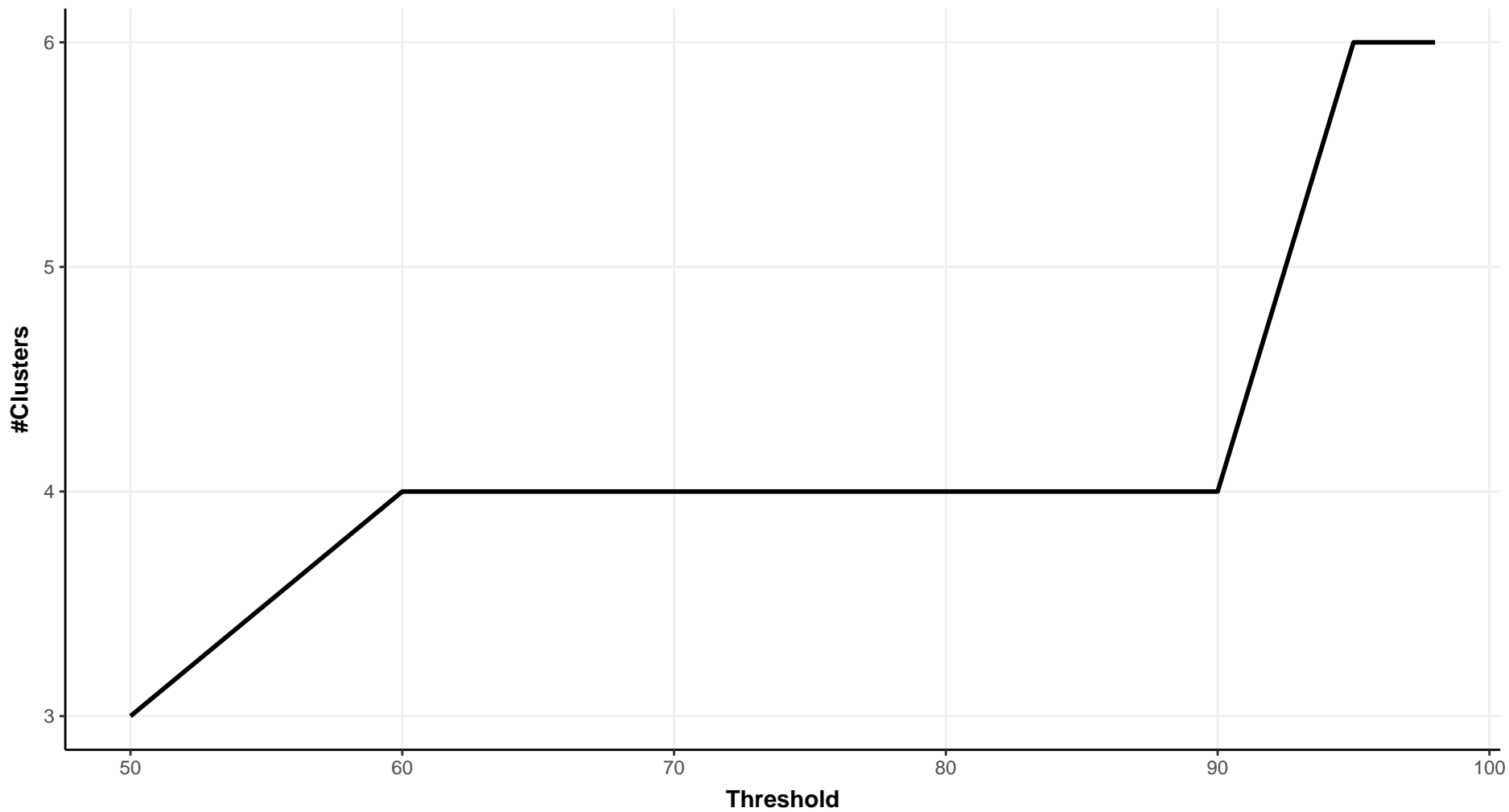
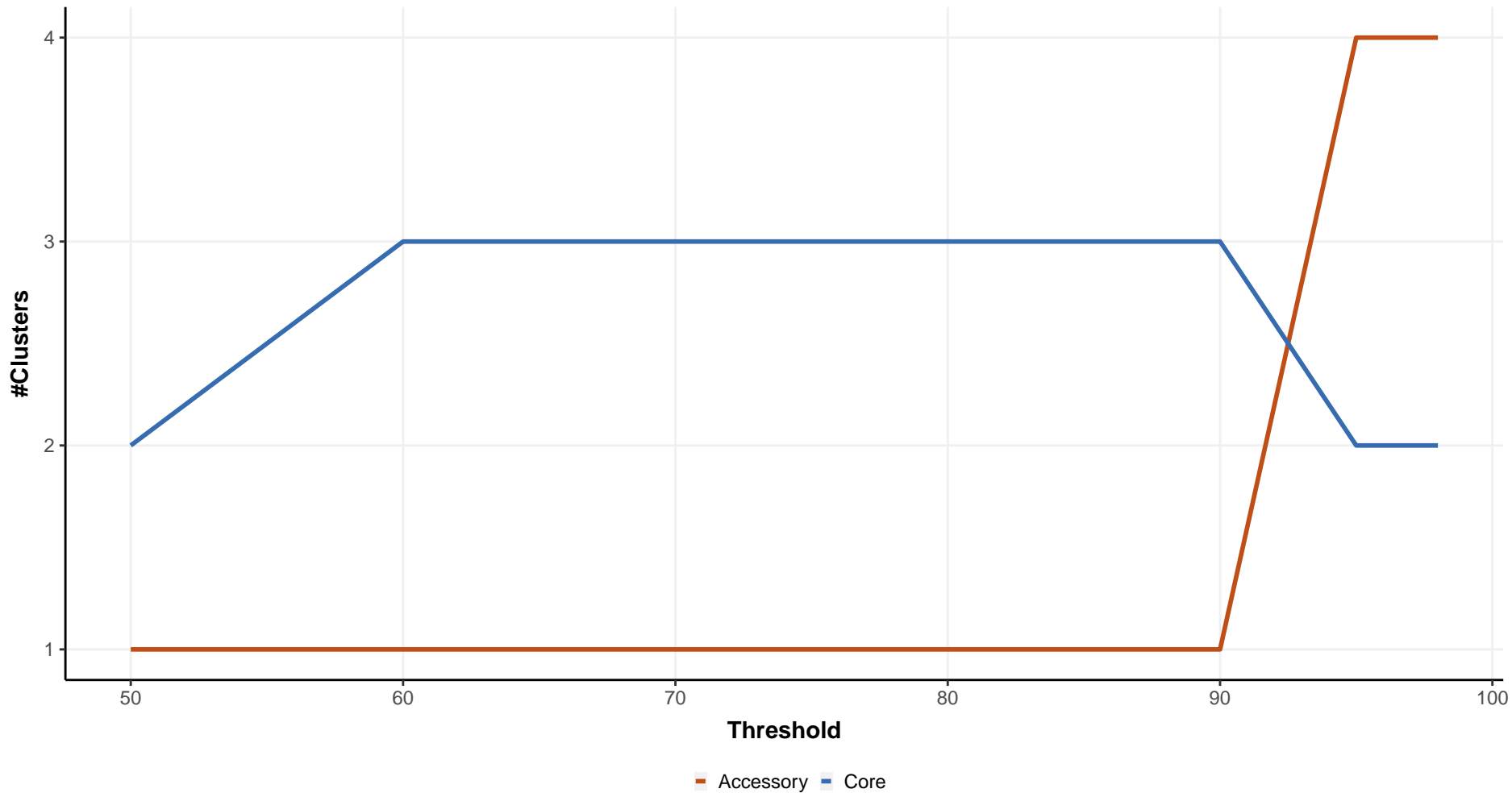


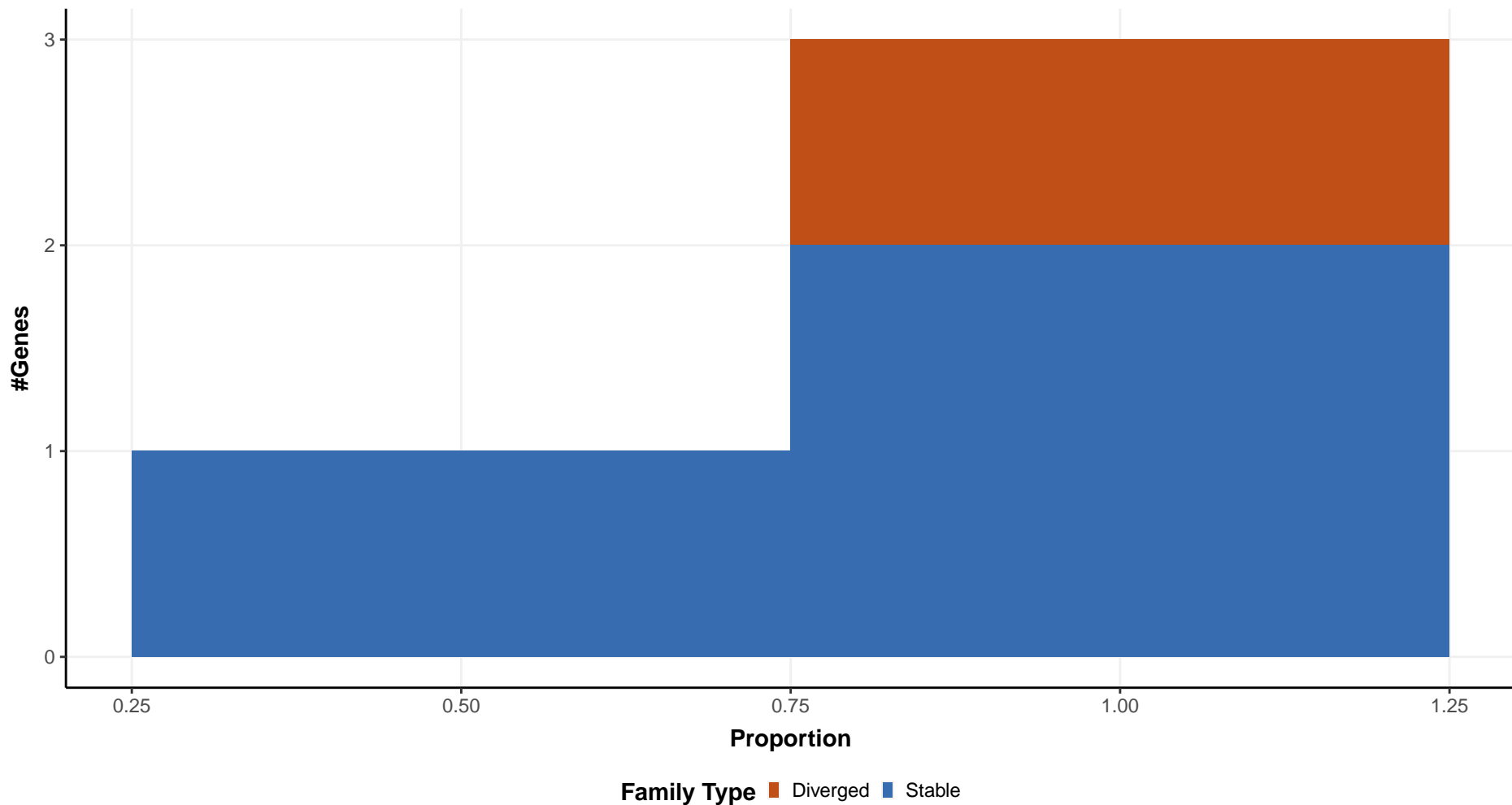
Number of Clusters per Threshold



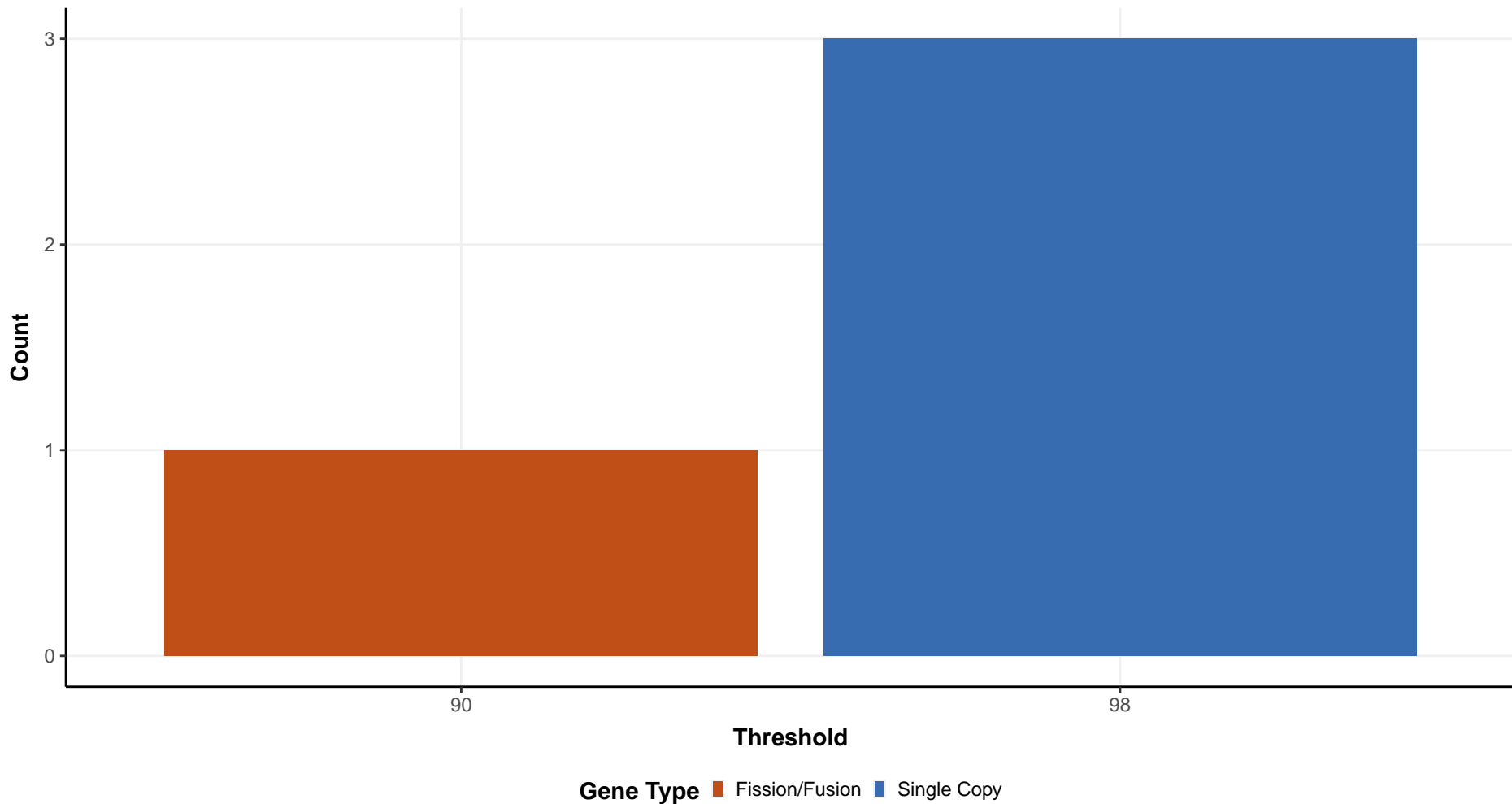
Core(>95%) versus accessory(<95%) alleles per threshold



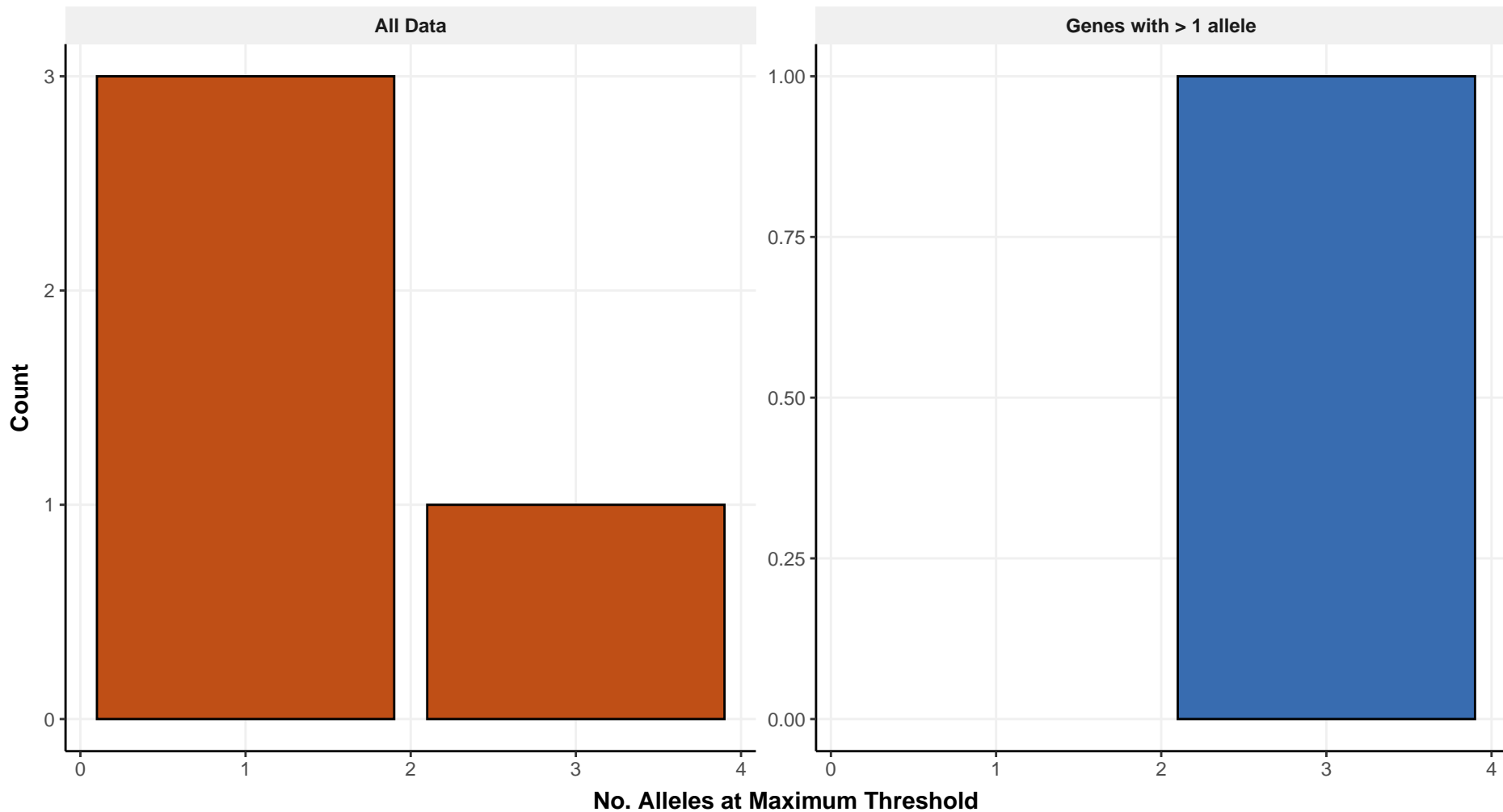
Frequency of gene presence/absence



Percentage identity of gene family at lowest threshold



Number of alleles at maximum %ID threshold



Unique alleles per threshold

