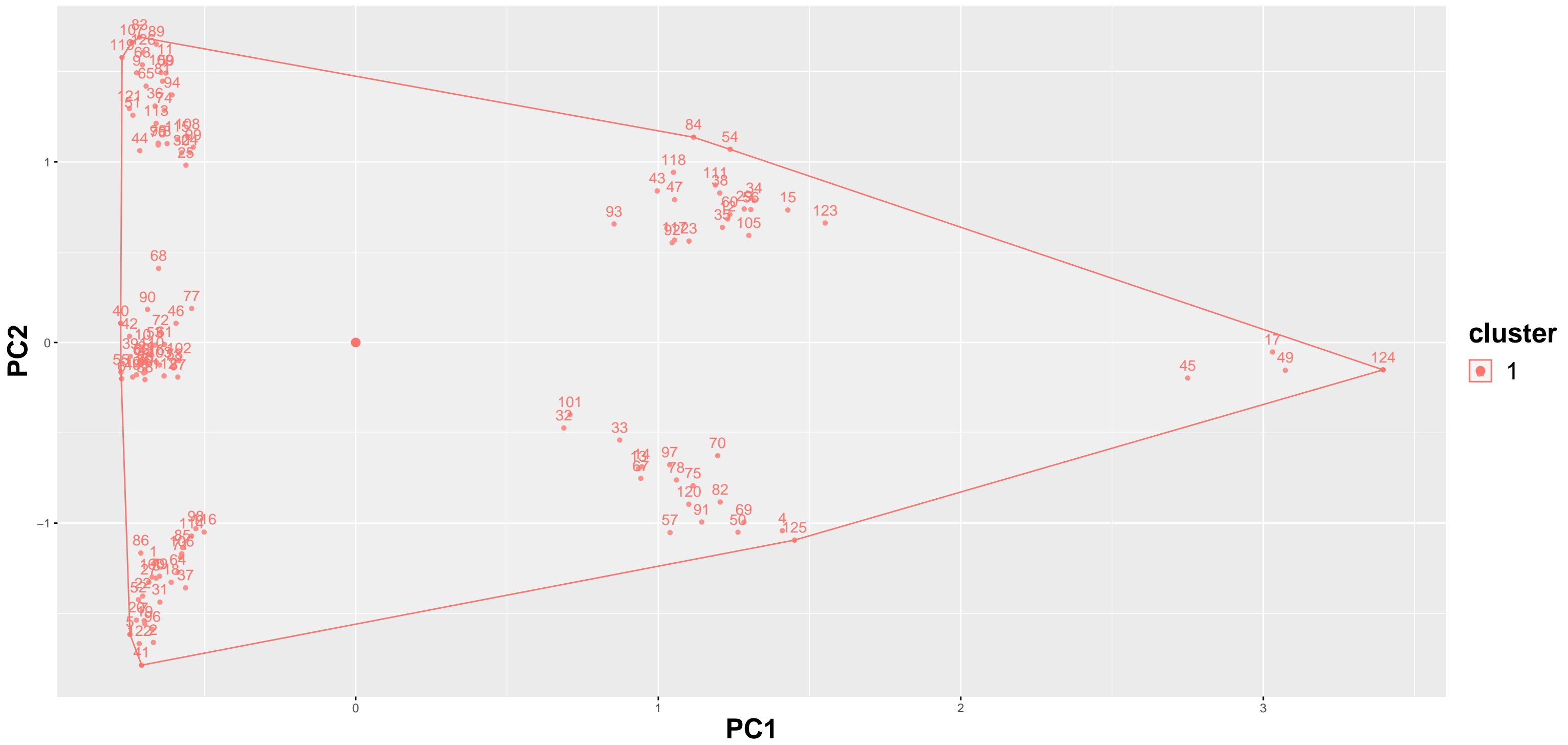
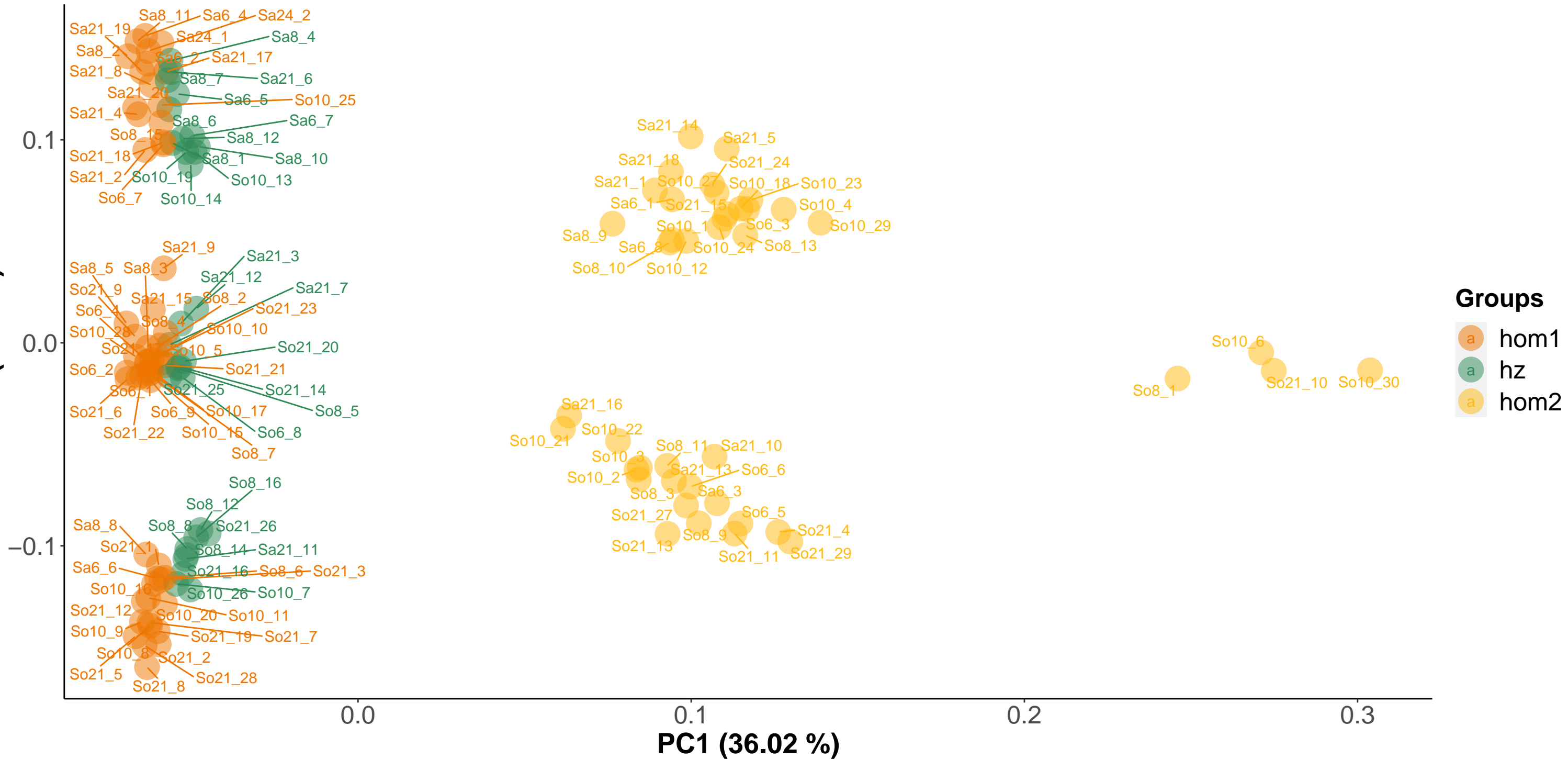


Best K-means clustering plot



PCA based on SNPs present in candidate structural variant

PC2 (9.79 %)



Heterozygosity between the Kmeans-based groups

