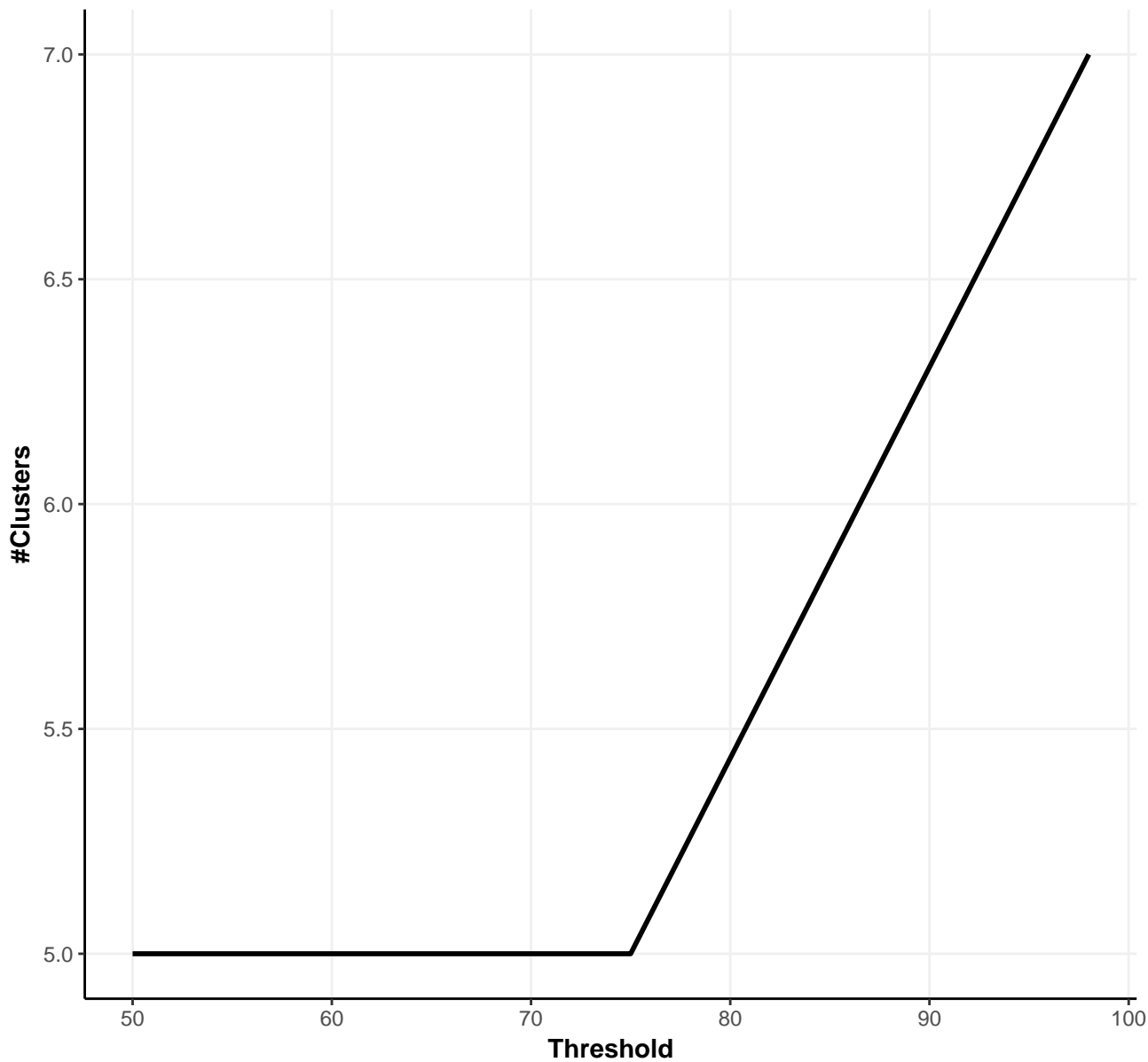
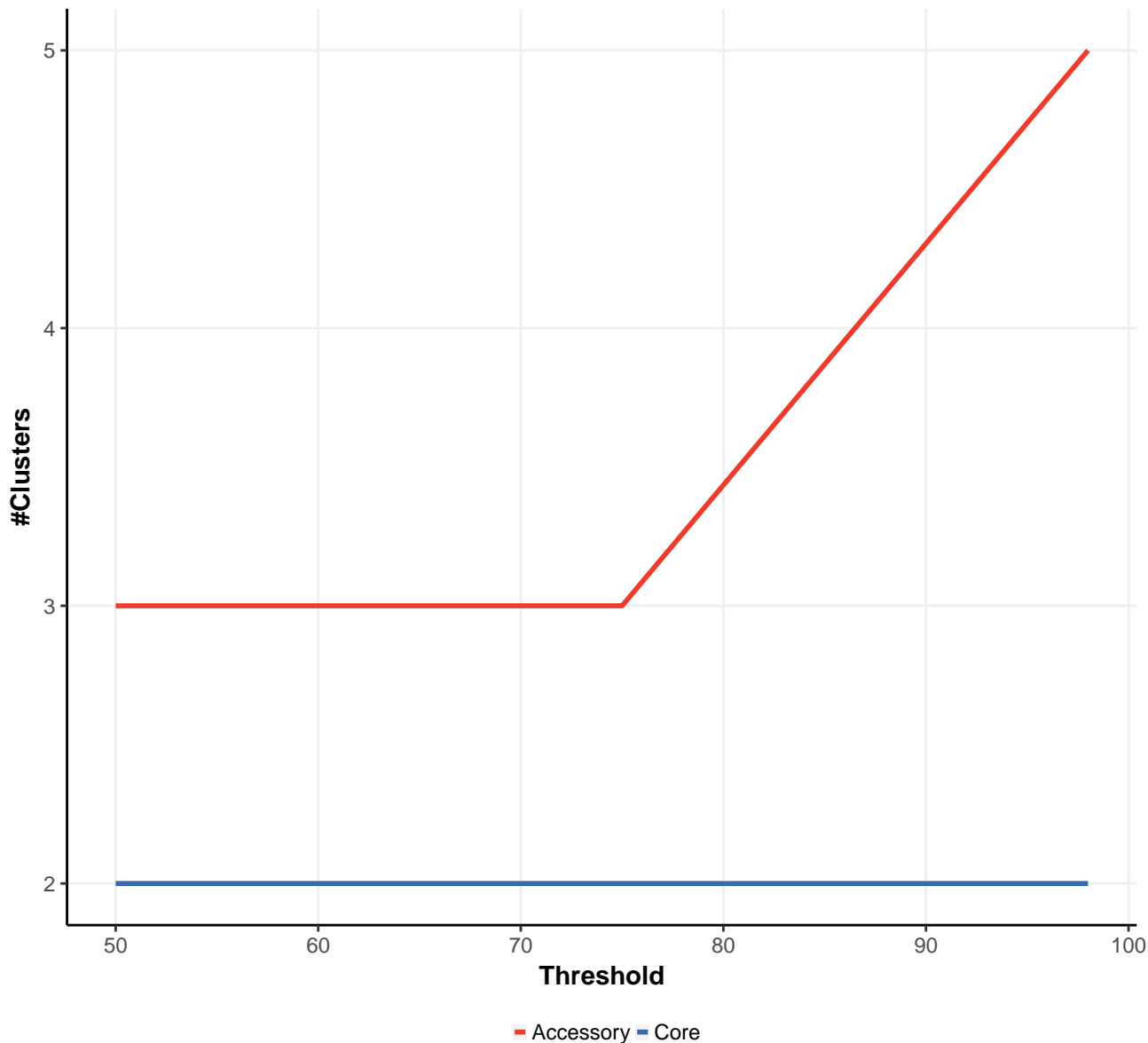


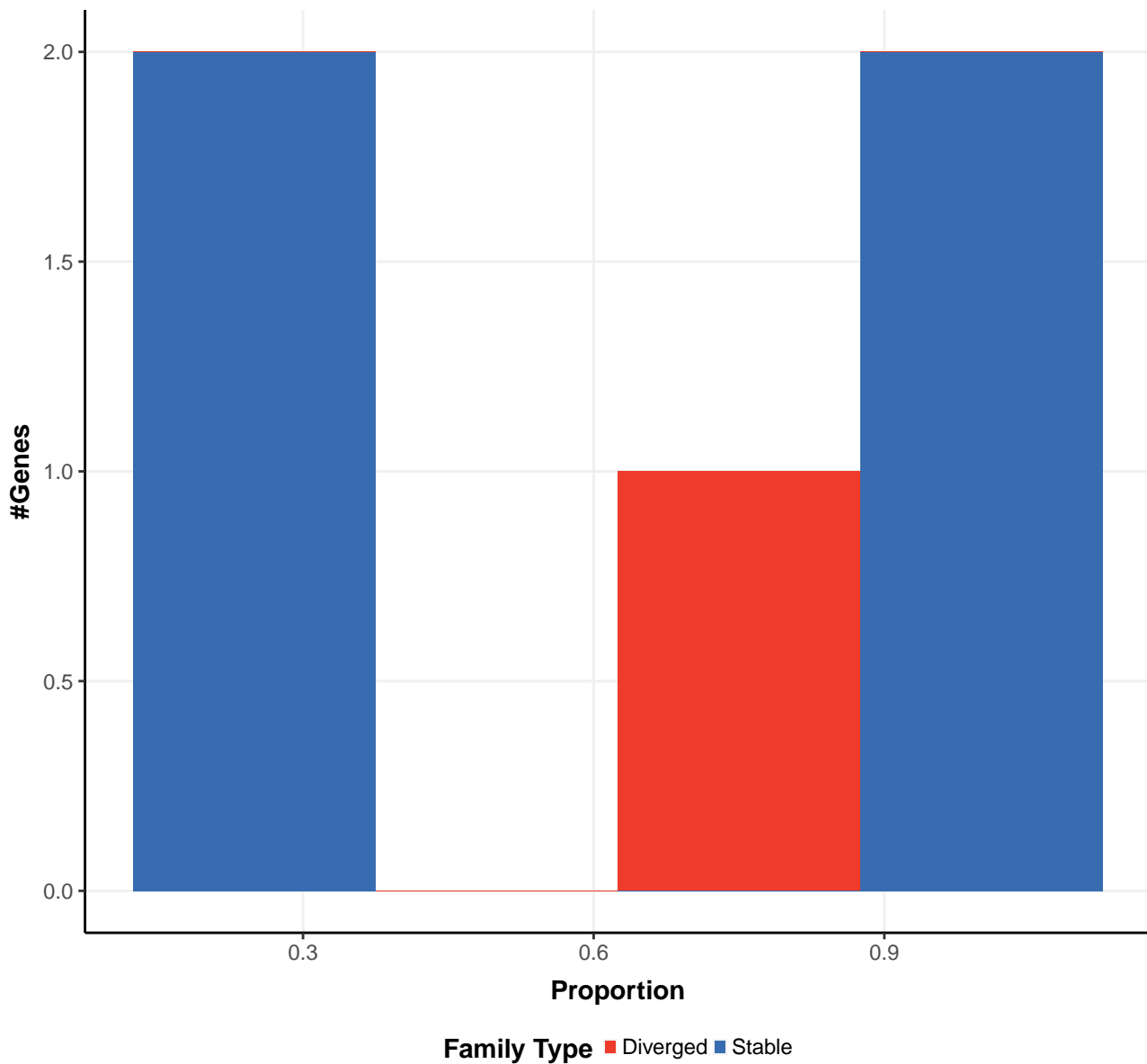
Number of Clusters per Threshold



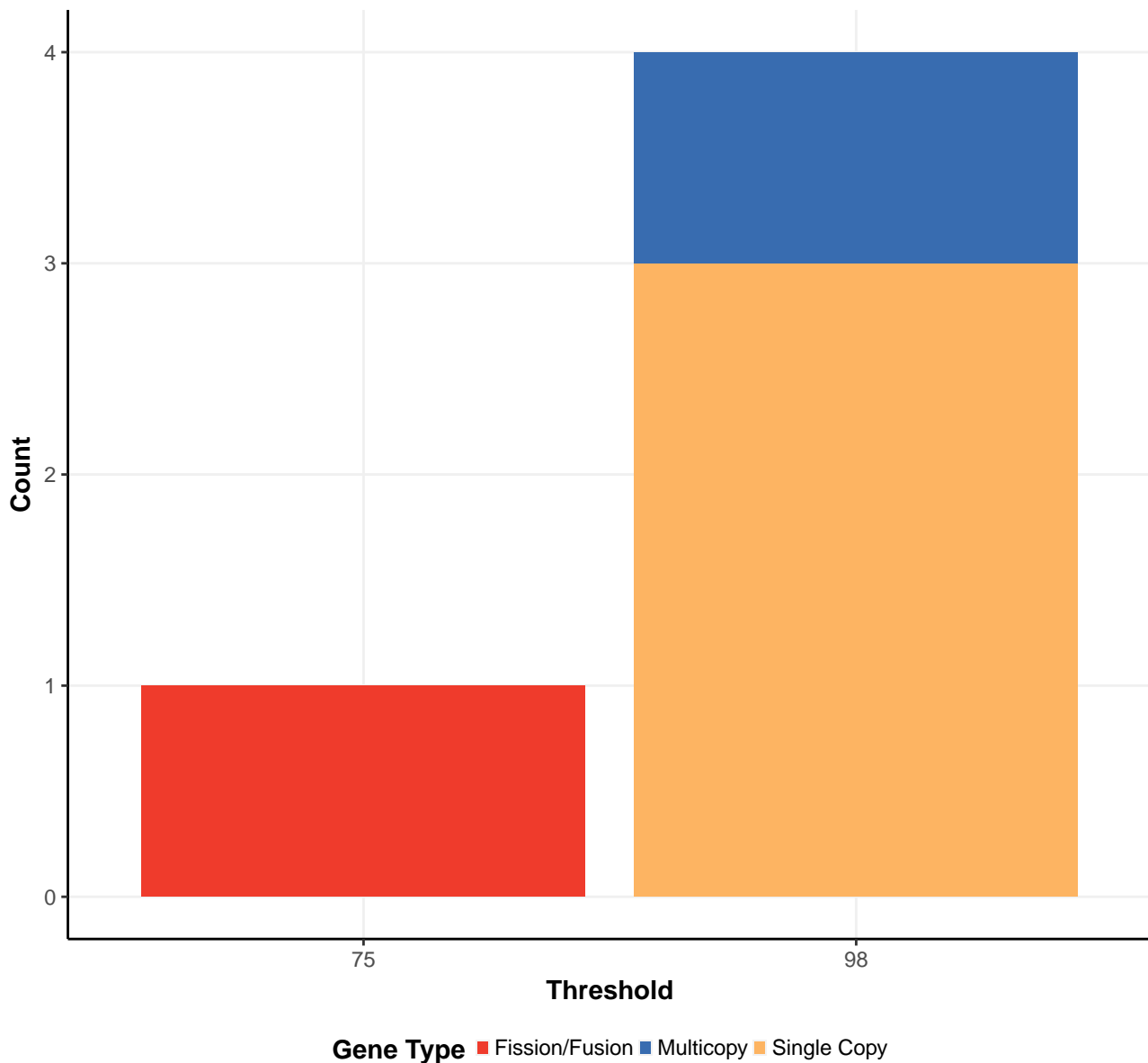
Core(>95%) versus accessory(<95%) alleles per threshold



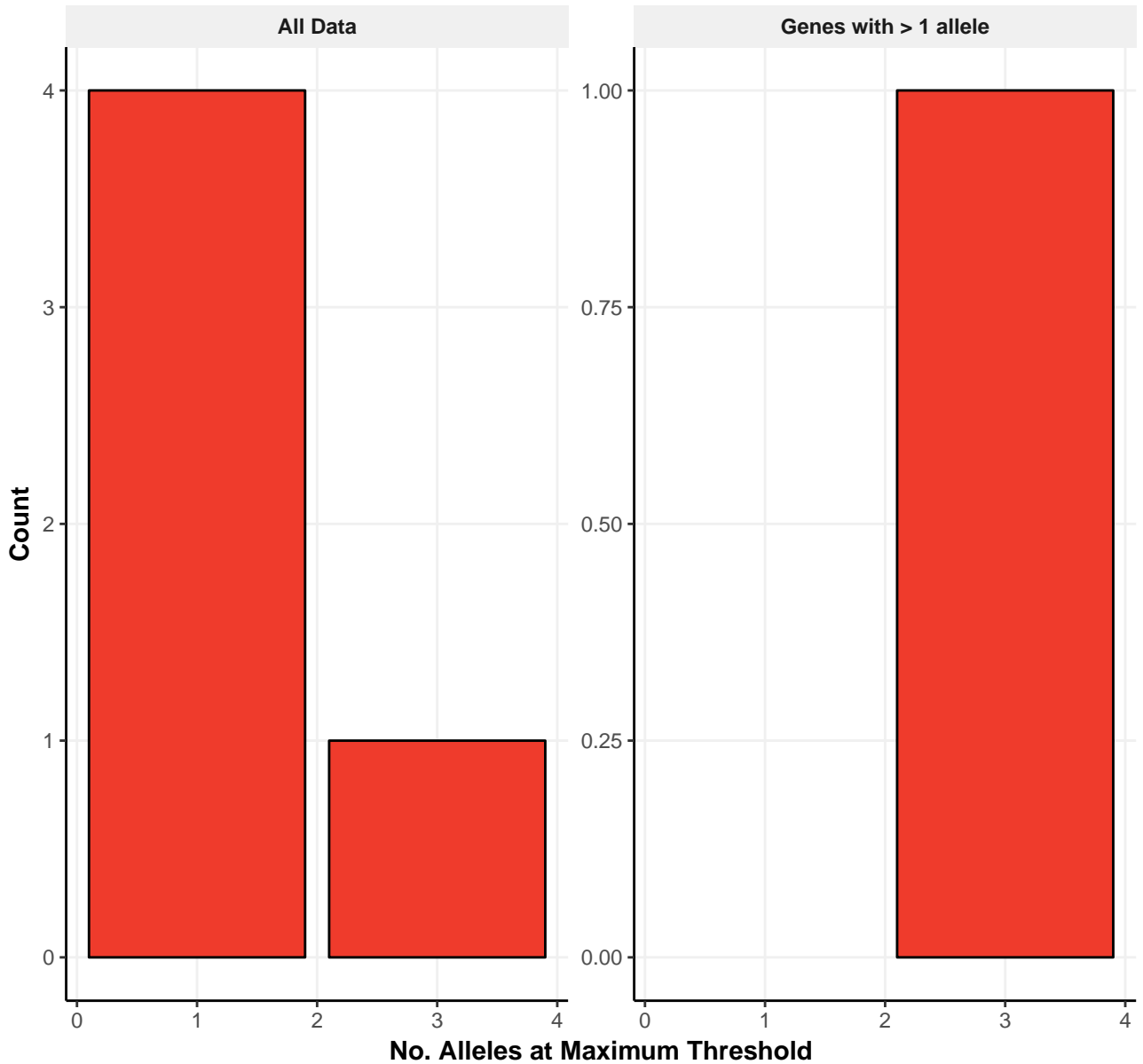
Frequency of gene presence/absence



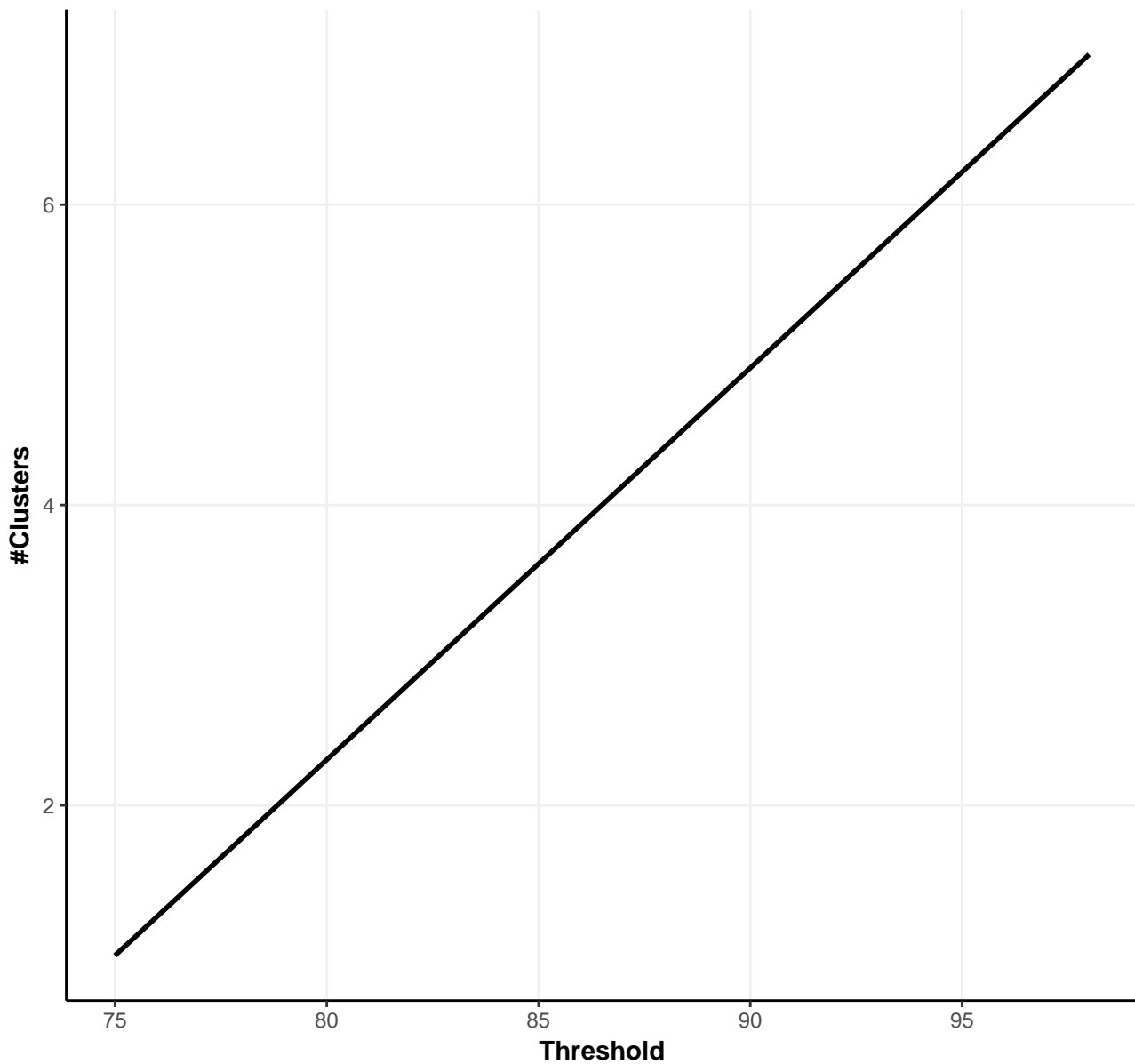
Percentage identity of gene family at lowest threshold



Number of alleles at maximum %ID threshold



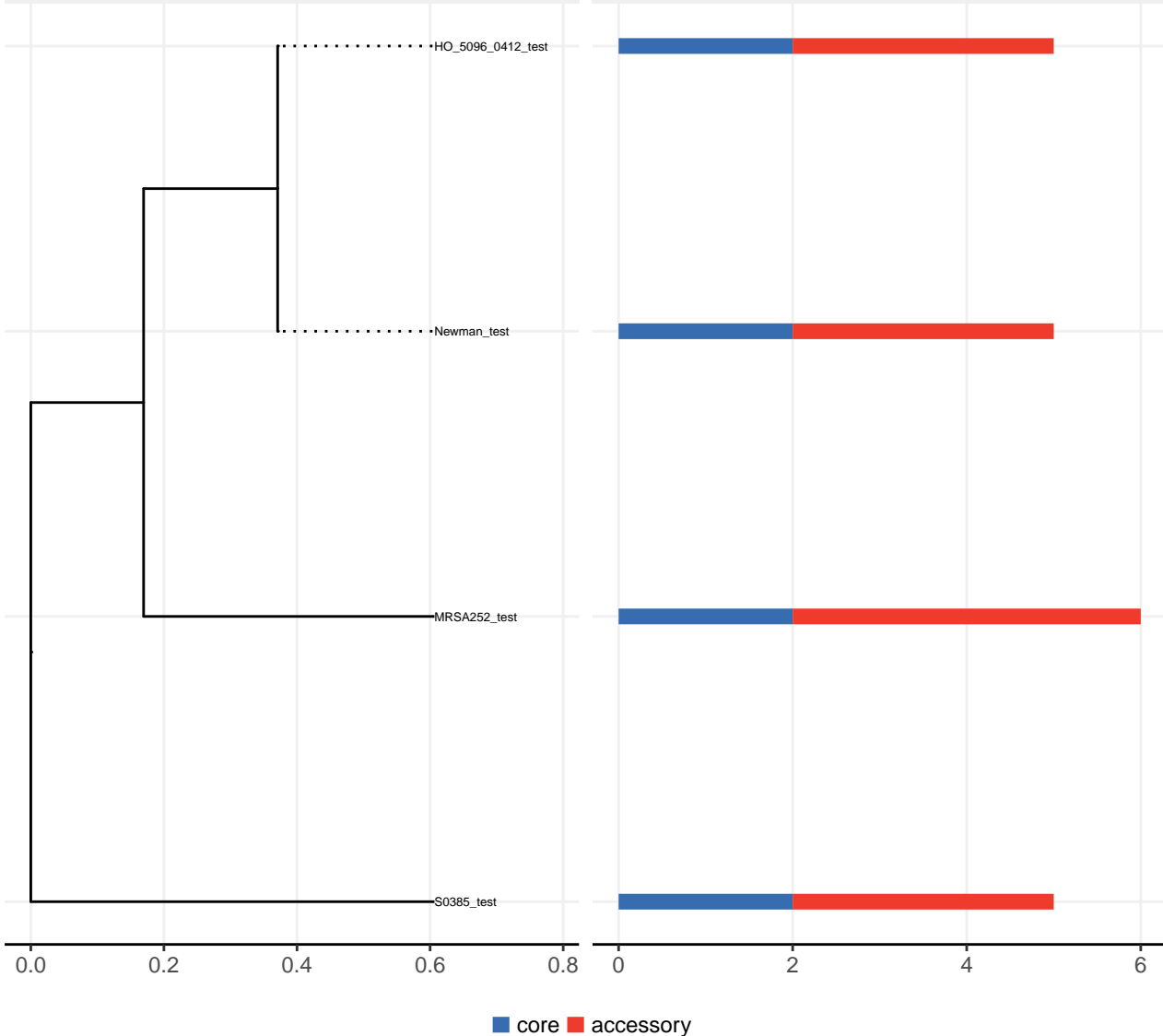
#Unique alleles per threshold



Number of gene families per sample

Tree

#Clusters



Pangenome coloured by minimum clustering threshold

