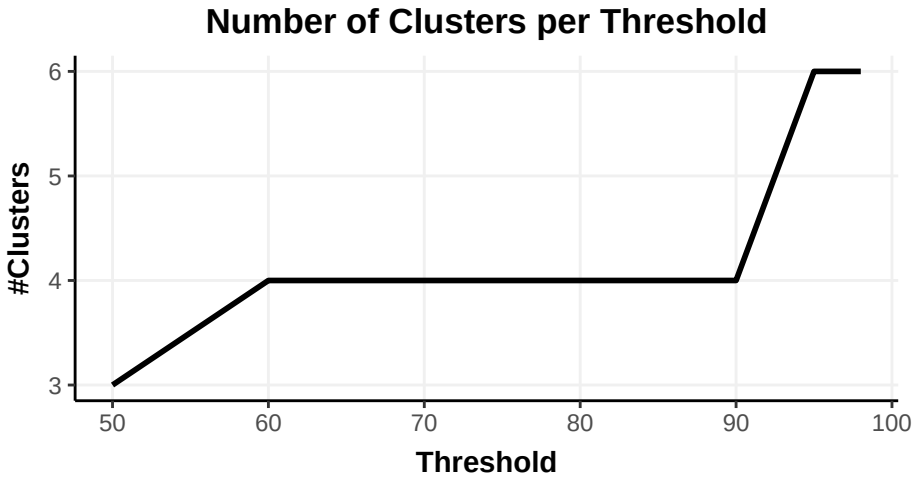
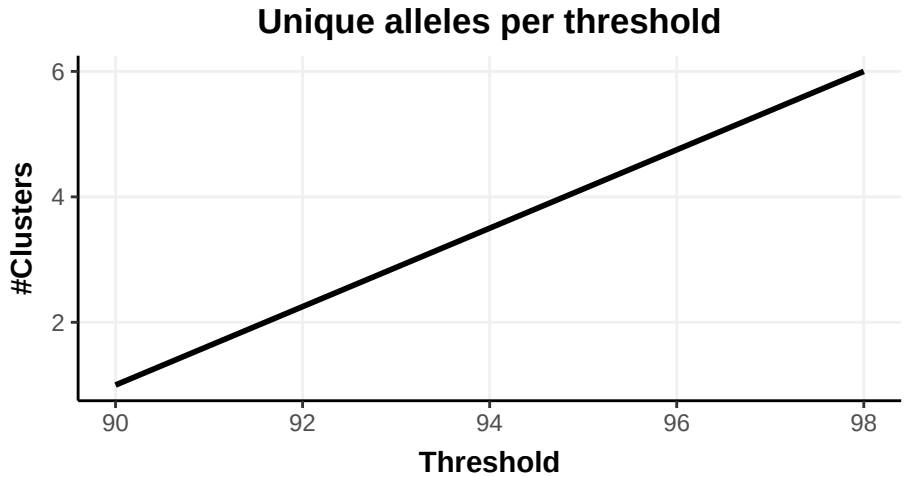
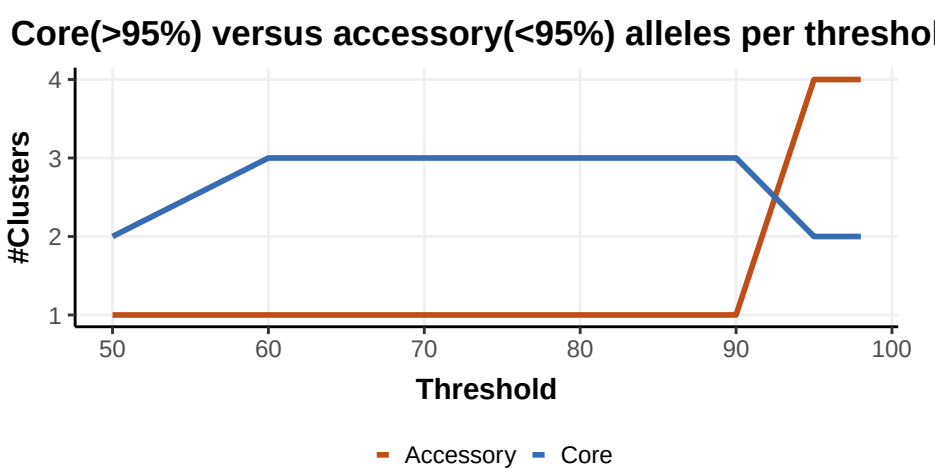
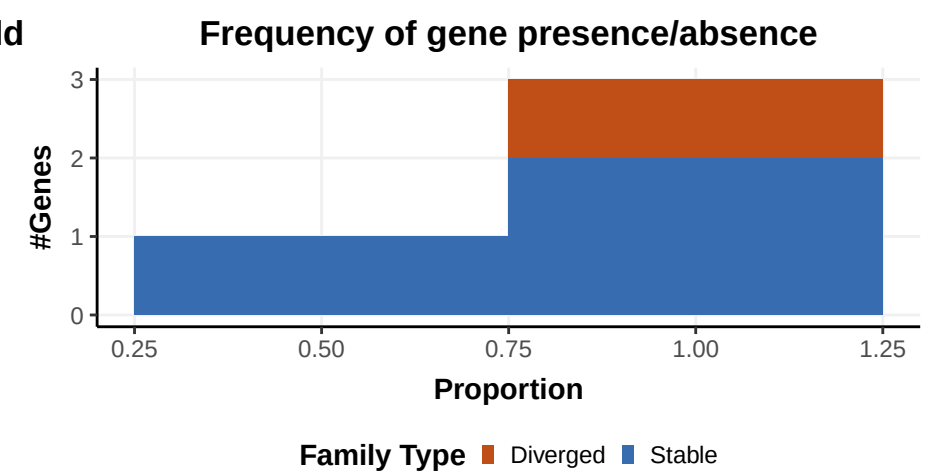
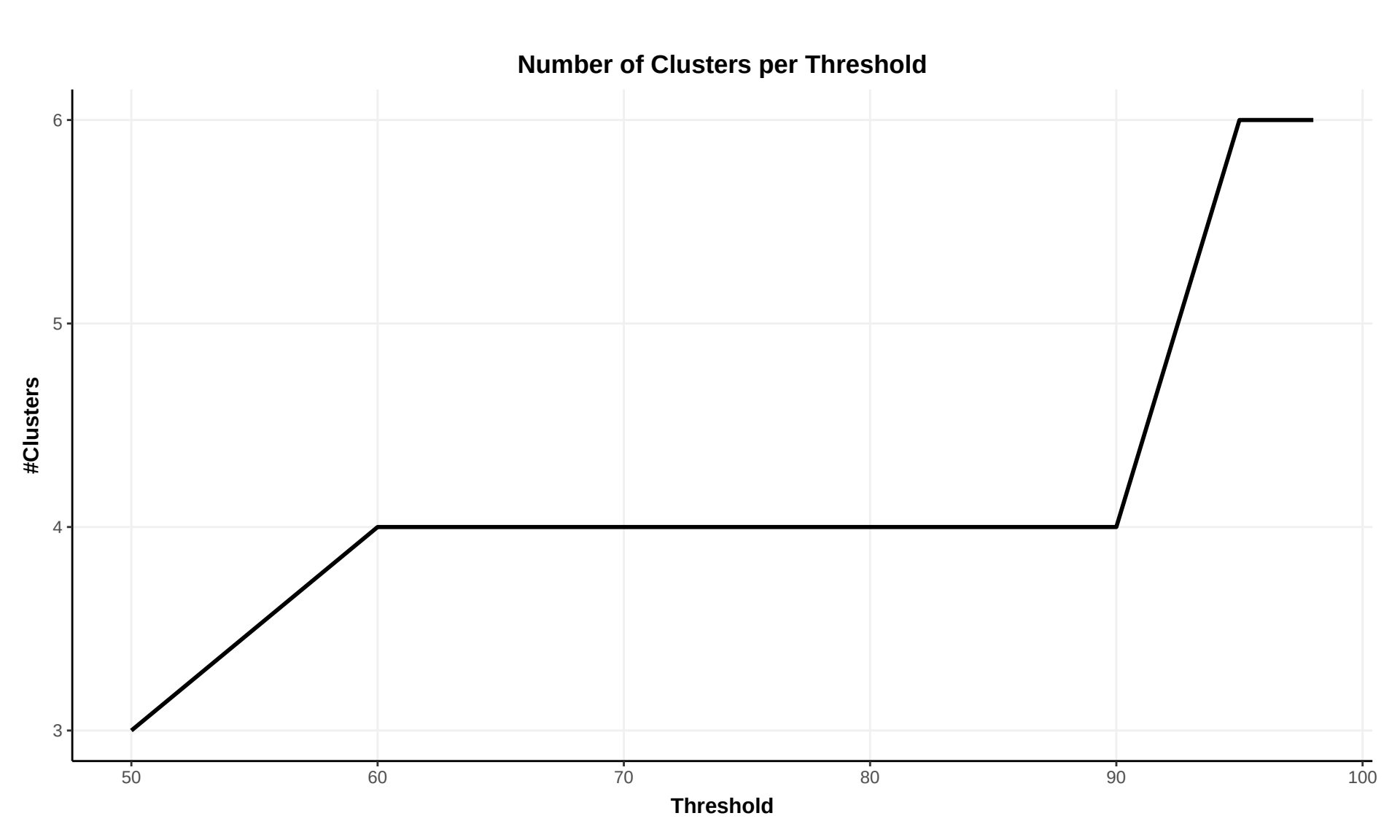
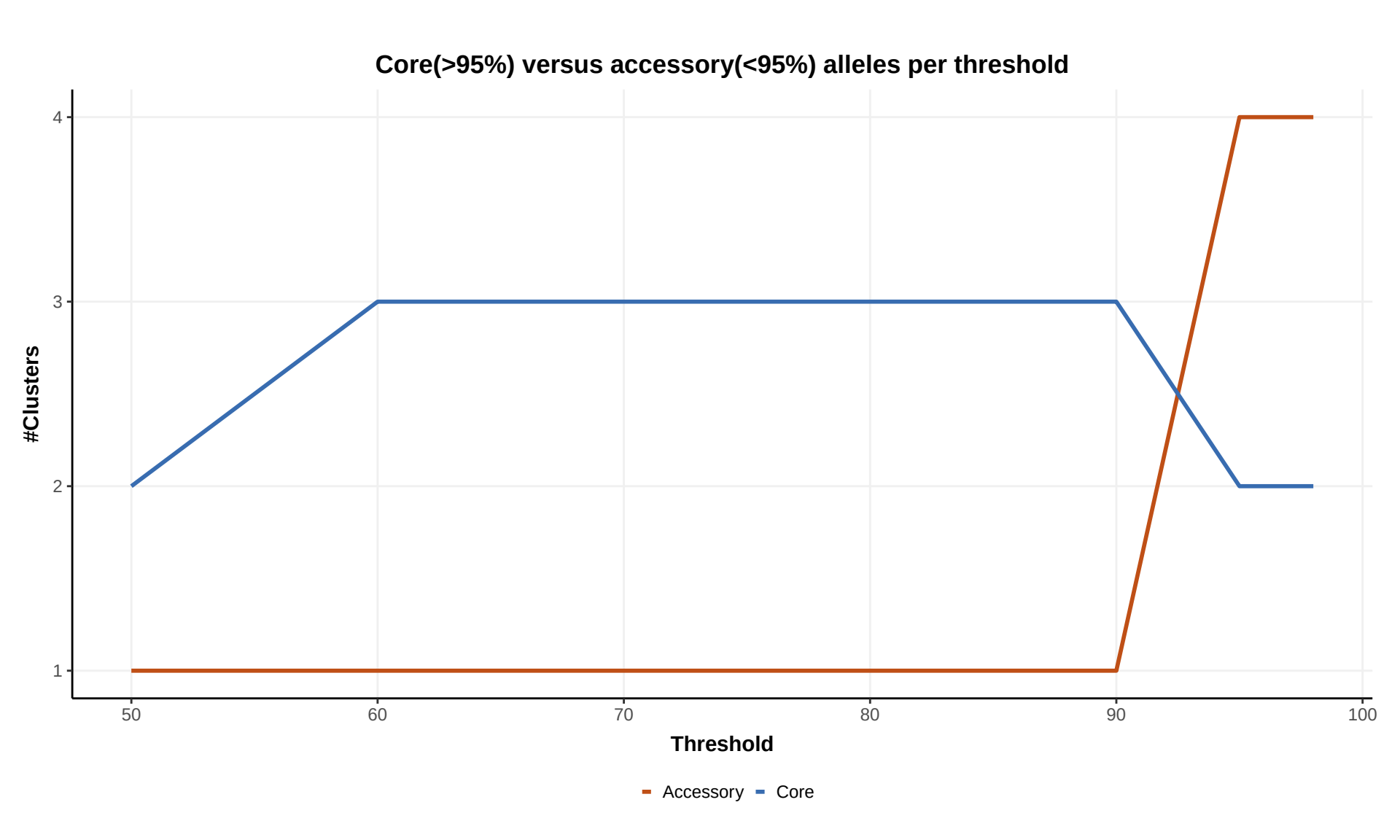

Number of Clusters per Threshold
#Clusters
6
5
4
3
50
60
70
80
90
100
Threshold

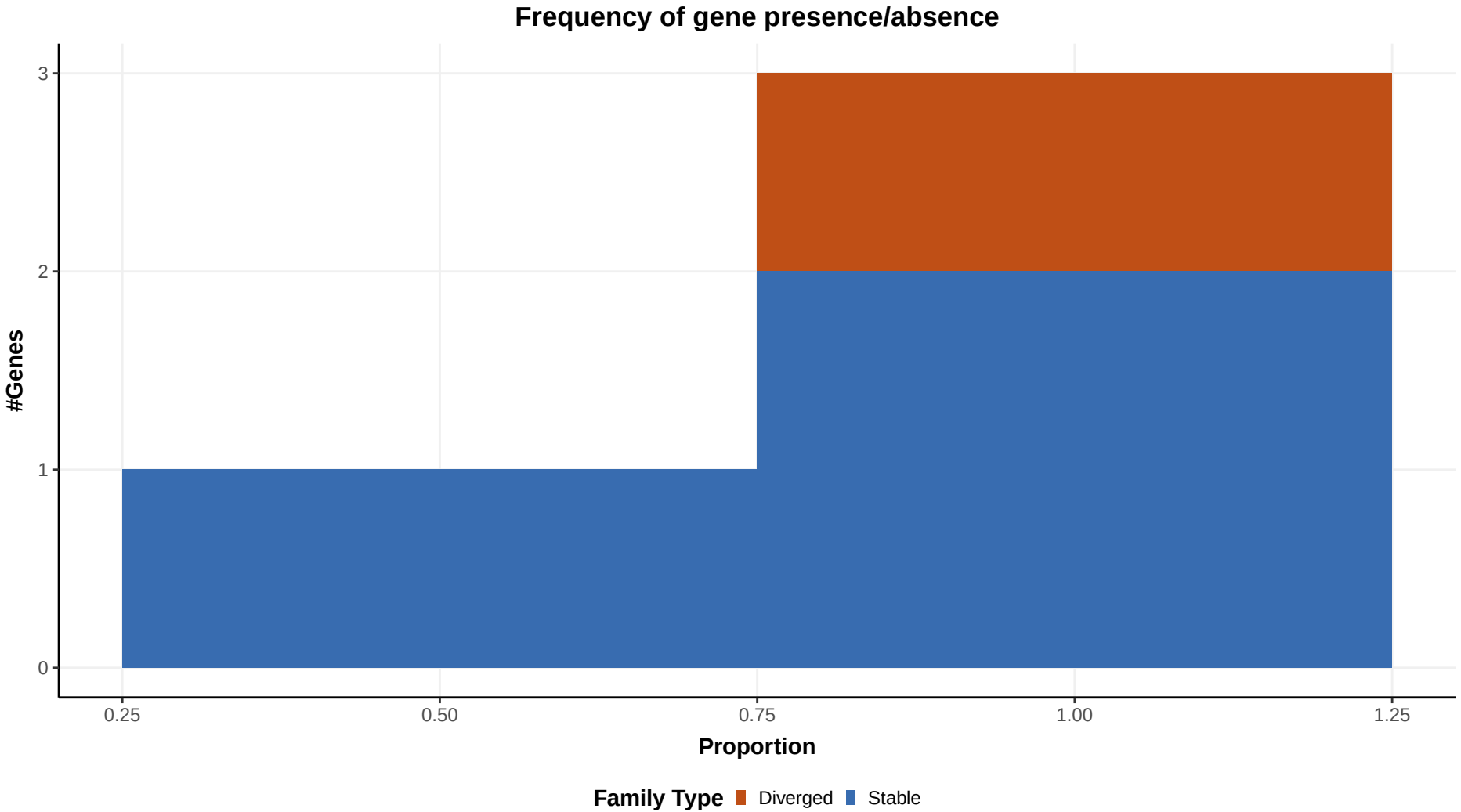

Unique alleles per threshold
#Clusters
6
4
2
90
92
94
96
98
Threshold

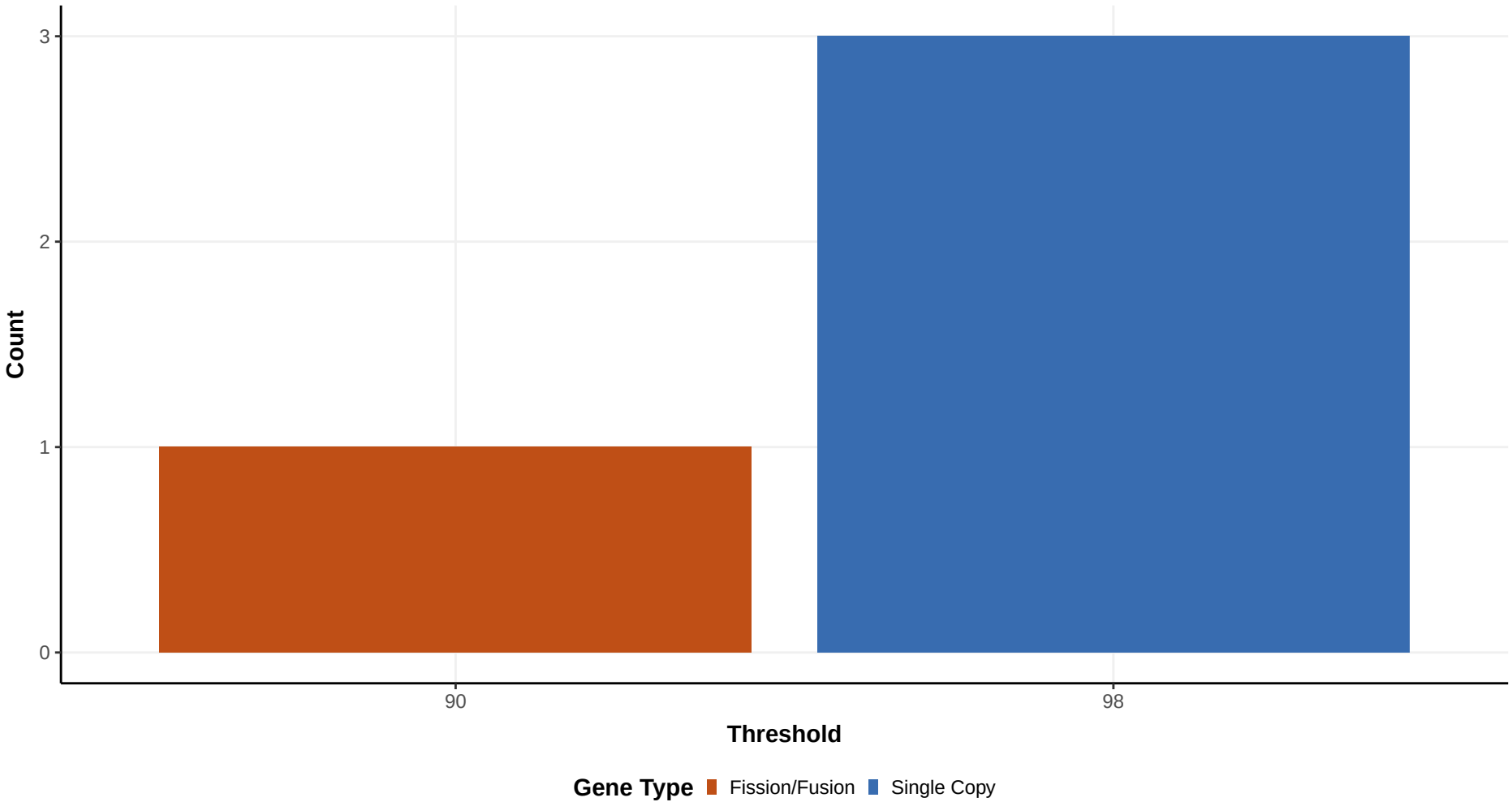

Core(>95%) versus accessory(<95%) alleles per threshold
#Clusters
4
3
2
1
50
60
70
80
90
100
Threshold
Accessory
Core

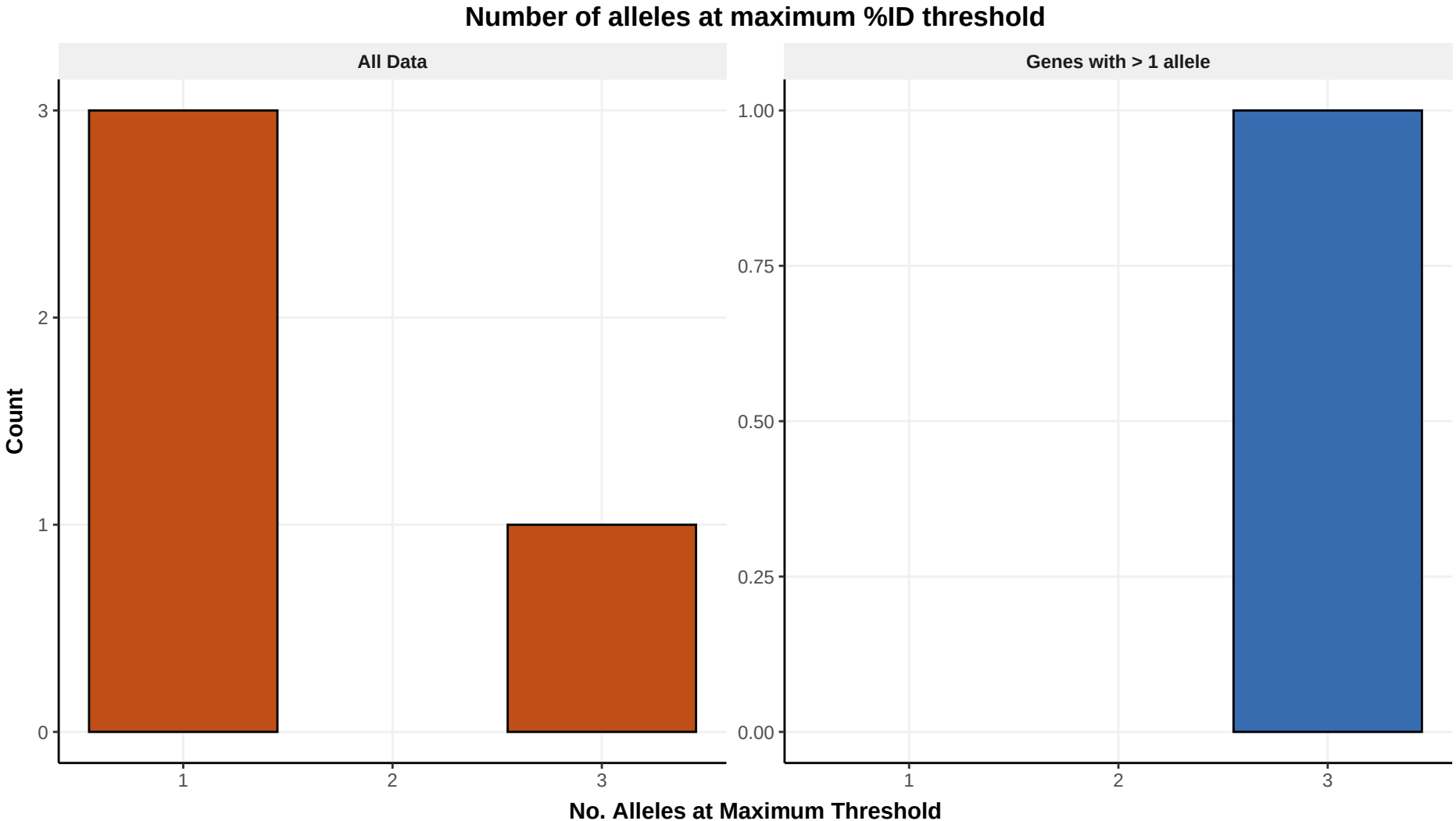

Frequency of gene presence/absence
#Genes
3
2
1
0
0.25
0.50
0.75
1.00
1.25
Proportion
Family Type
Diverged
Stable

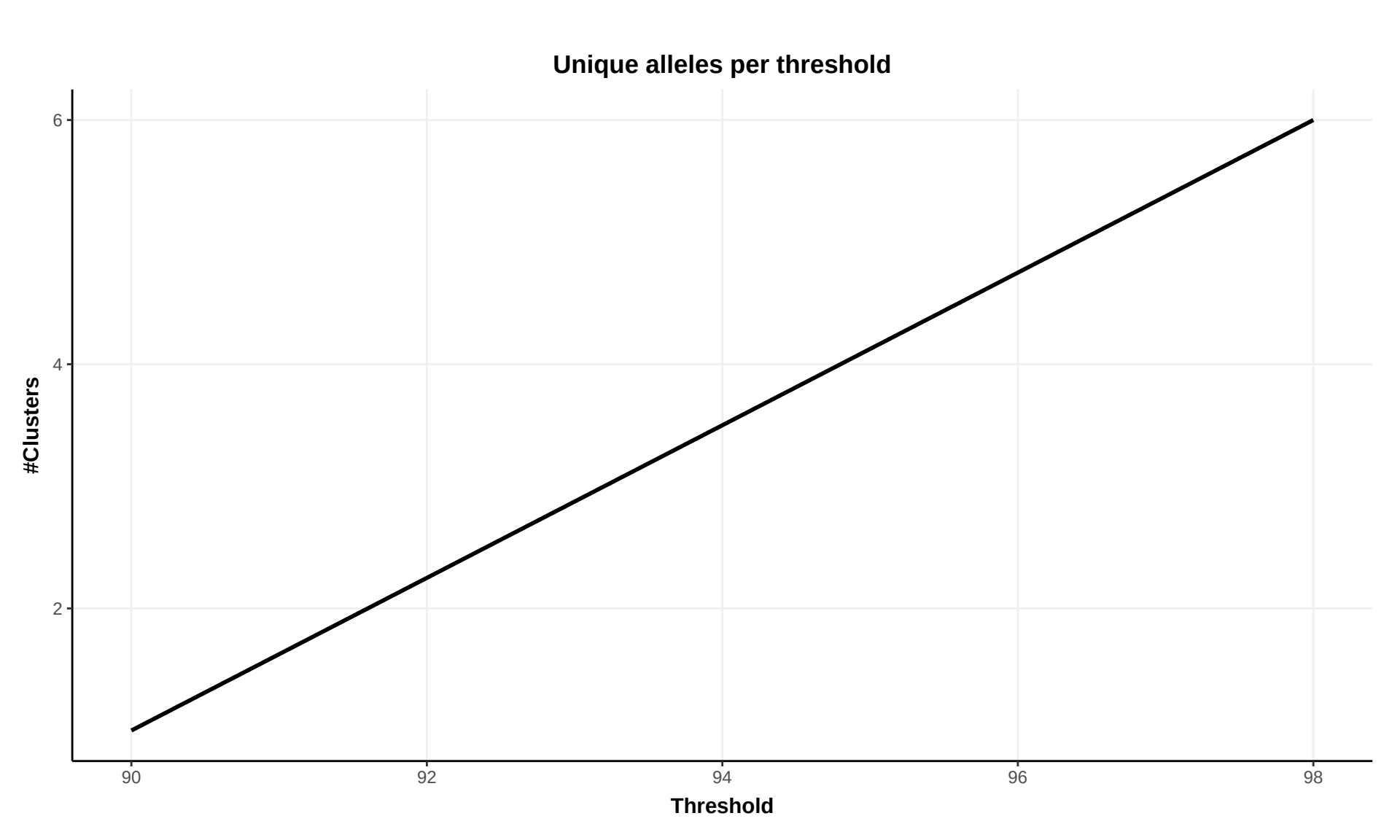

Number of Clusters per Threshold
#Clusters
6
5
4
3
50
60
70
80
90
100
Threshold

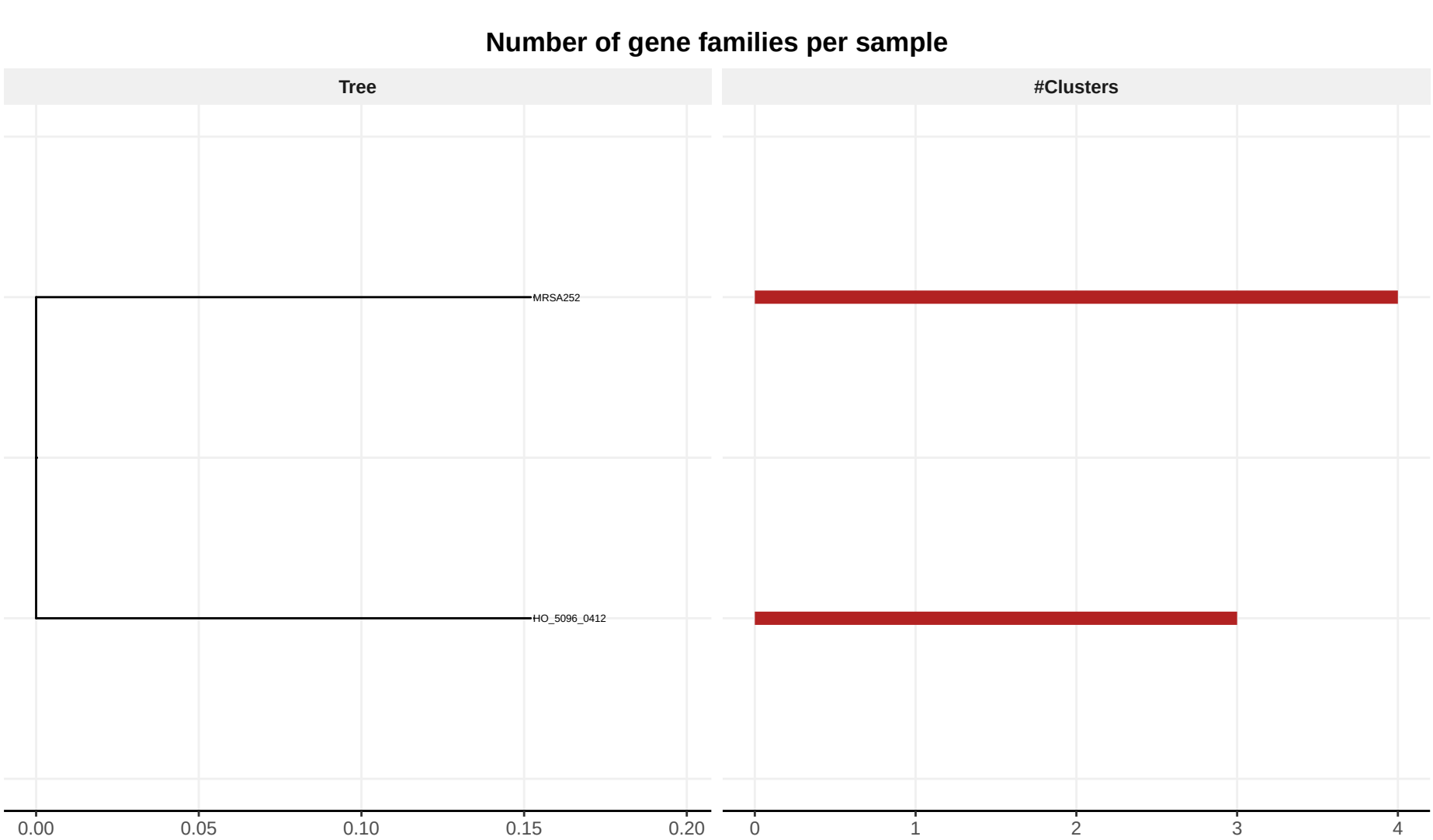

Core(>95%) versus accessory(<95%) alleles per threshold
#Clusters
4
3
2
1
50
60
70
80
90
100
Threshold
Accessory
Core

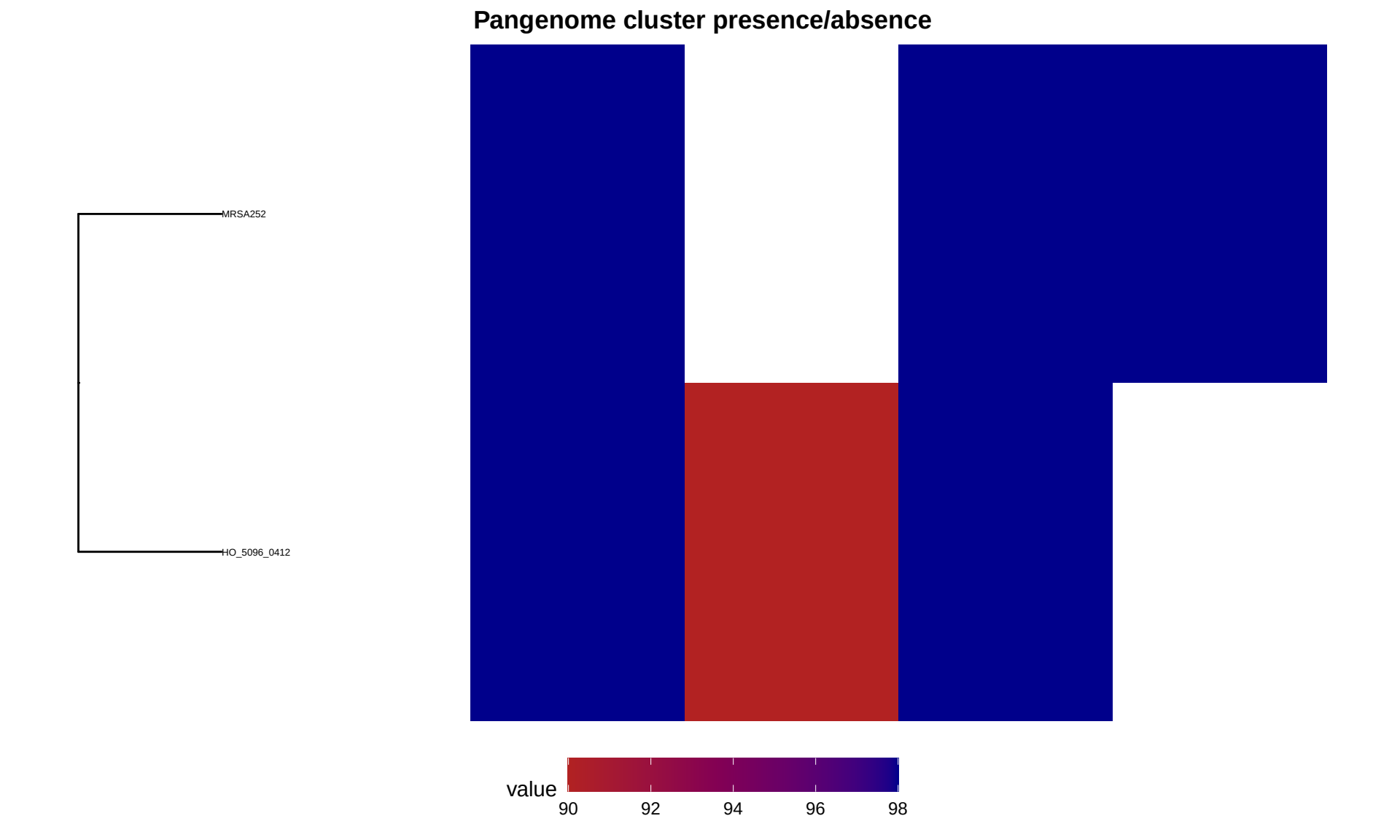

Frequency of gene presence/absence
#Genes
3
2
1
0
0.25
0.50
0.75
1.00
1.25
Proportion
Family Type
Diverged
Stable


Percentage identity of gene family at lowest threshold
Count
3
2
1
0
90
98
Threshold
Gene Type
Fission/Fusion
Single Copy


Number of alleles at maximum %ID threshold
All Data
Genes with > 1 allele
Count
3
2
1
0
1.00
0.75
0.50
0.25
0.00
1
2
3
No. Alleles at Maximum Threshold


Unique alleles per threshold
#Clusters
6
4
2
90
92
94
96
98
Threshold


Number of gene families per sample
Tree
#Clusters
MRSA252
HO_5096_0412
0.00
0.05
0.10
0.15
0.20
0
1
2
3
4


Pangenome cluster presence/absence
MRSA252
HO_5096_0412
value
90
92
94
96
98