Best K−means clustering plot
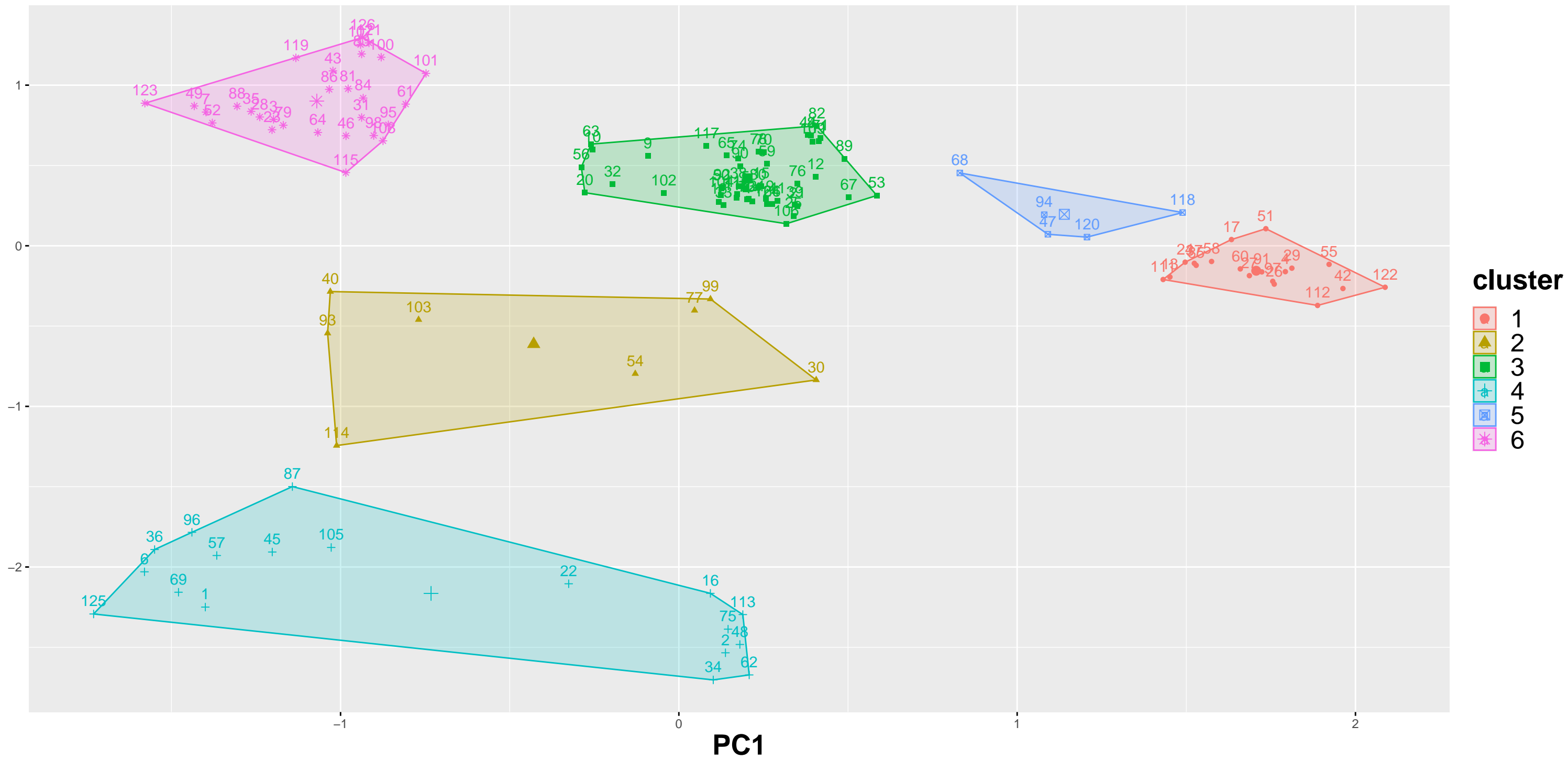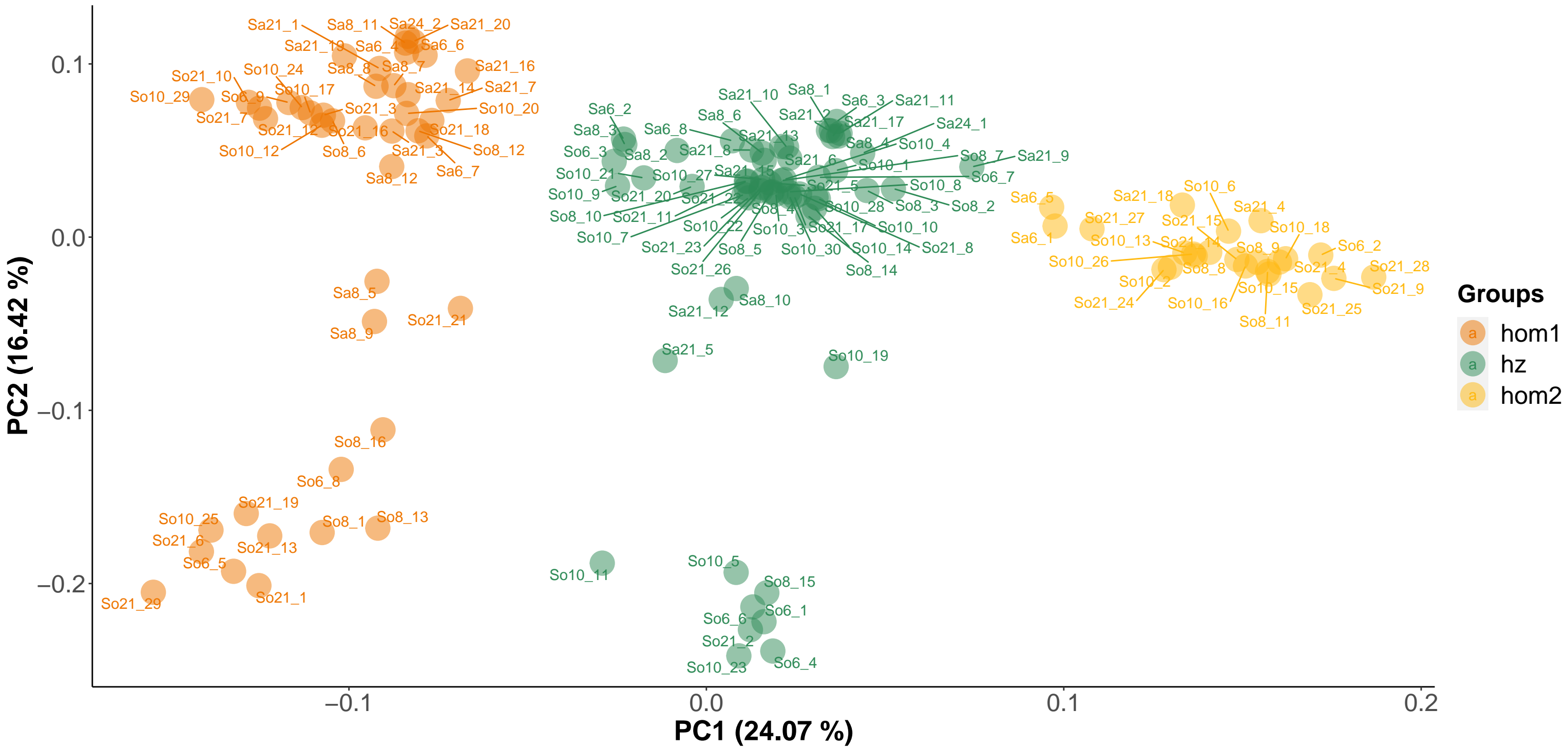
PCA based on SNPs present in candidate structural variant

Heterozygosity between the Kmeans−based groups