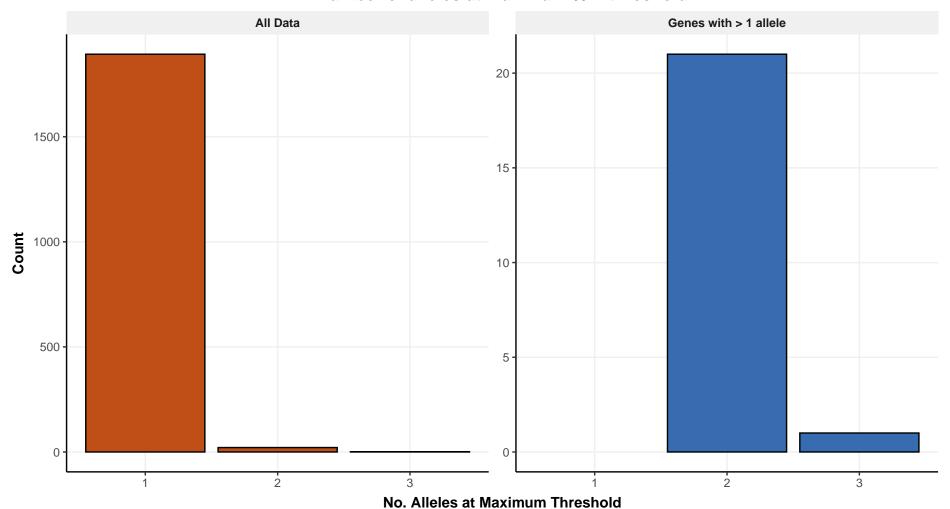
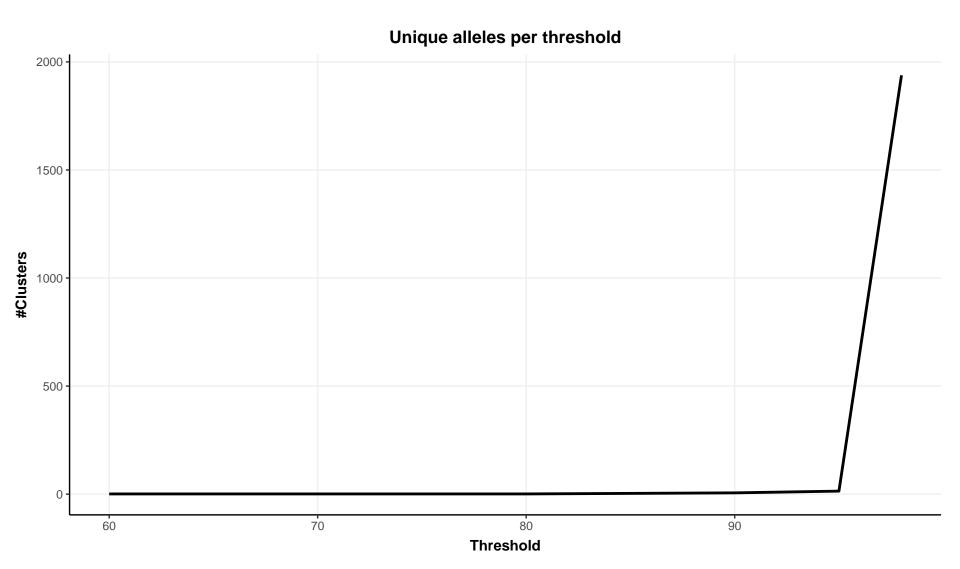
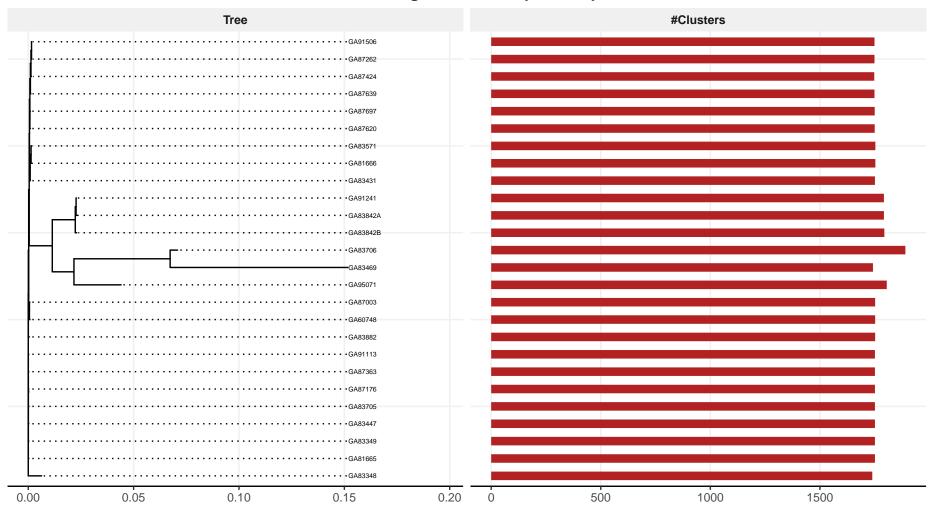


Number of alleles at maximum %ID threshold

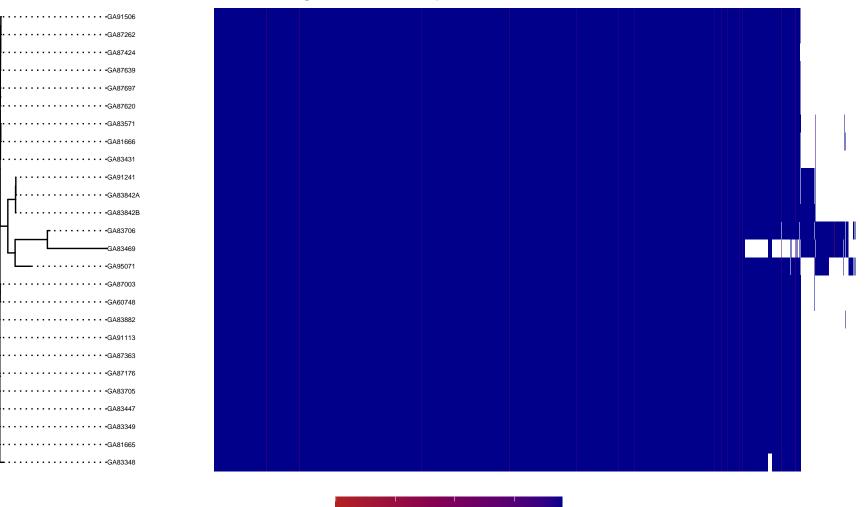




Number of gene families per sample



Pangenome cluster presence/absence



value

70

80

90