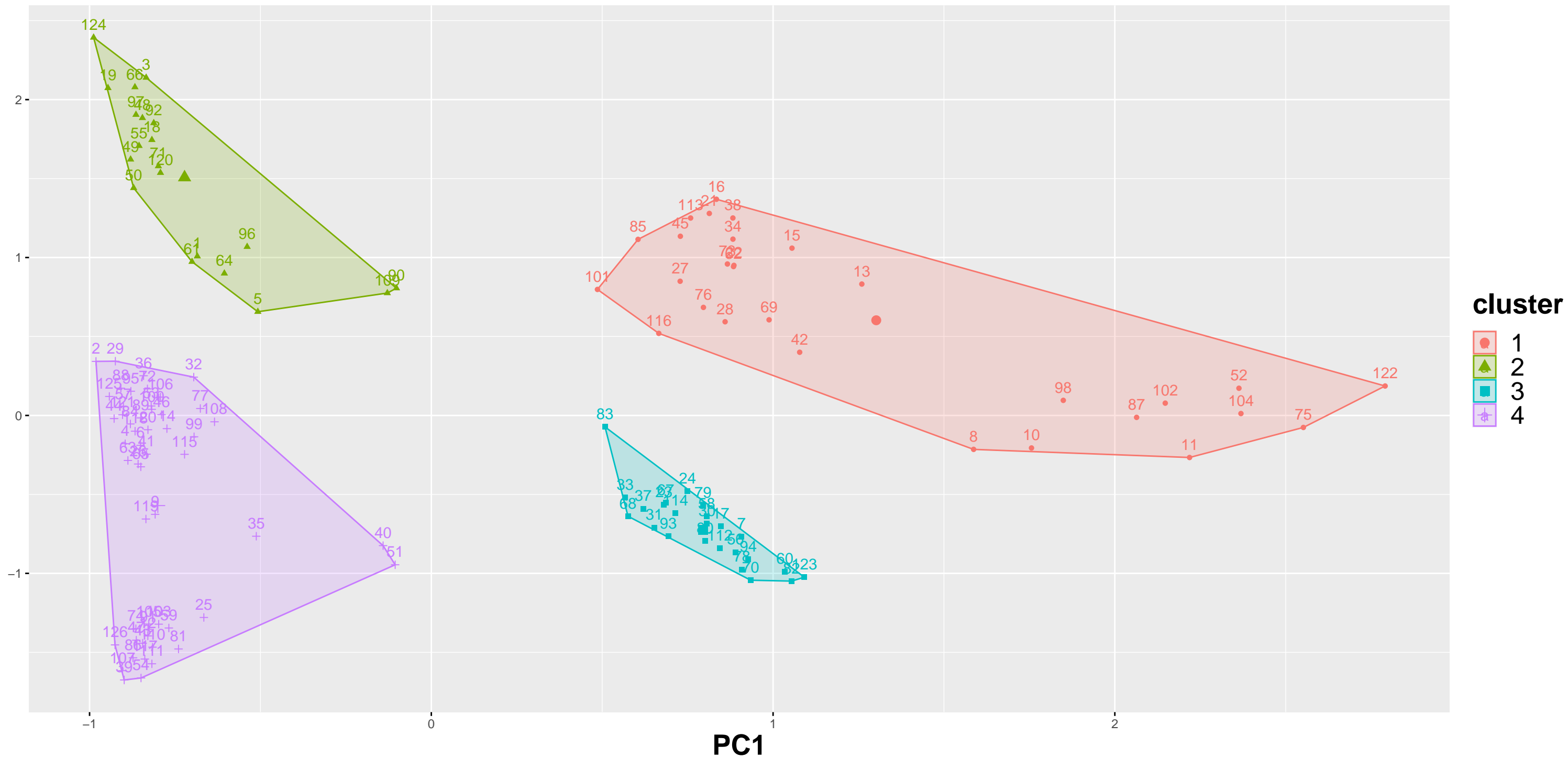
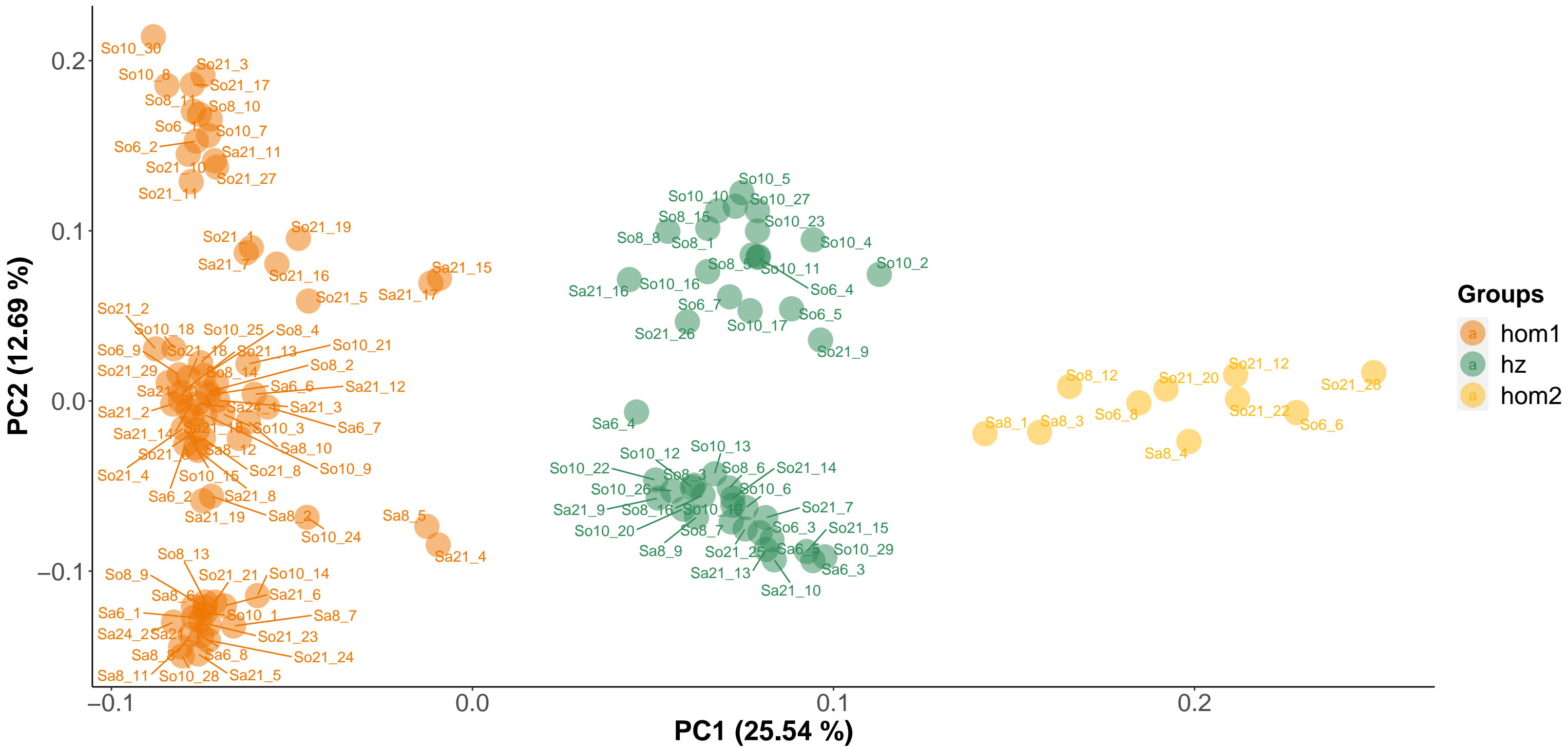


Best K-means clustering plot

PC2



PCA based on SNPs present in candidate structural variant



Heterozygosity between the Kmeans-based groups

Proportion of heterozygous sites

