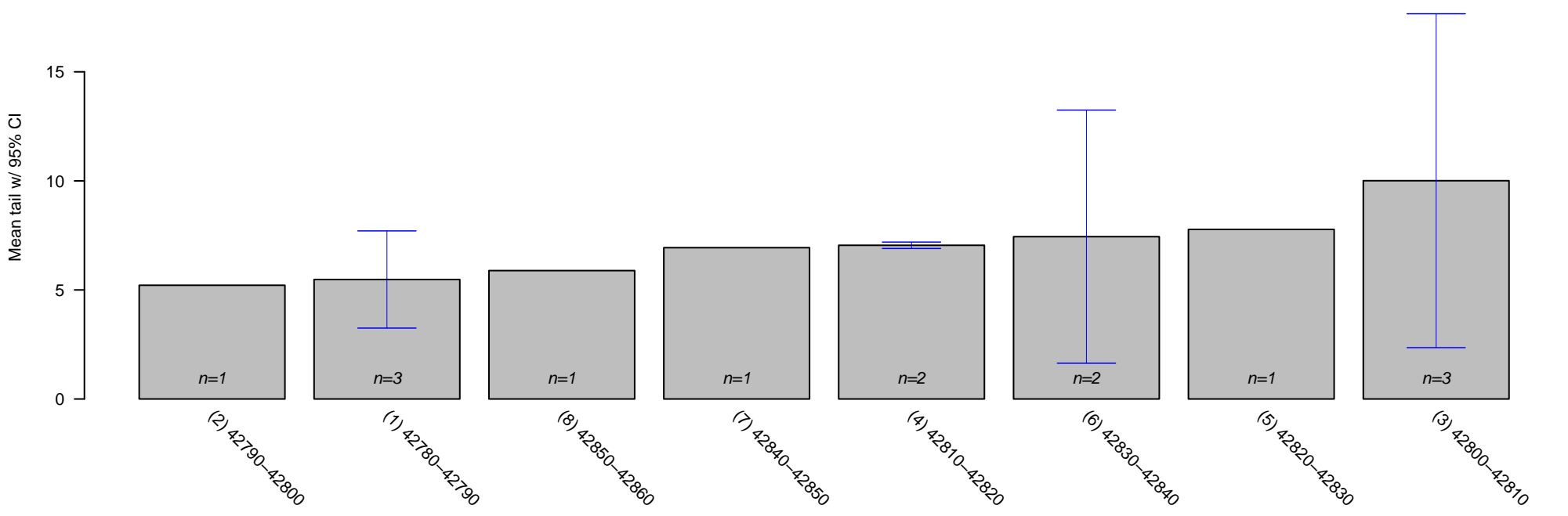
Consent.Date



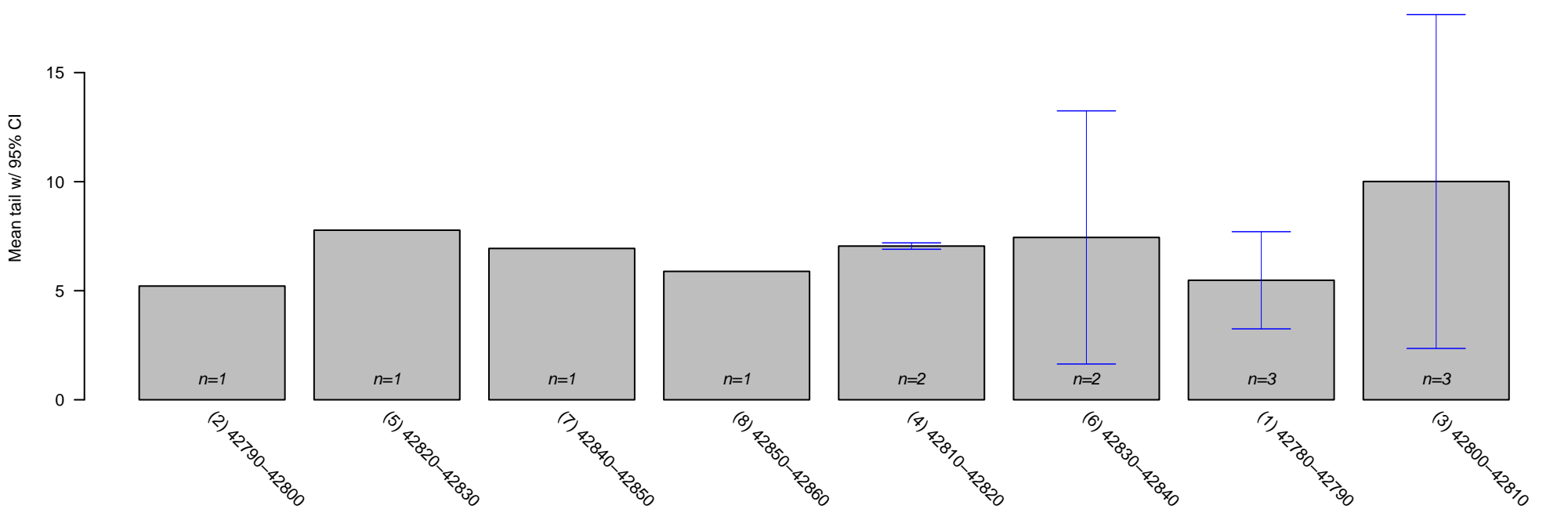
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Consent.Date: Ordered by Diversity

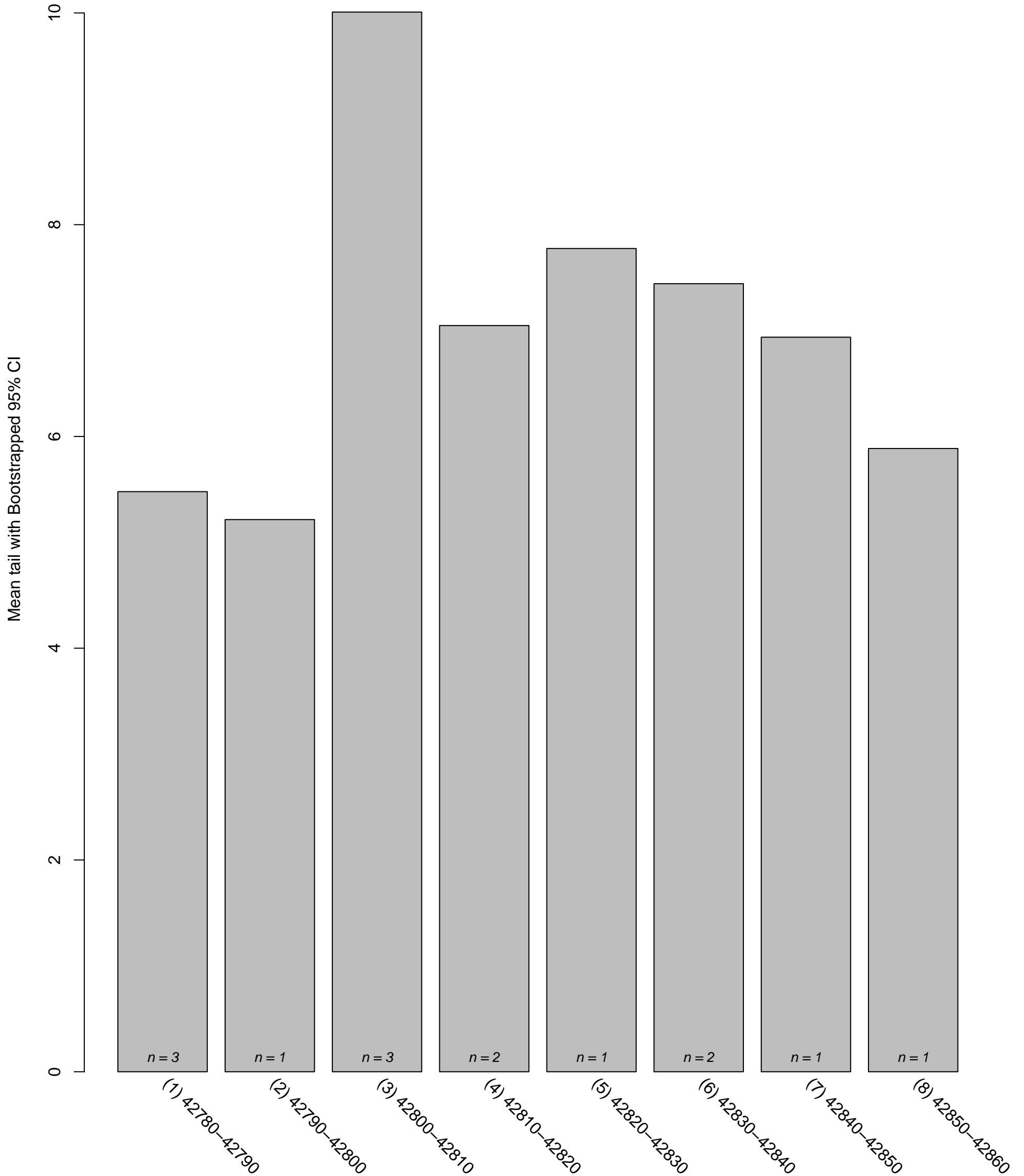


Consent.Date: Ordered by Sample Size

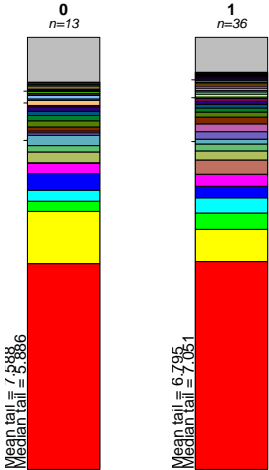


Consent.Date: Ordered by Category

with Wilcoxon rank sum test (difference between group means) p-values

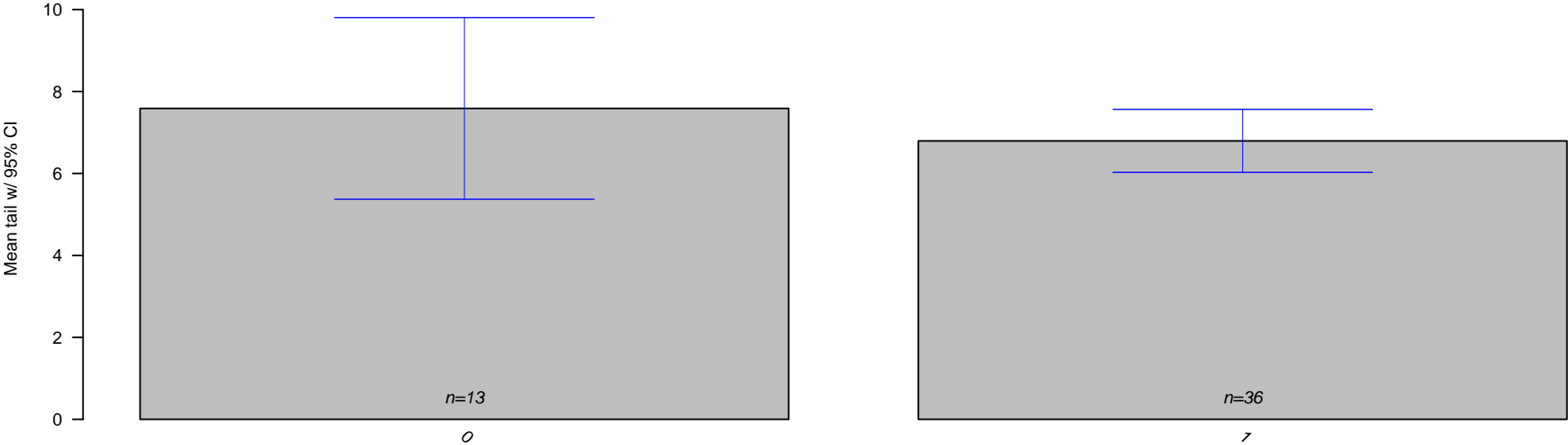


isFemale

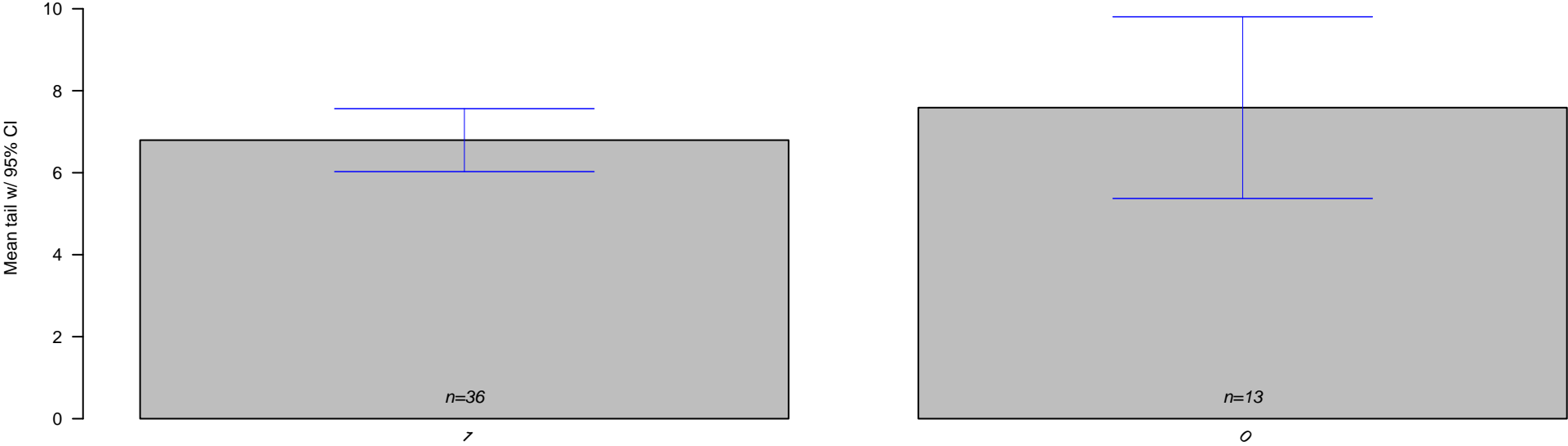


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

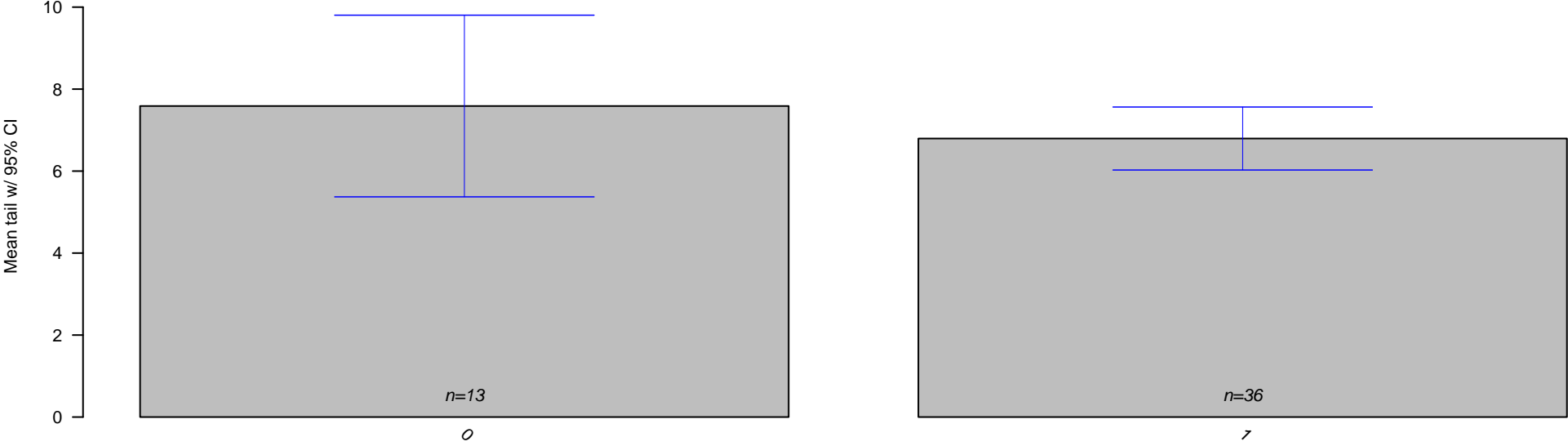
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isFemale: Ordered by Diversity

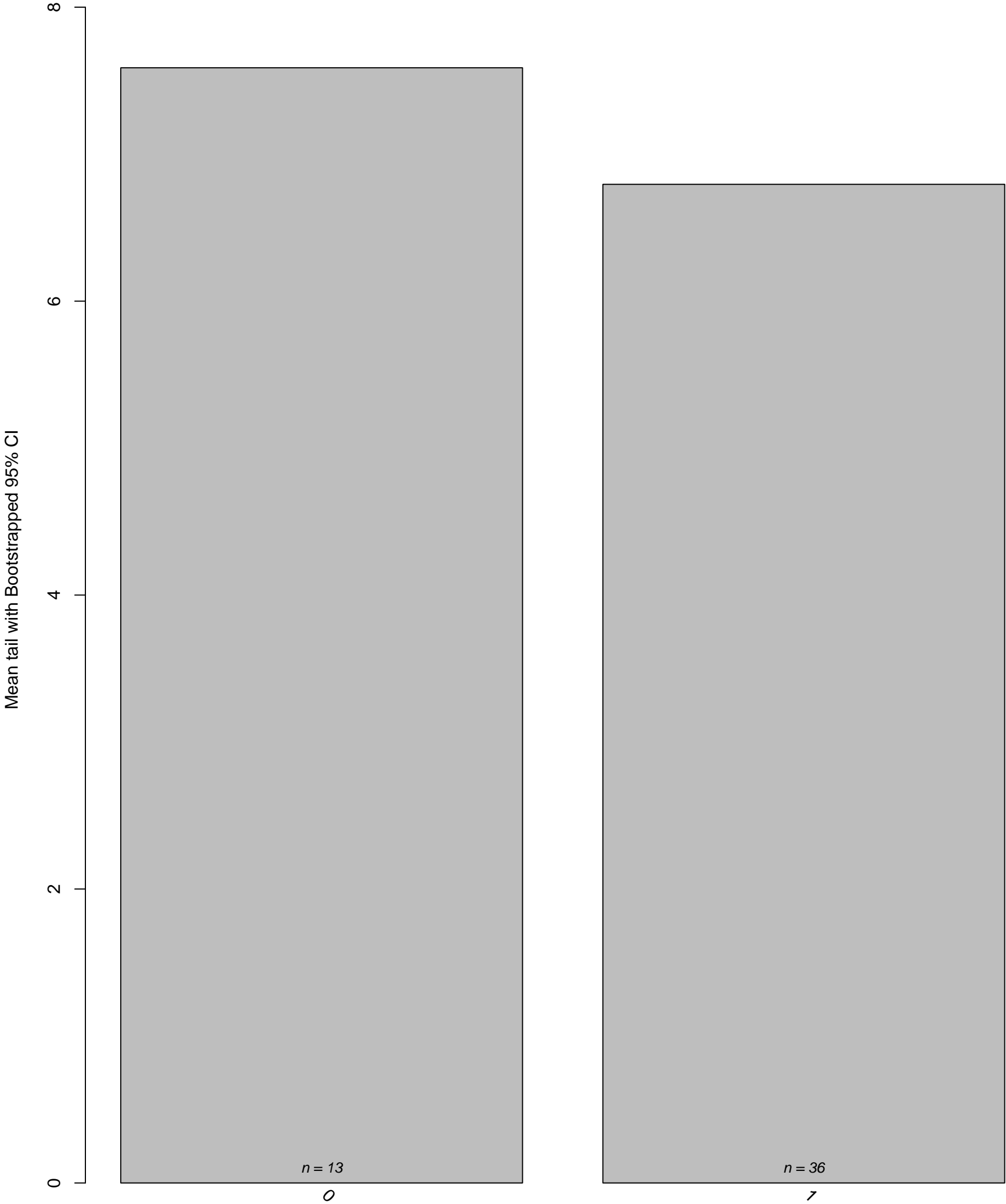


isFemale: Ordered by Sample Size

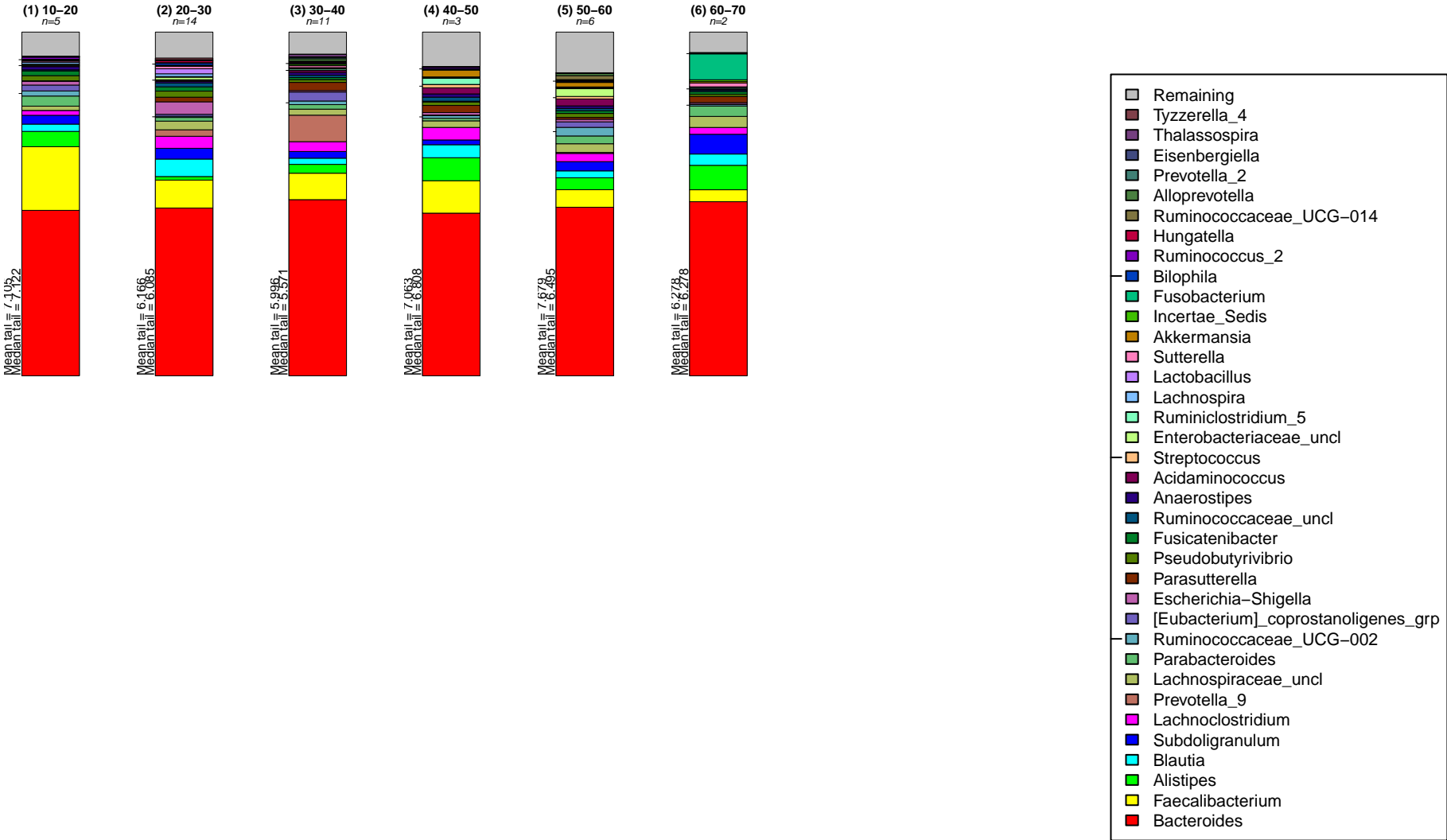


isFemale: Ordered by Category

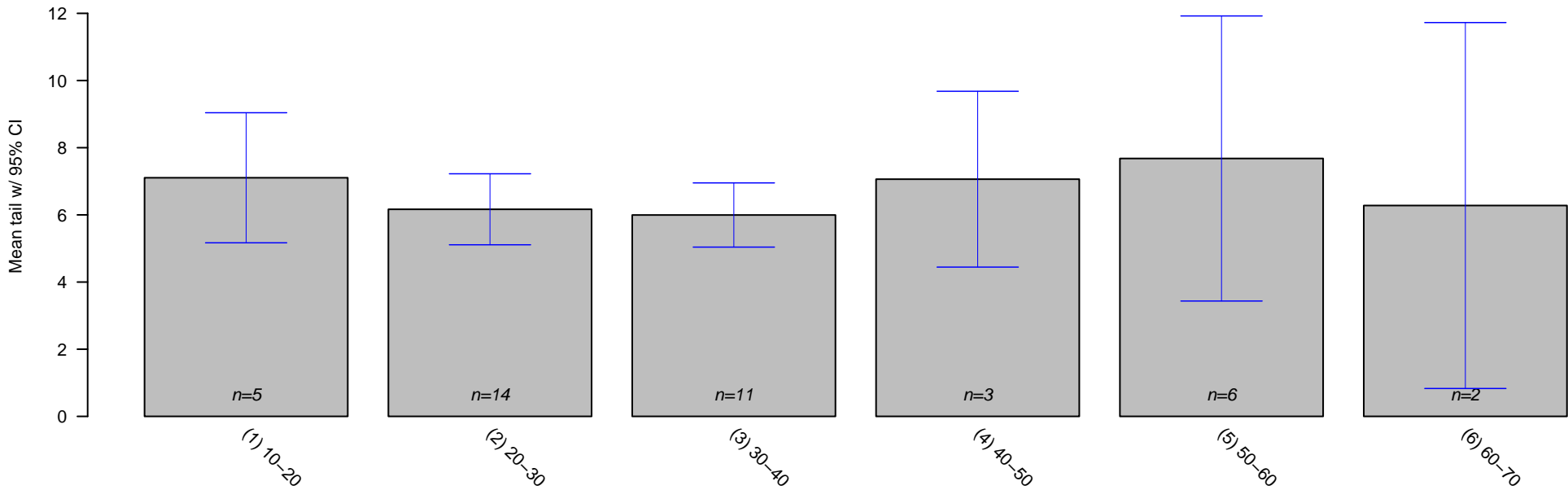
with Wilcoxon rank sum test (difference between group means) *p*-values



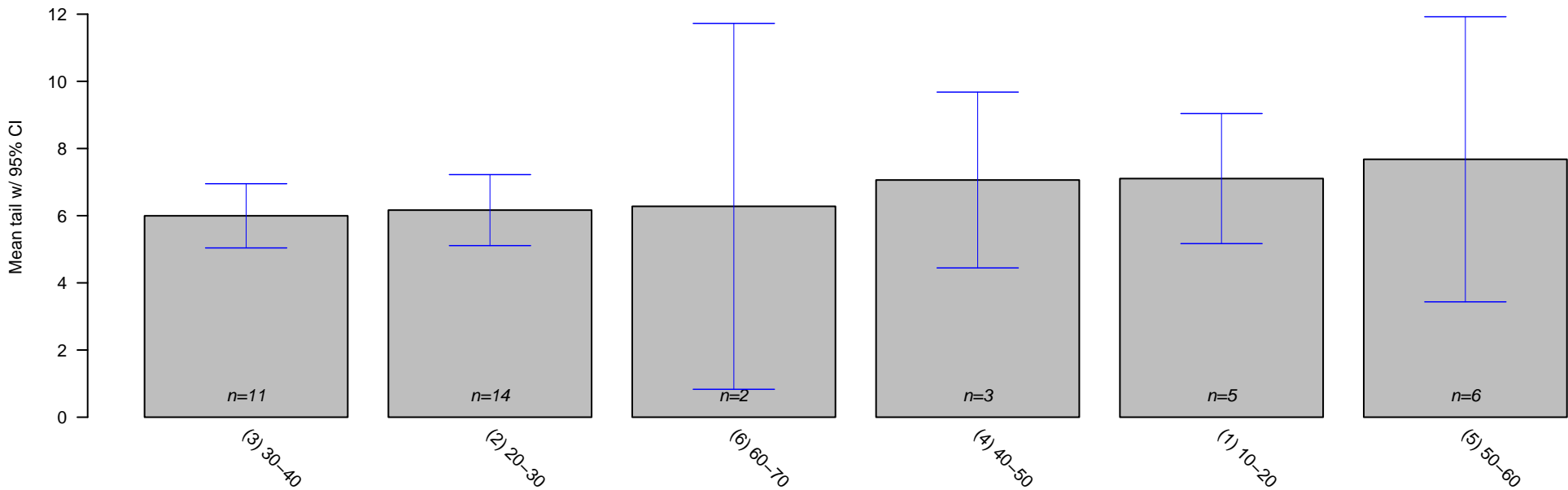
Age



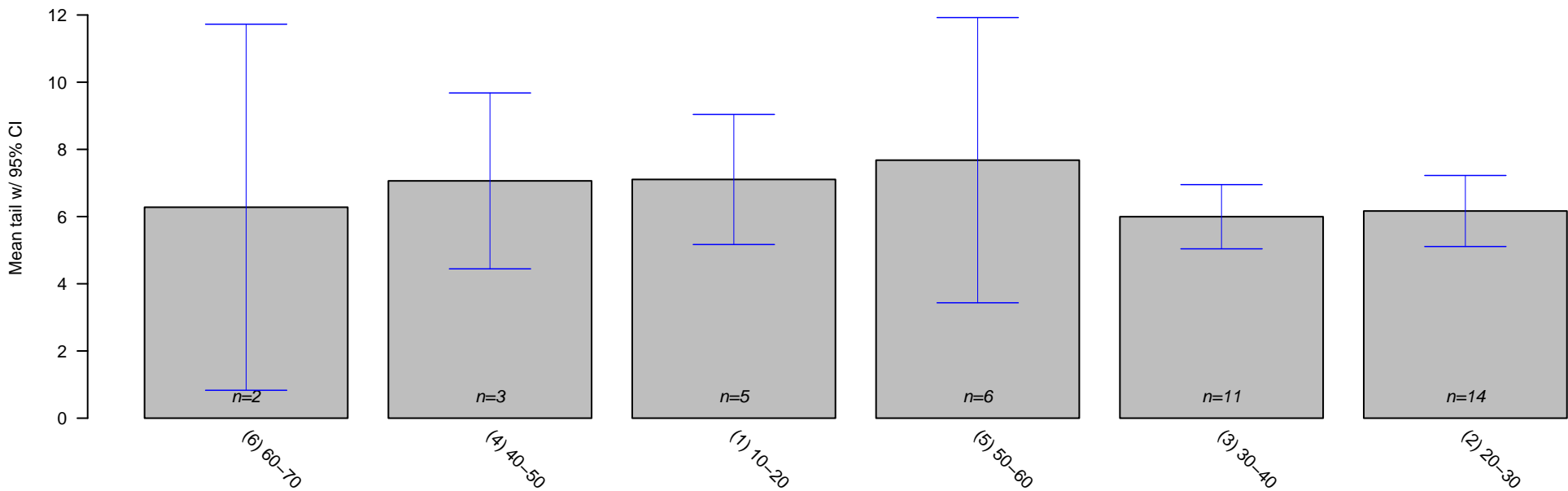
Age: Ordered by Category



Age: Ordered by Diversity

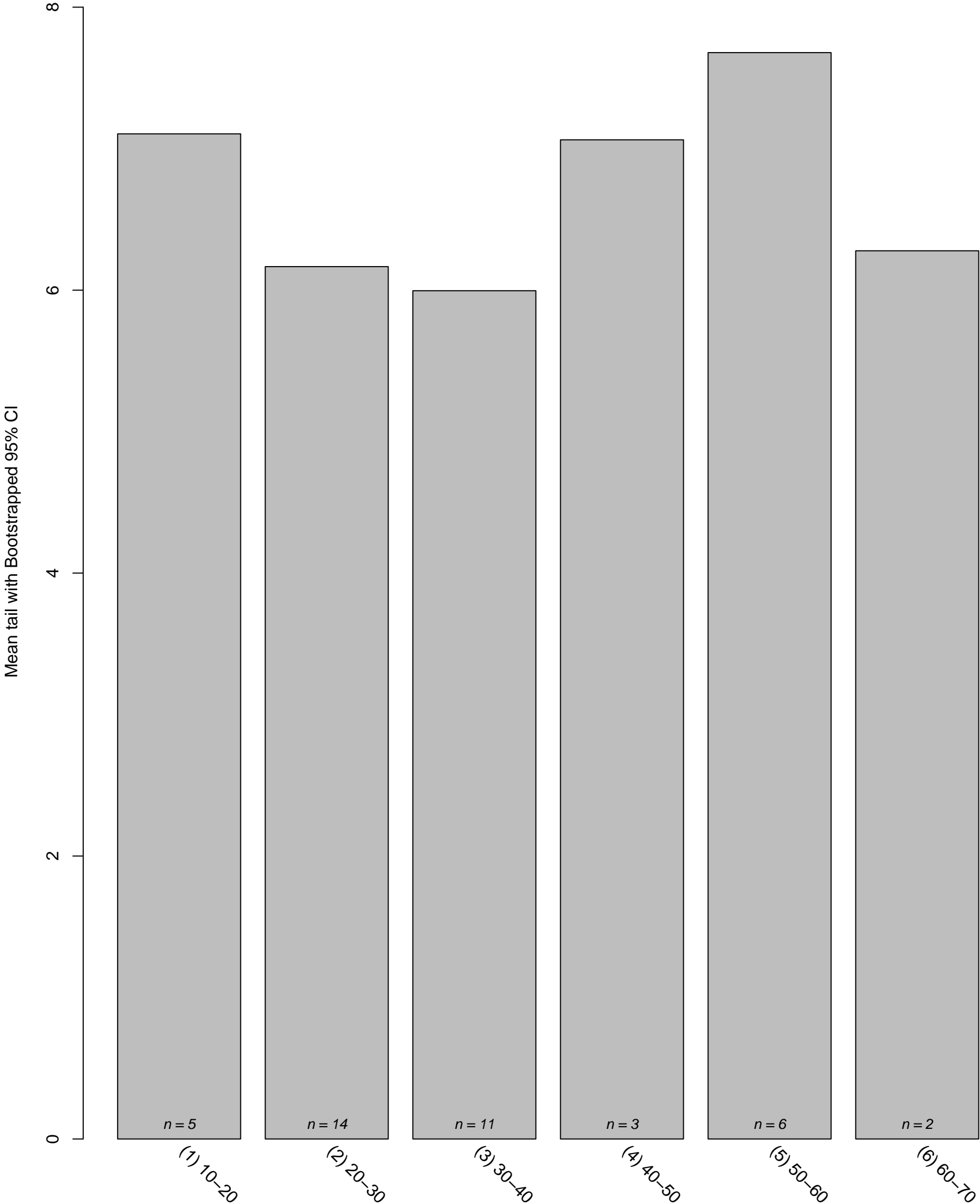


Age: Ordered by Sample Size

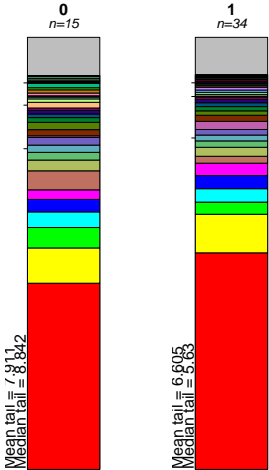


Age: Ordered by Category

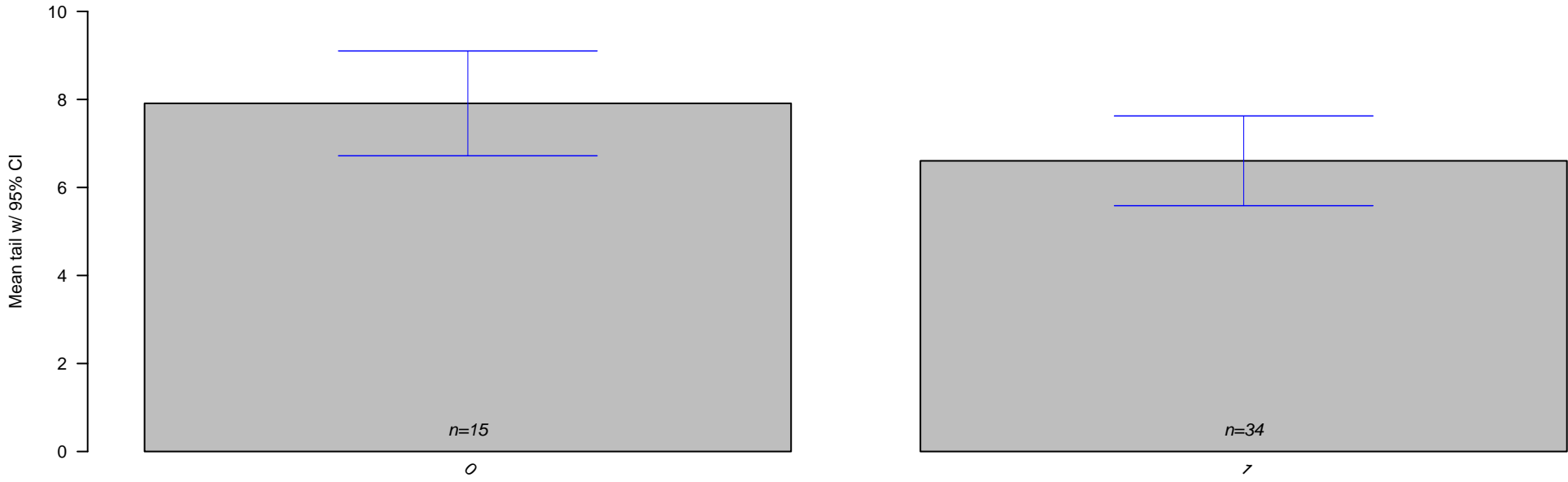
with Wilcoxon rank sum test (difference between group means) p-values



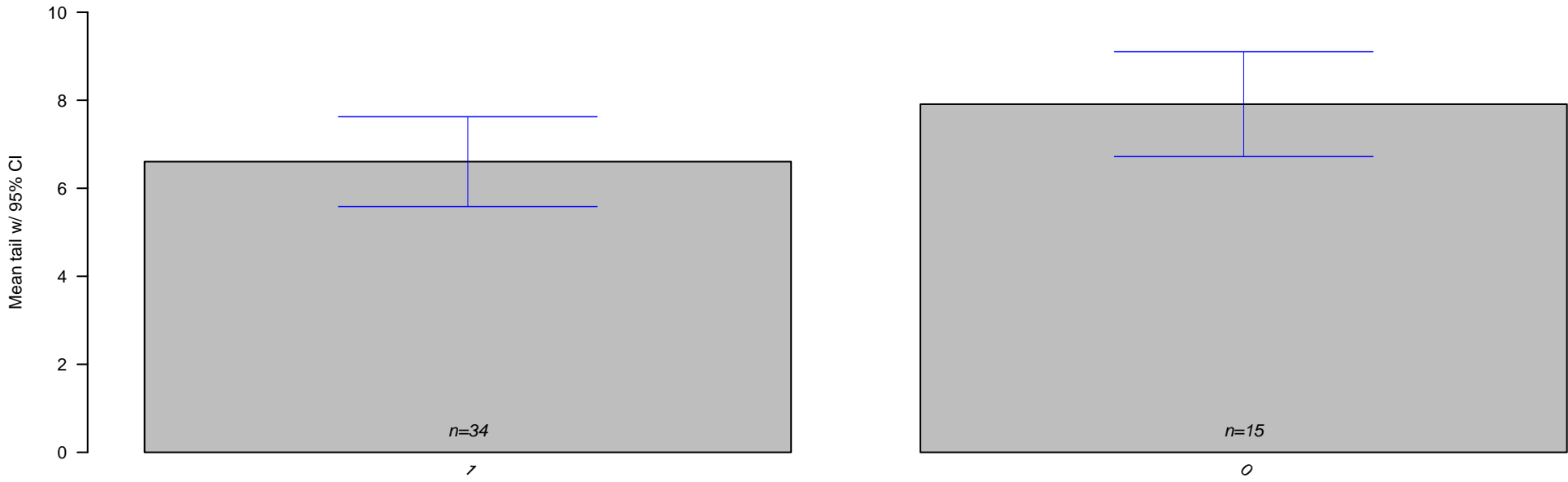
isCase



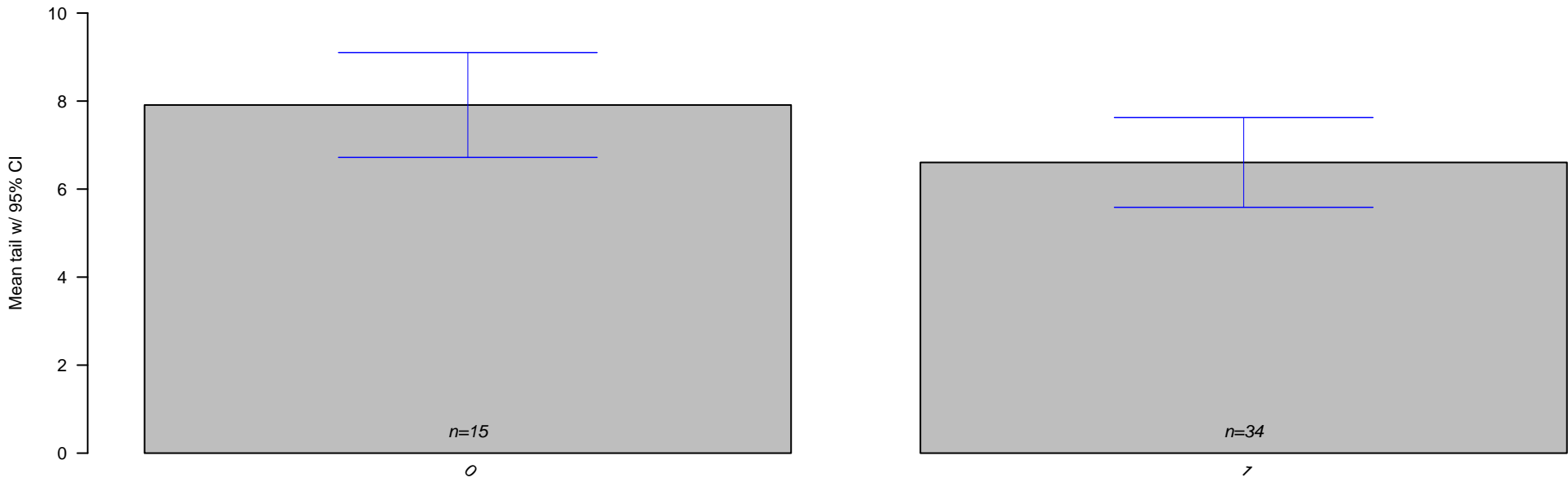
isCase: Ordered by Category



isCase: Ordered by Diversity

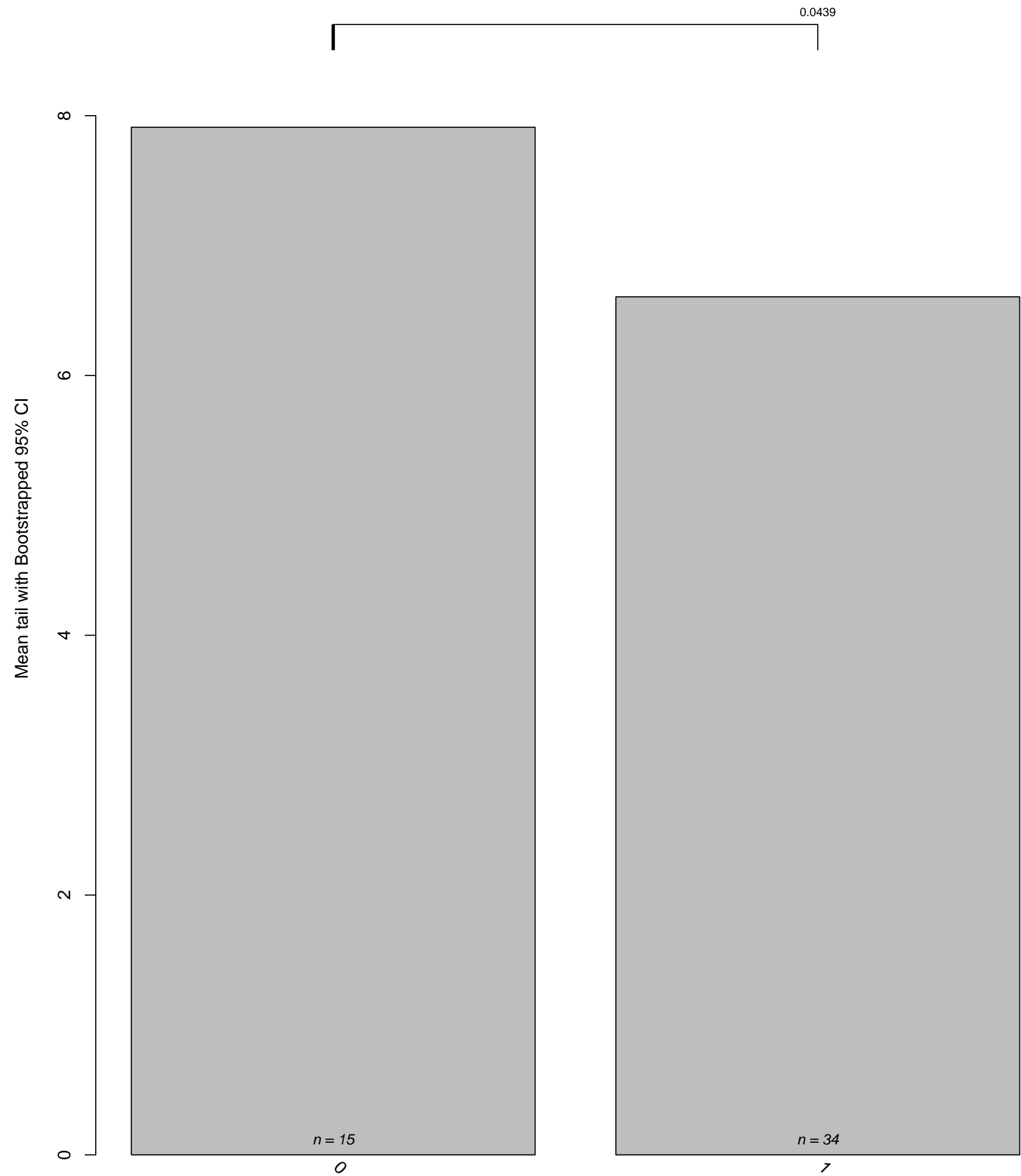


isCase: Ordered by Sample Size

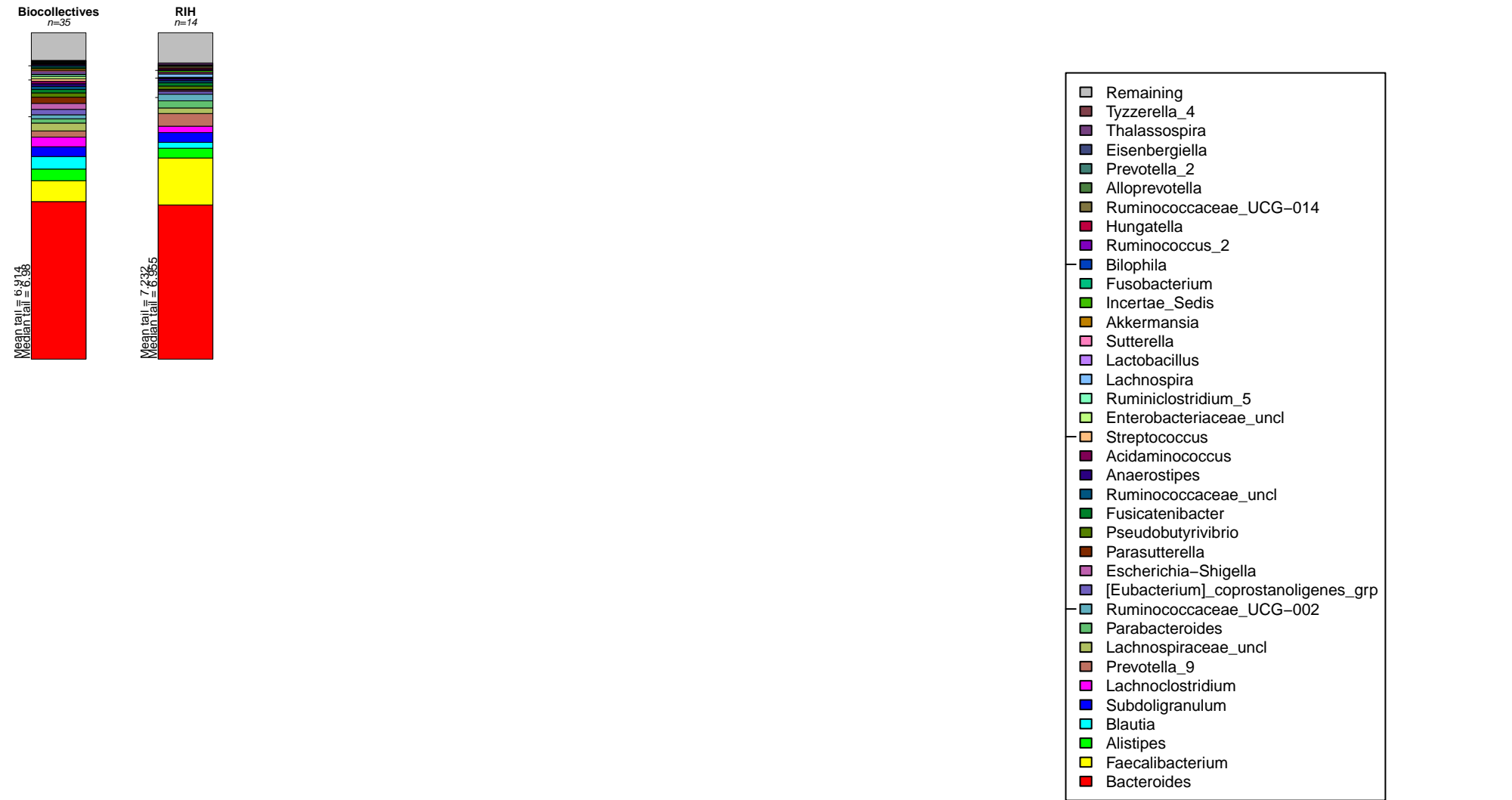


isCase: Ordered by Category

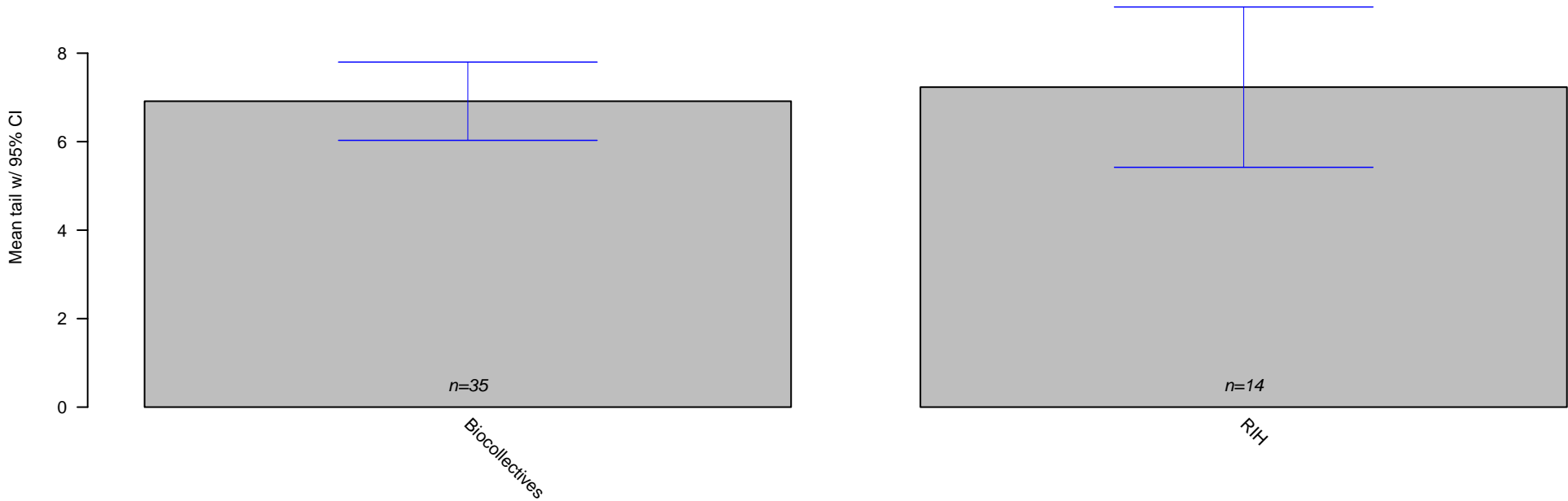
with Wilcoxon rank sum test (difference between group means) *p*-values



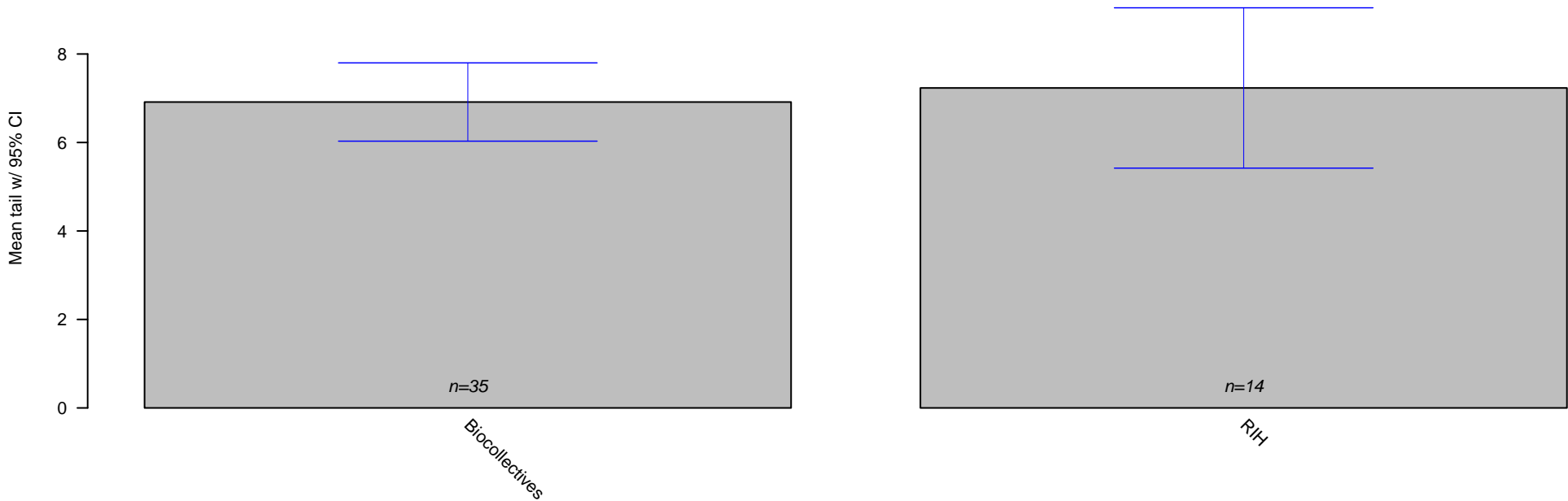
Origin



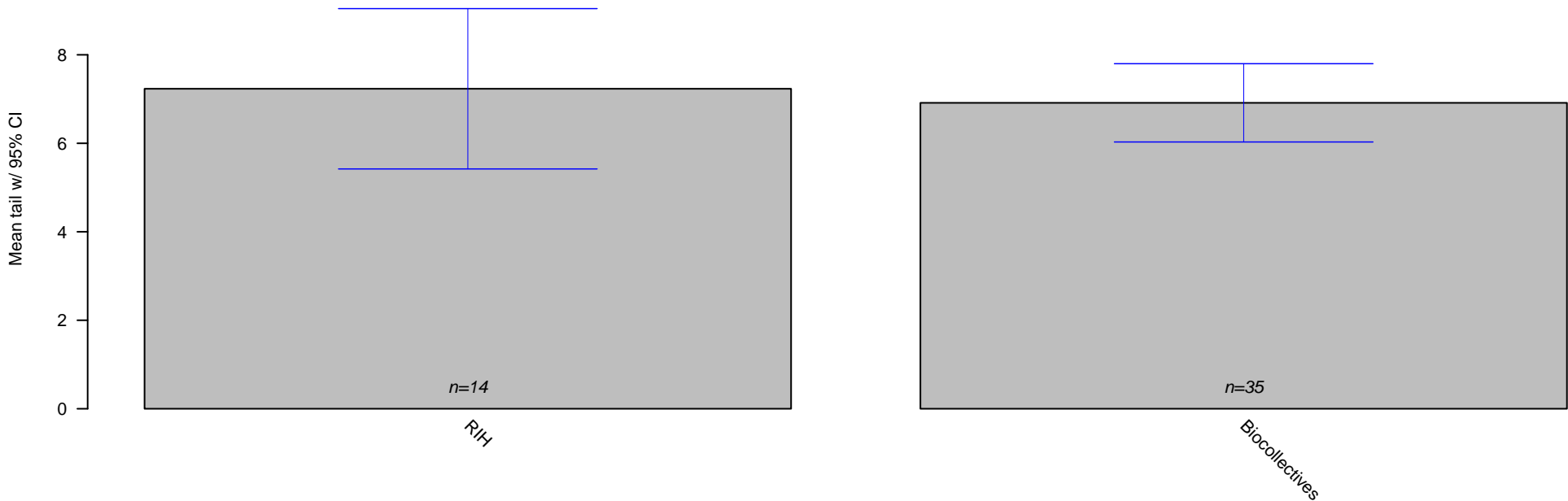
Origin: Ordered by Category



Origin: Ordered by Diversity

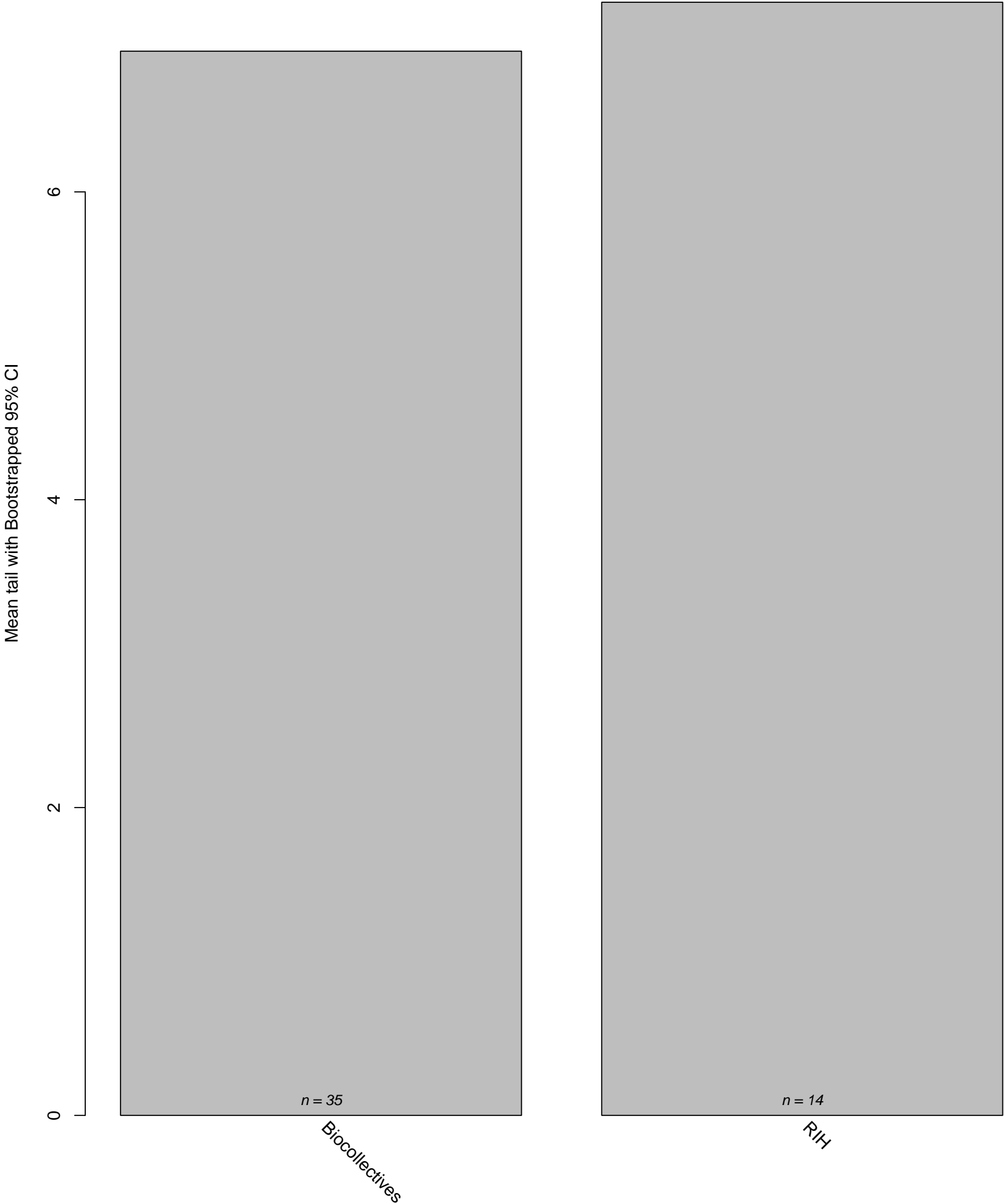


Origin: Ordered by Sample Size

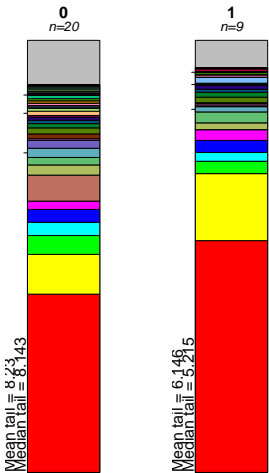


Origin: Ordered by Category

with Wilcoxon rank sum test (difference between group means) *p*-values

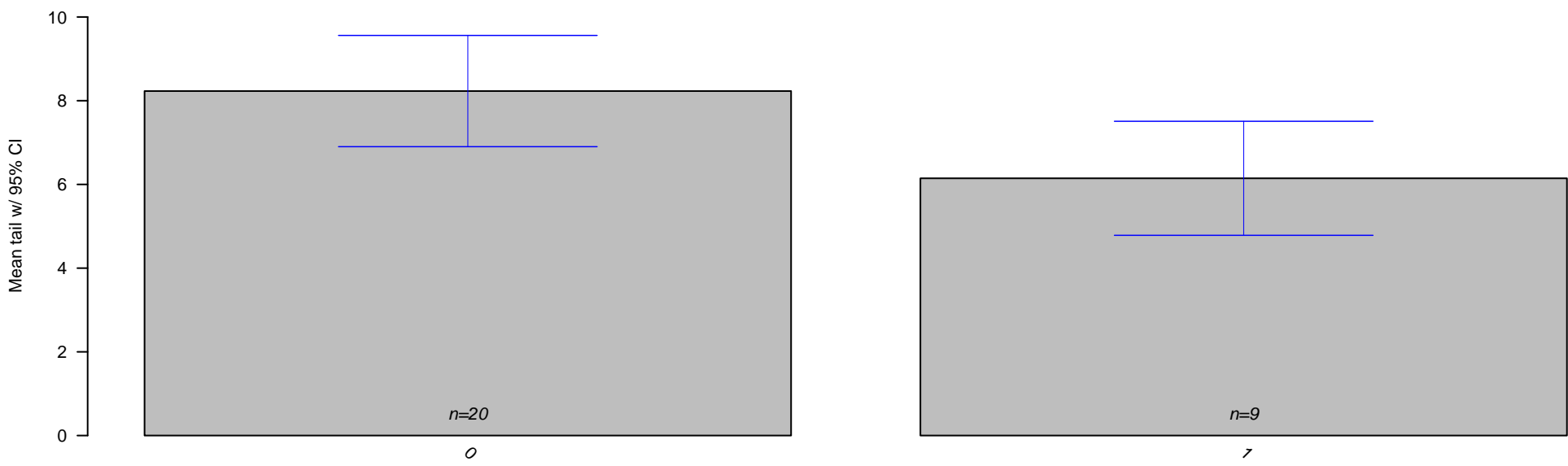


hadHydrea

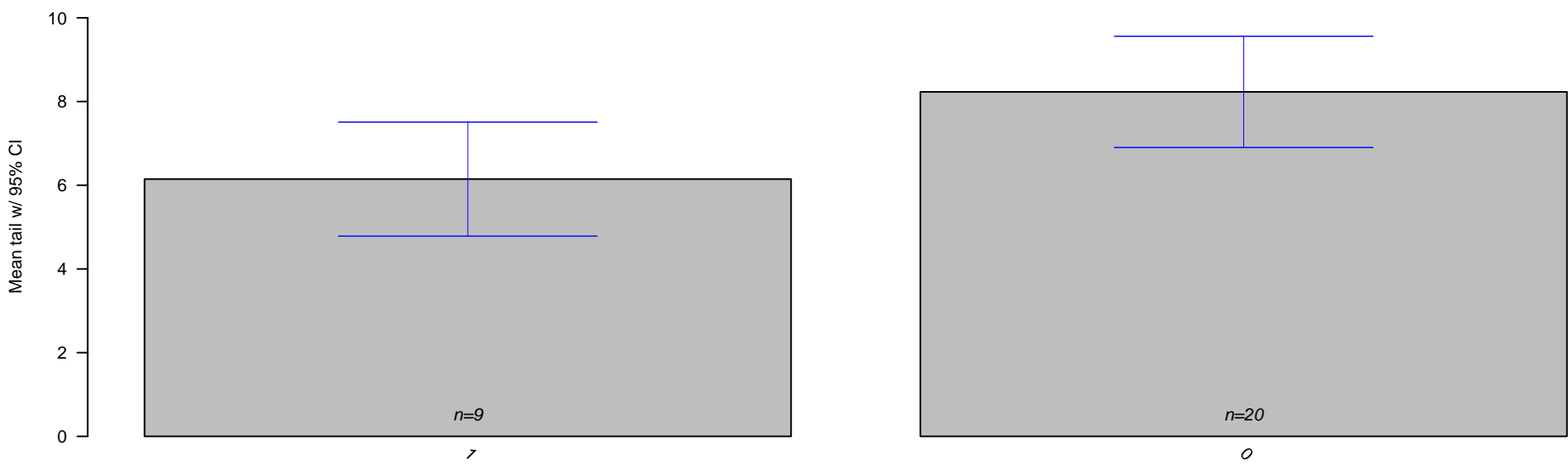


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

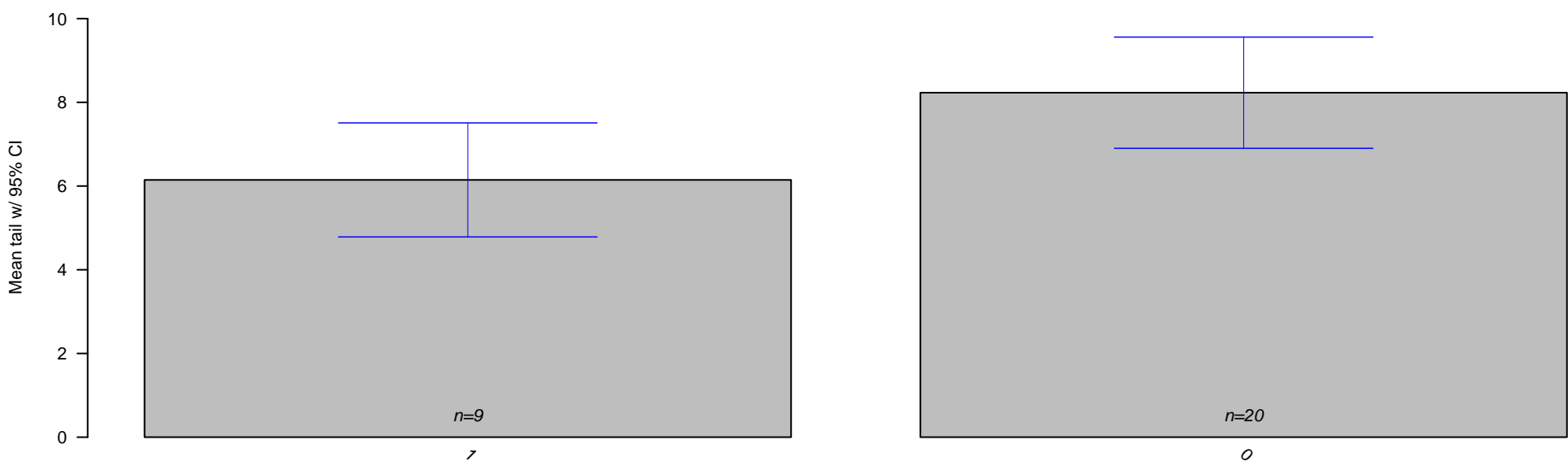
hadHydrea: Ordered by Category



hadHydrea: Ordered by Diversity

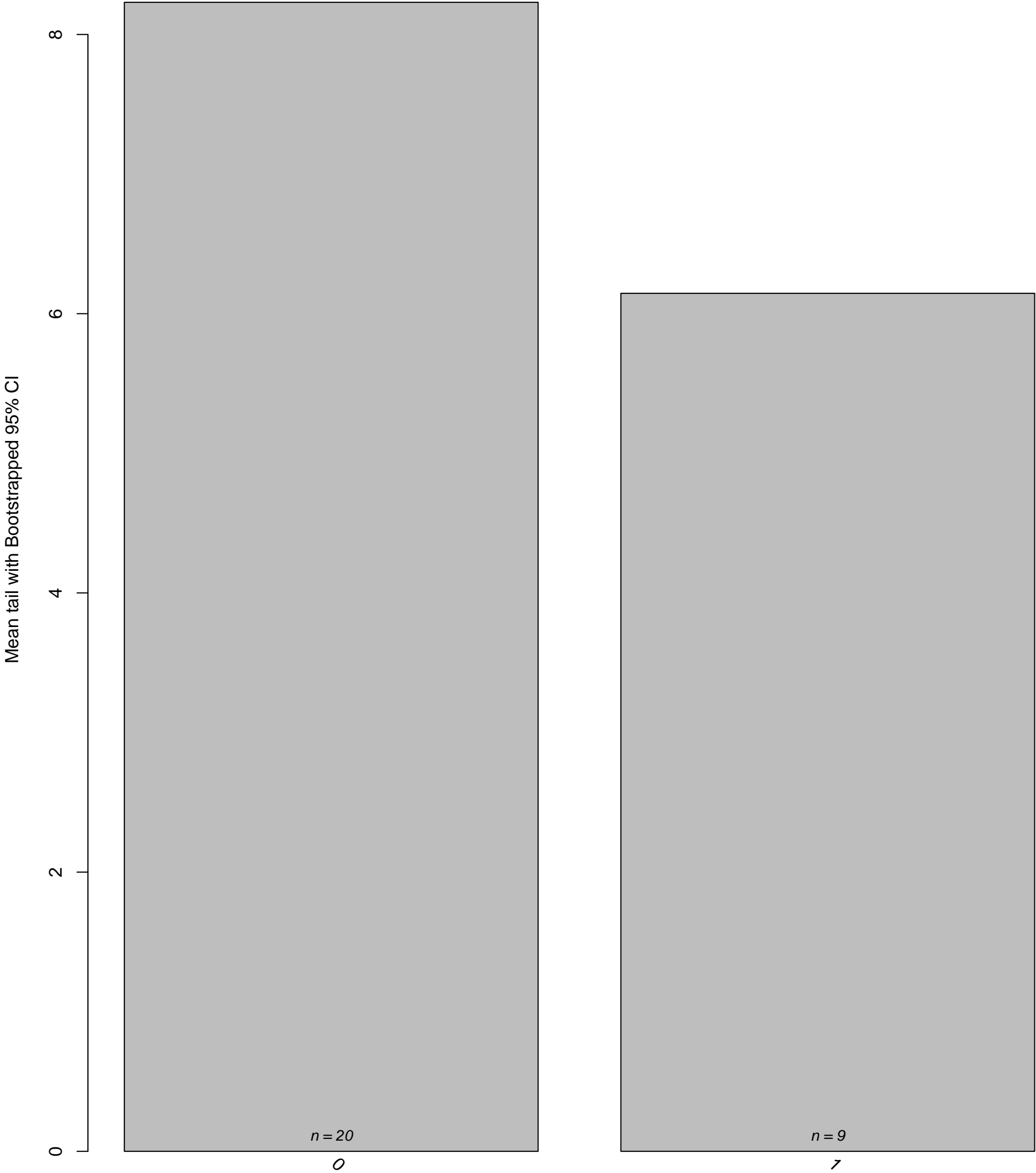


hadHydrea: Ordered by Sample Size

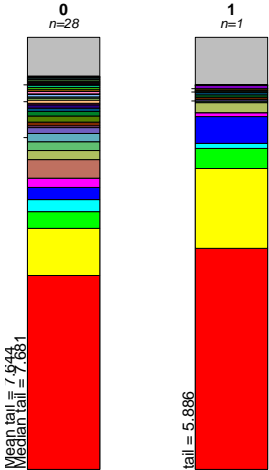


hadHydrea: Ordered by Category

with Wilcoxon rank sum test (difference between group means) *p*-values

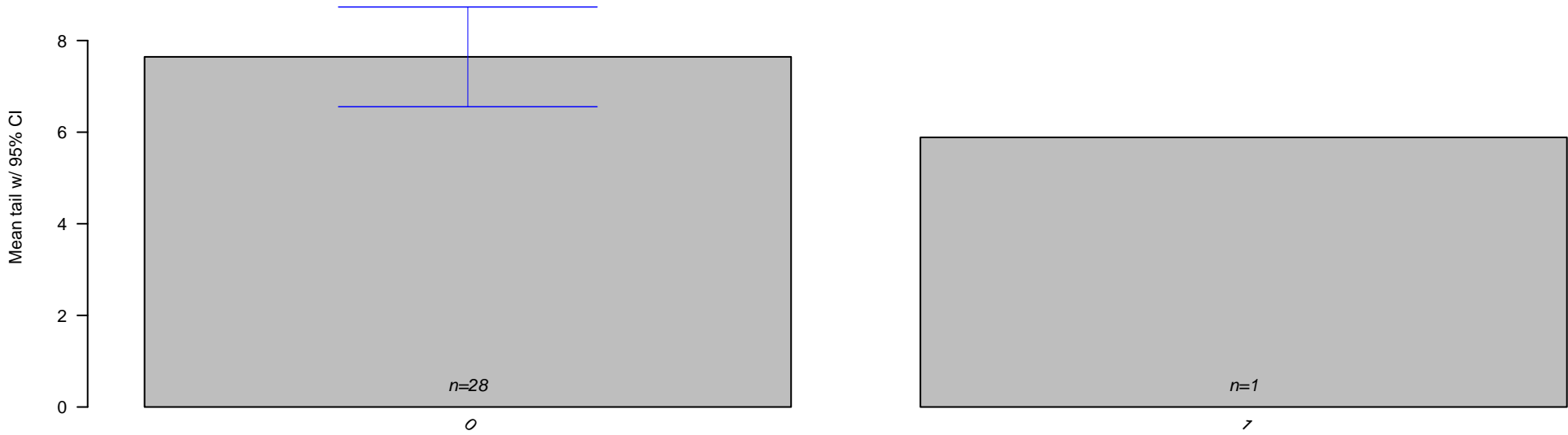


hadExch

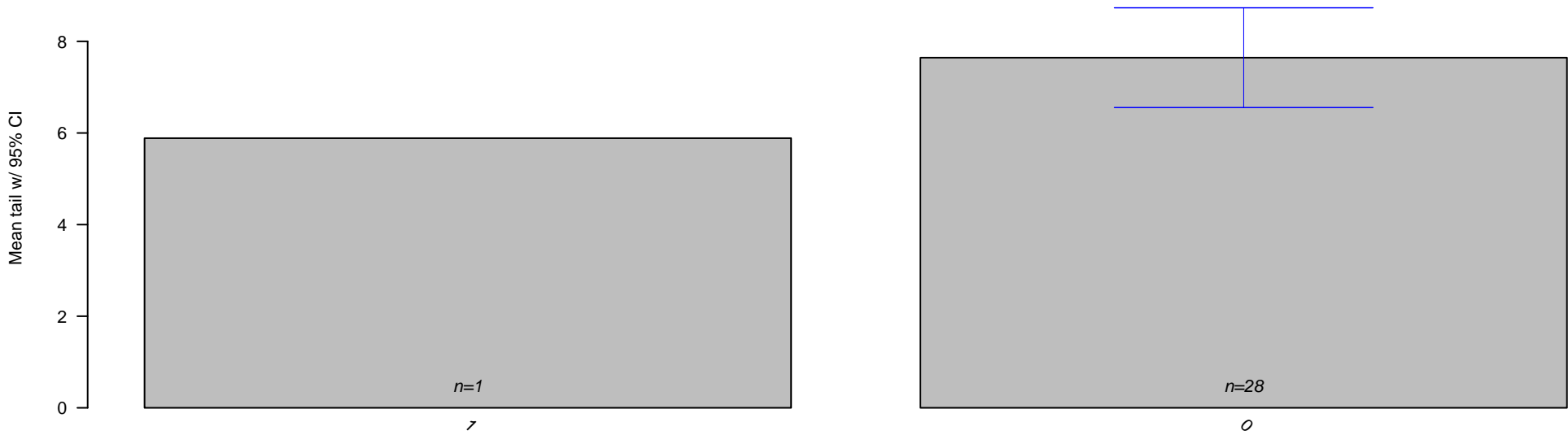


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

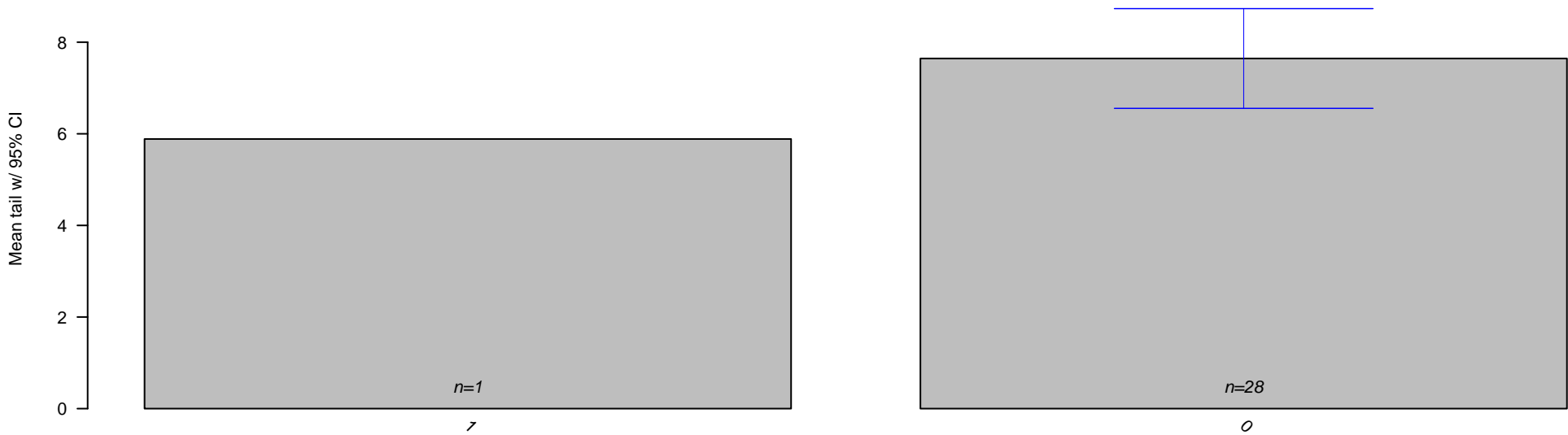
hadExch: Ordered by Category



hadExch: Ordered by Diversity

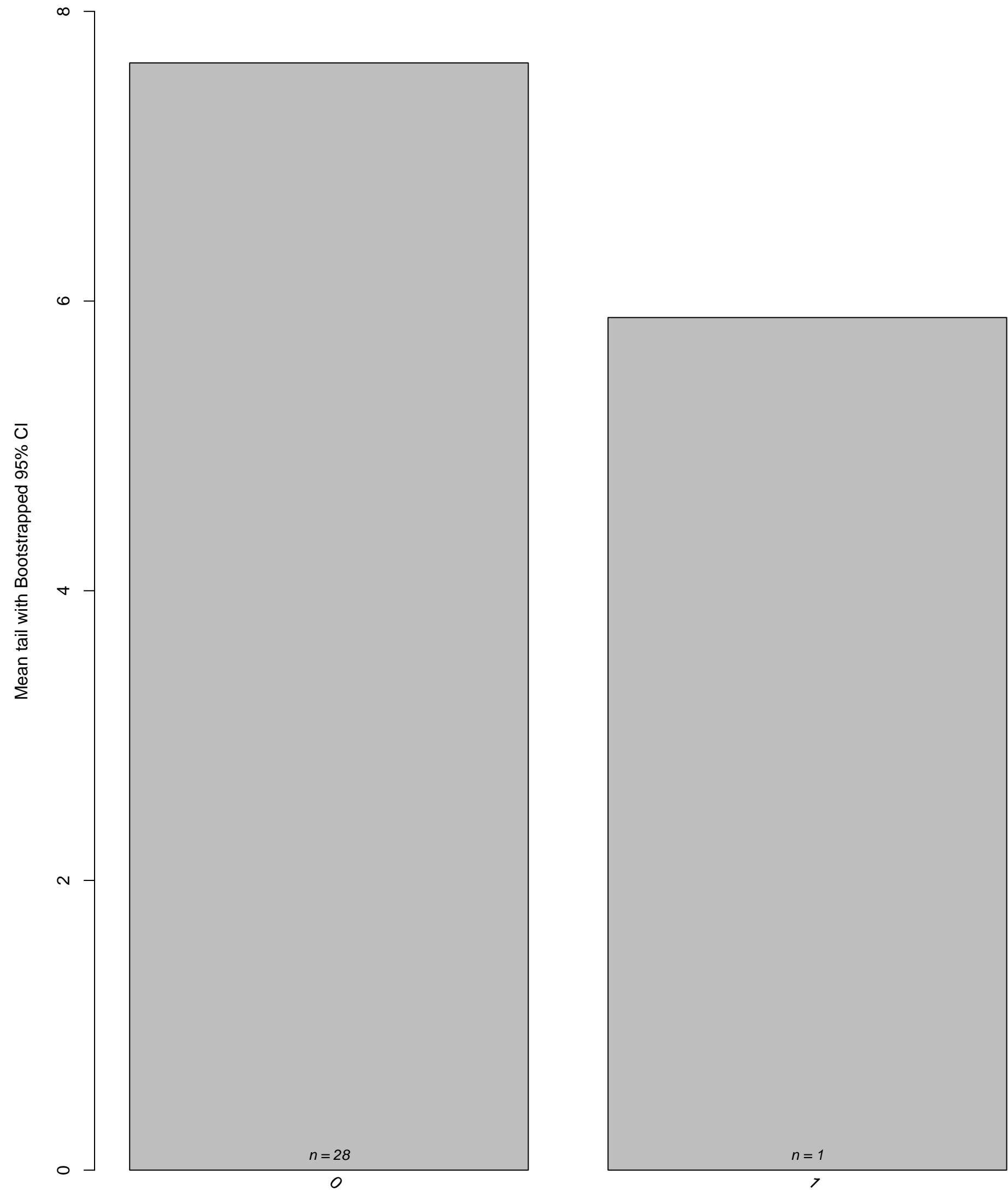


hadExch: Ordered by Sample Size

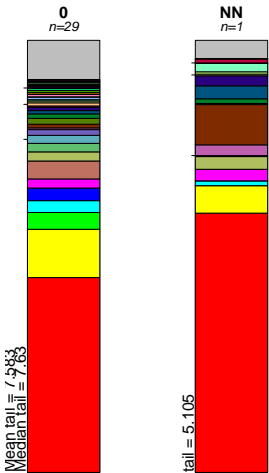


hadExch: Ordered by Category

with Wilcoxon rank sum test (difference between group means) *p*-values

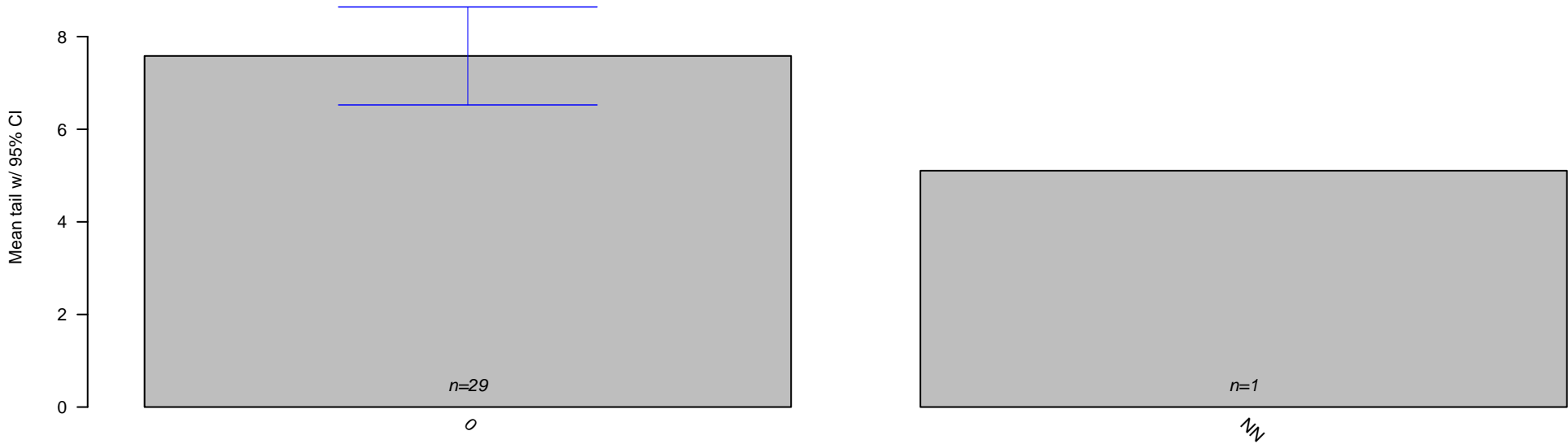


onPCN

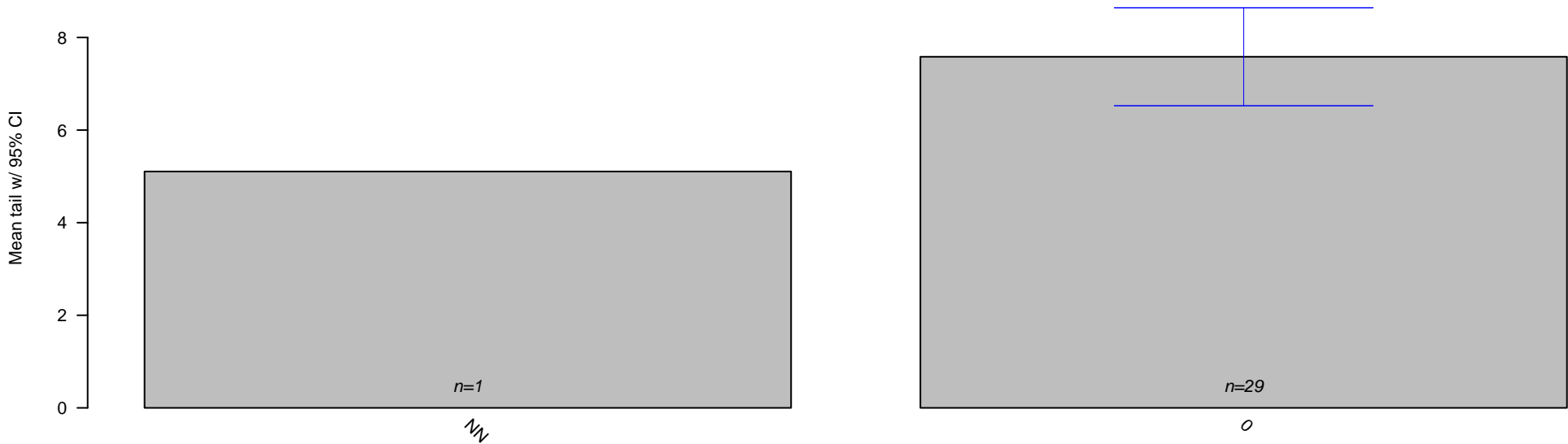


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

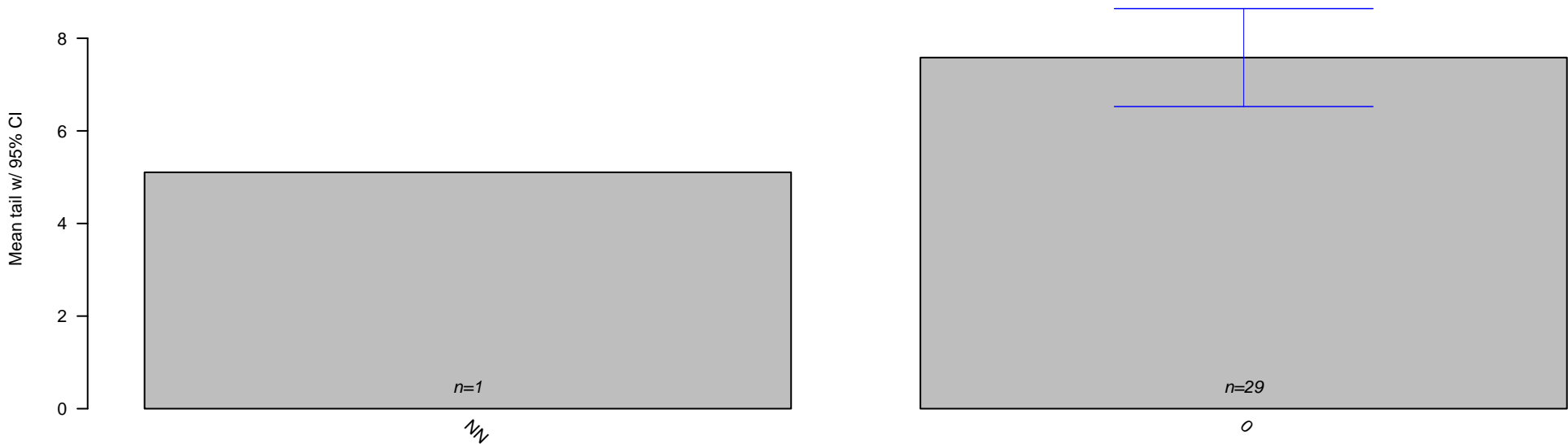
onPCN: Ordered by Category



onPCN: Ordered by Diversity

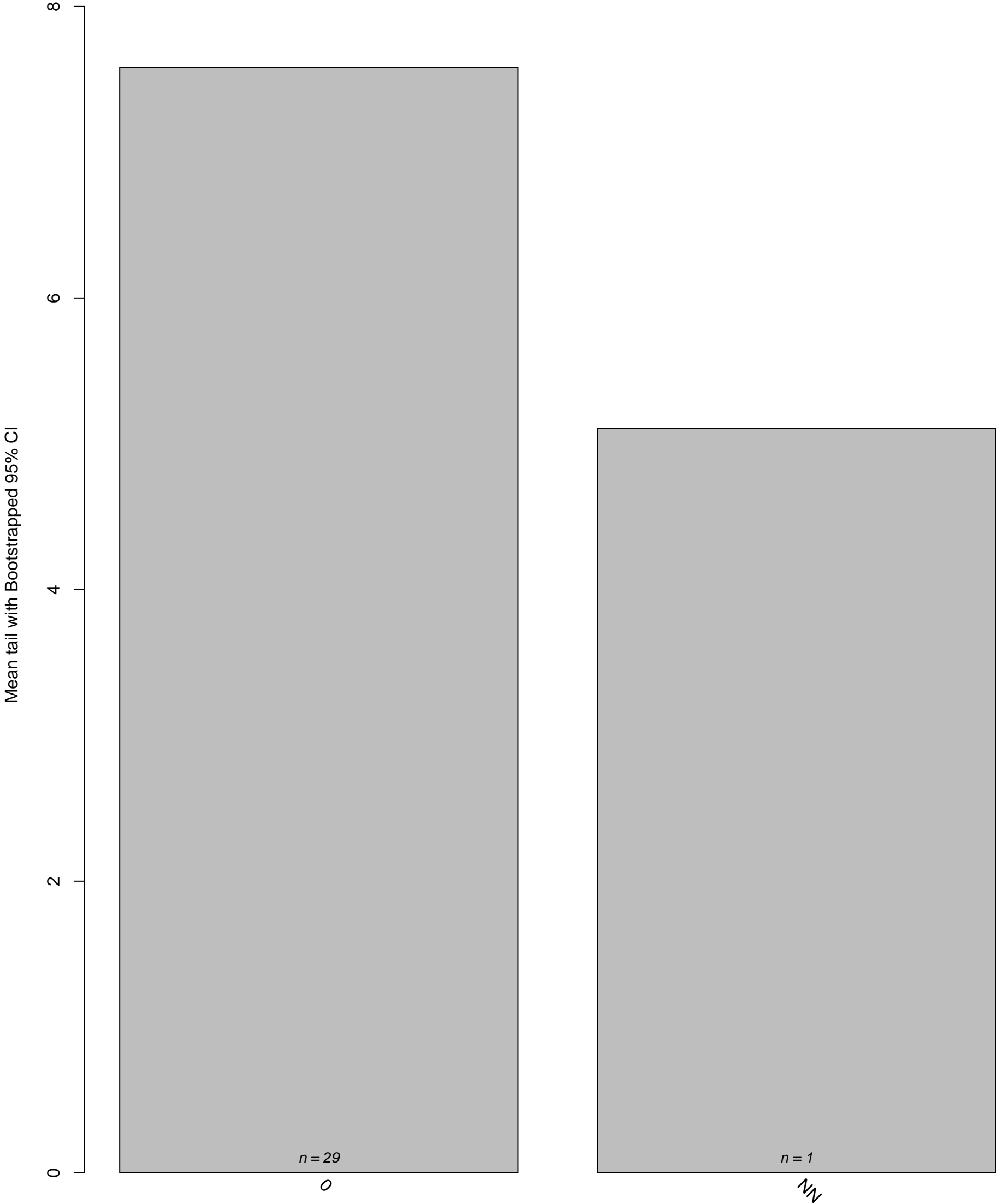


onPCN: Ordered by Sample Size

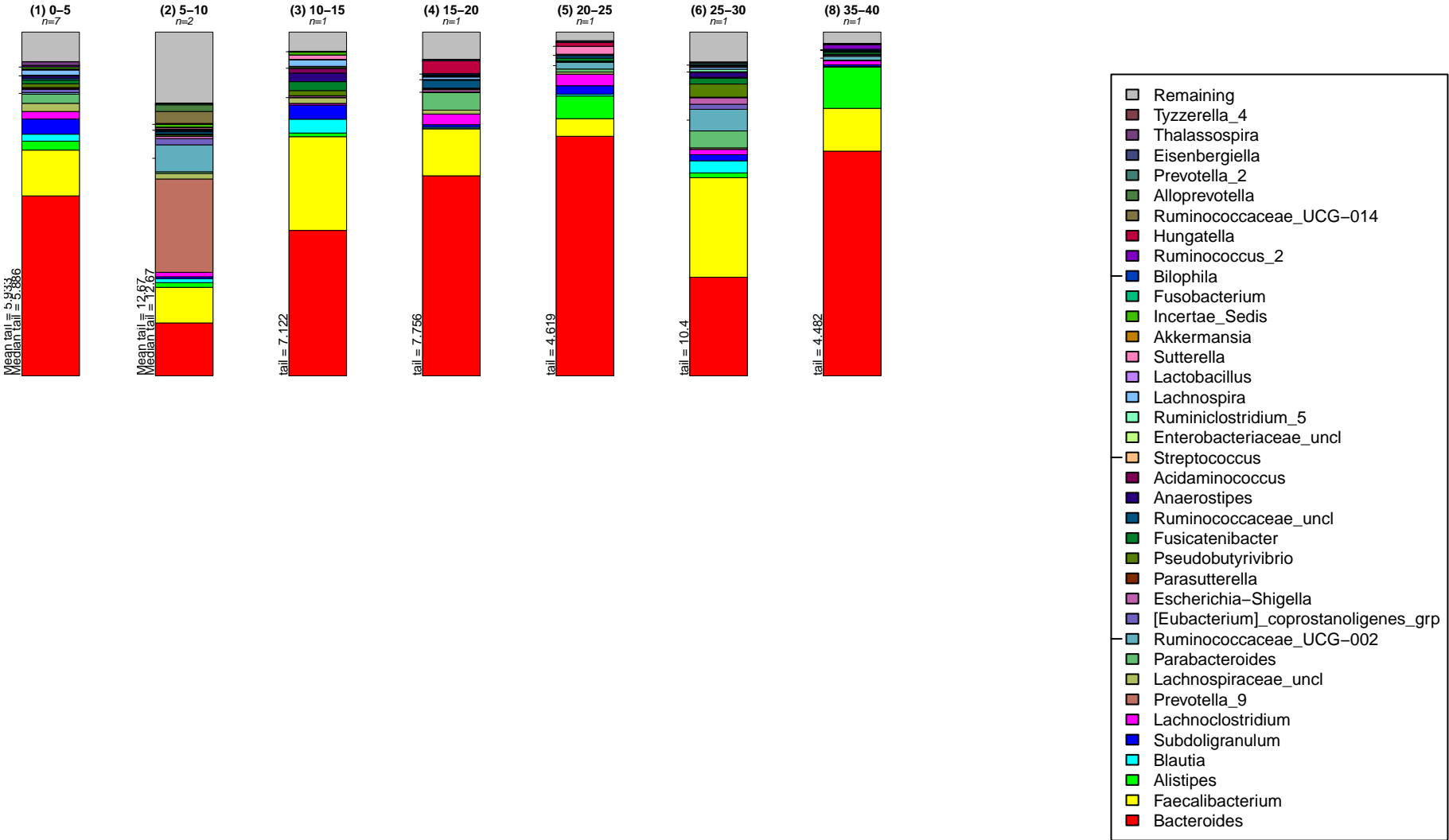


onPCN: Ordered by Category

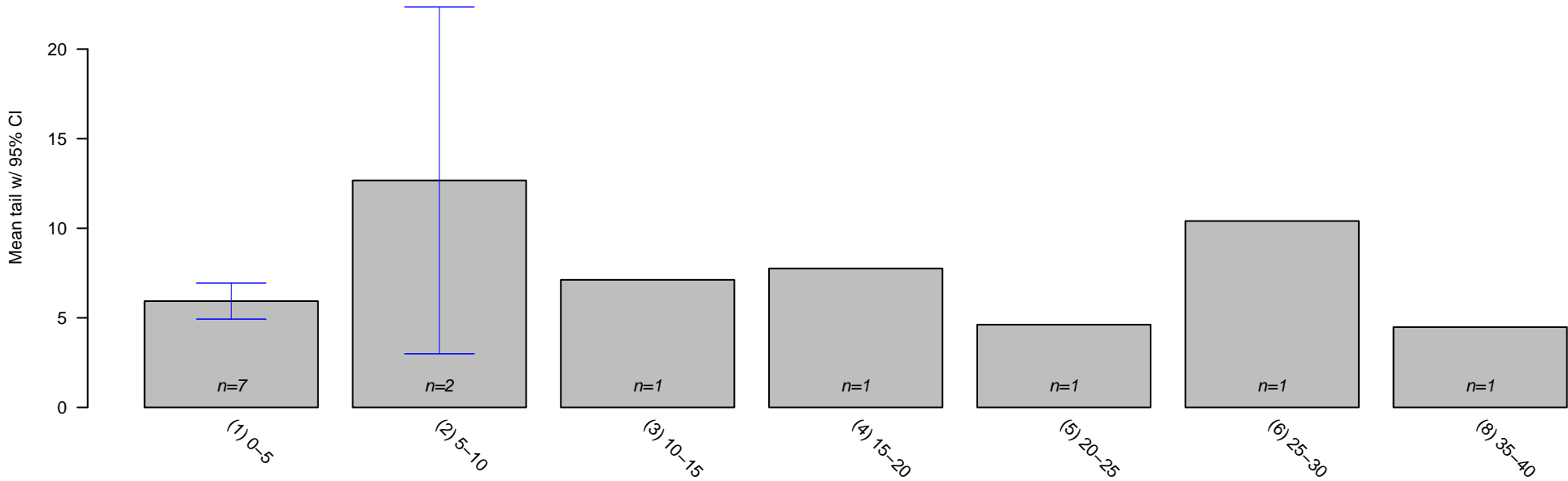
with Wilcoxon rank sum test (difference between group means) p-values



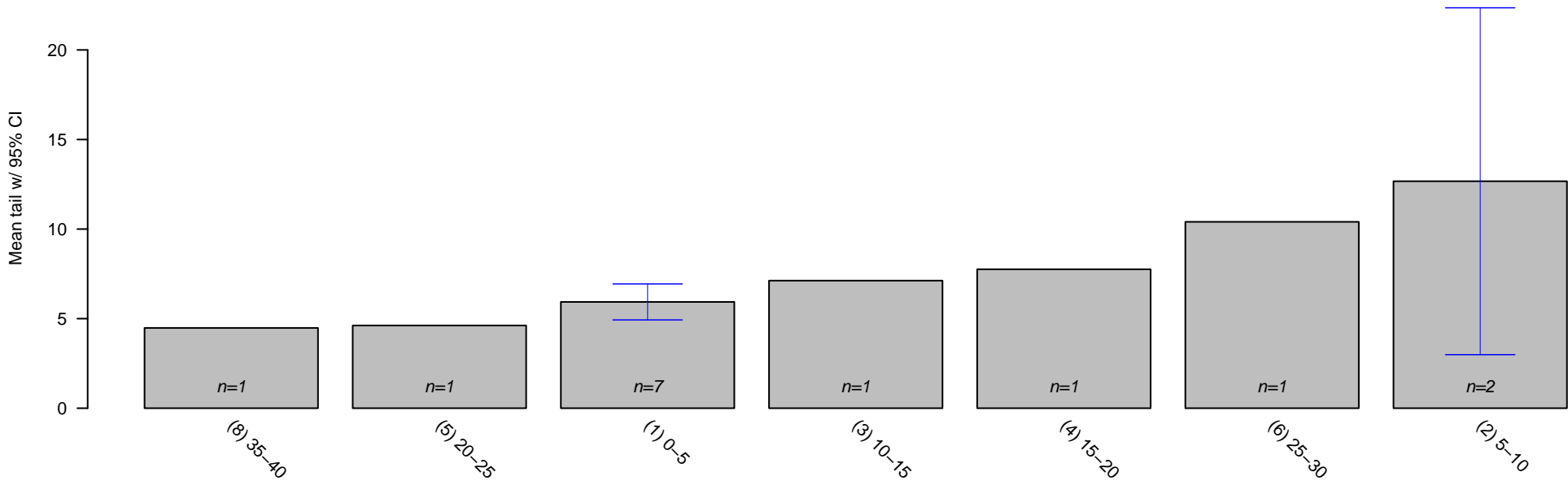
HbF



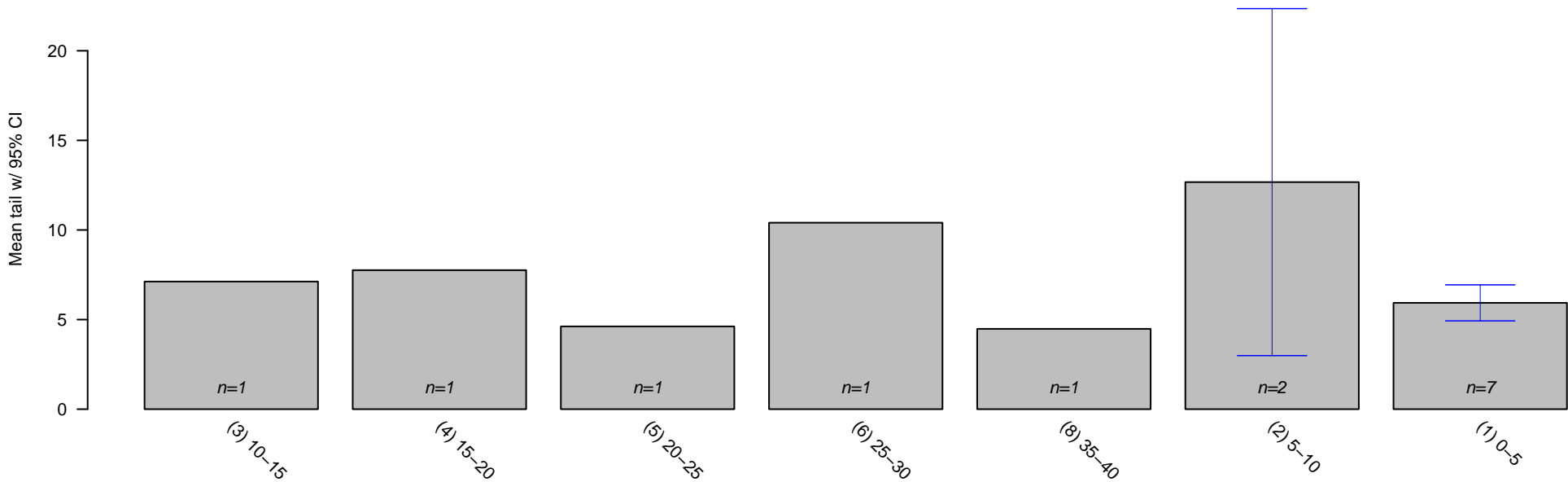
HbF: Ordered by Category



HbF: Ordered by Diversity

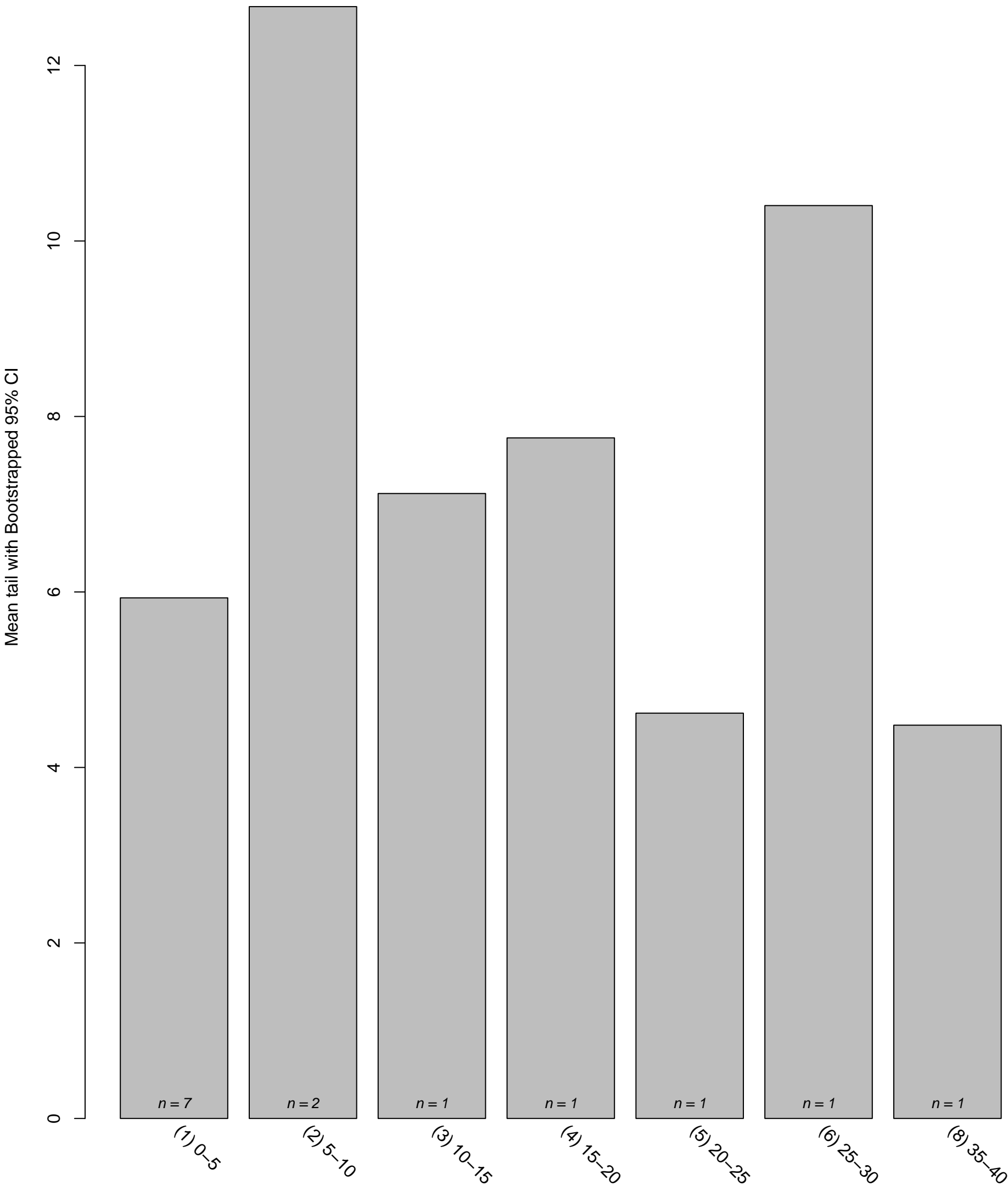


HbF: Ordered by Sample Size

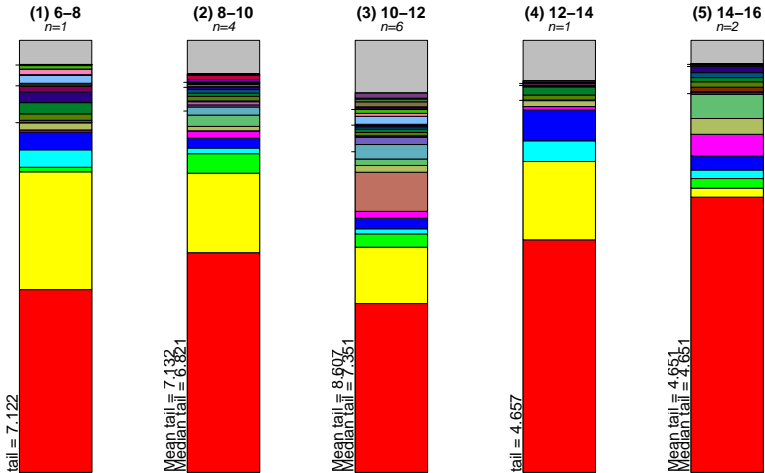


HbF: Ordered by Category

with Wilcoxon rank sum test (difference between group means) *p*-values

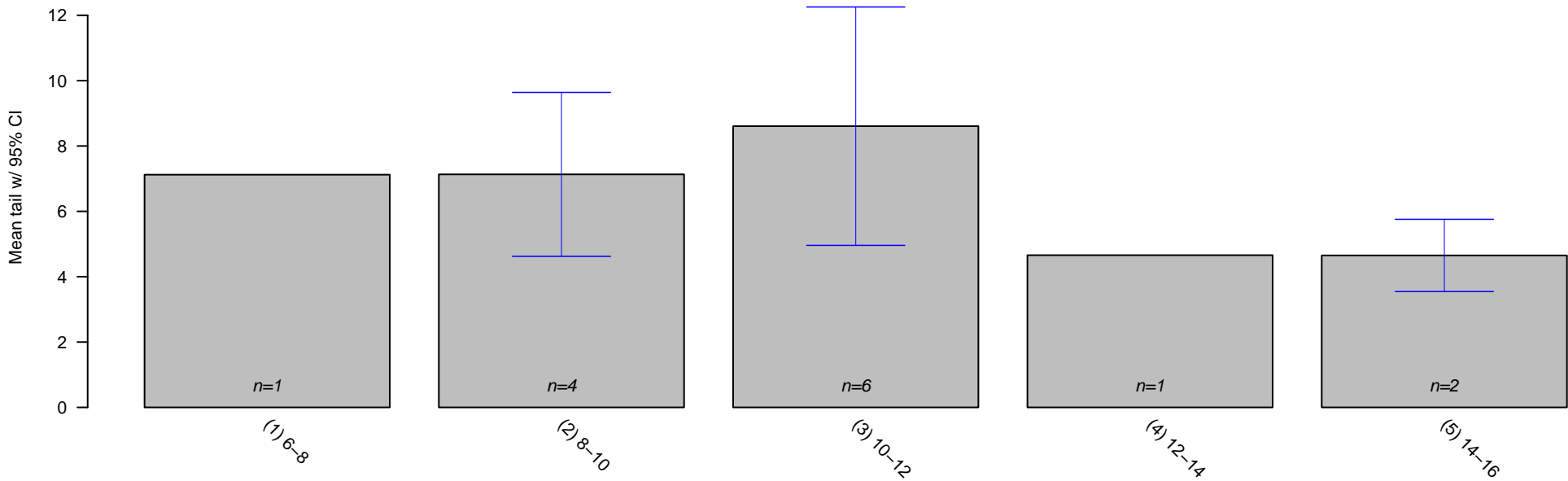


Hb

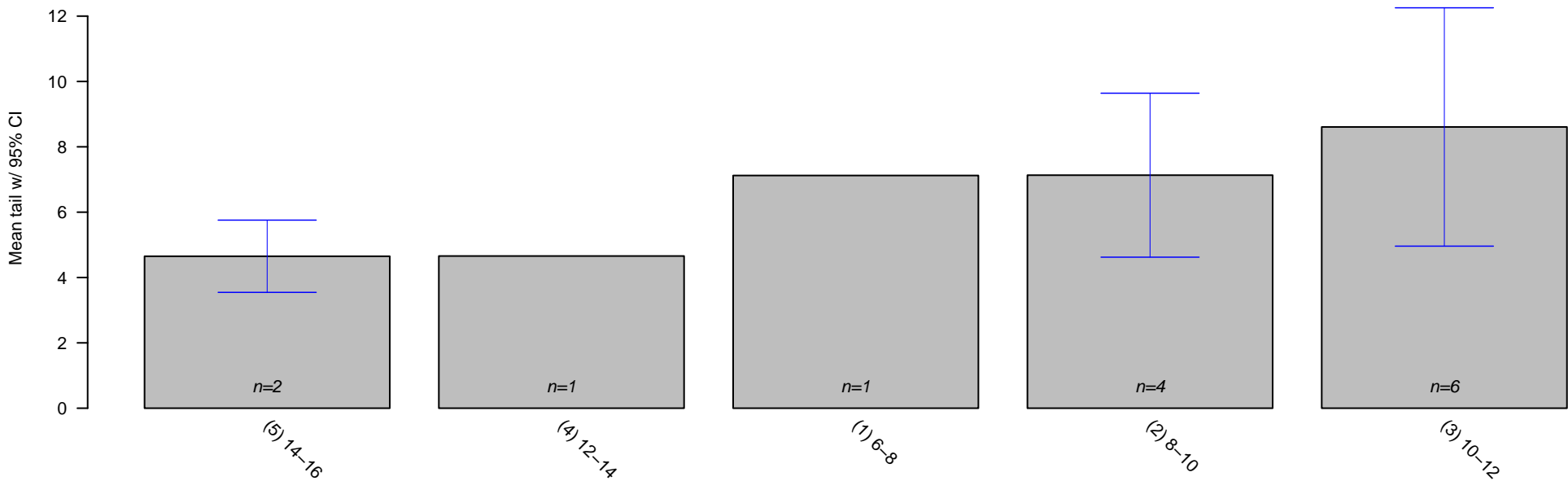


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

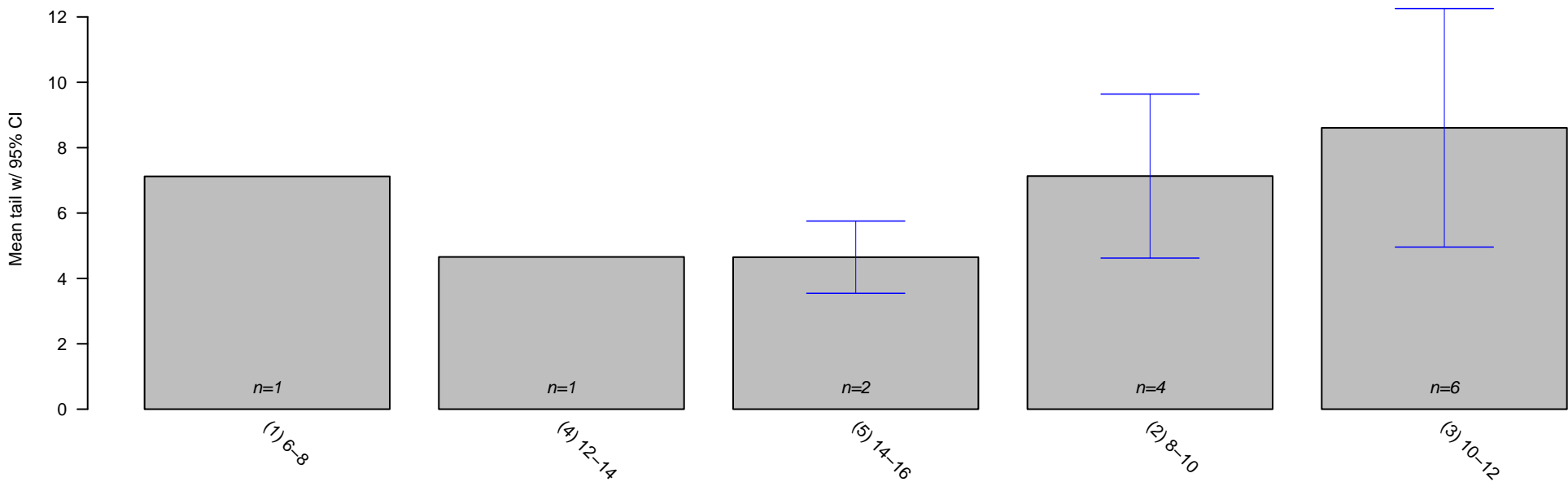
Hb: Ordered by Category



Hb: Ordered by Diversity

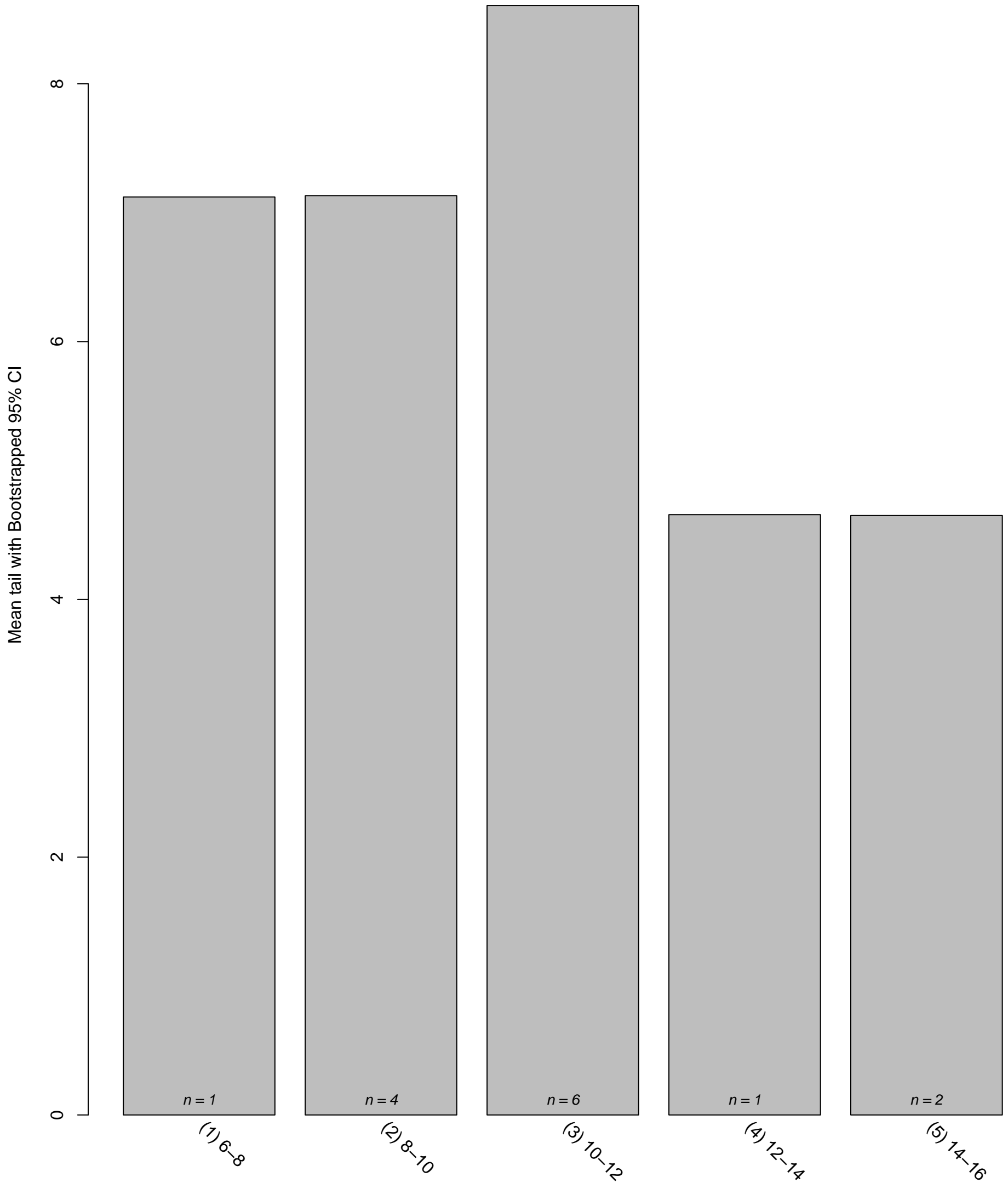


Hb: Ordered by Sample Size

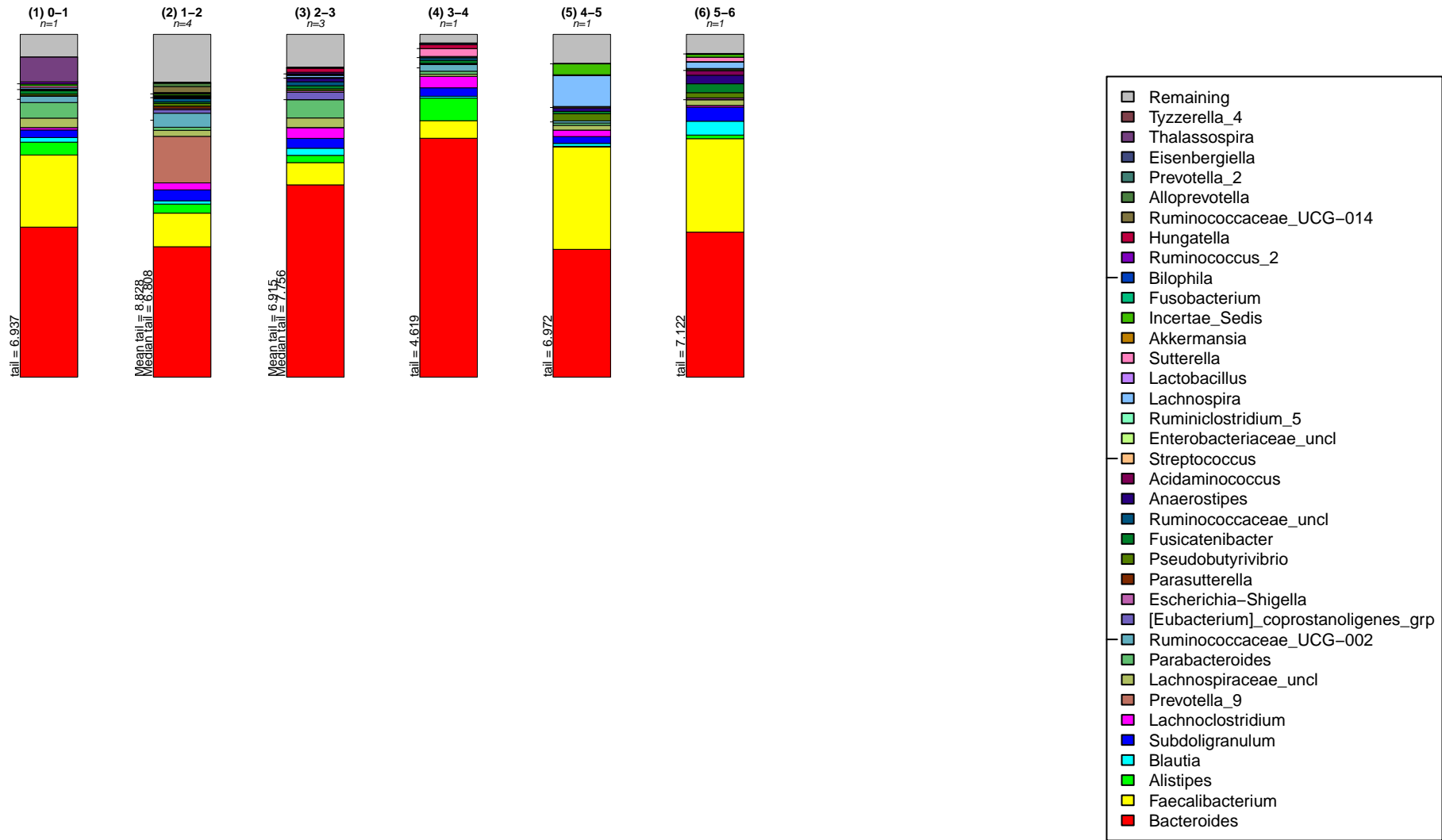


Hb: Ordered by Category

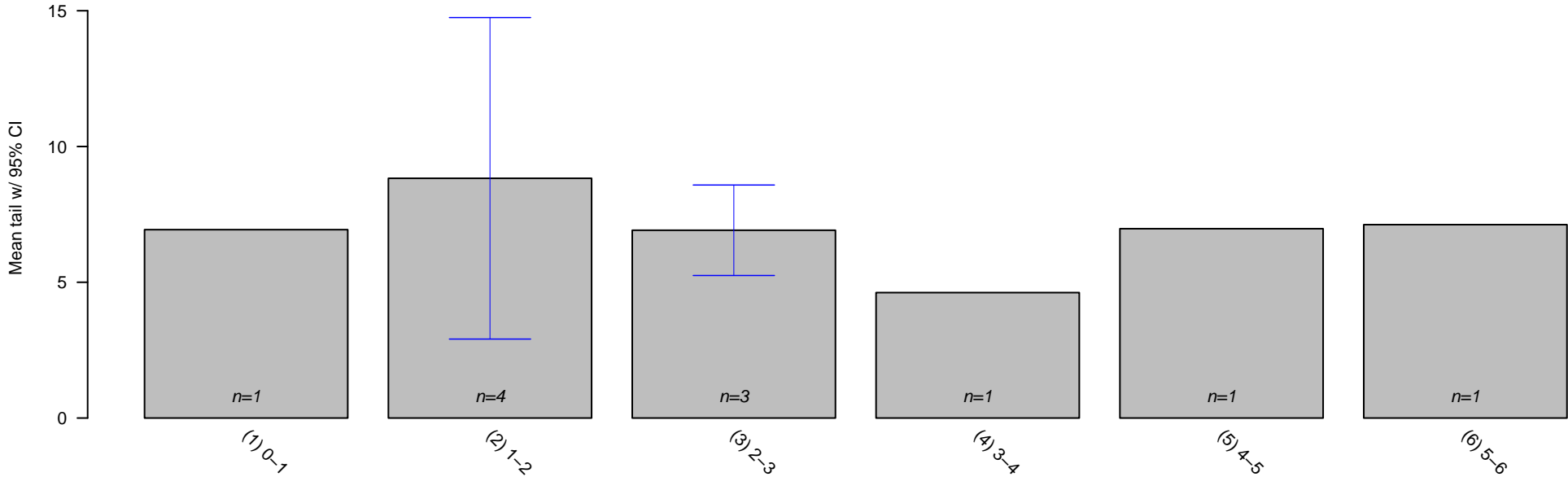
with Wilcoxon rank sum test (difference between group means) *p*-values



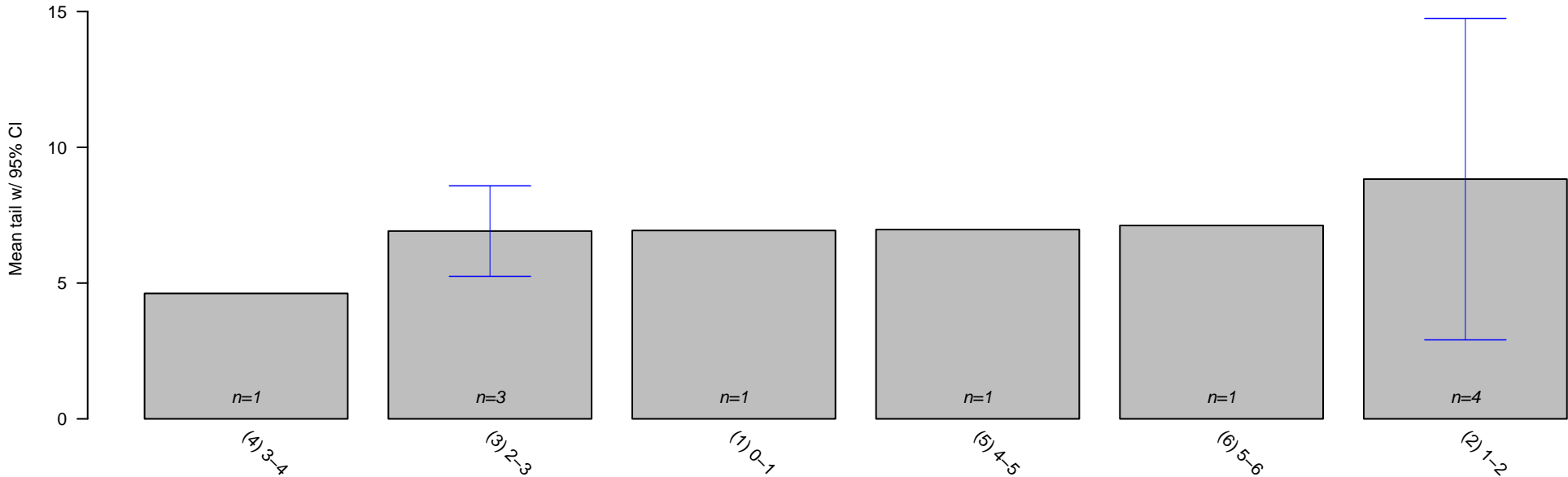
Bilirubin



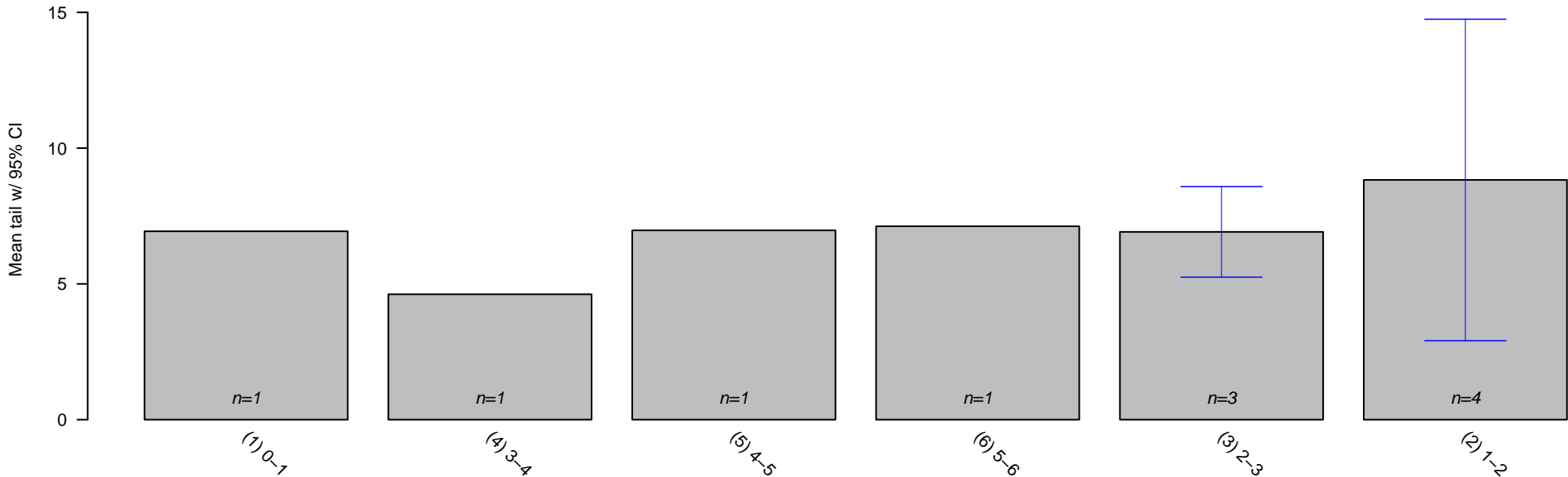
Bilirubin: Ordered by Category



Bilirubin: Ordered by Diversity

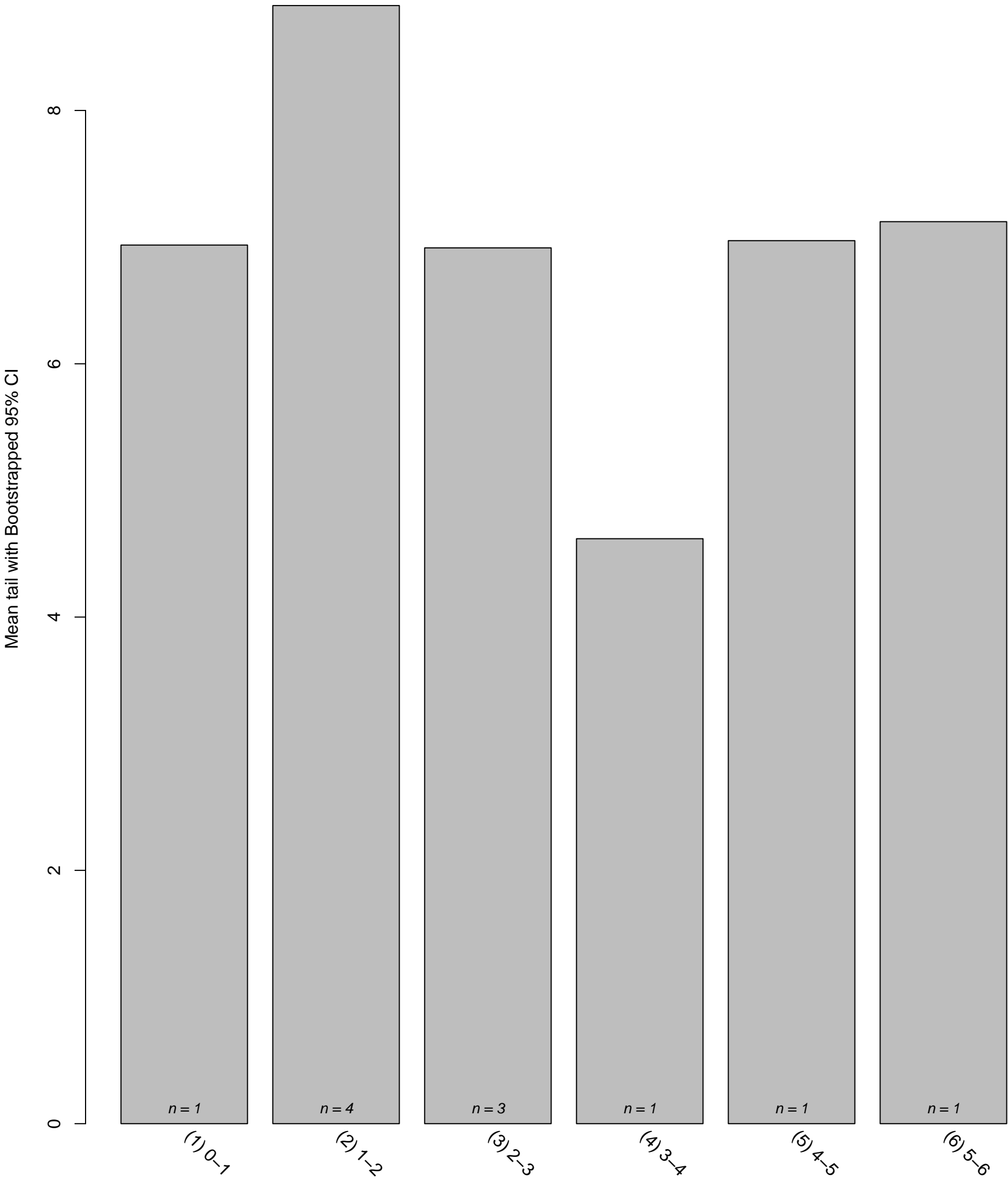


Bilirubin: Ordered by Sample Size

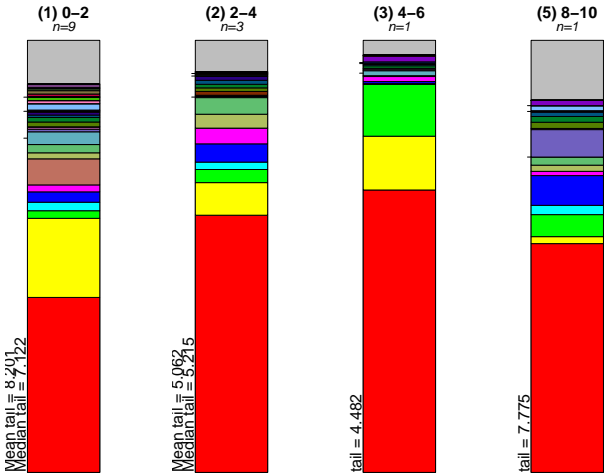


Bilirubin: Ordered by Category

with Wilcoxon rank sum test (difference between group means) p-values

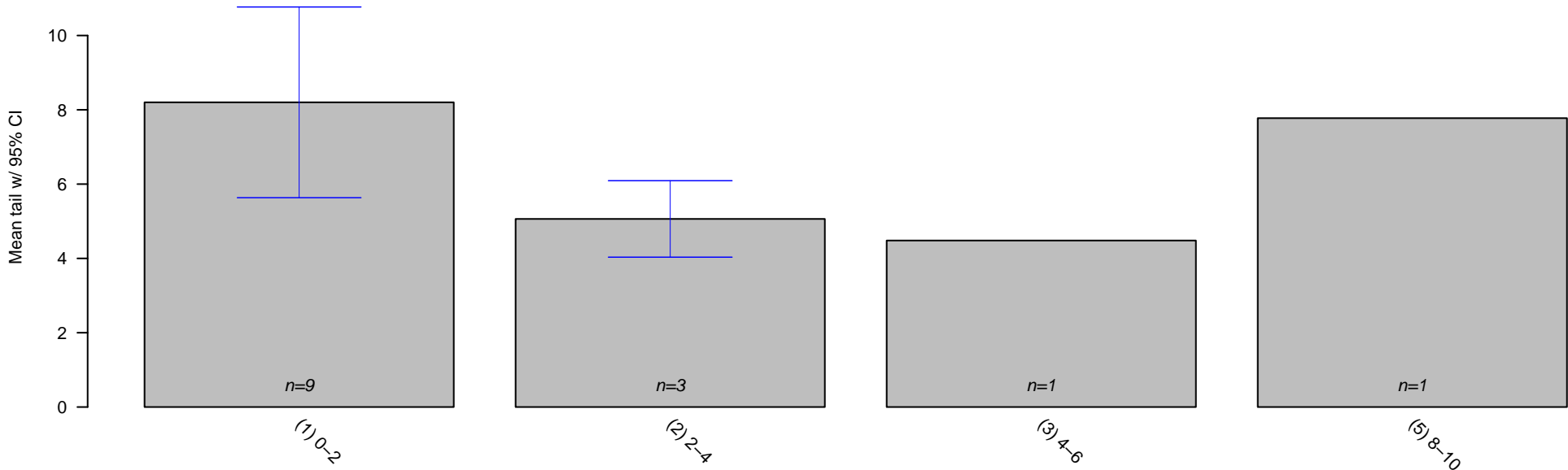


Admit

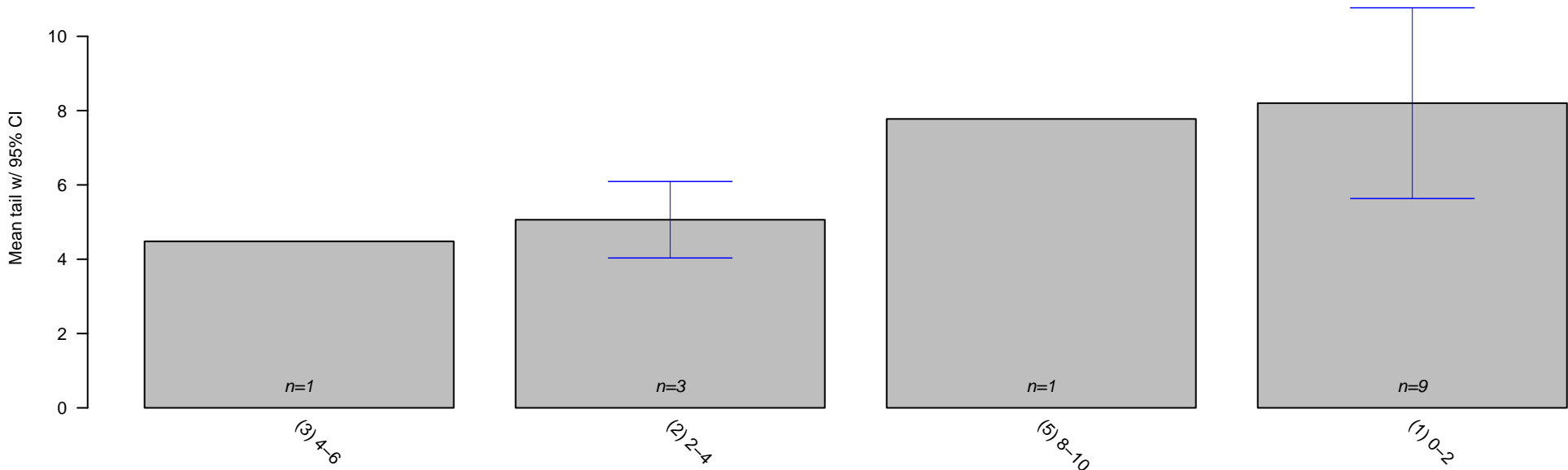


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

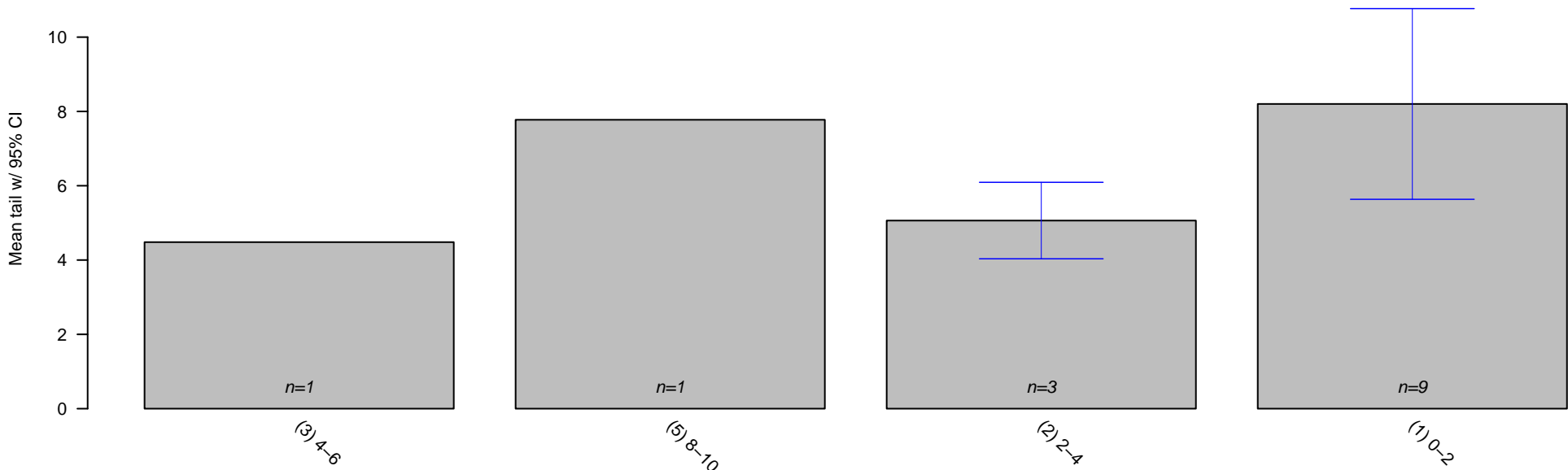
Admit: Ordered by Category



Admit: Ordered by Diversity

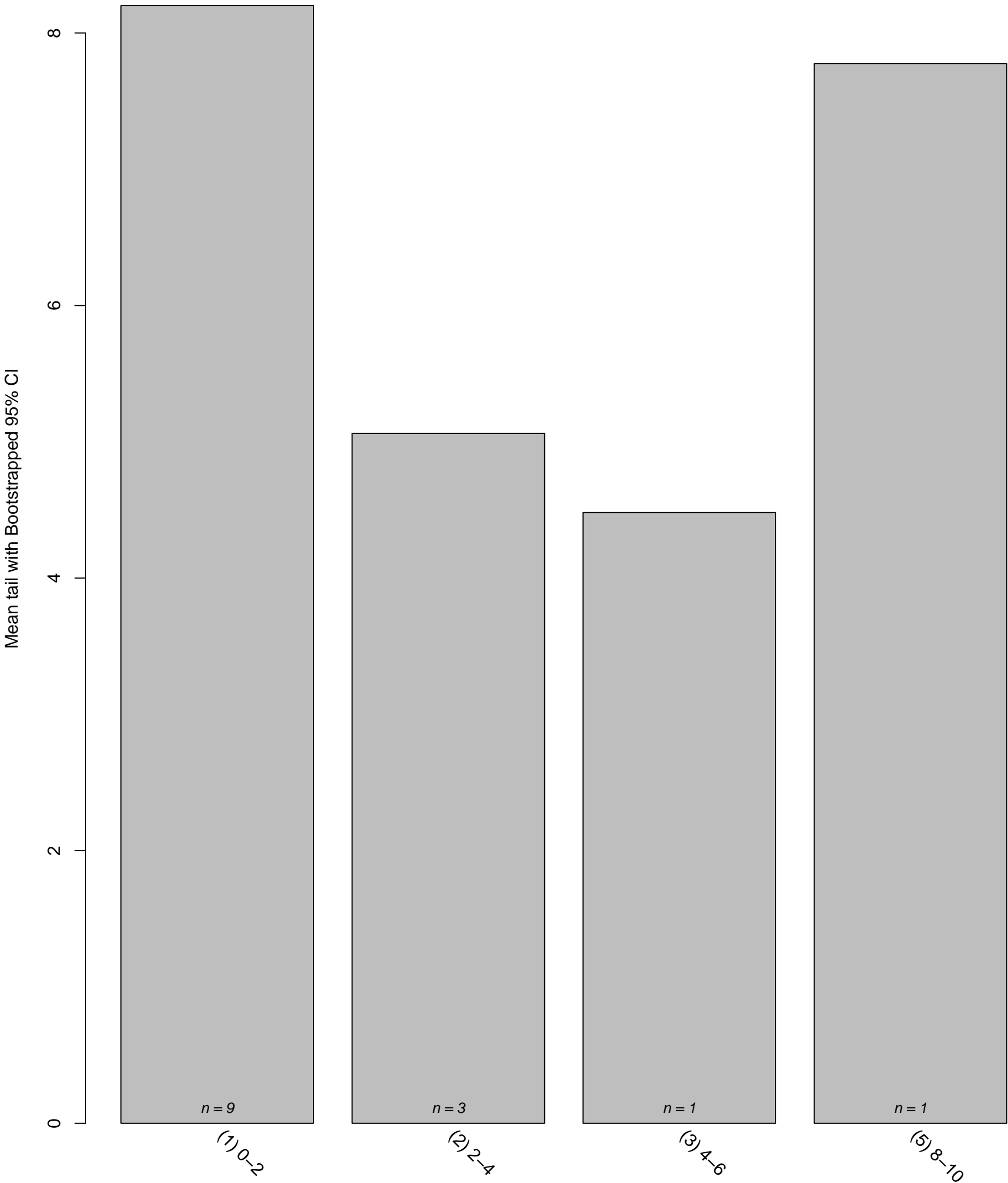


Admit: Ordered by Sample Size

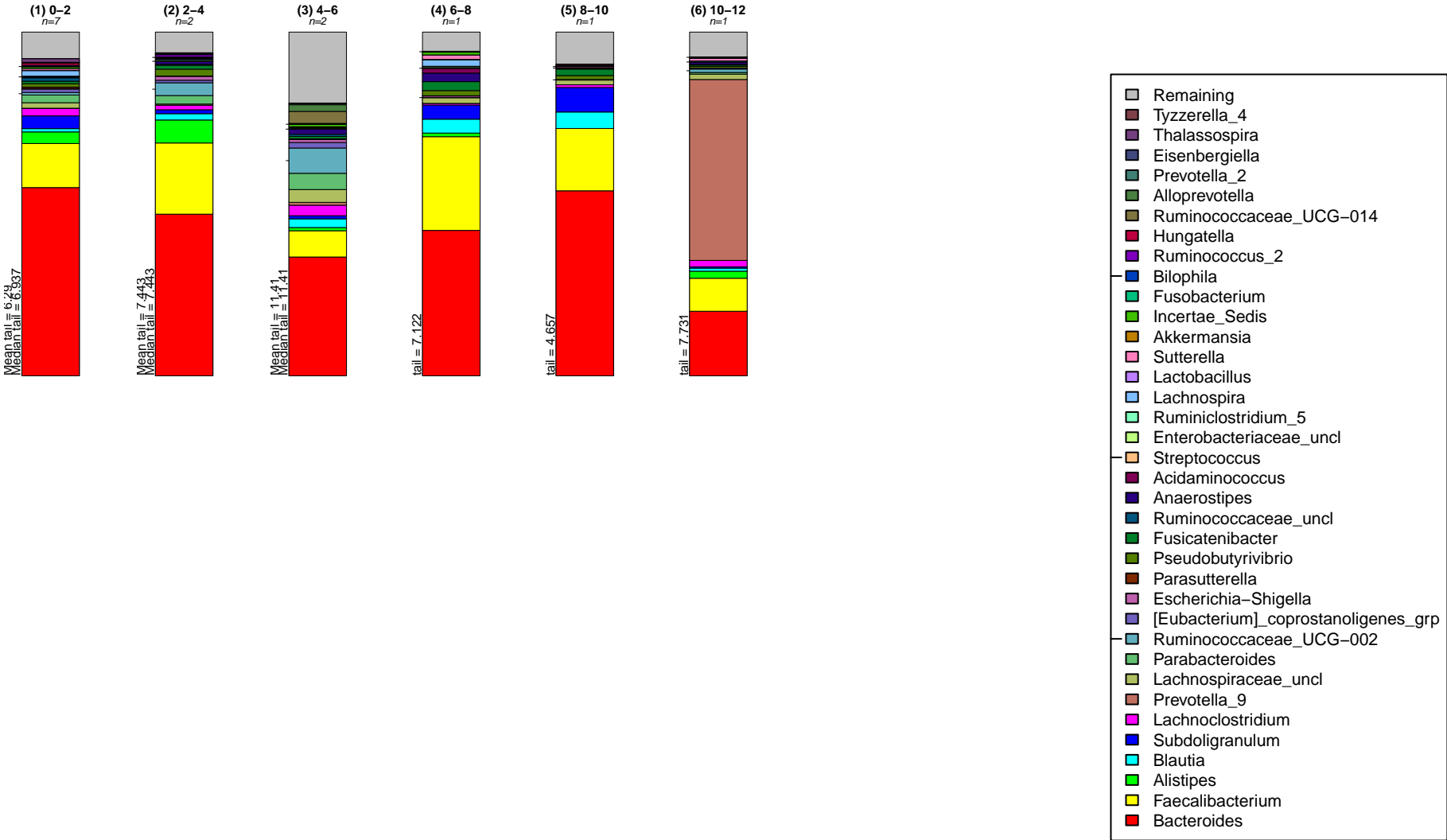


Admit: Ordered by Category

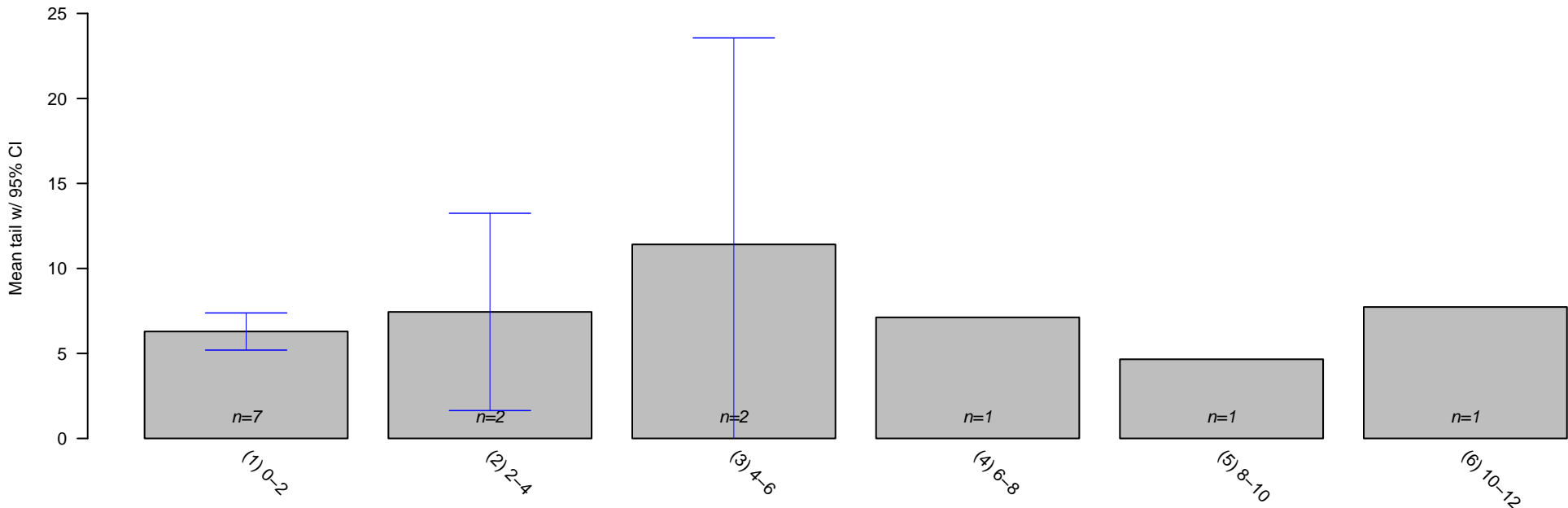
with Wilcoxon rank sum test (difference between group means) p-values



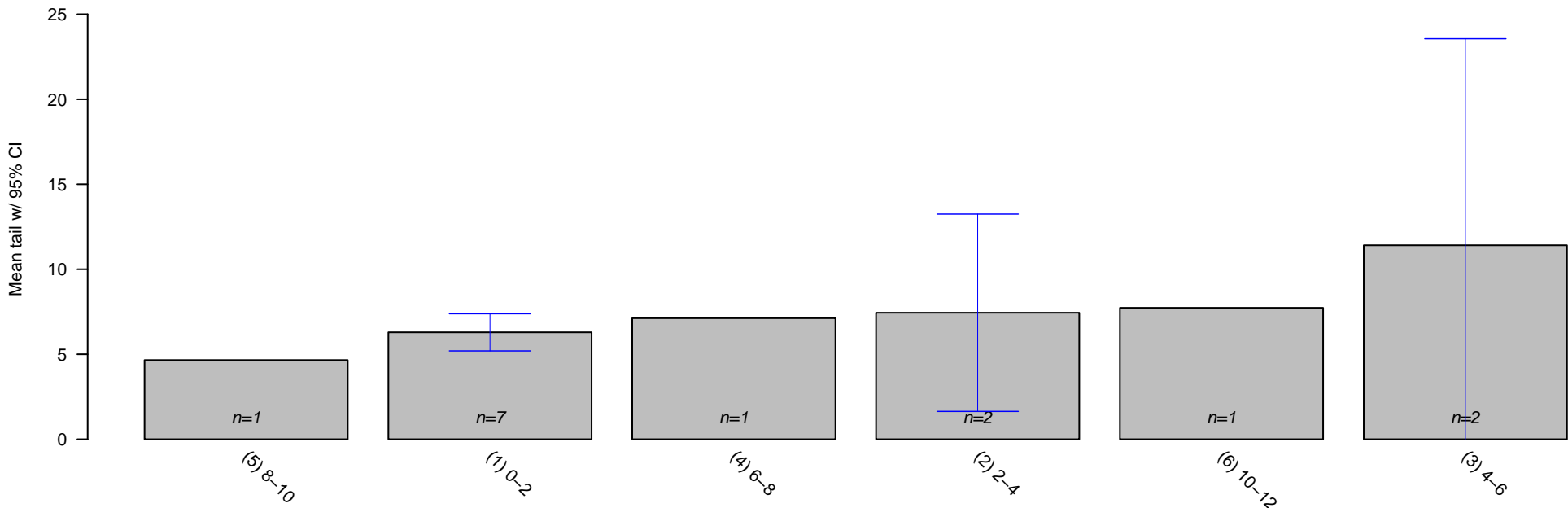
LastVOC



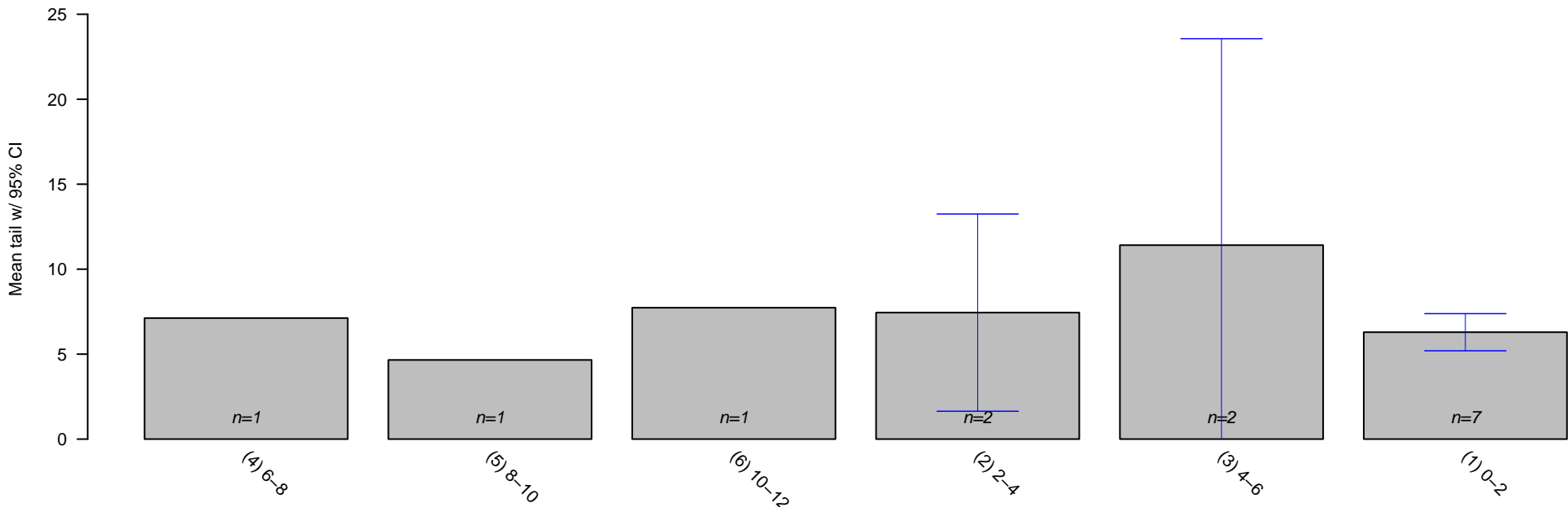
LastVOC: Ordered by Category



LastVOC: Ordered by Diversity

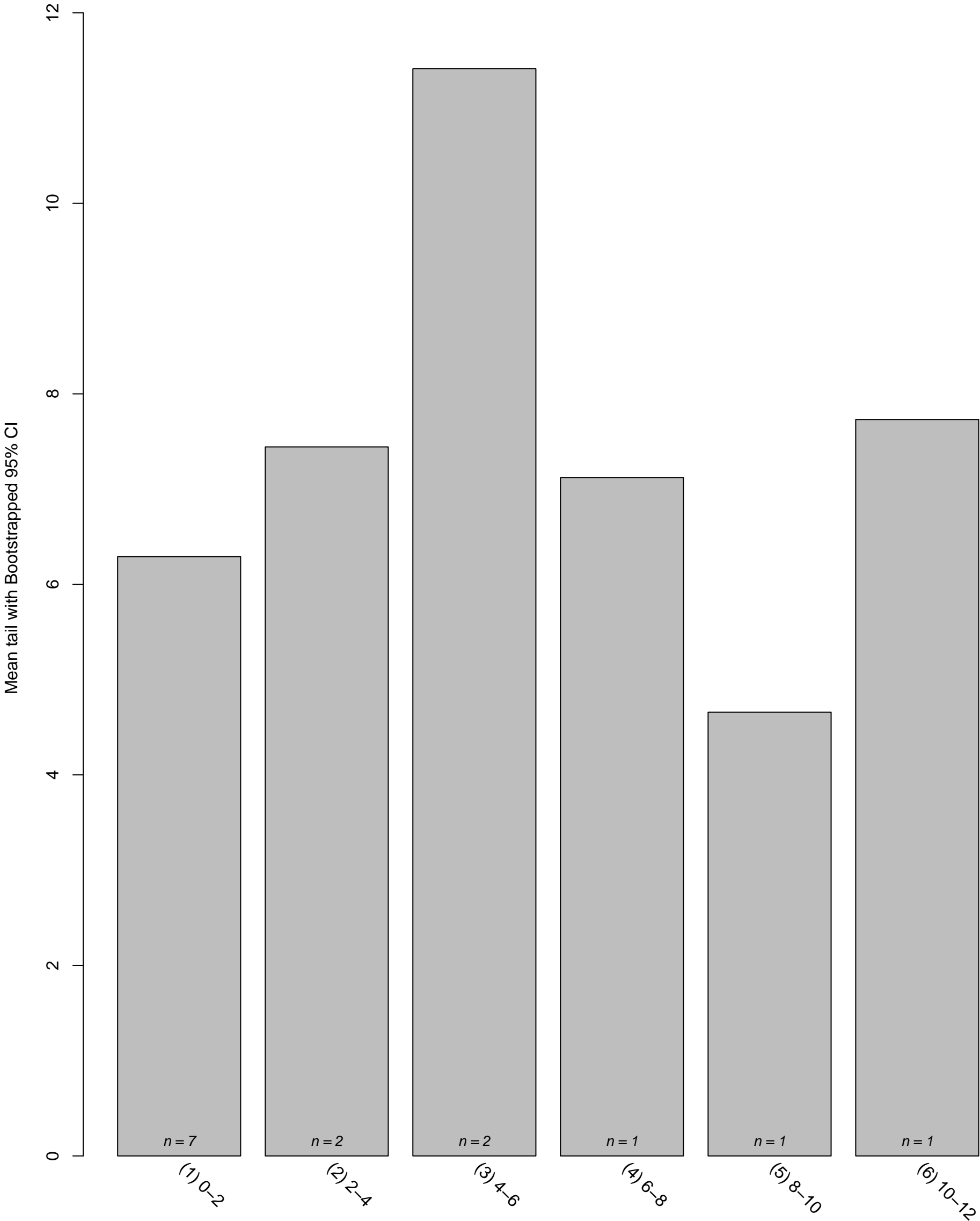


LastVOC: Ordered by Sample Size

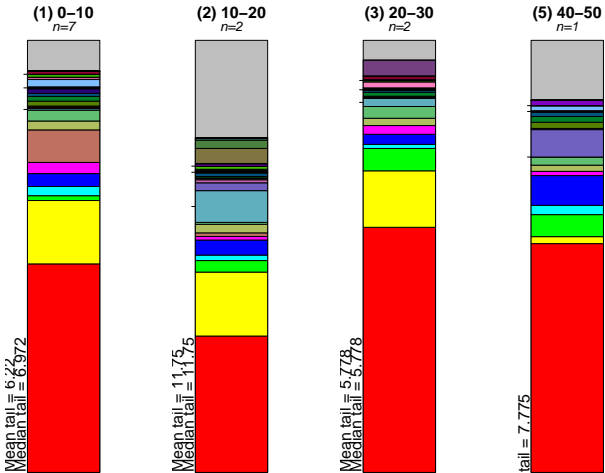


LastVOC: Ordered by Category

with Wilcoxon rank sum test (difference between group means) p-values

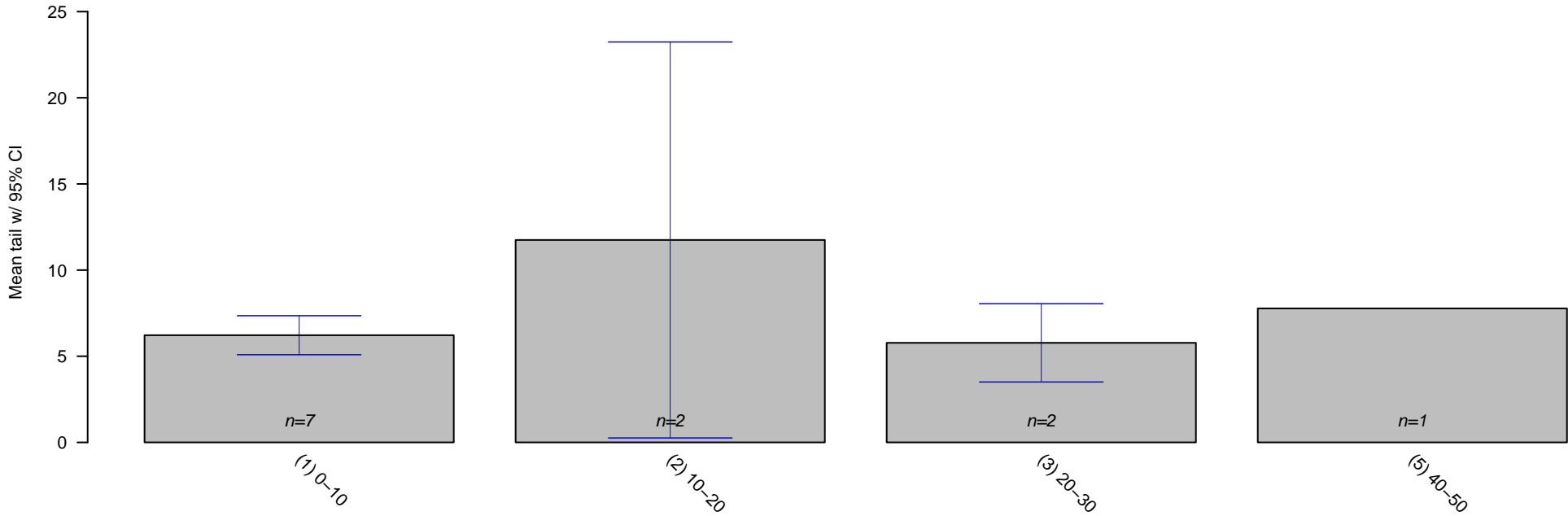


ESR

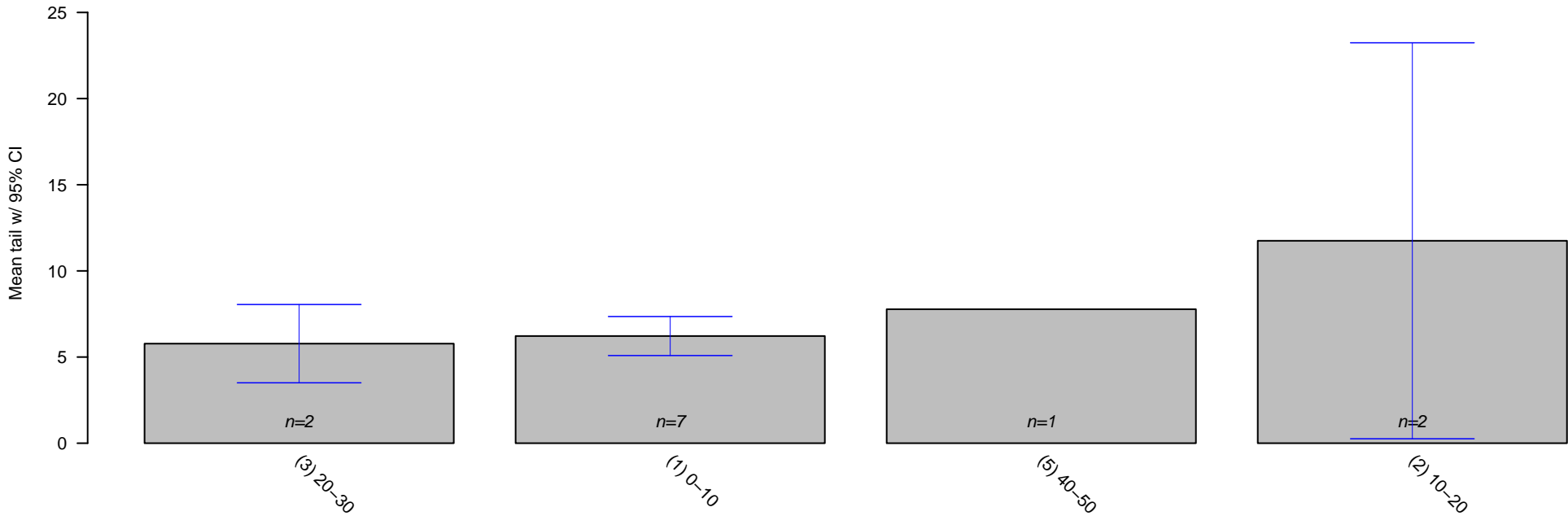


- Remaining
- Tyzzarella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

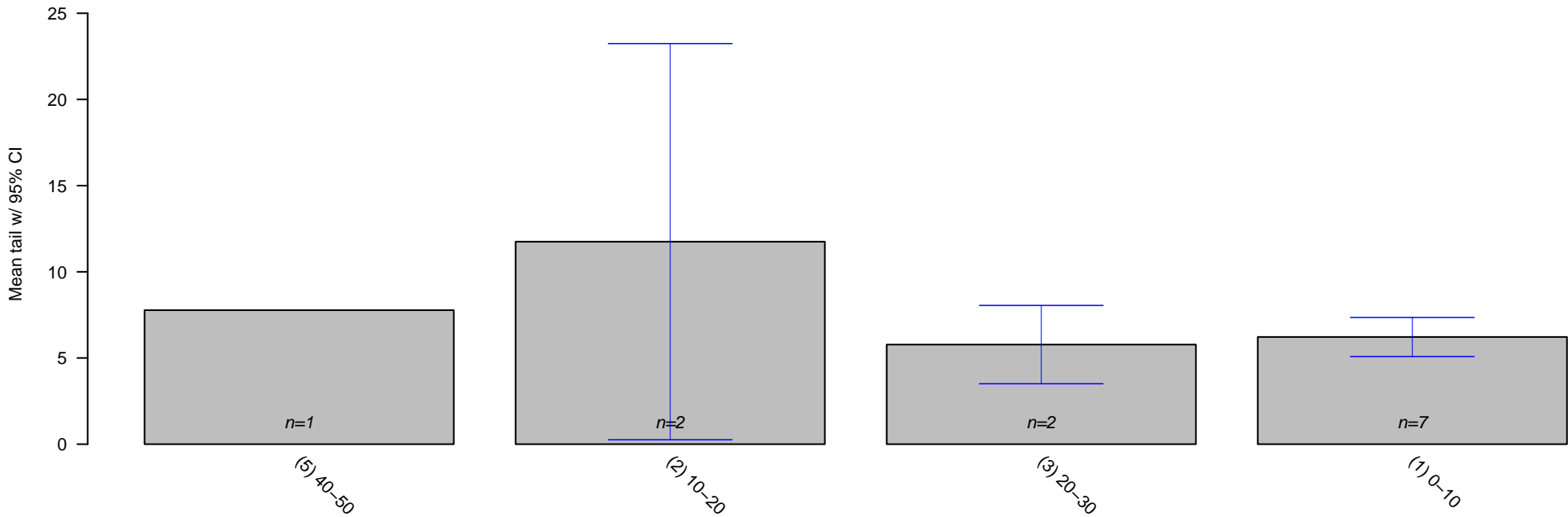
ESR: Ordered by Category



ESR: Ordered by Diversity

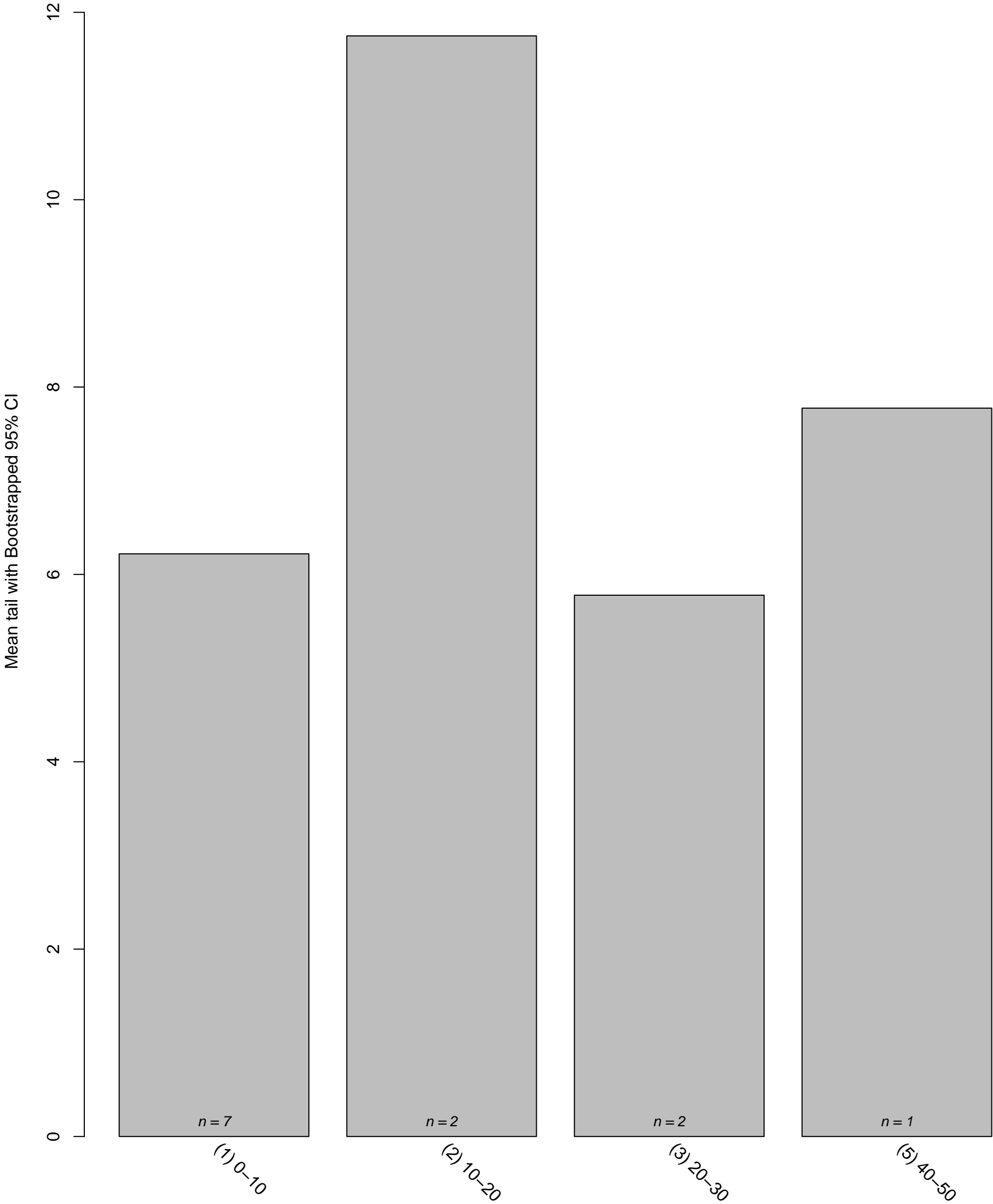


ESR: Ordered by Sample Size

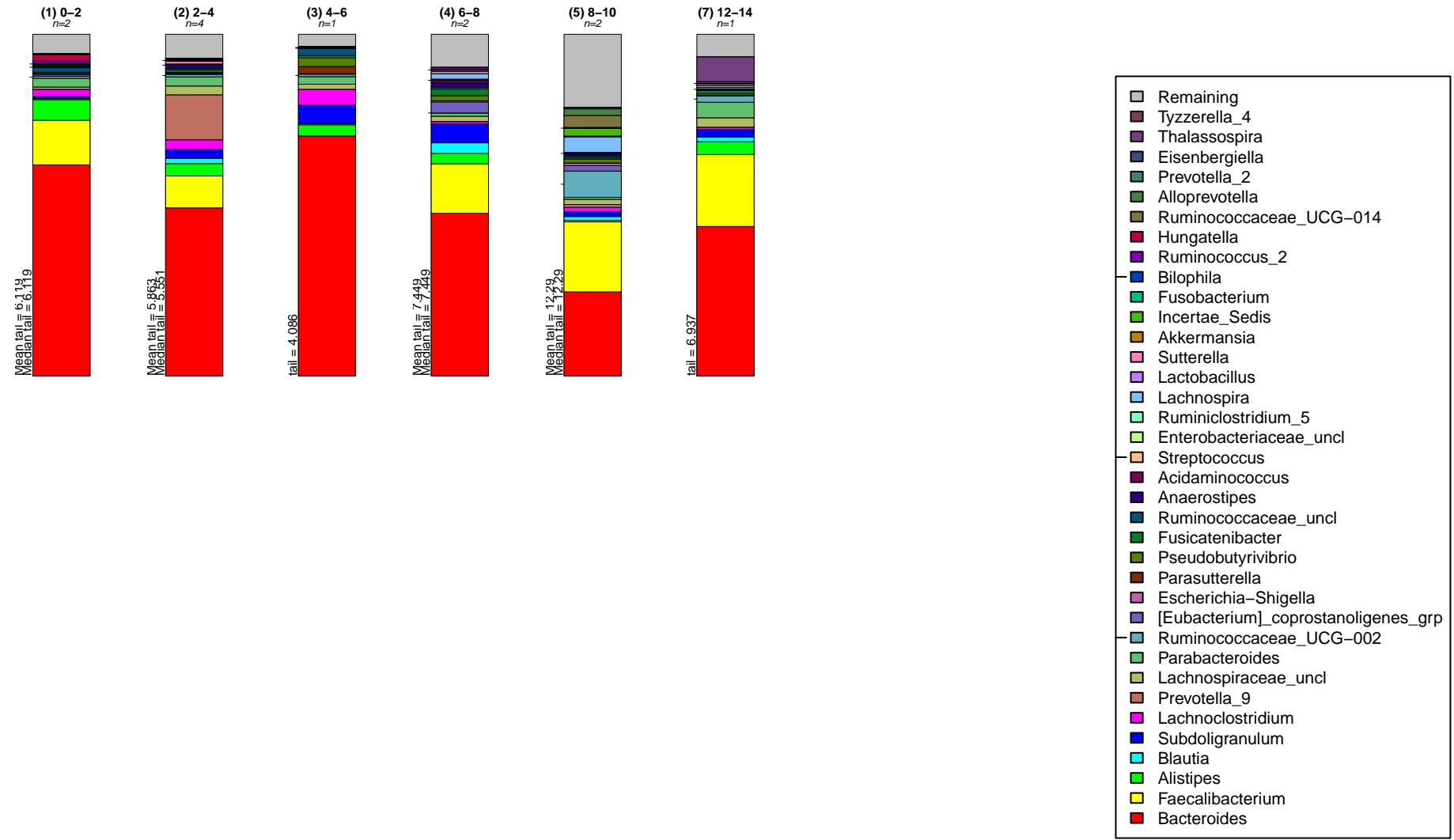


ESR: Ordered by Category

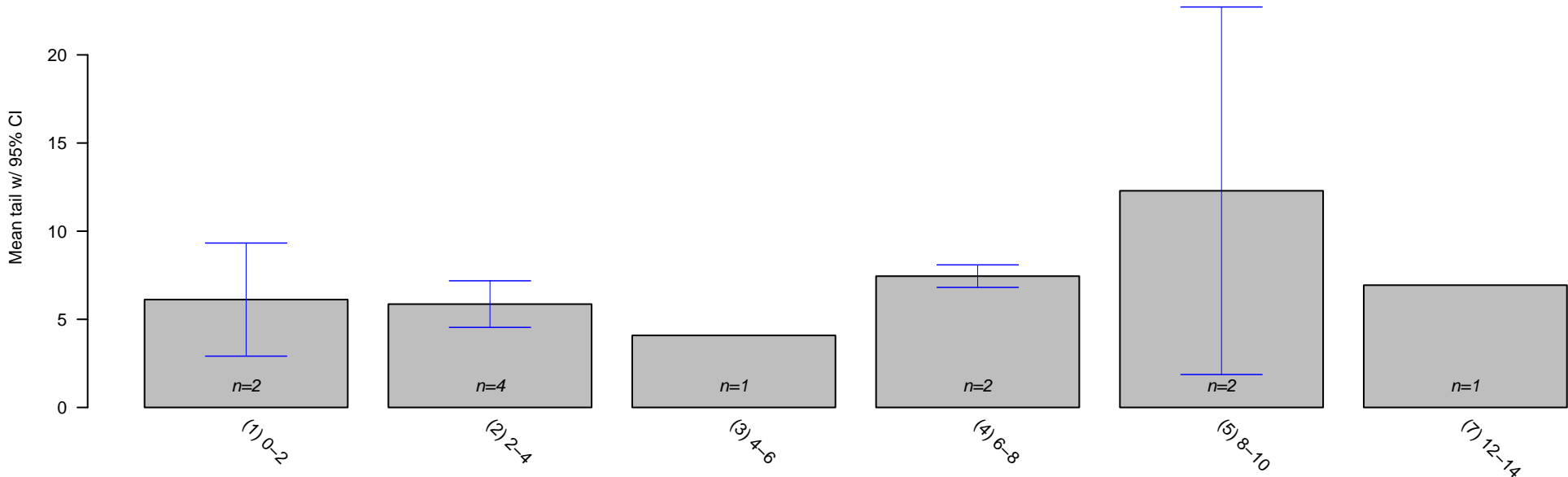
with Wilcoxon rank sum test (difference between group means) *p*-values



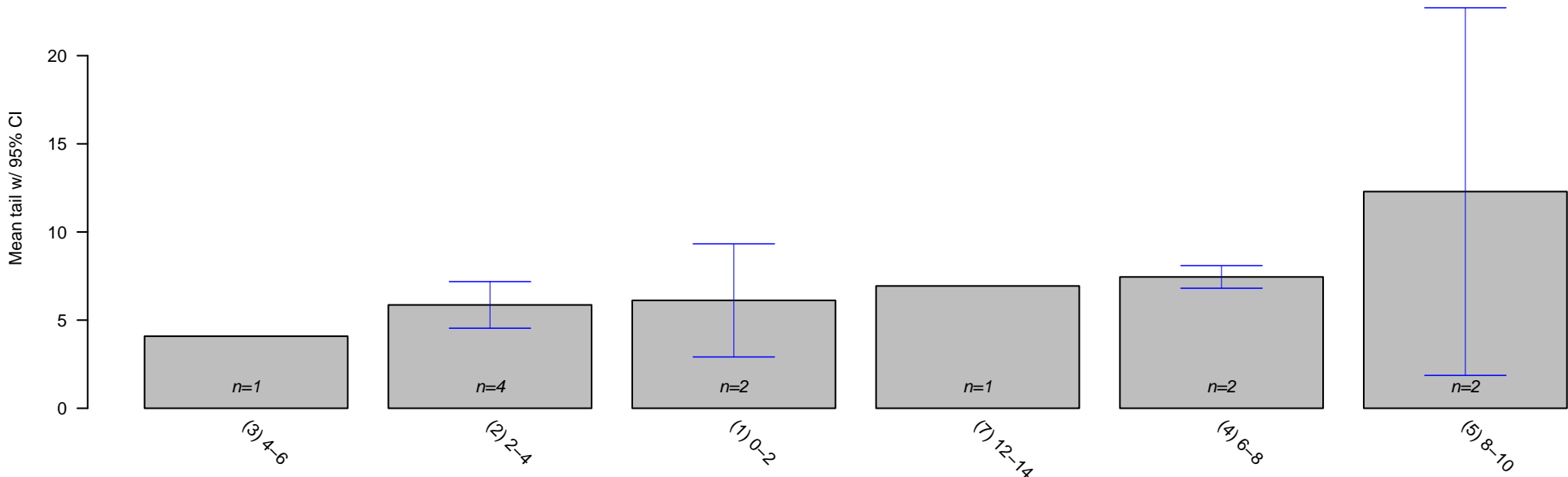
CRP



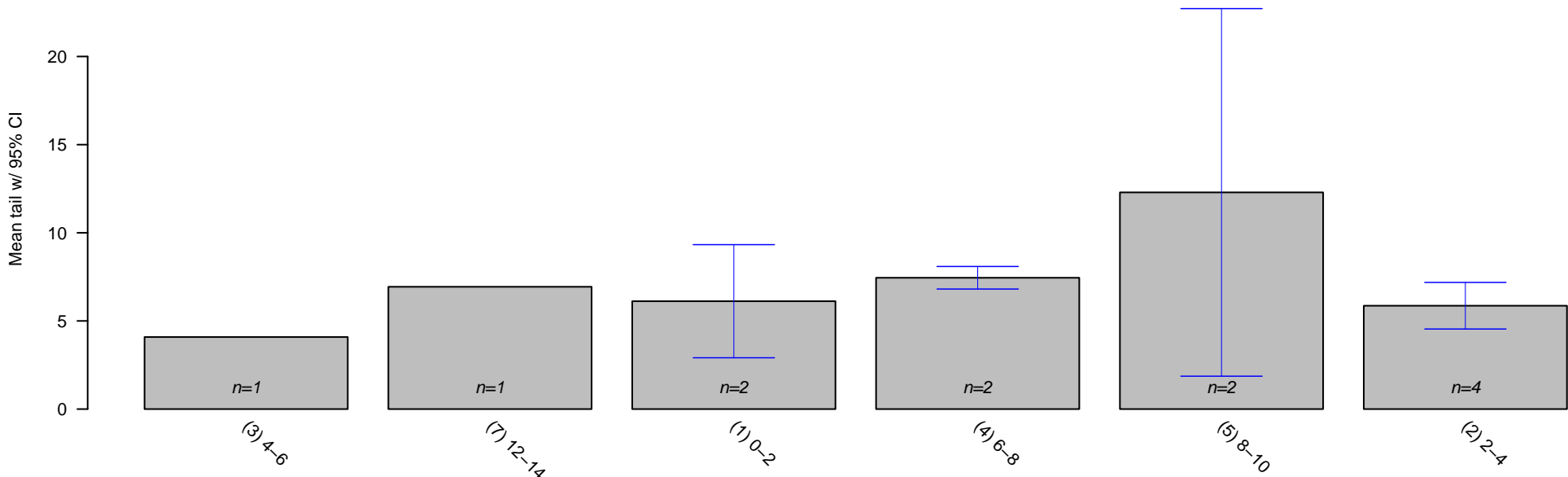
CRP: Ordered by Category



CRP: Ordered by Diversity

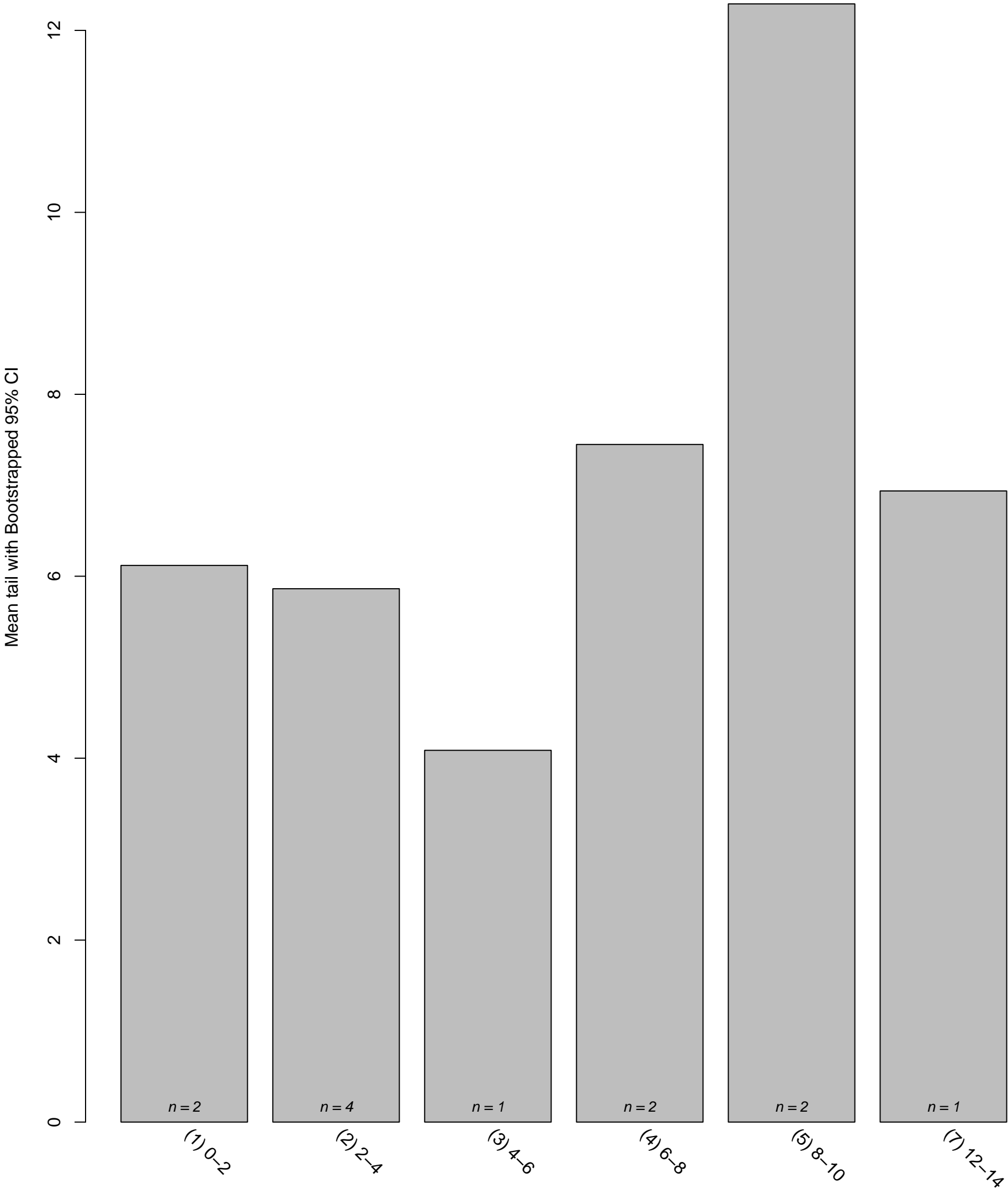


CRP: Ordered by Sample Size

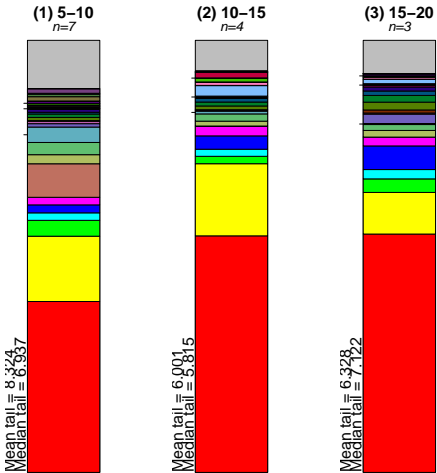


CRP: Ordered by Category

with Wilcoxon rank sum test (difference between group means) p-values

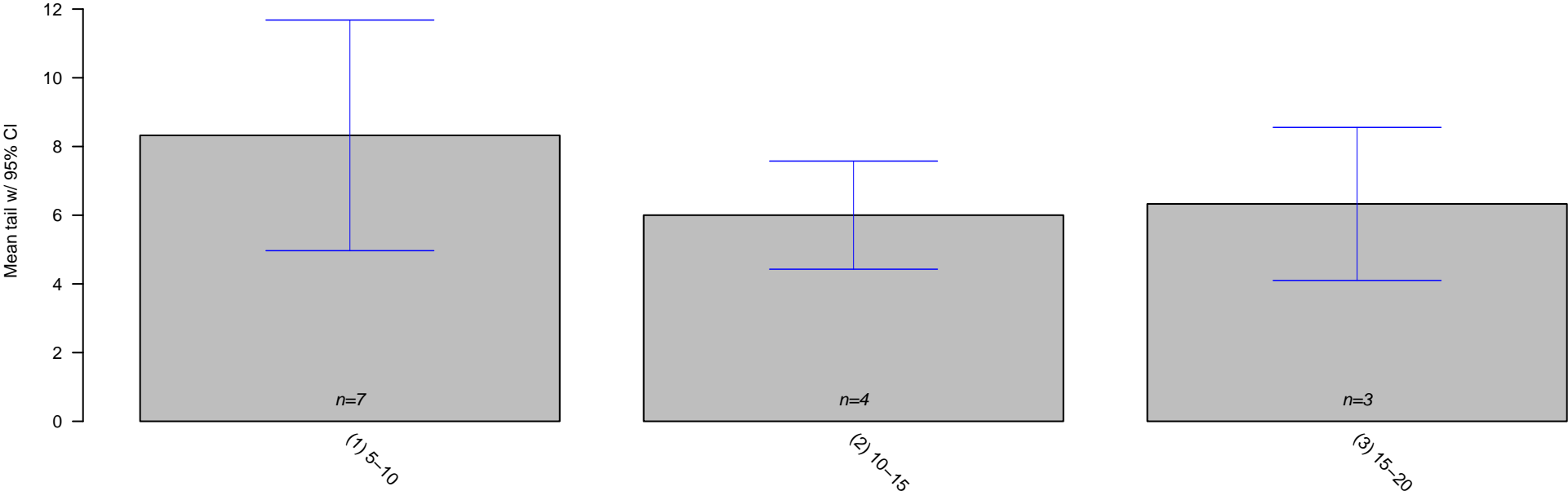


WBC

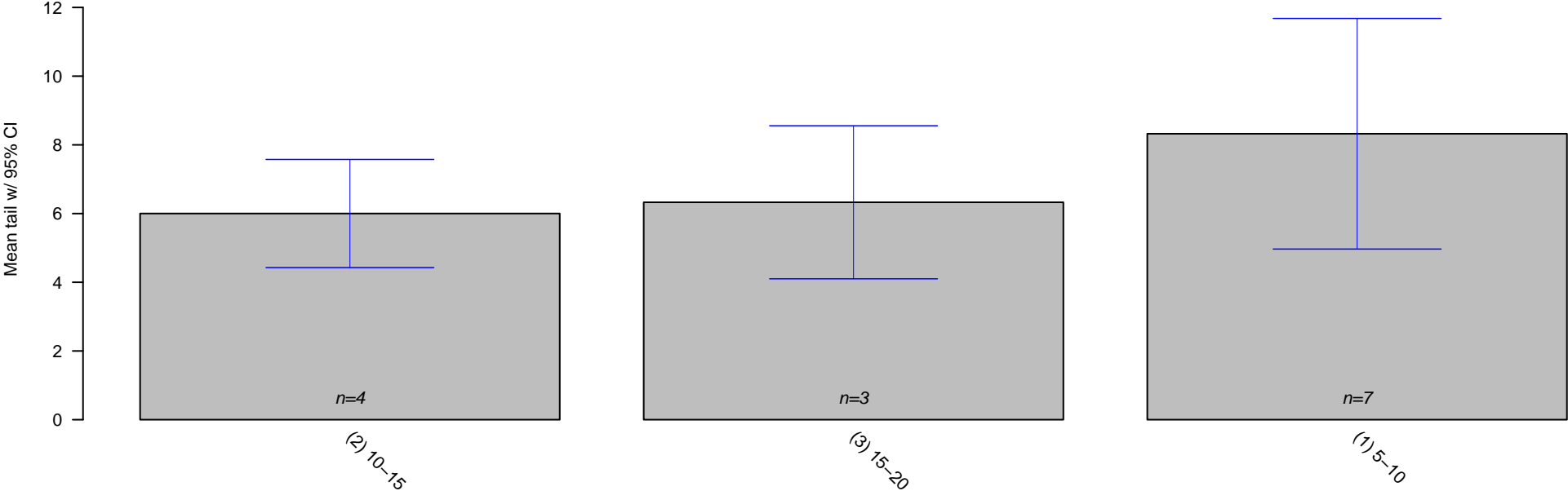


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

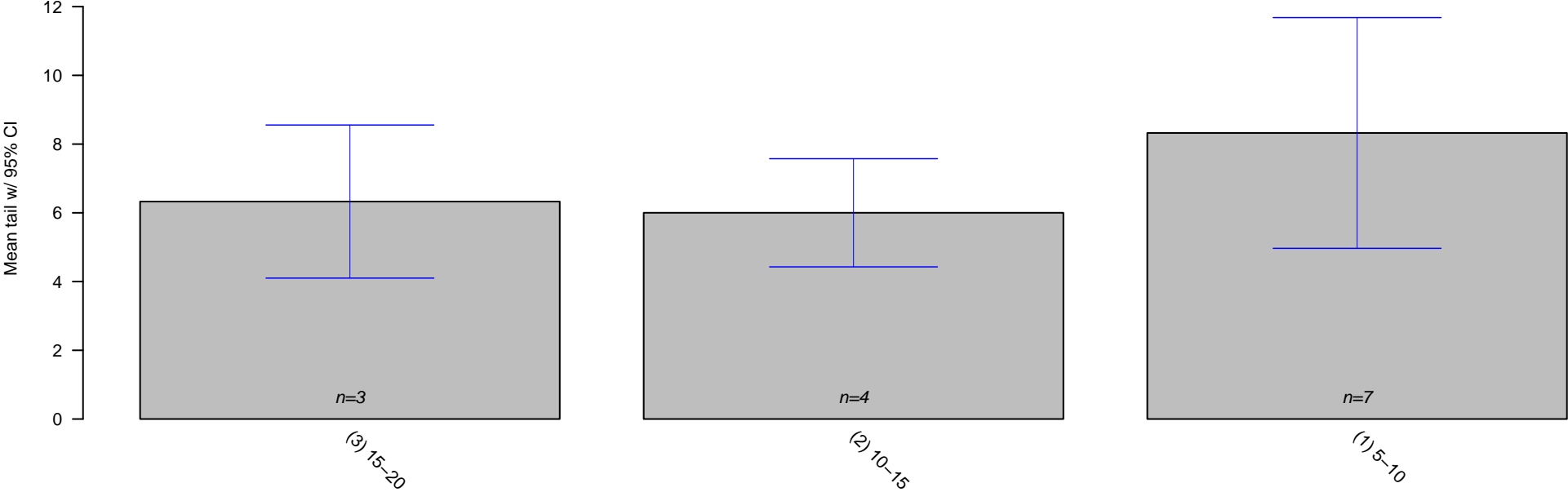
WBC: Ordered by Category



WBC: Ordered by Diversity

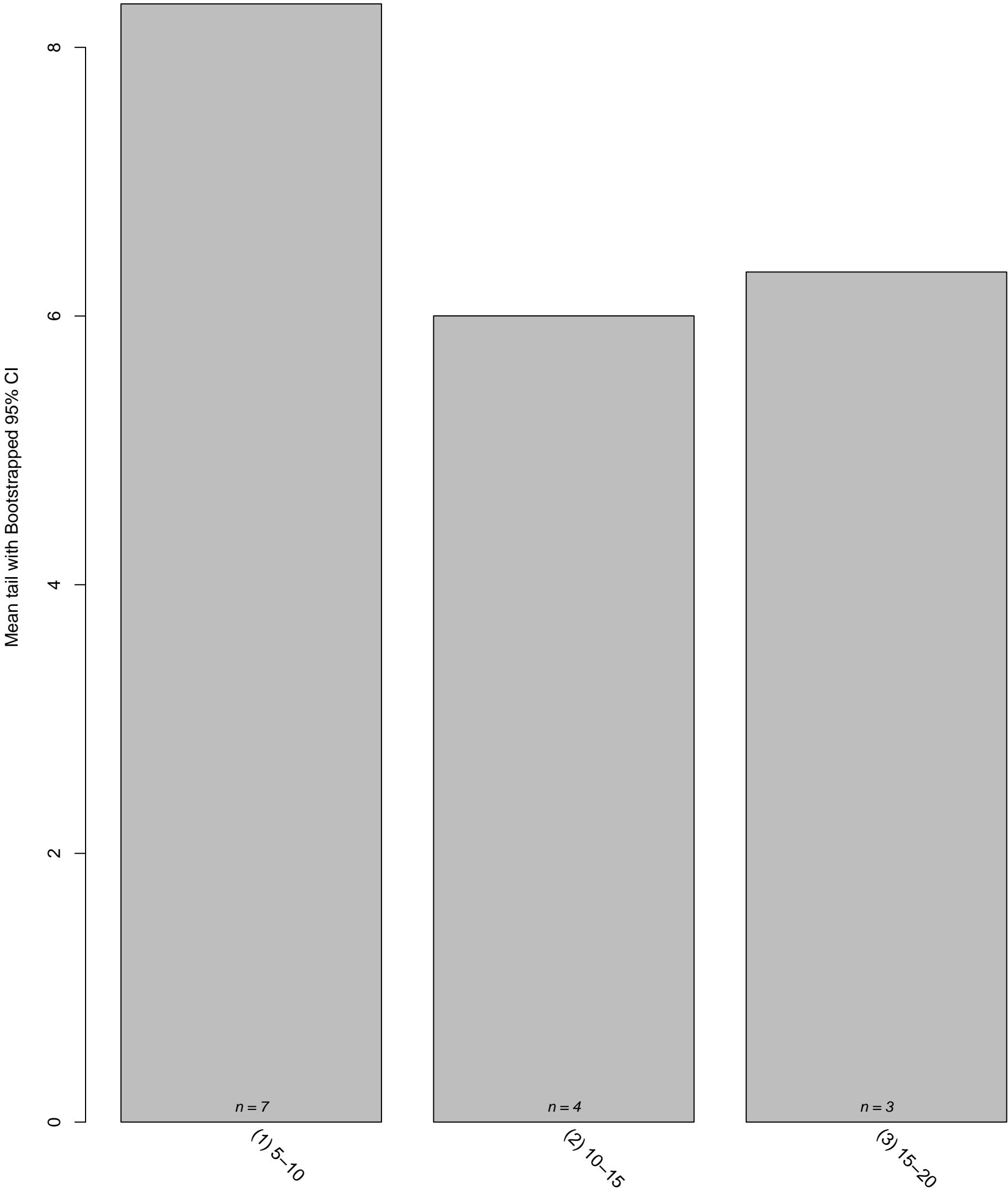


WBC: Ordered by Sample Size

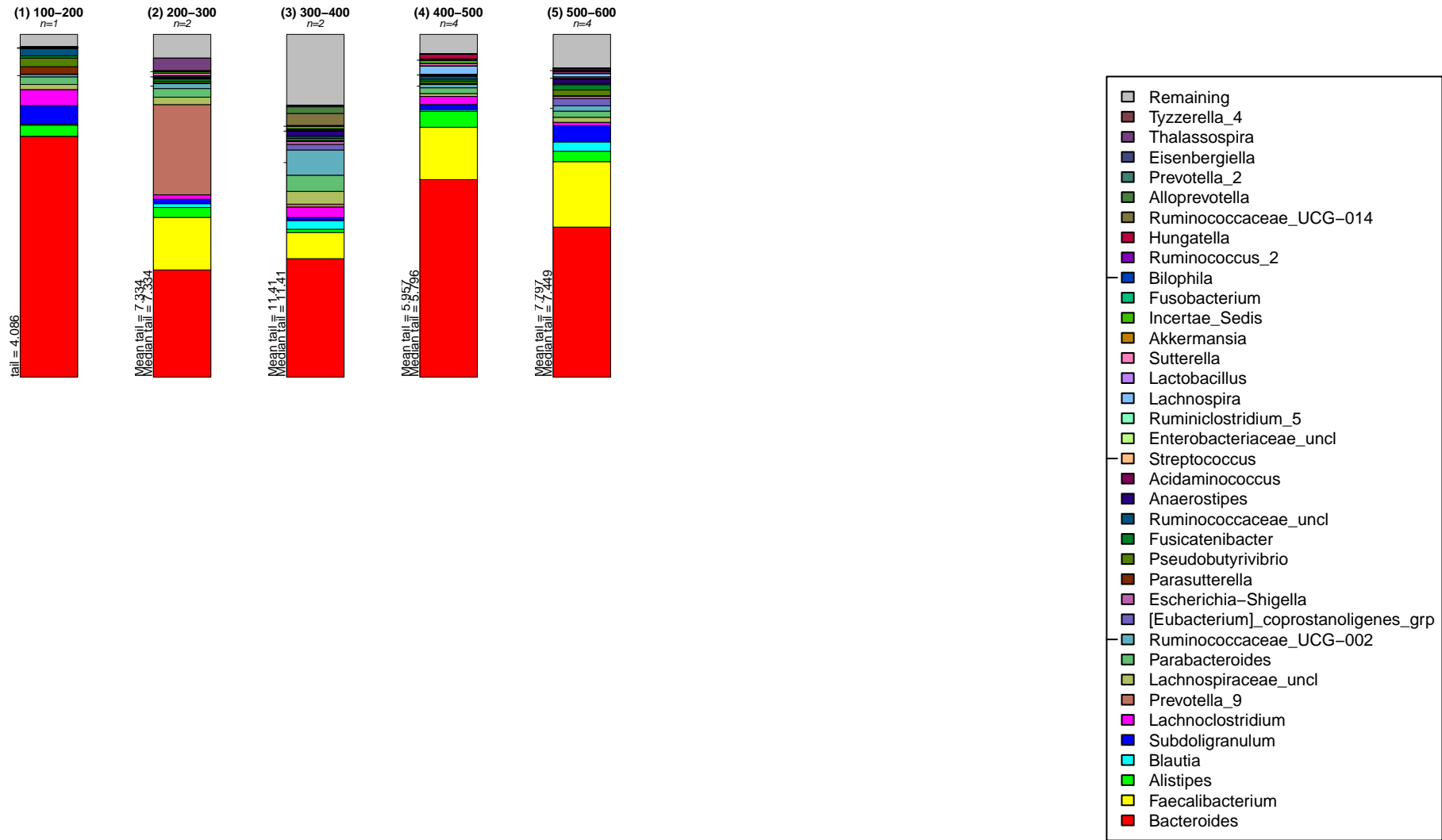


WBC: Ordered by Category

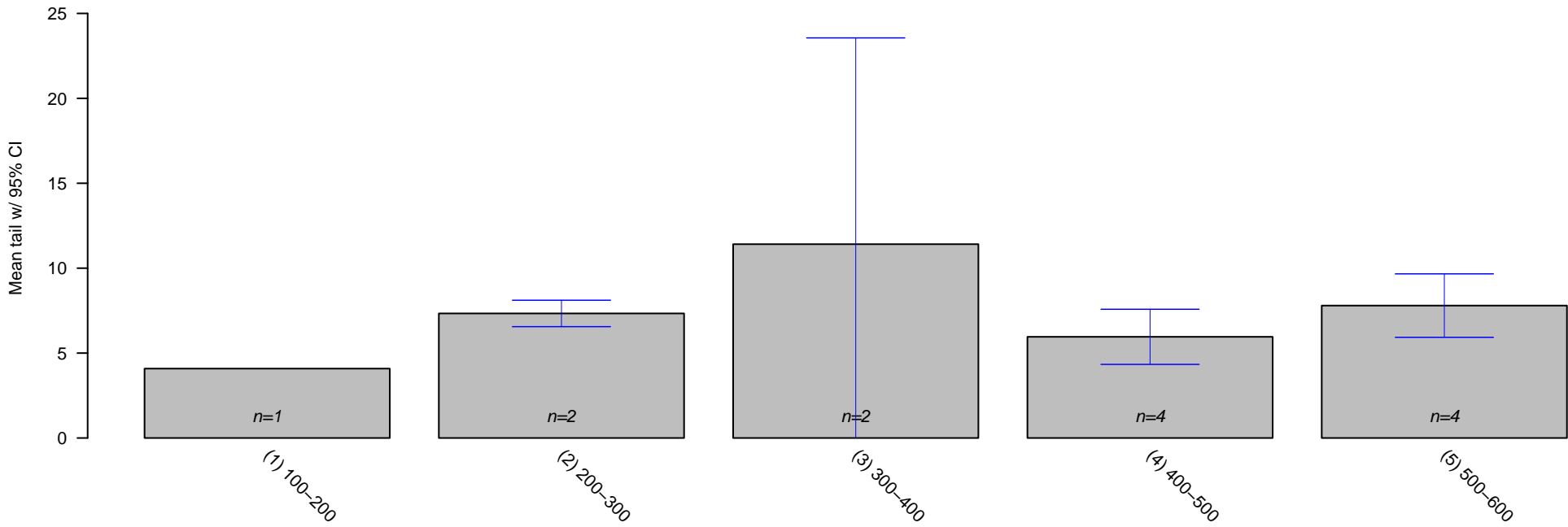
with Wilcoxon rank sum test (difference between group means) *p*-values



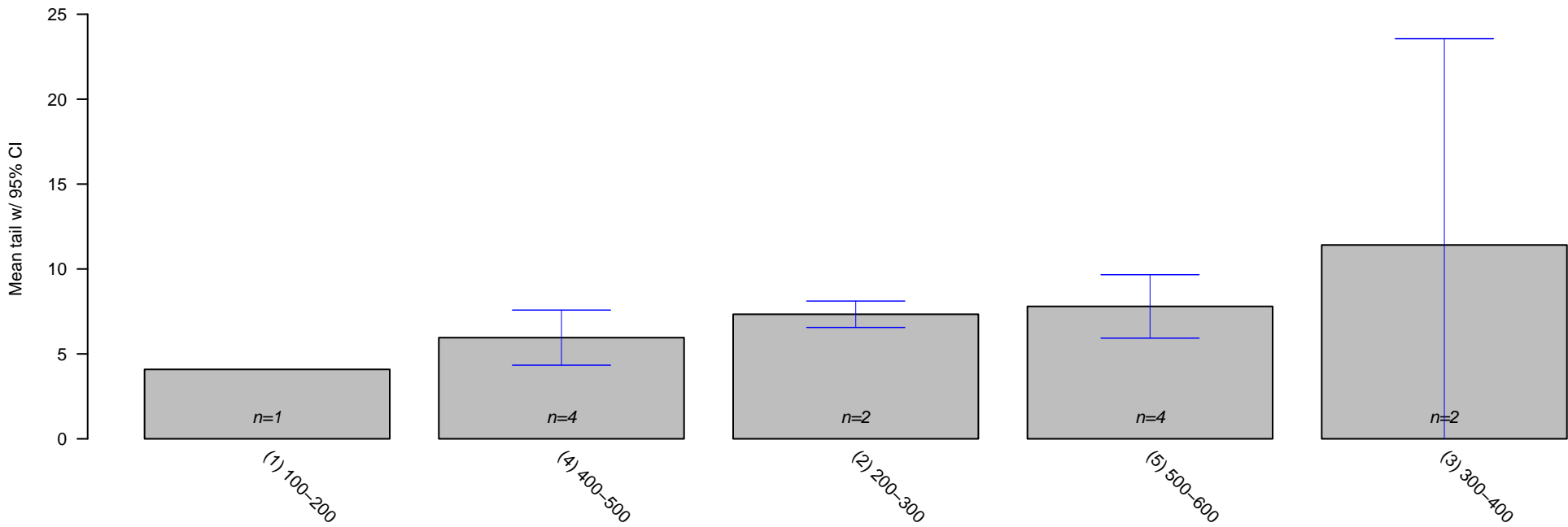
LDH



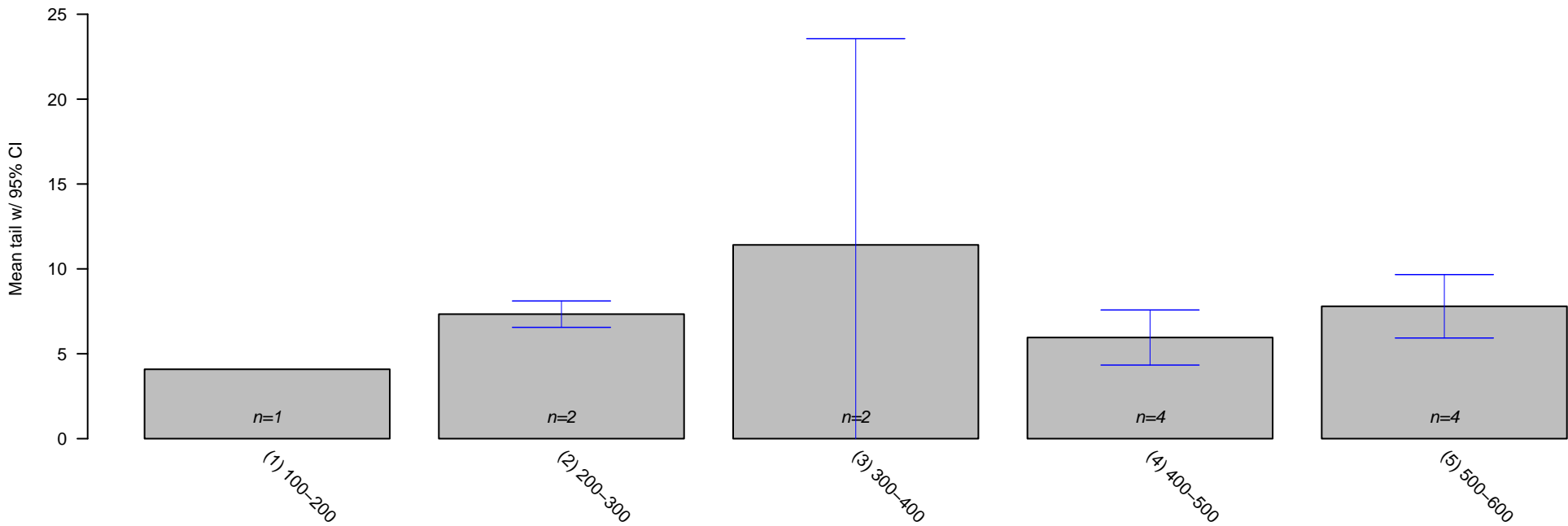
LDH: Ordered by Category



LDH: Ordered by Diversity

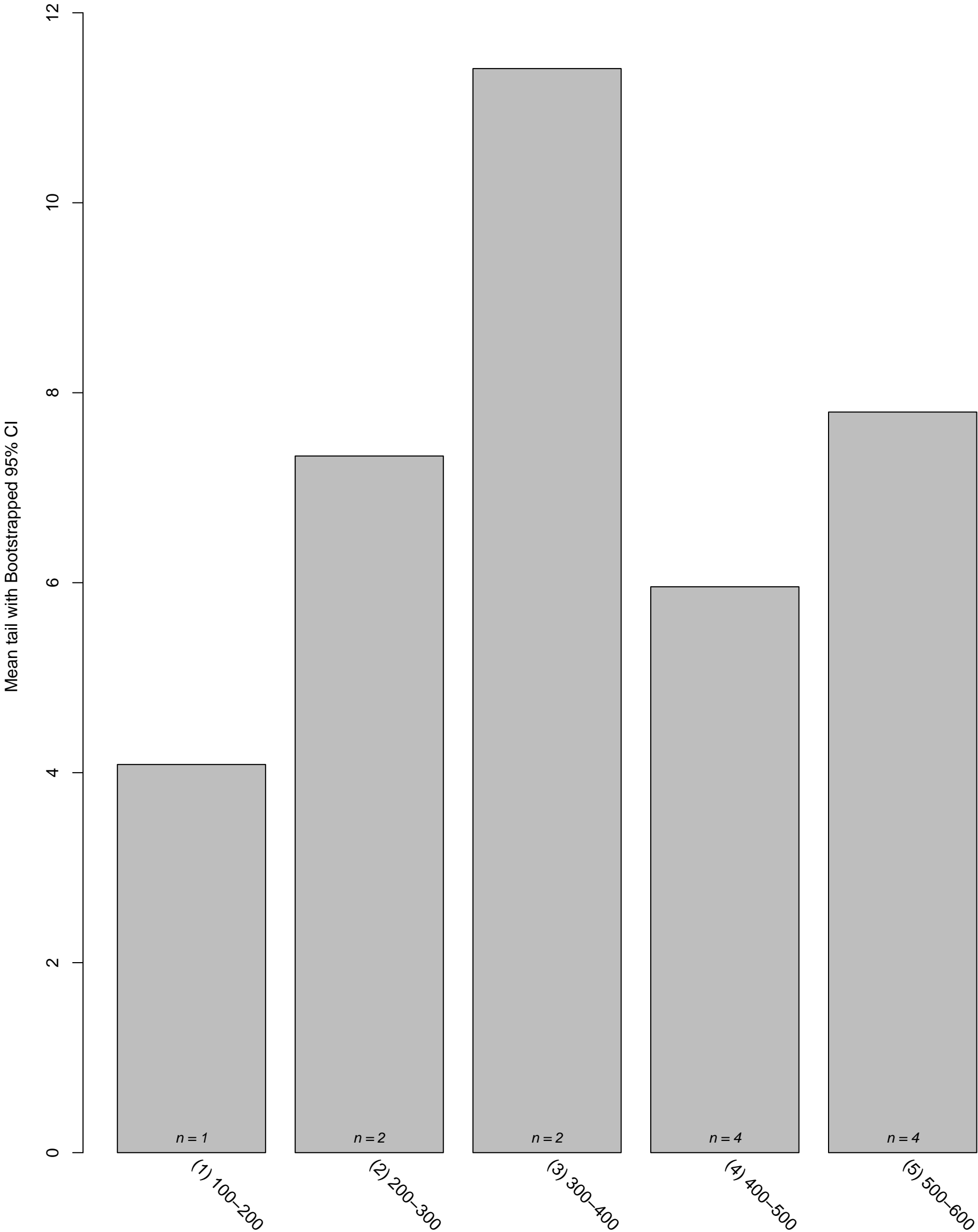


LDH: Ordered by Sample Size

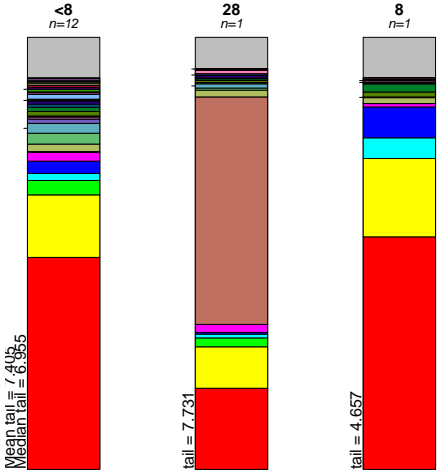


LDH: Ordered by Category

with Wilcoxon rank sum test (difference between group means) *p*-values

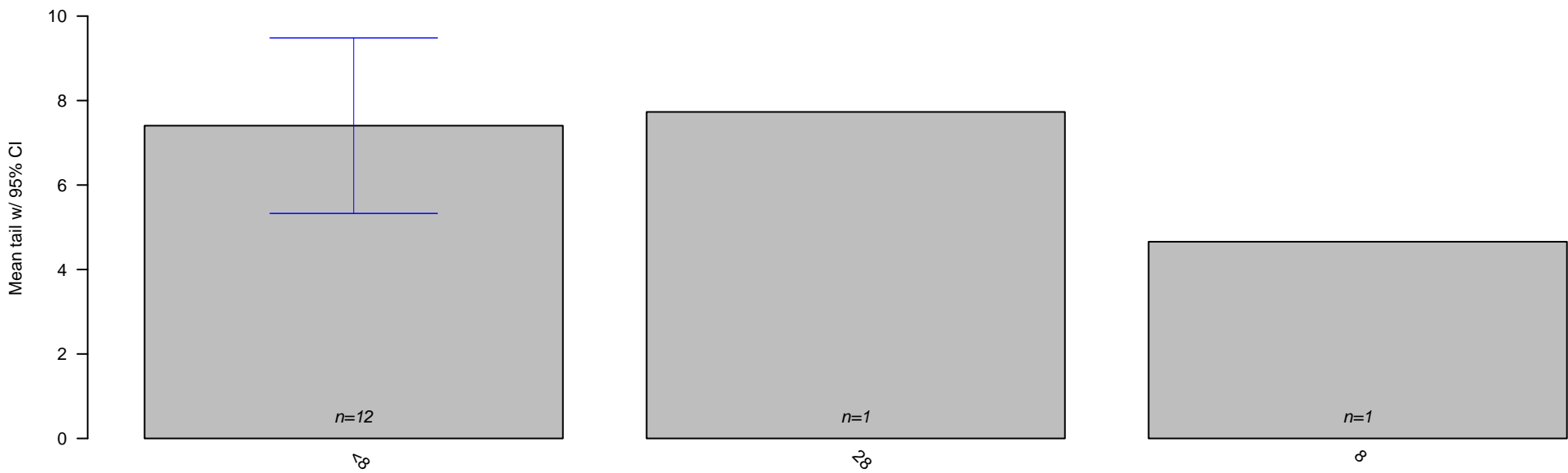


Hapto

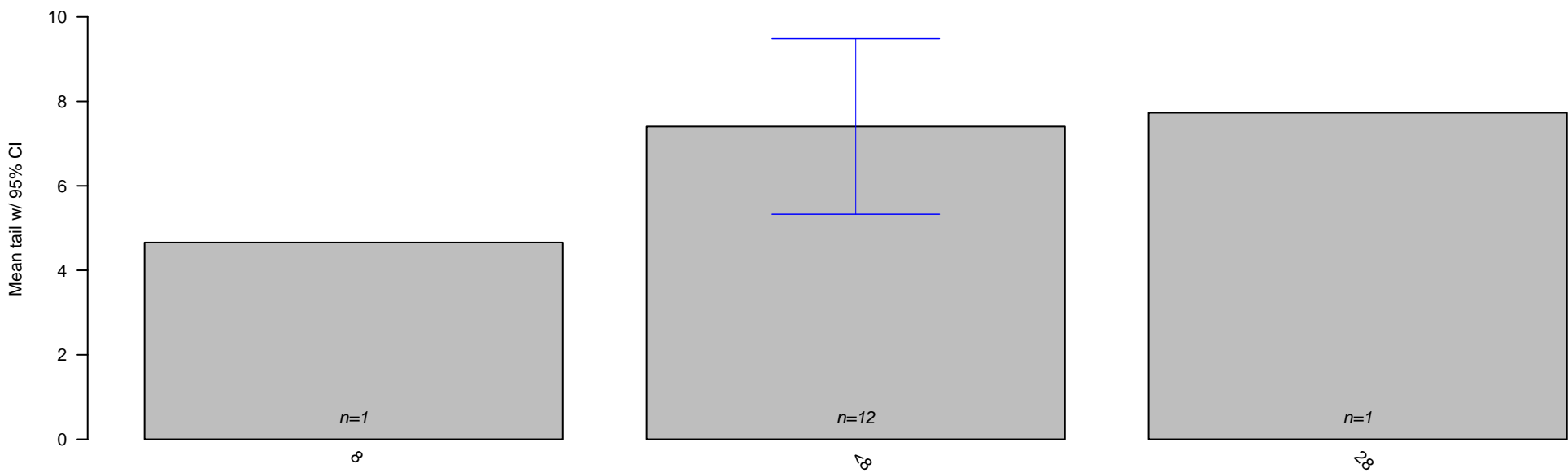


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

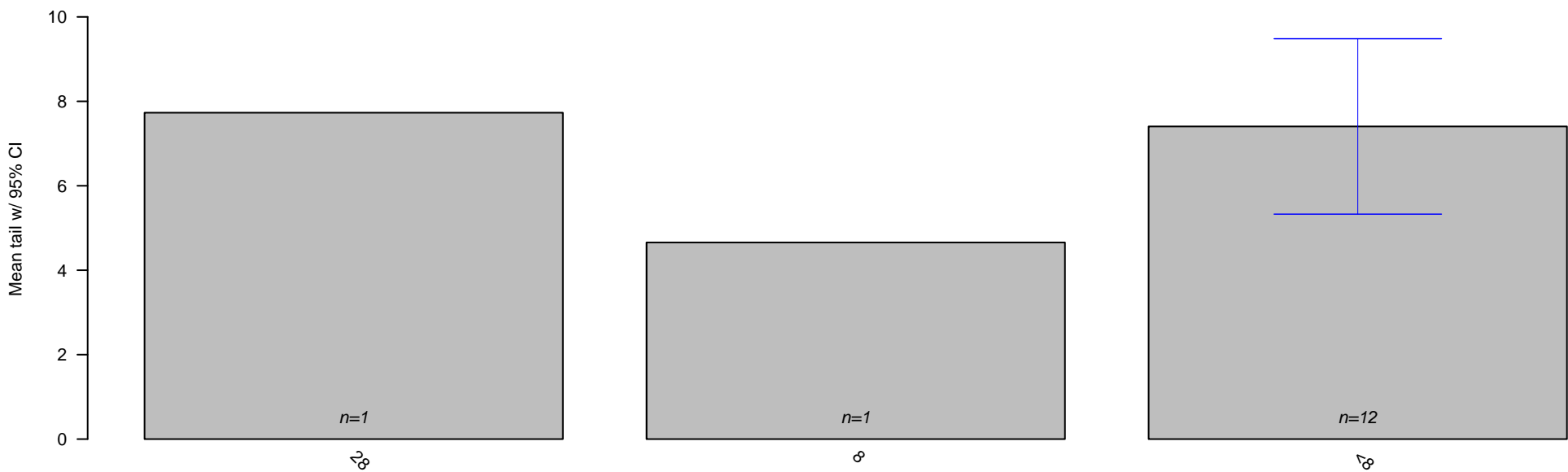
Hapto: Ordered by Category



Hapto: Ordered by Diversity

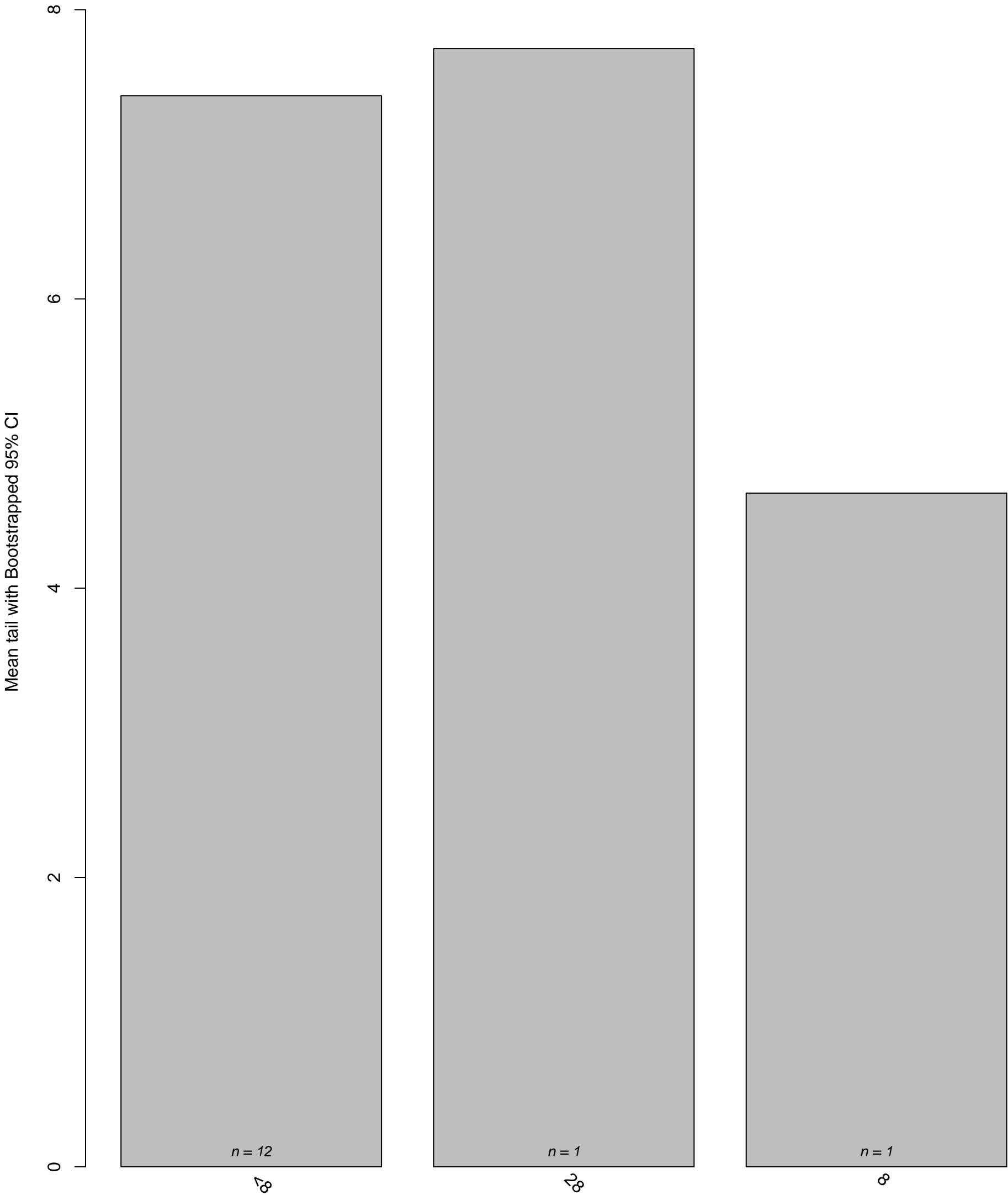


Hapto: Ordered by Sample Size

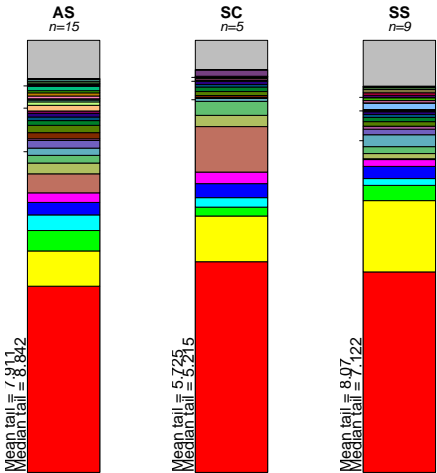


Hapto: Ordered by Category

with Wilcoxon rank sum test (difference between group means) *p*-values

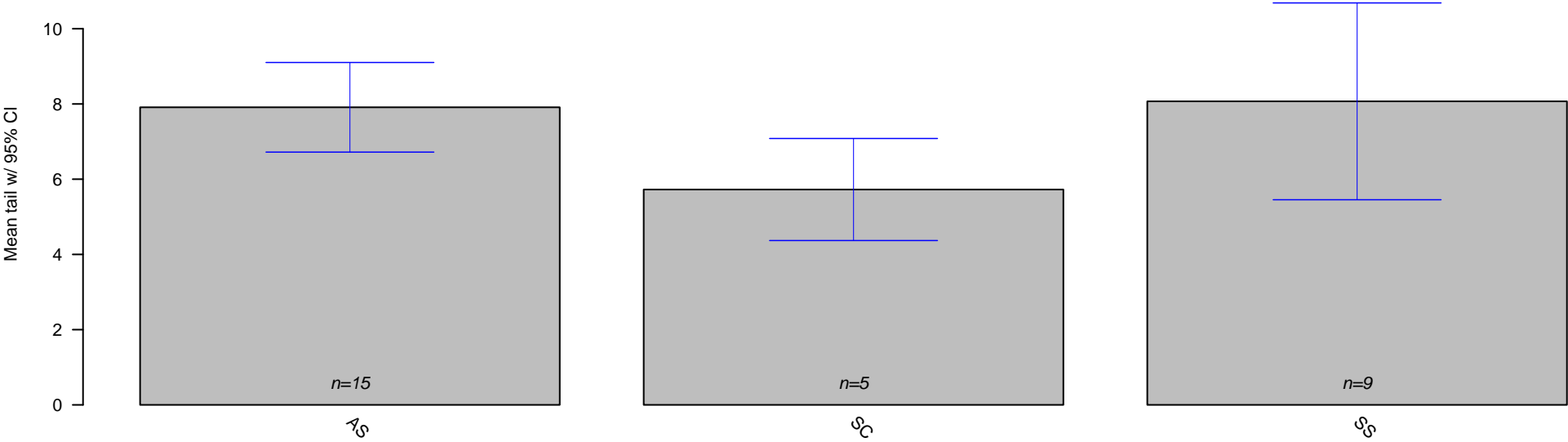


Type

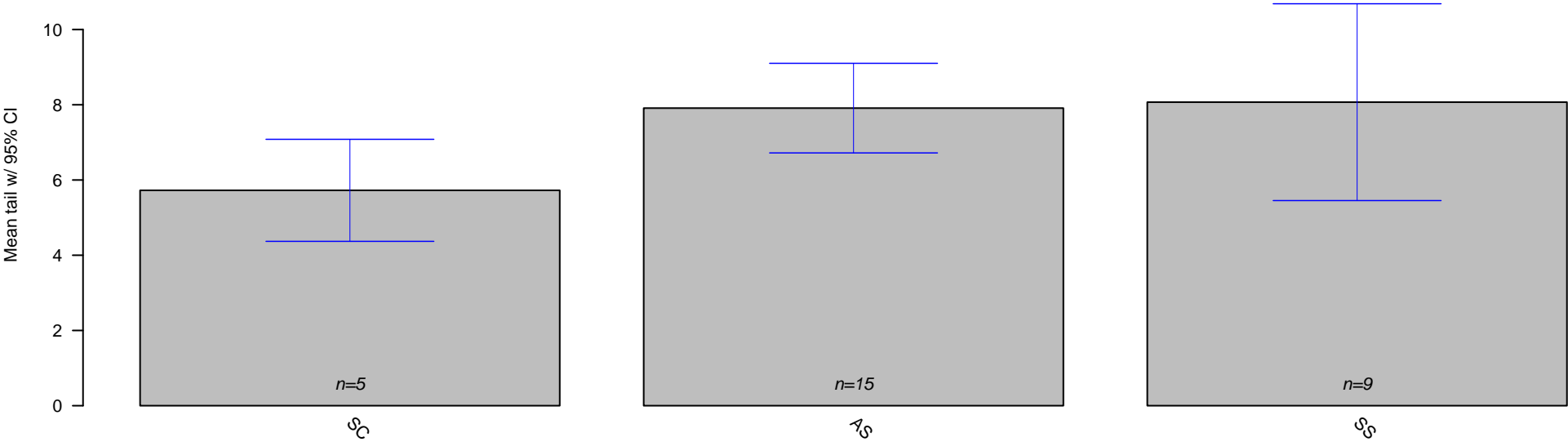


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

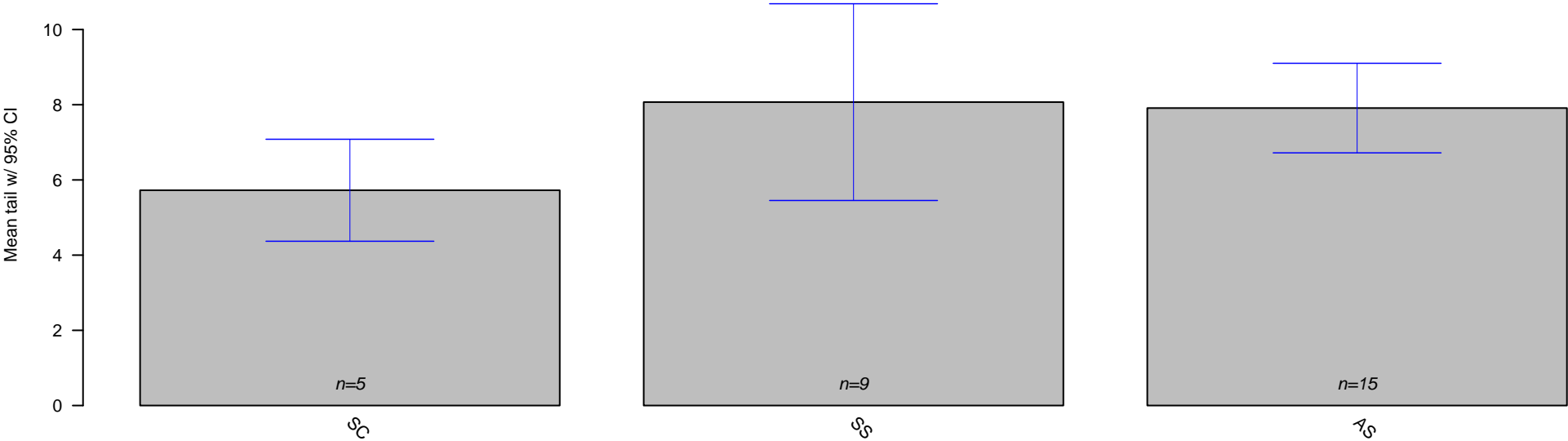
Type: Ordered by Category



Type: Ordered by Diversity

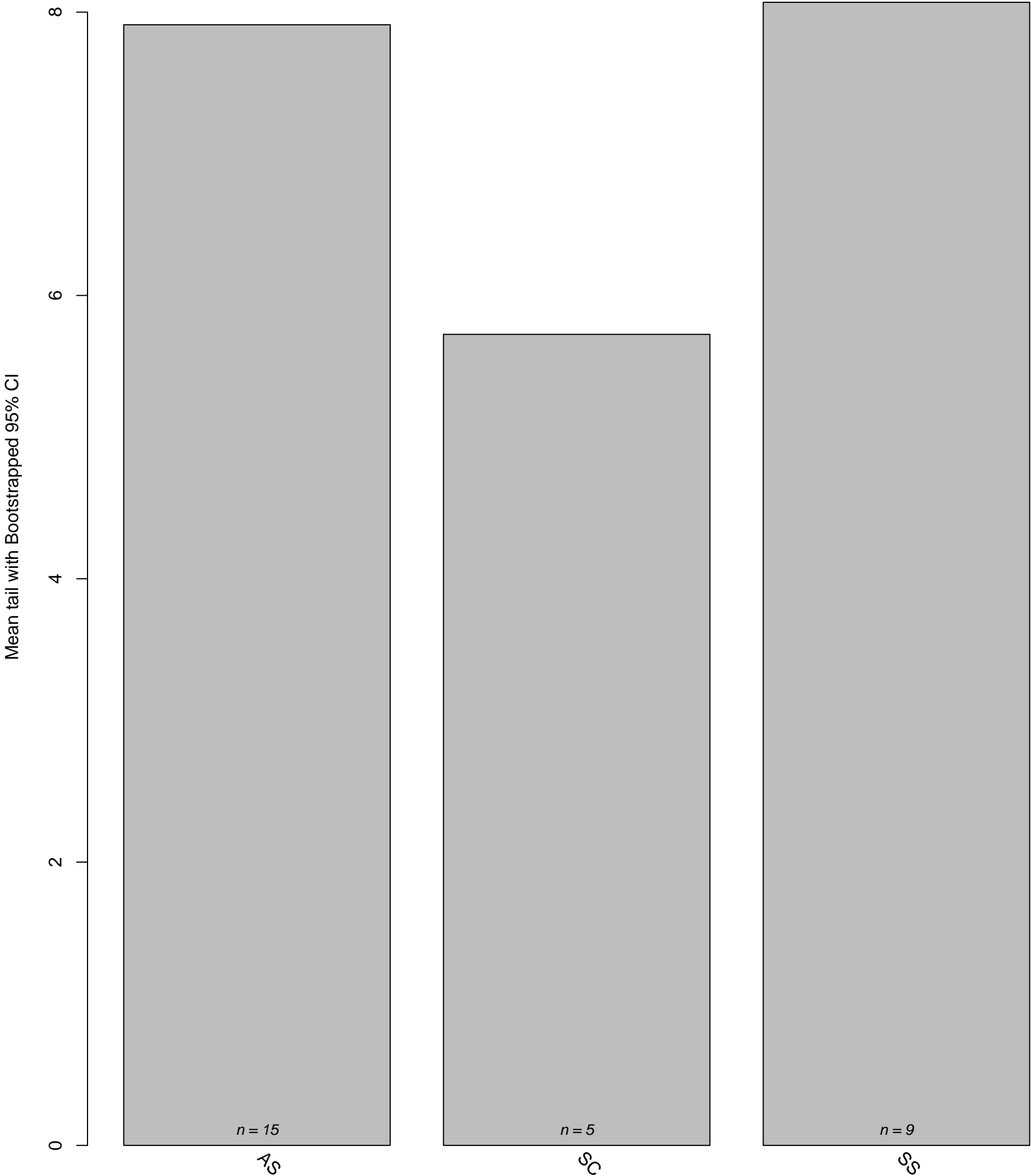


Type: Ordered by Sample Size

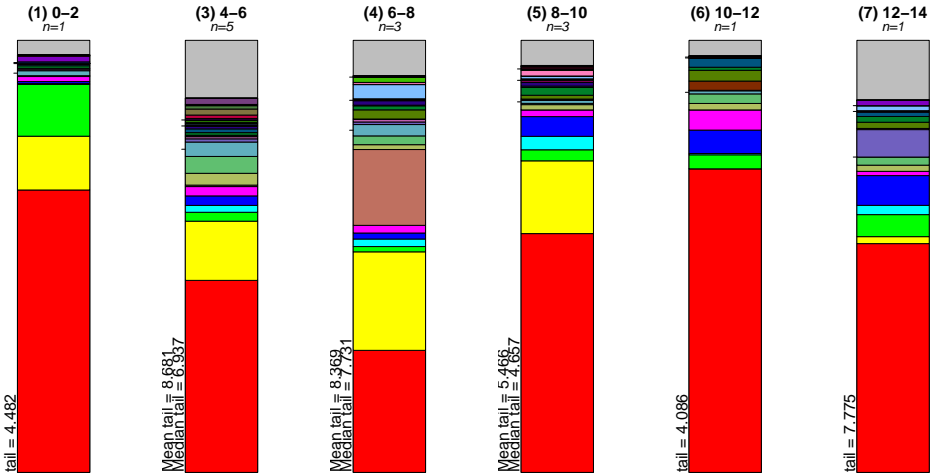


Type: Ordered by Category

with Wilcoxon rank sum test (difference between group means) p -values

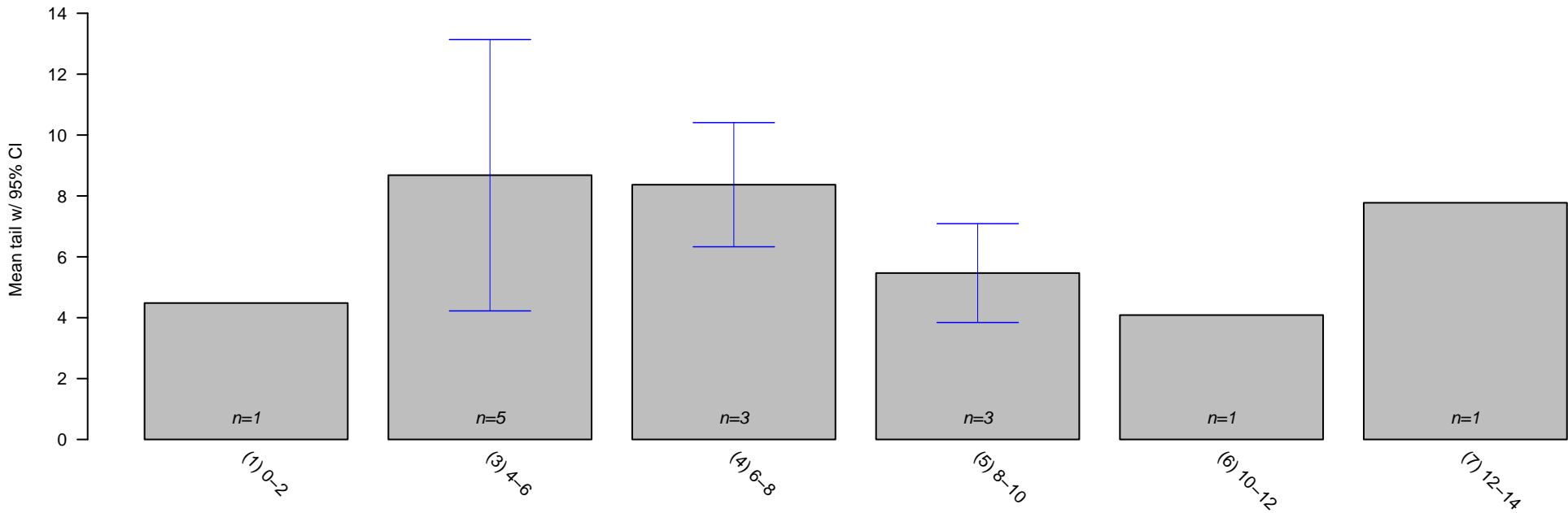


Neut

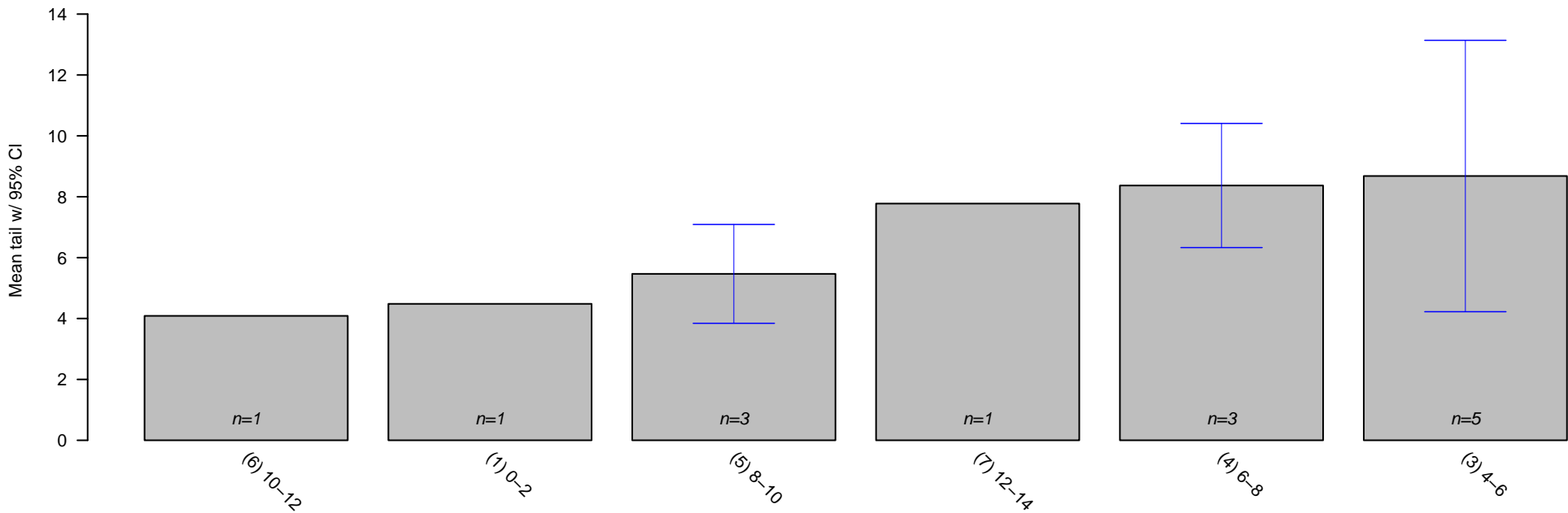


- Remaining
- Tyzzereella_4
- Thalassospira
- Eisenbergiella
- Prevotella_2
- Alloprevotella
- Ruminococcaceae_UCG-014
- Hungatella
- Ruminococcus_2
- Bilophila
- Fusobacterium
- Incertae_Sedis
- Akkermansia
- Sutterella
- Lactobacillus
- Lachnospira
- Ruminiclostridium_5
- Enterobacteriaceae_uncl
- Streptococcus
- Acidaminococcus
- Anaerostipes
- Ruminococcaceae_uncl
- Fusicatenibacter
- Pseudobutyrvibrio
- Parasutterella
- Escherichia-Shigella
- [Eubacterium]_coprostanoligenes_grp
- Ruminococcaceae_UCG-002
- Parabacteroides
- Lachnospiraceae_uncl
- Prevotella_9
- Lachnoclostridium
- Subdoligranulum
- Blautia
- Alistipes
- Faecalibacterium
- Bacteroides

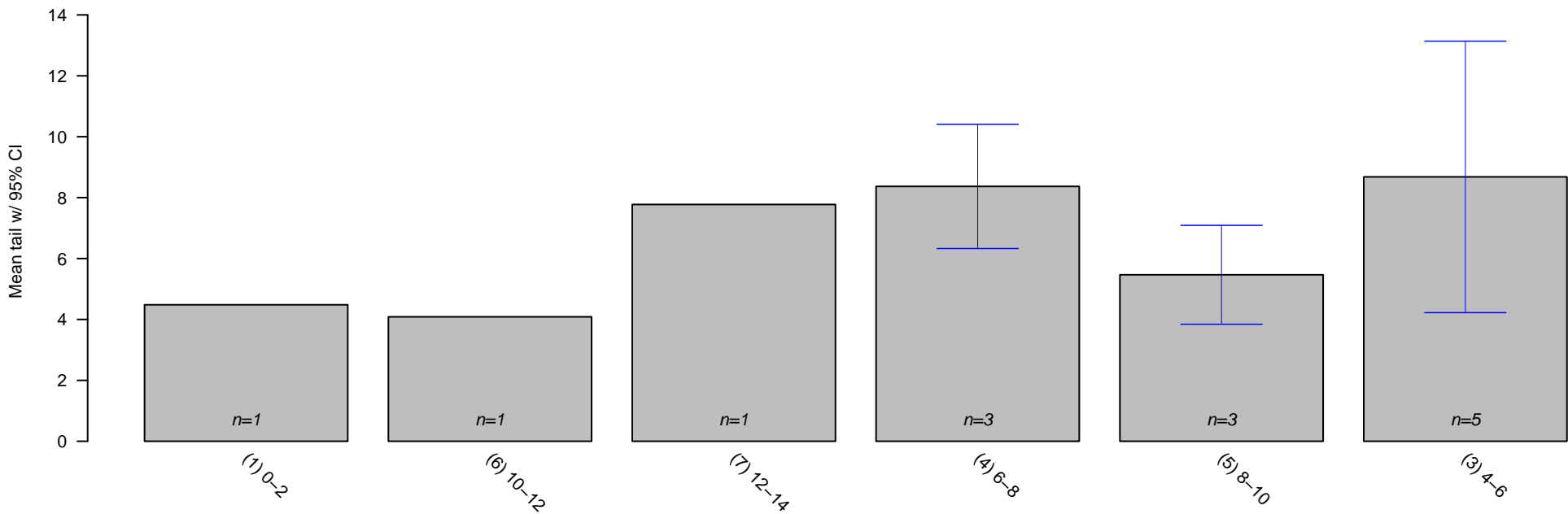
Neut: Ordered by Category



Neut: Ordered by Diversity

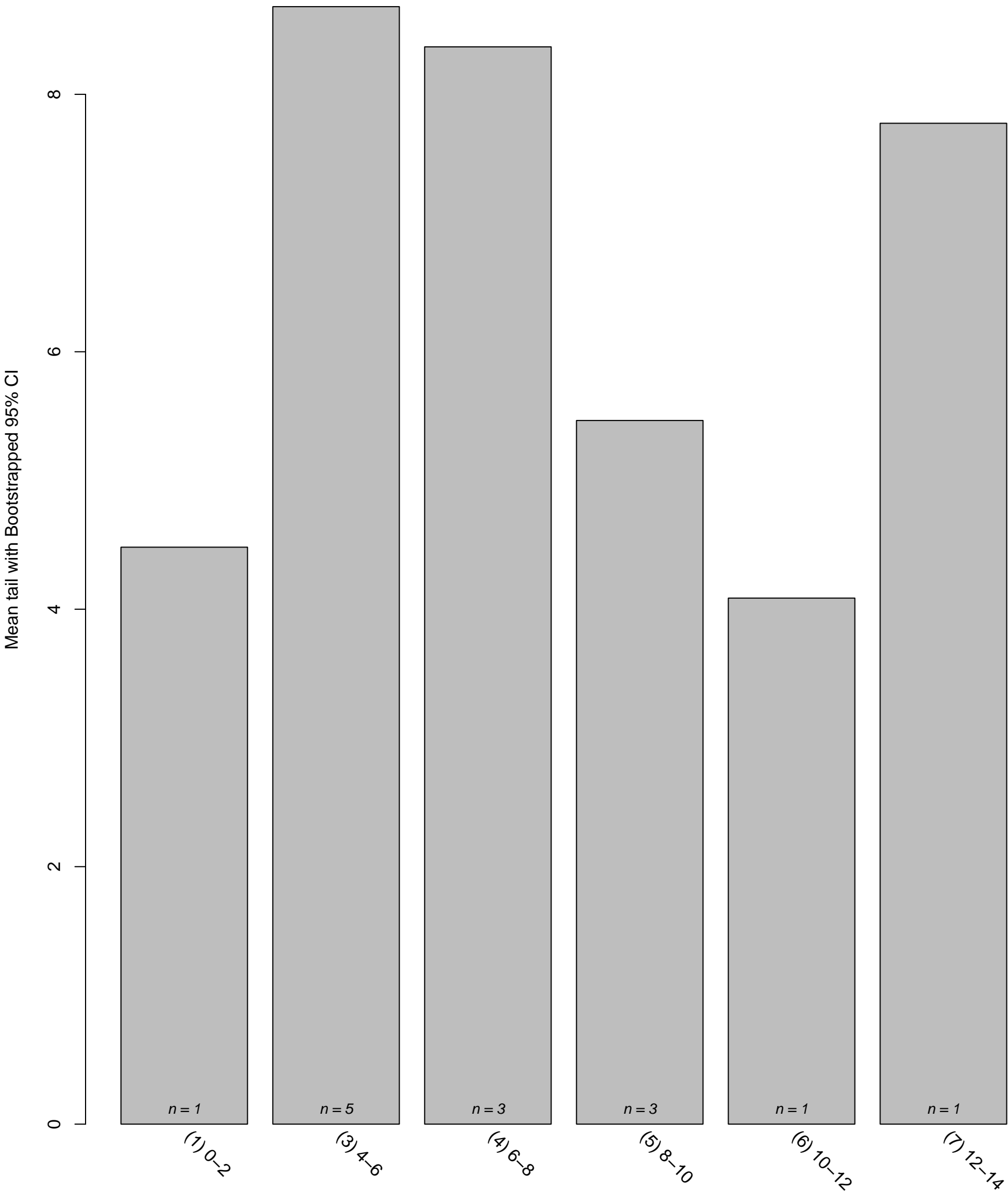


Neut: Ordered by Sample Size

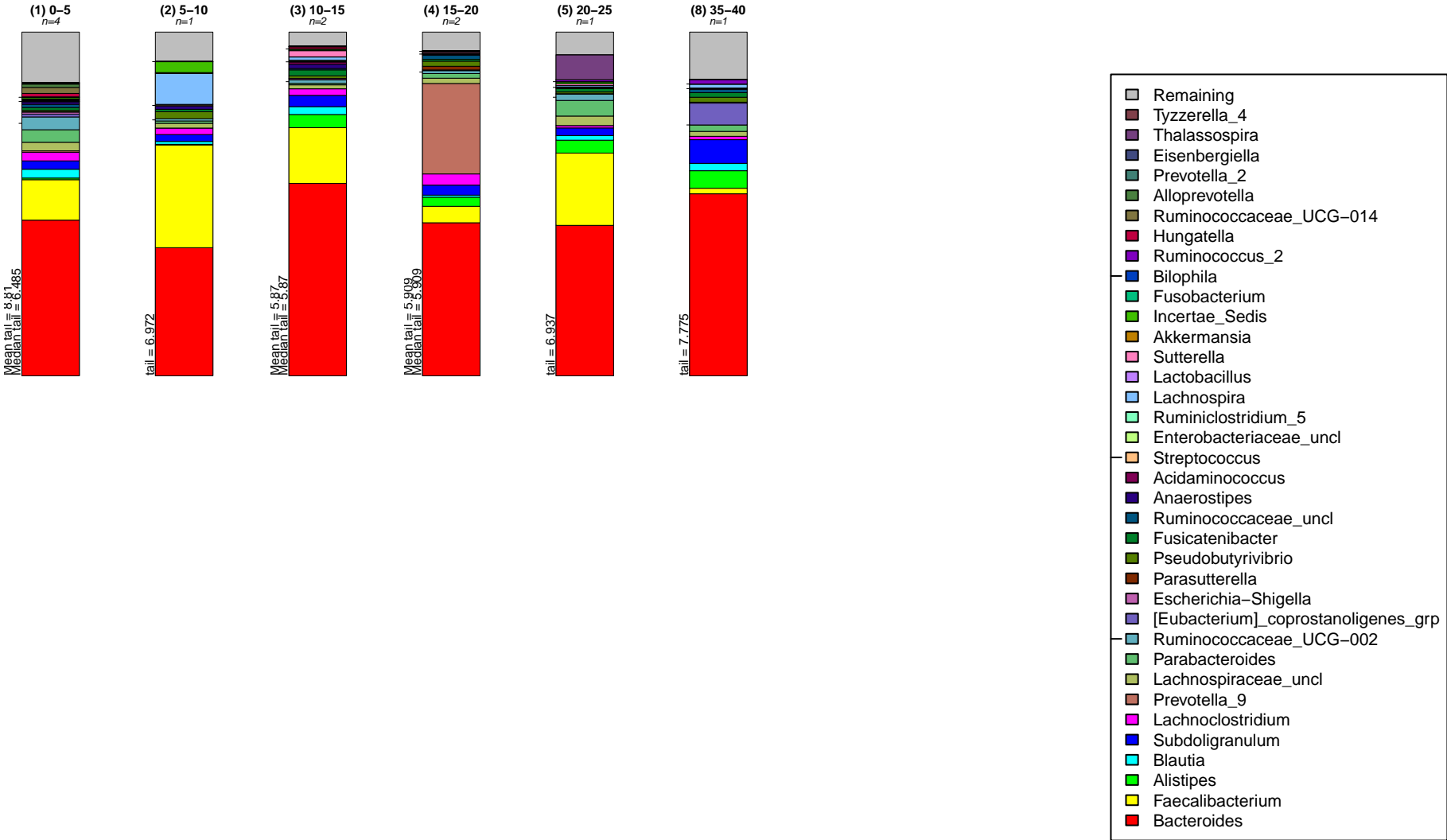


Neut: Ordered by Category

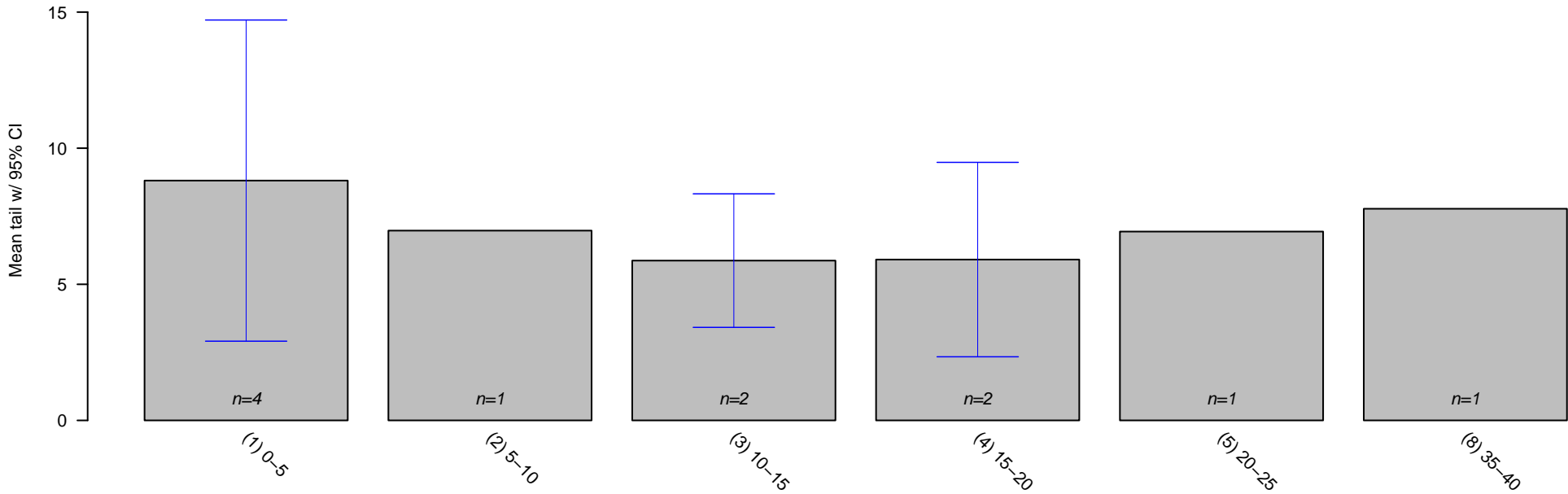
with Wilcoxon rank sum test (difference between group means) p-values



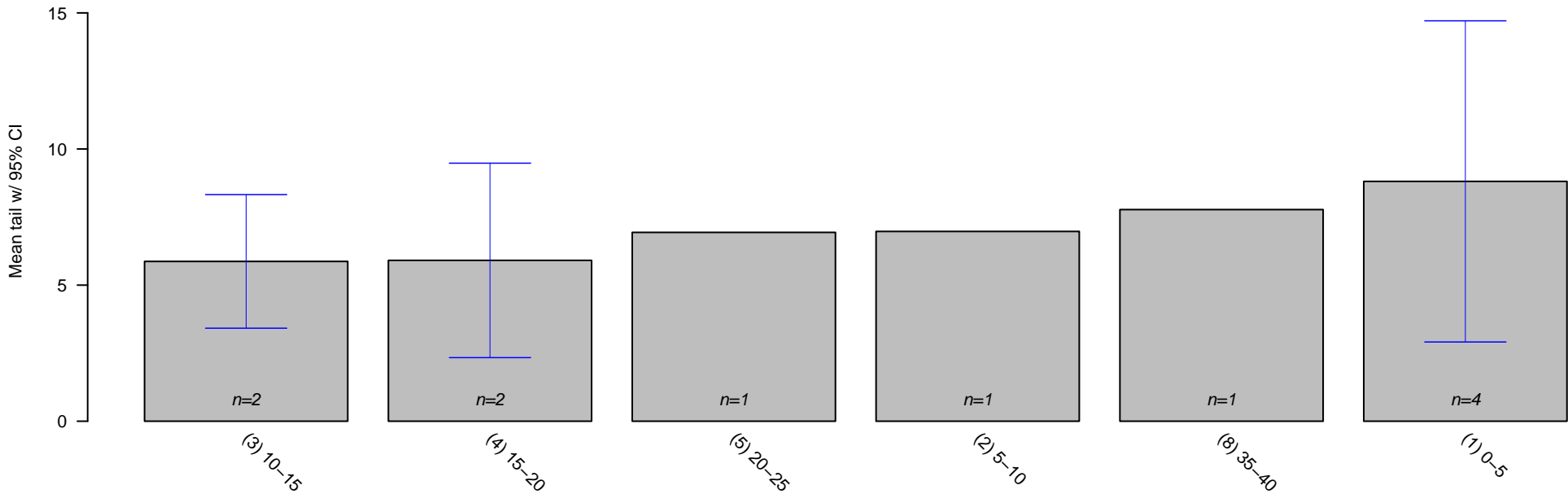
SCChi_CD16



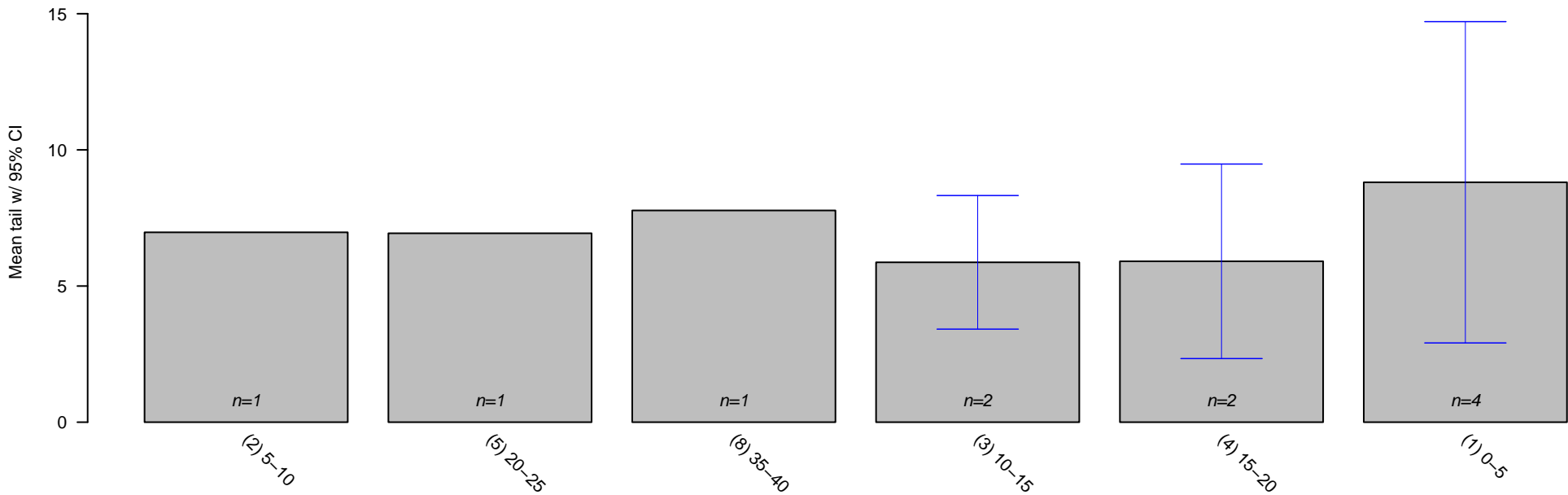
SCChi_CD16: Ordered by Category



SCChi_CD16: Ordered by Diversity

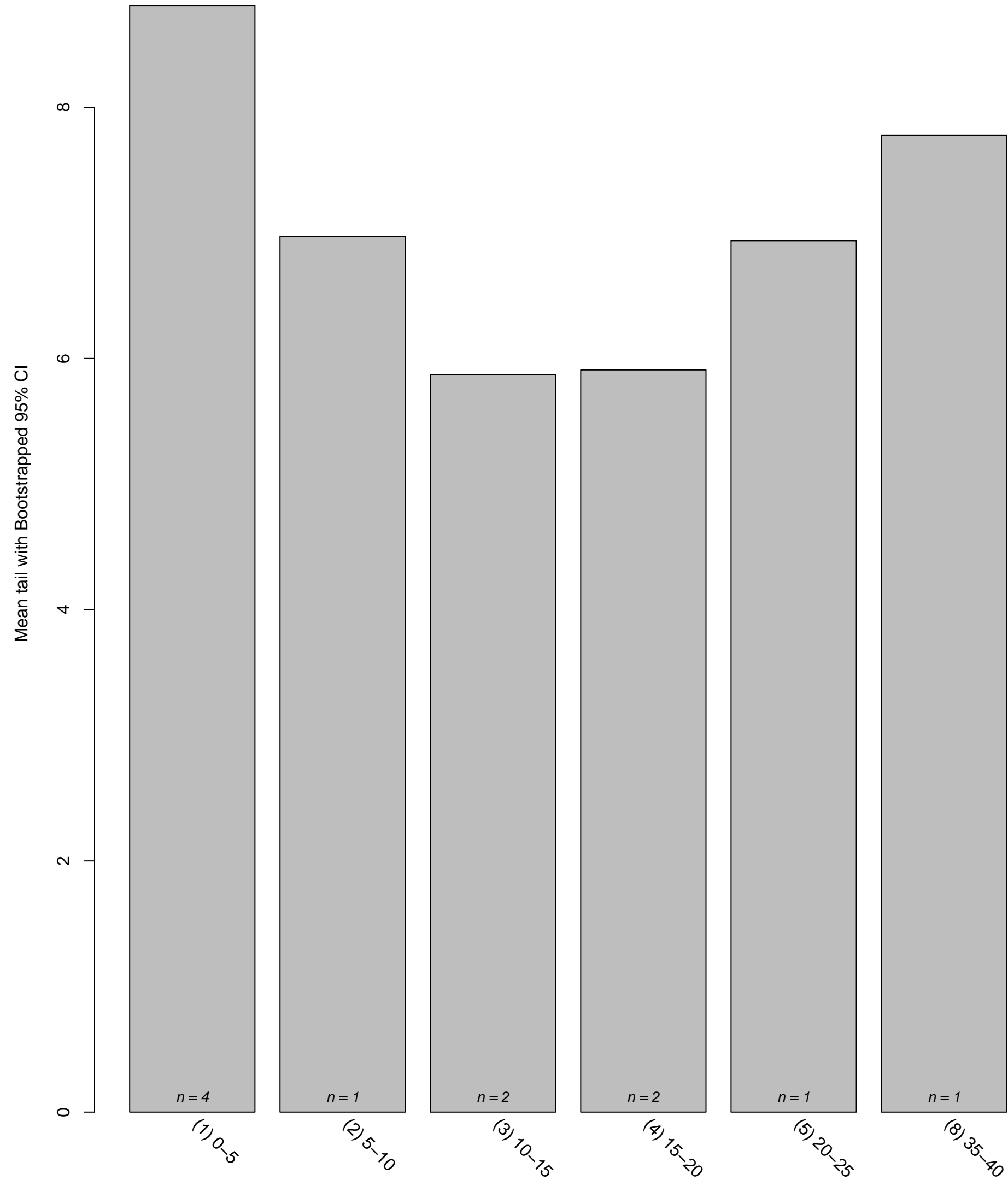


SCChi_CD16: Ordered by Sample Size

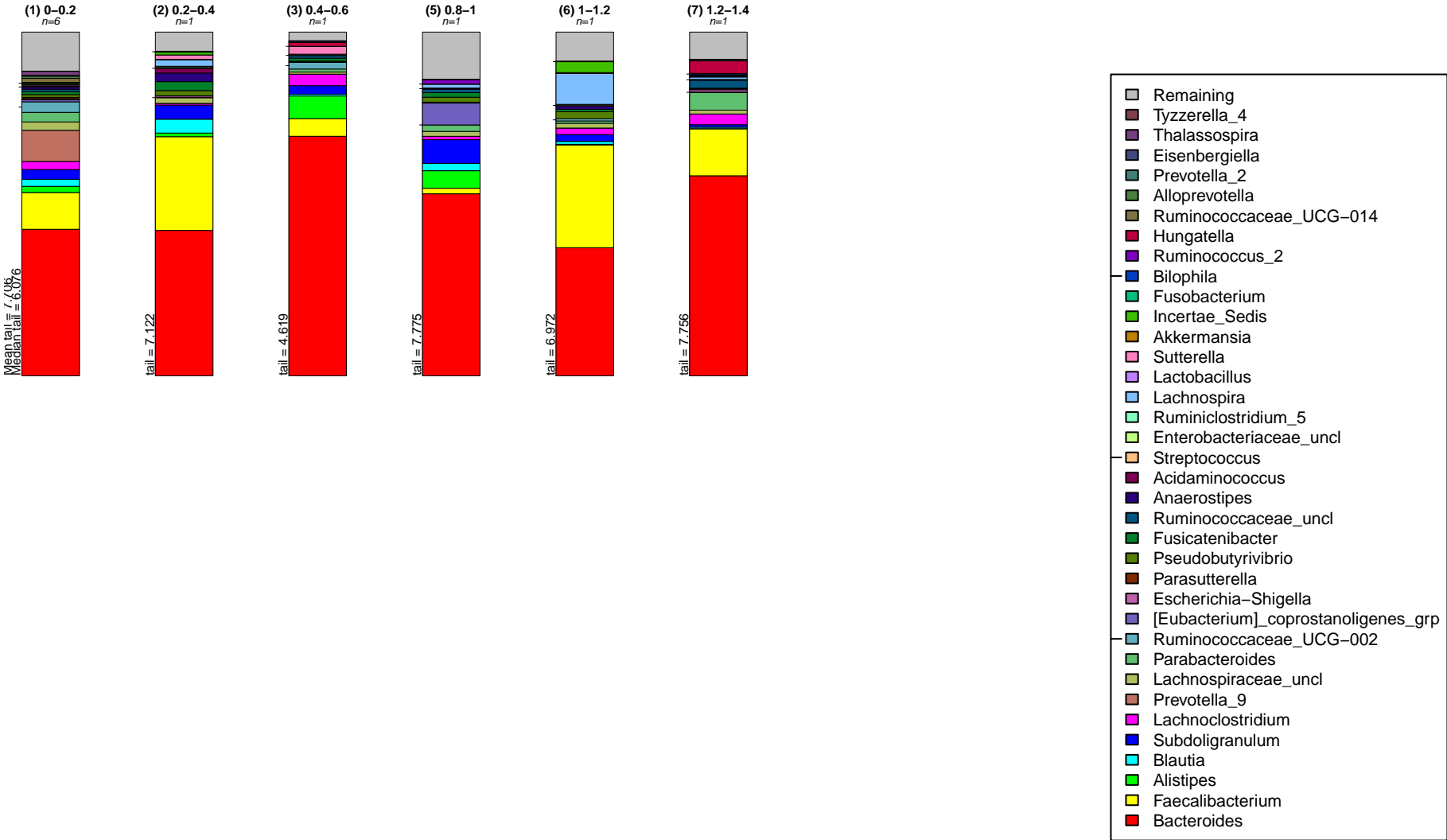


SCChi_CD16: Ordered by Category

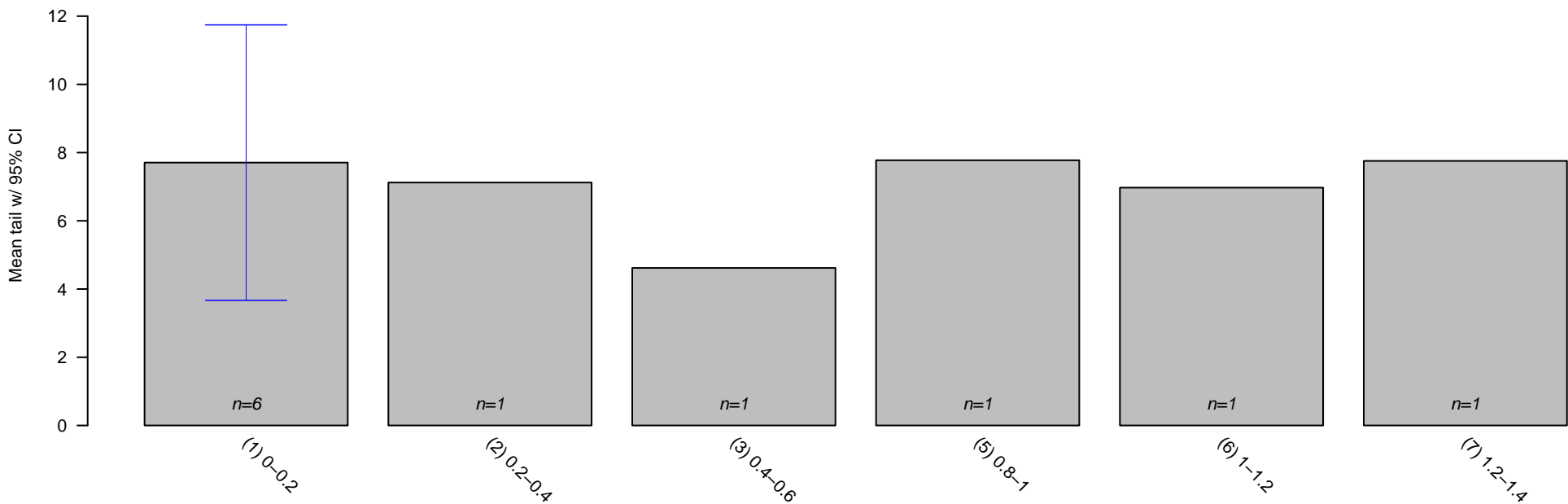
with Wilcoxon rank sum test (difference between group means) p-values



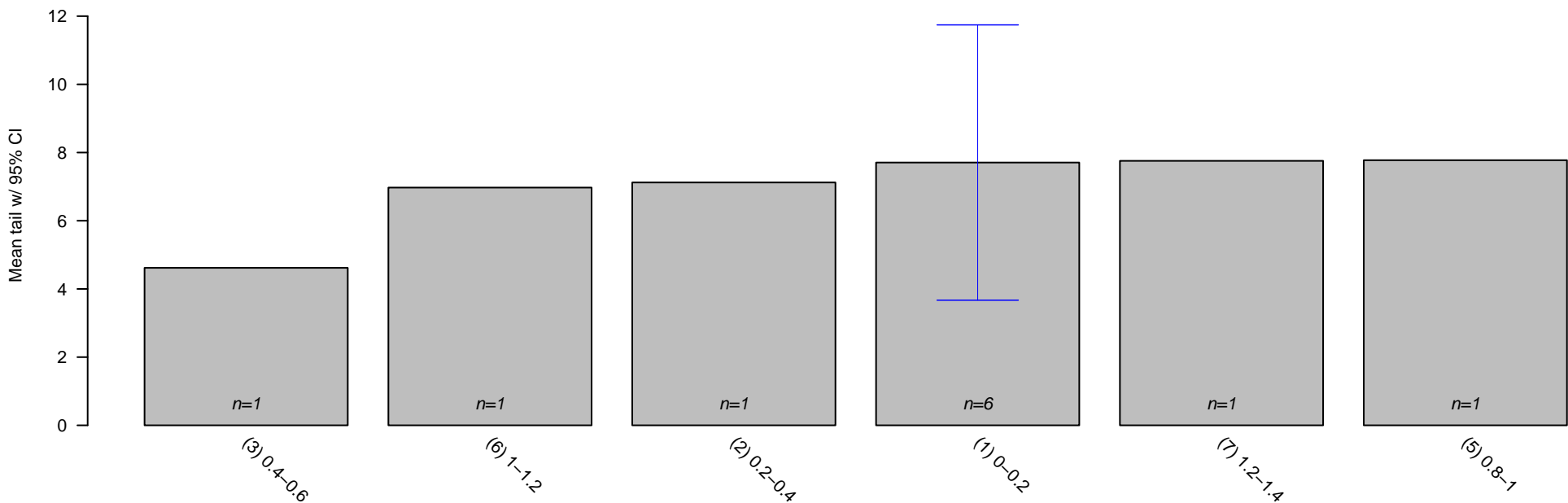
CD62lo_CD11b



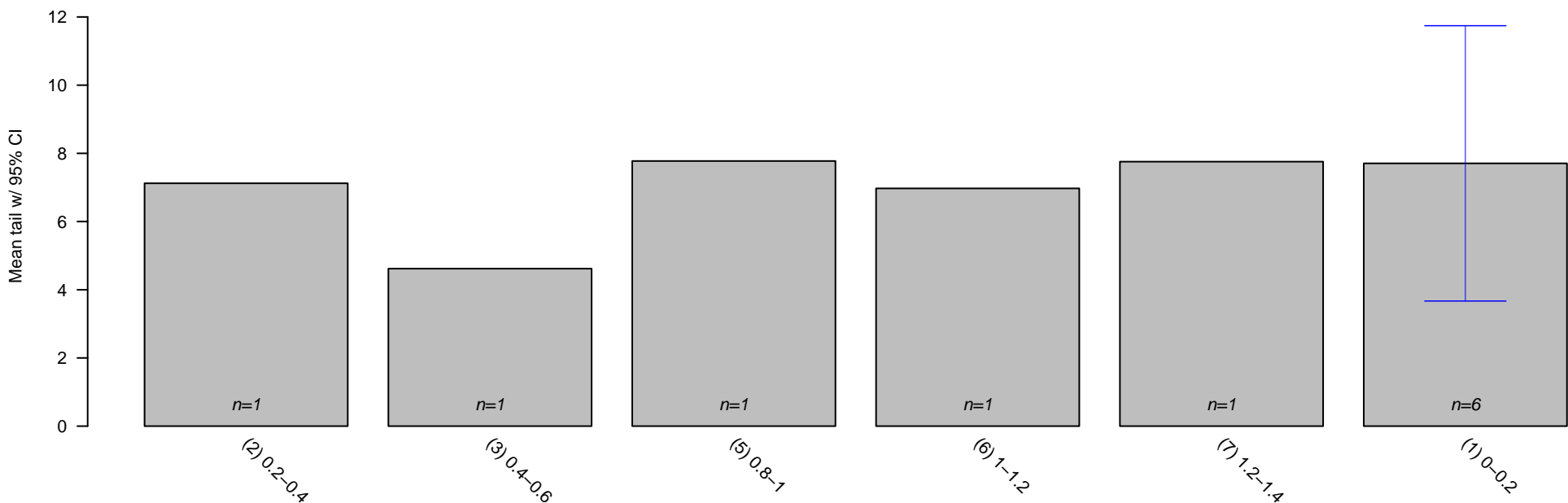
CD62lo_CD11b: Ordered by Category



CD62lo_CD11b: Ordered by Diversity

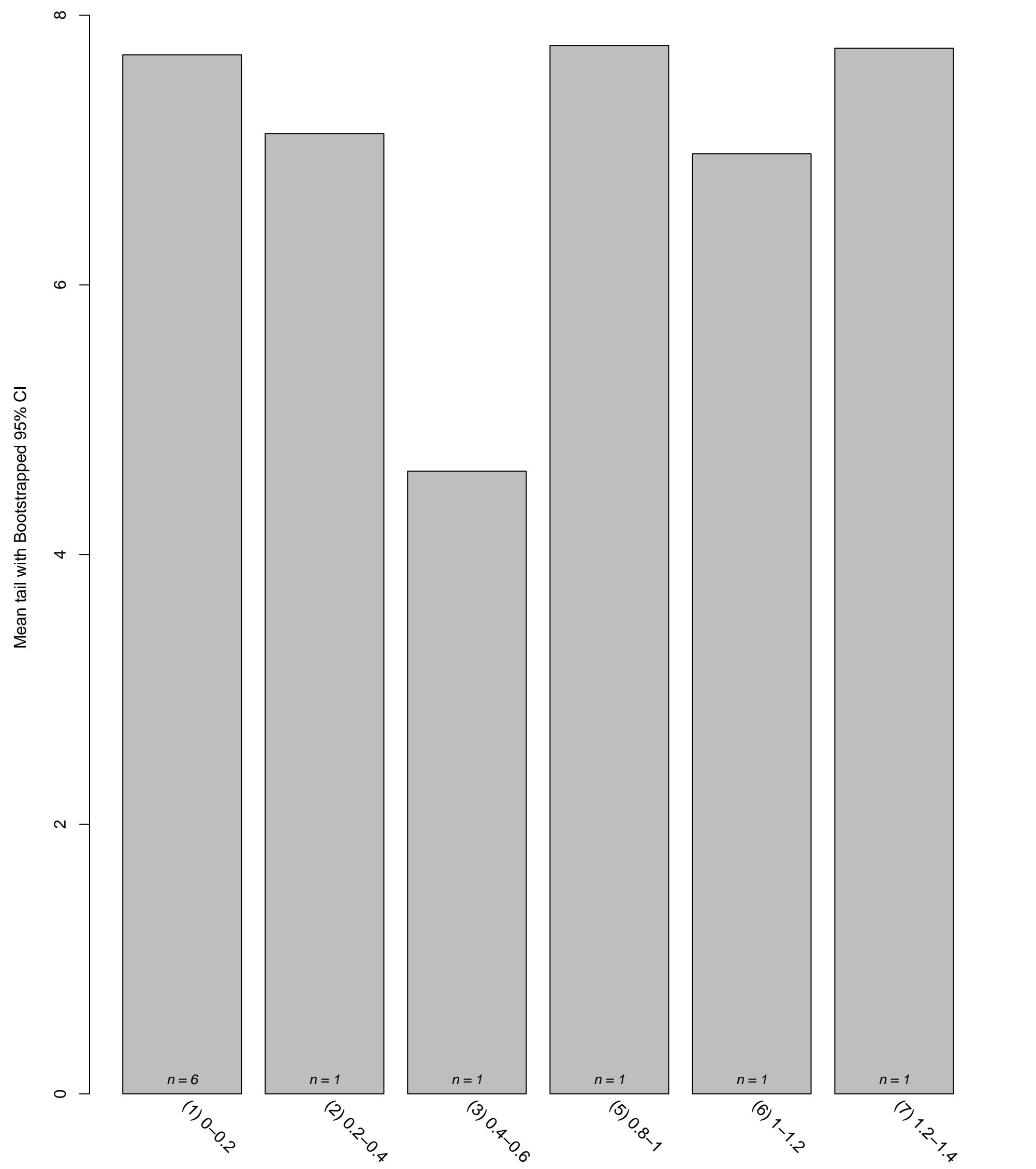


CD62lo_CD11b: Ordered by Sample Size



CD62lo_CD11b: Ordered by Category

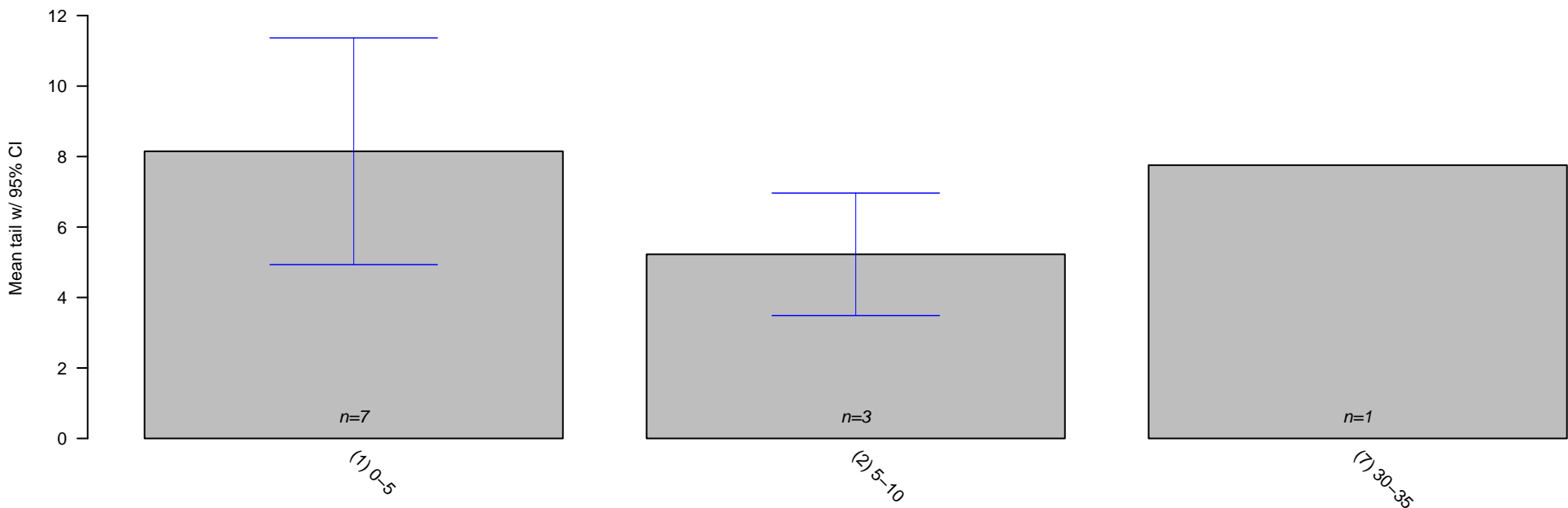
with Wilcoxon rank sum test (difference between group means) p-values



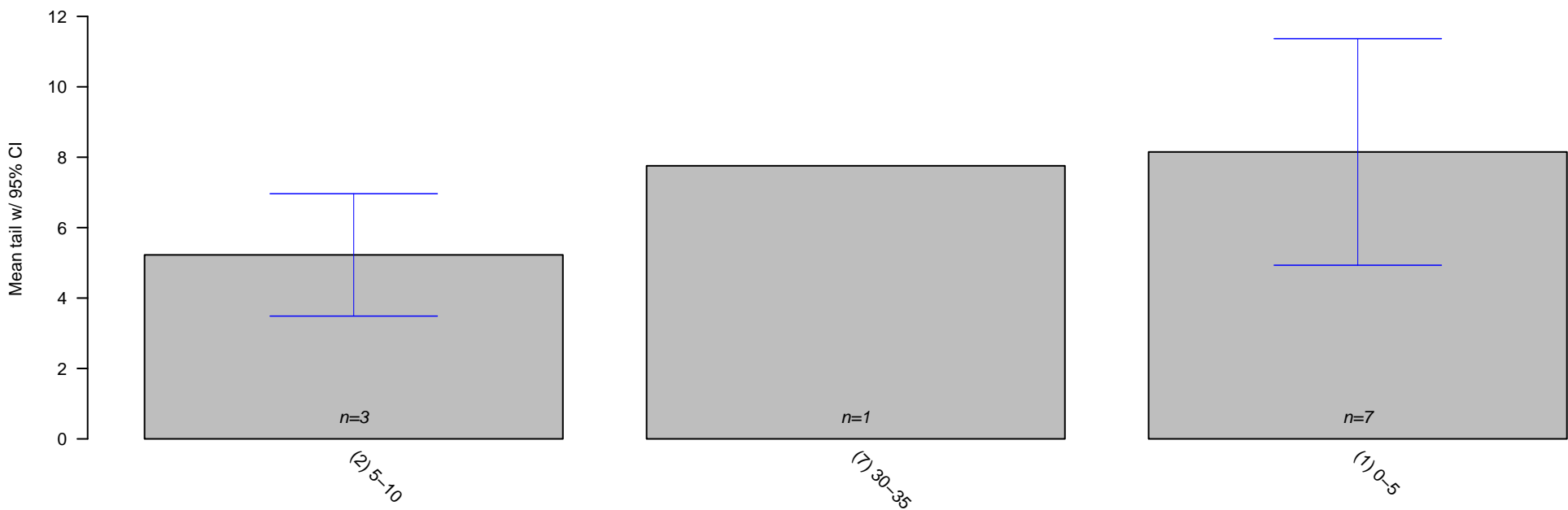
CD62lo_CD184



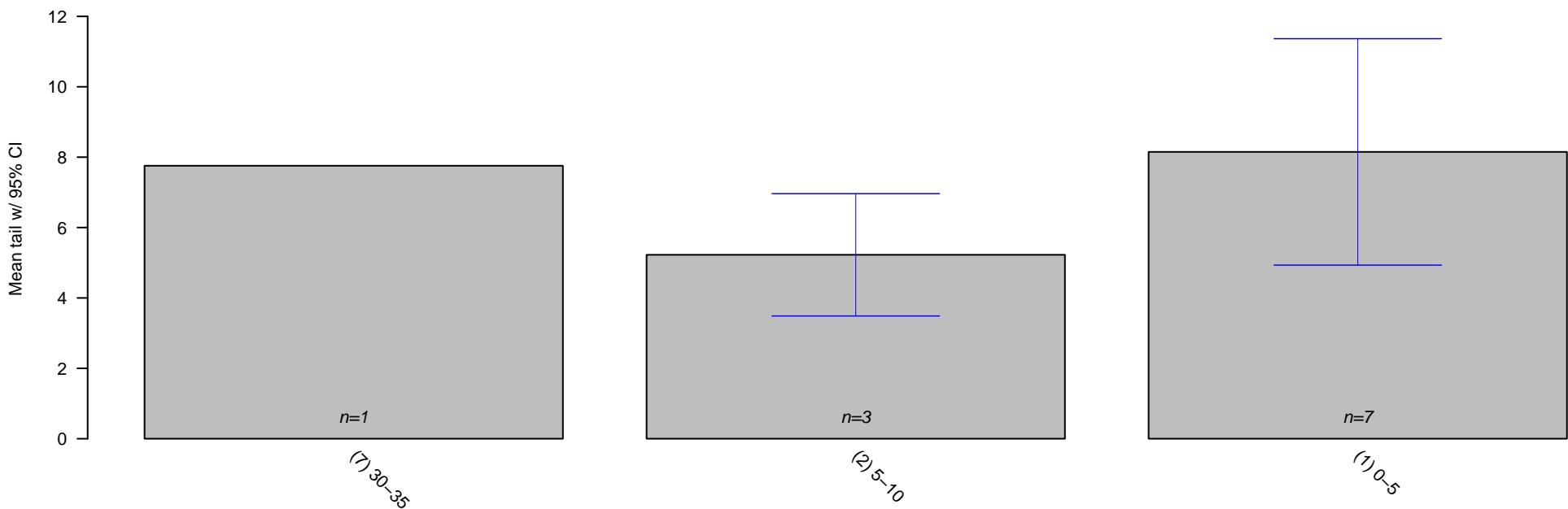
CD62lo_CD184: Ordered by Category



CD62lo_CD184: Ordered by Diversity



CD62lo_CD184: Ordered by Sample Size



CD62lo_CD184: Ordered by Category

with Wilcoxon rank sum test (difference between group means) p-values

