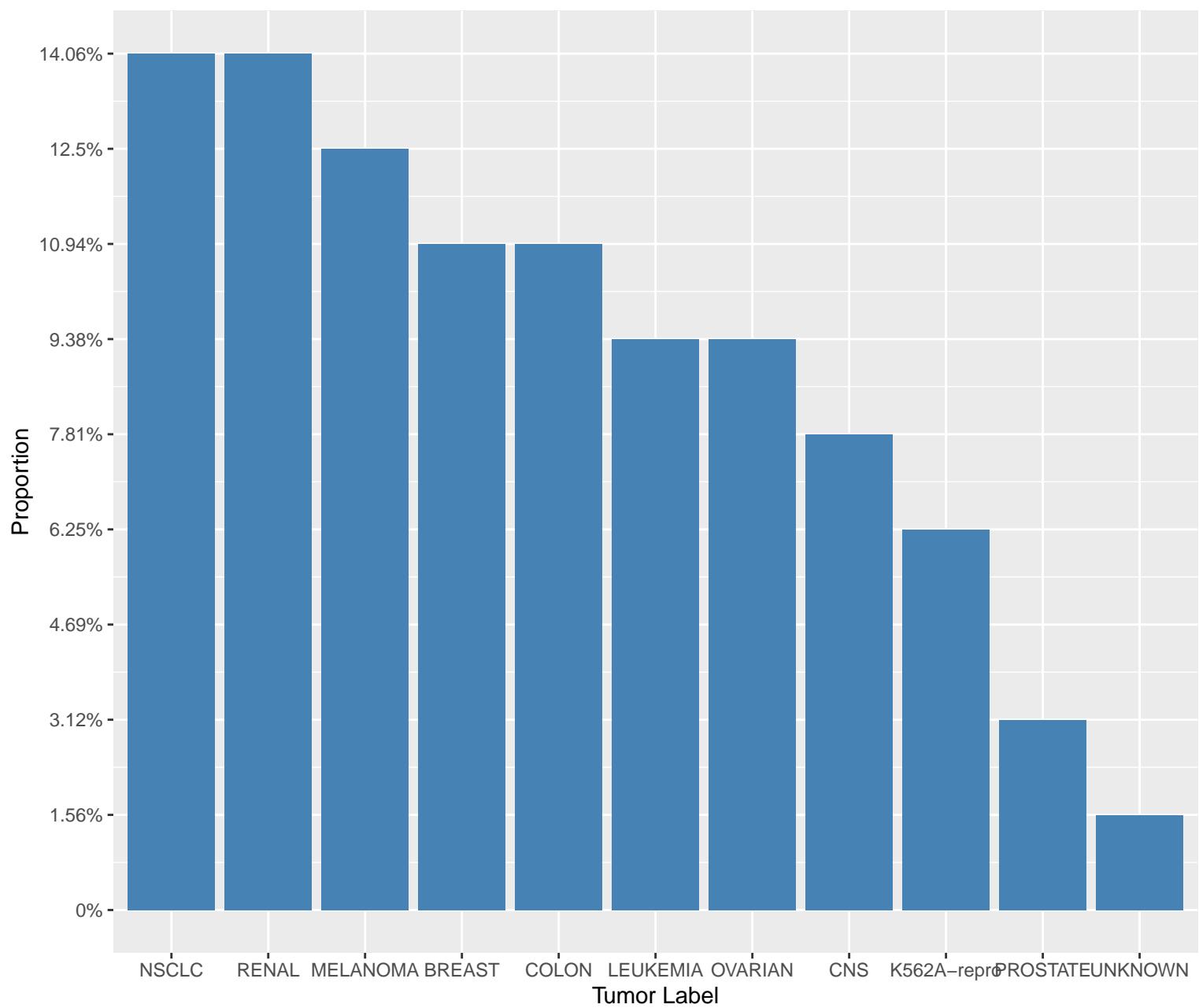
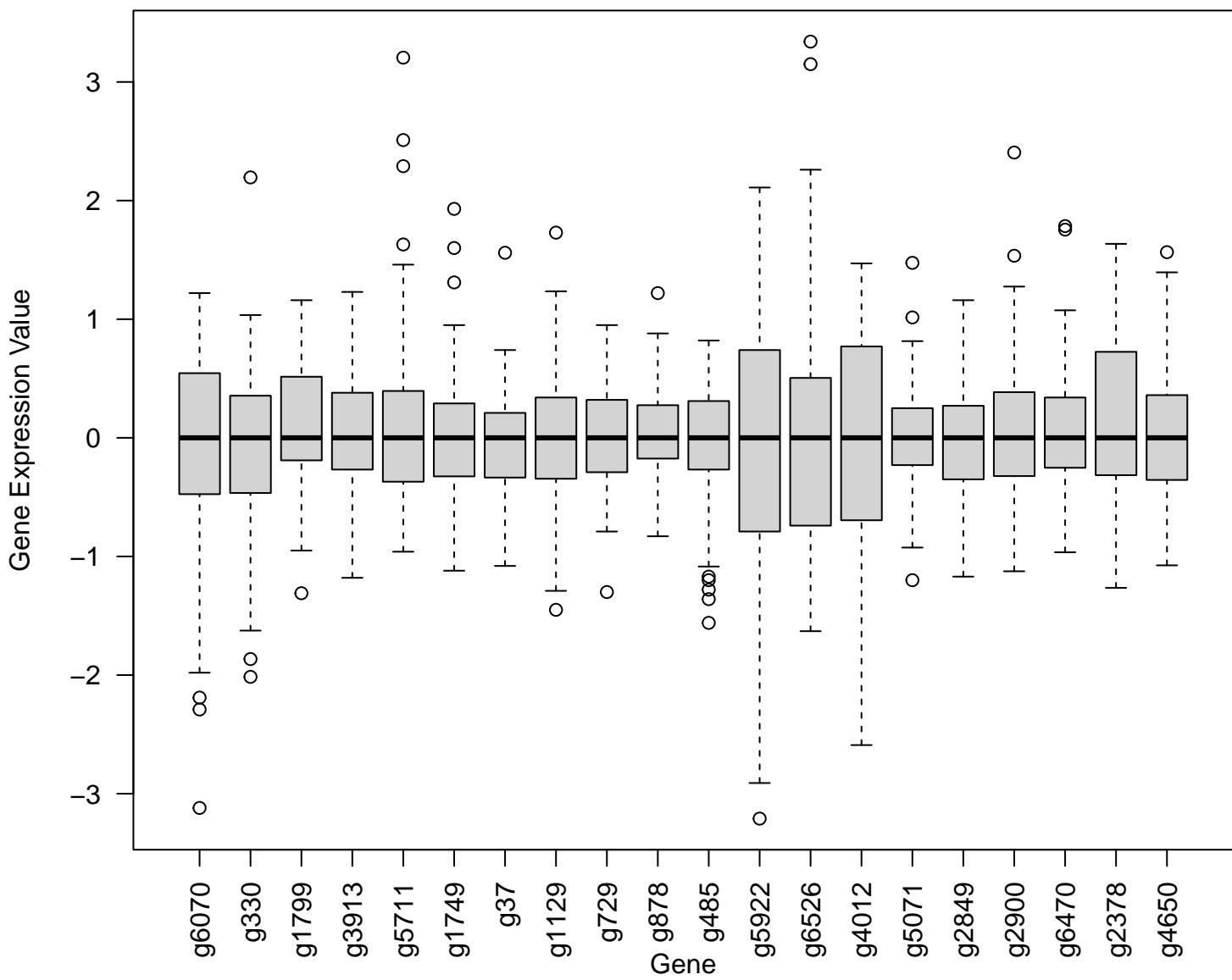


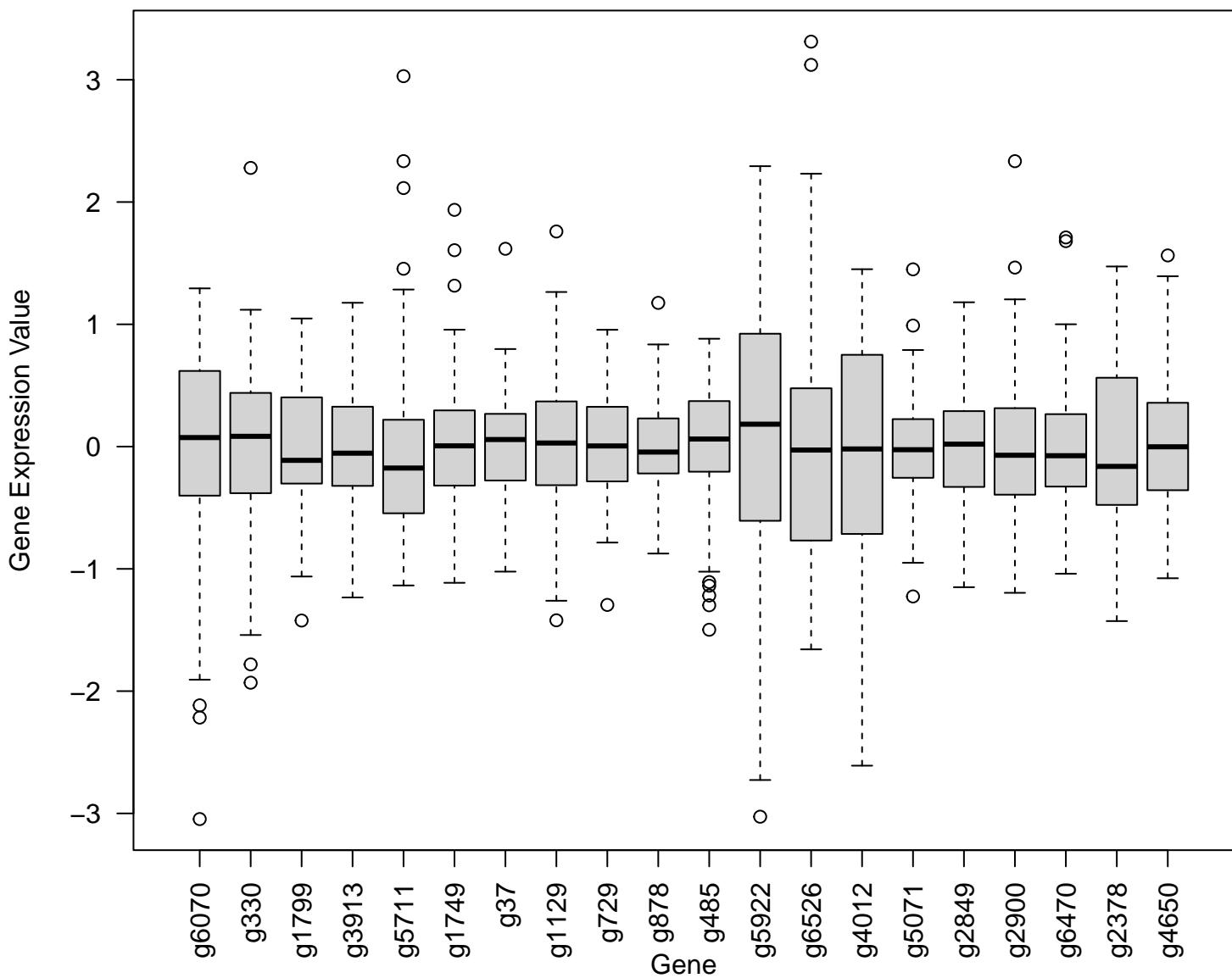
Tumor Label Distribution



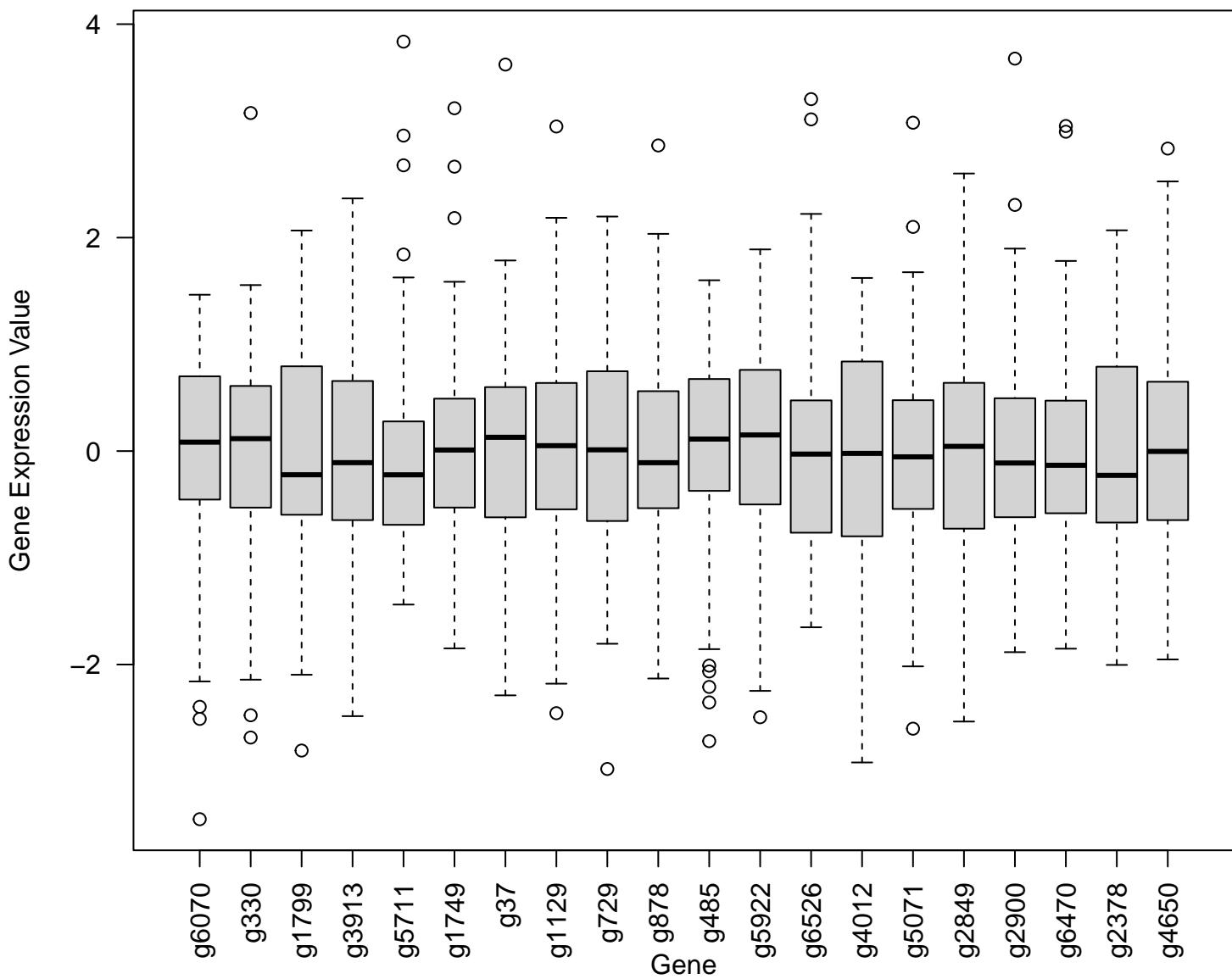
Subset Gene Expression marginalised over patients

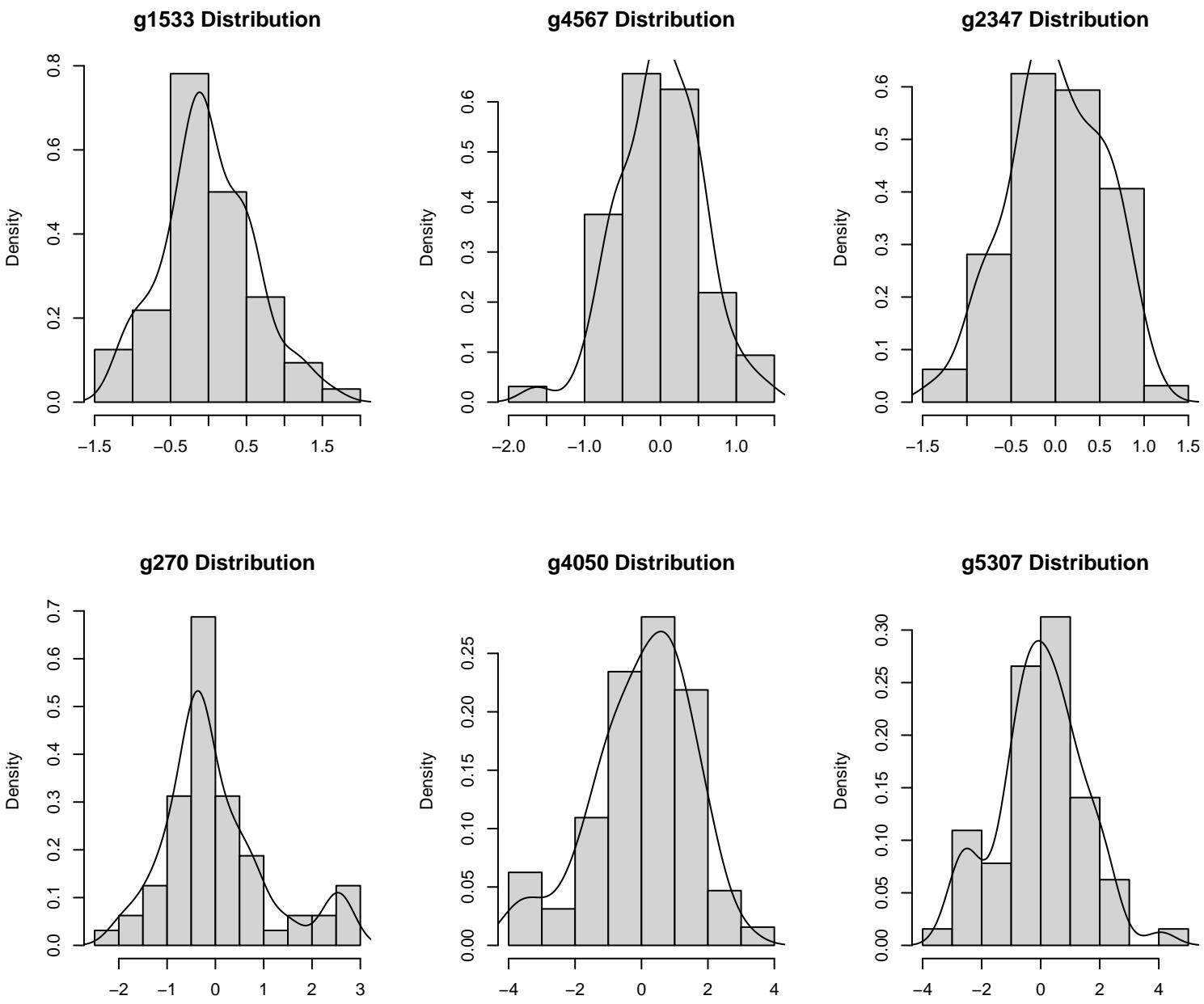


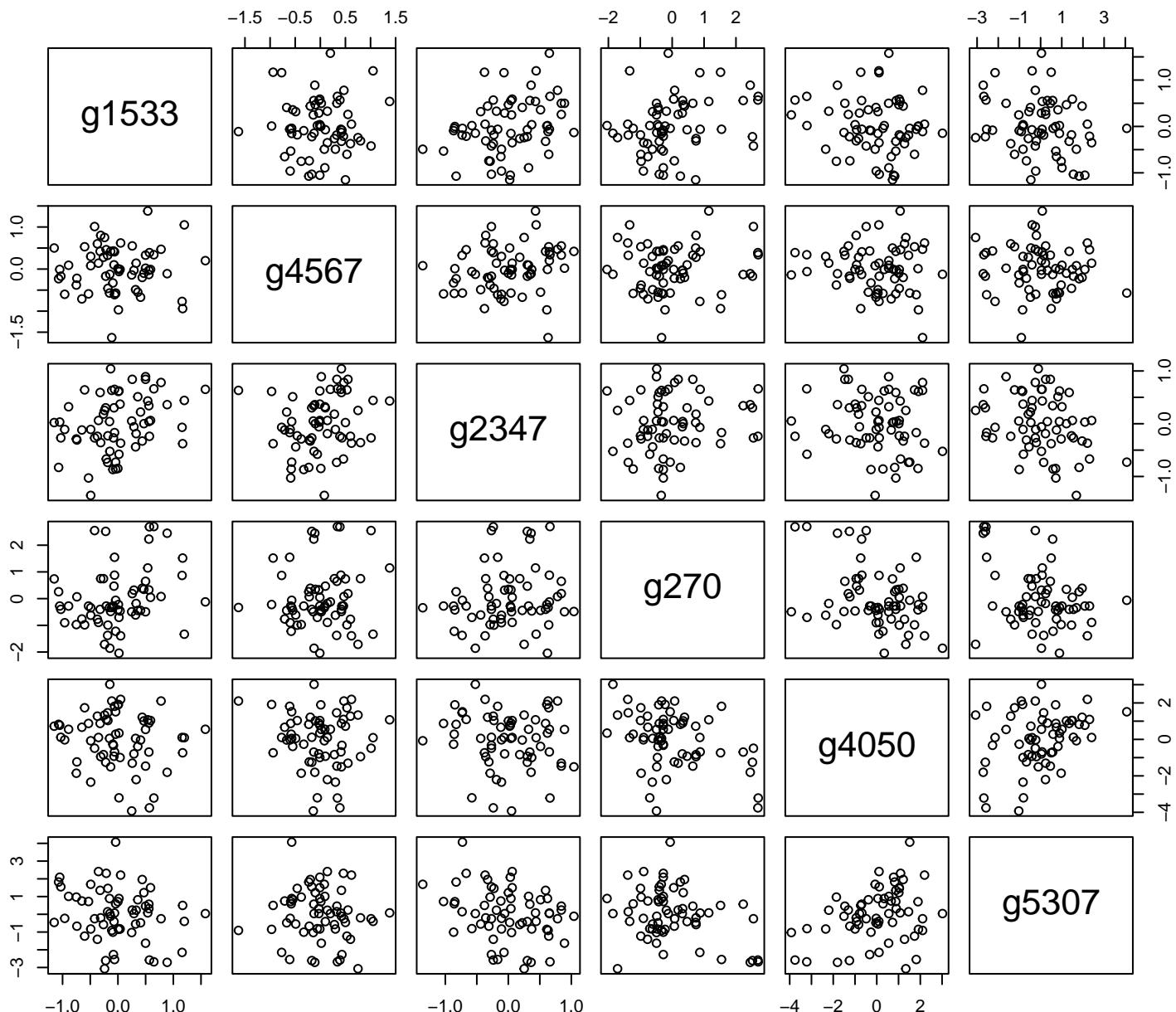
Subset Gene Expression marginalised over patients (centered)

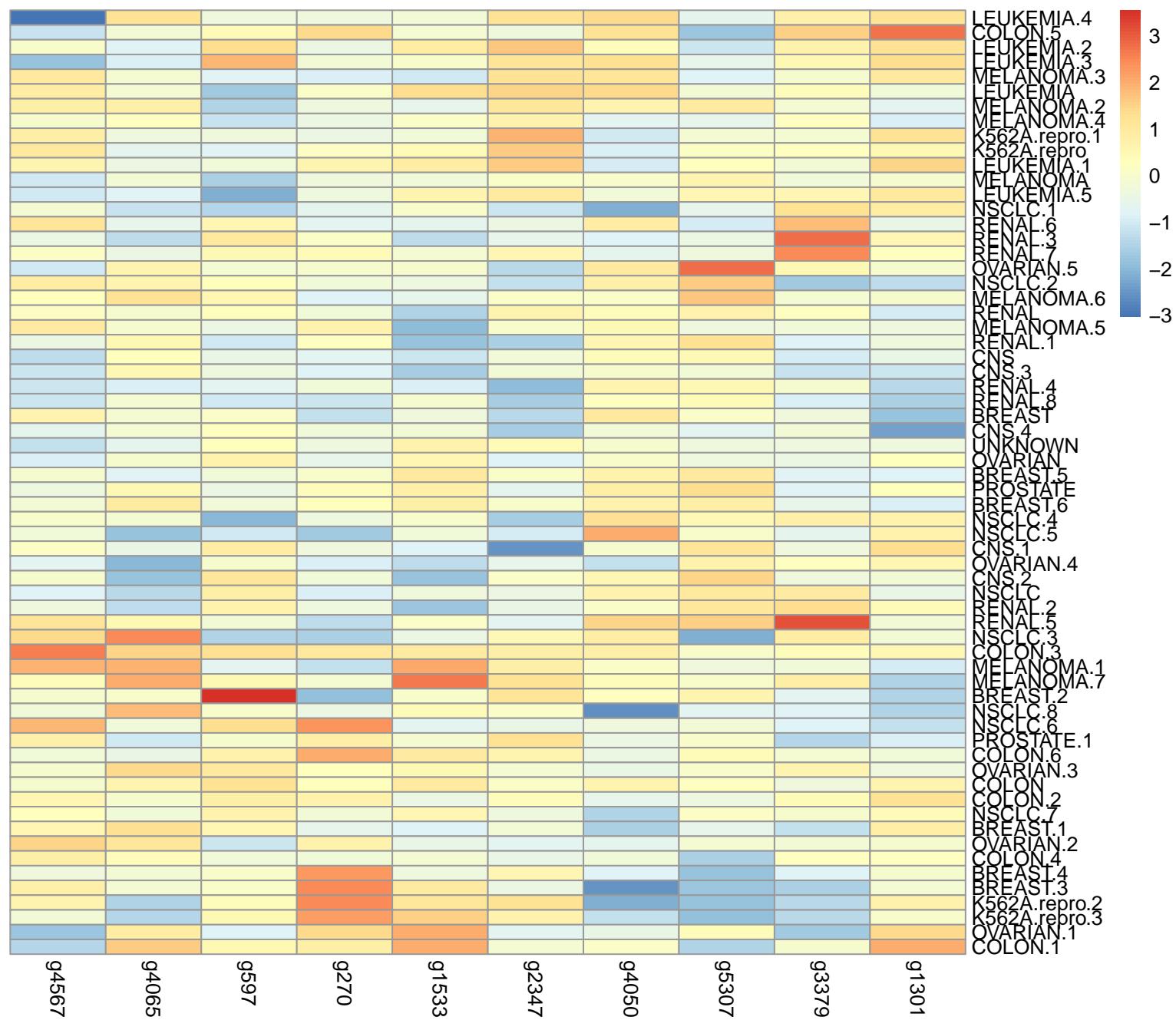


Subset Gene Expression marginalised over patients (centered and scaled)

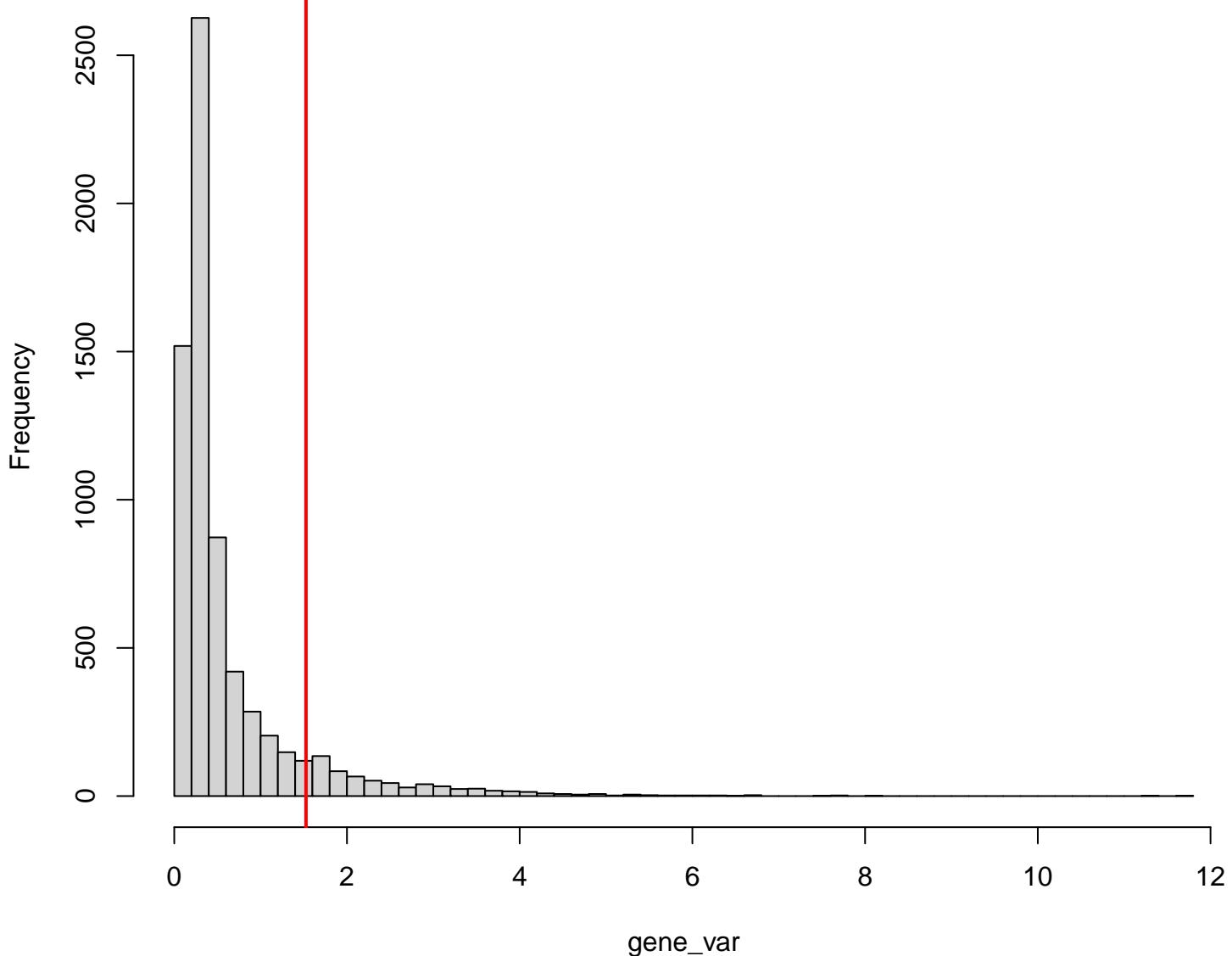




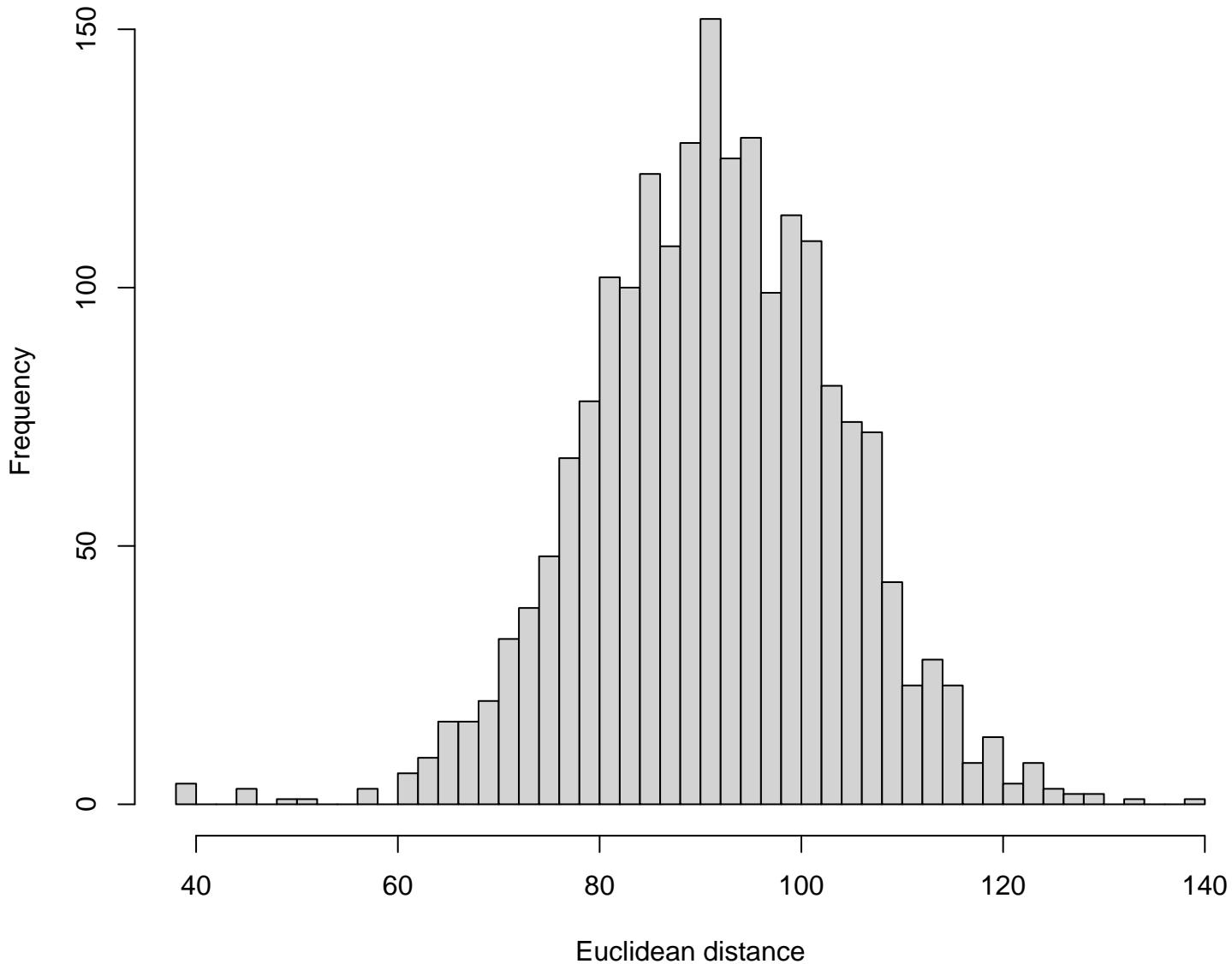




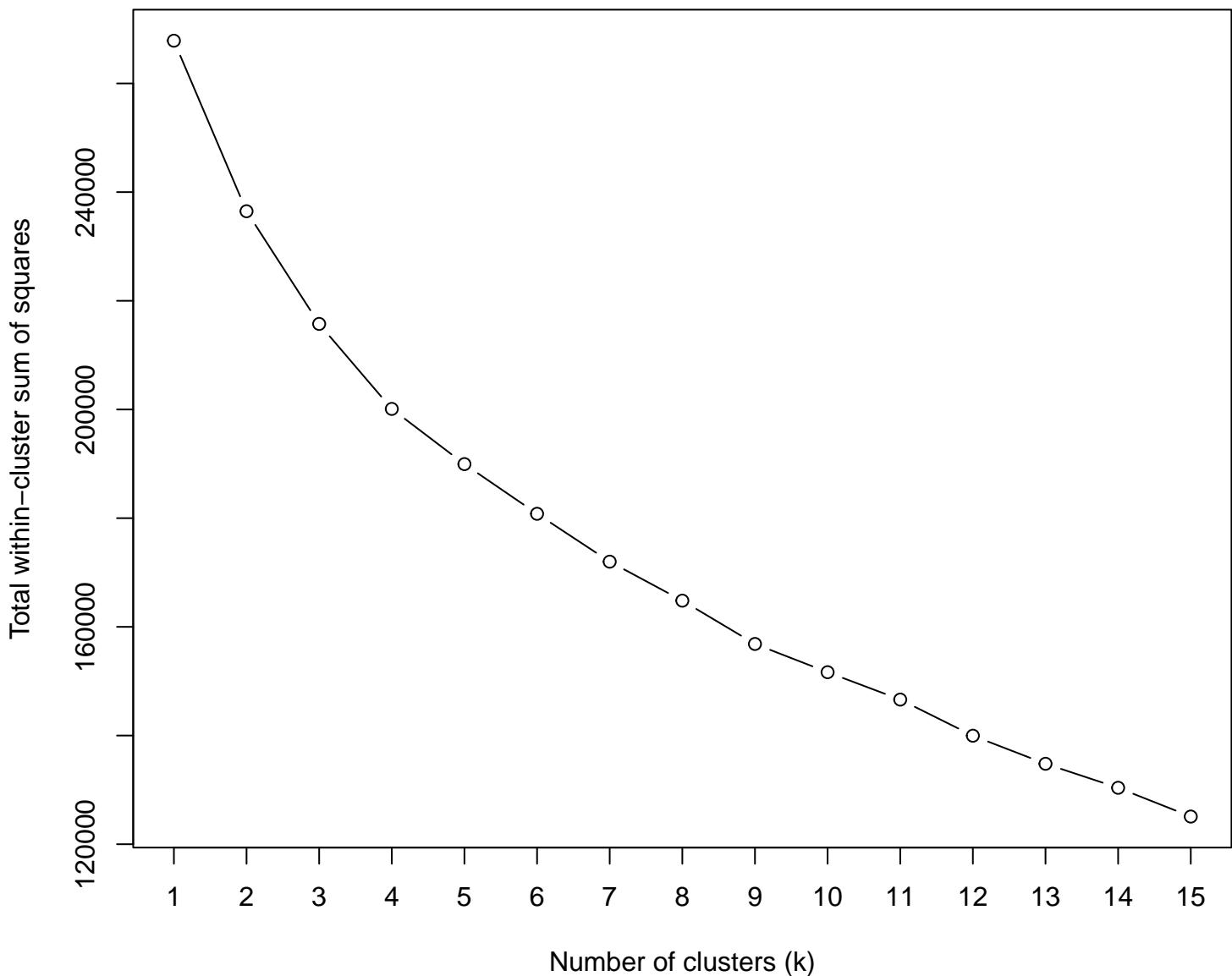
Gene-wise variance distribution



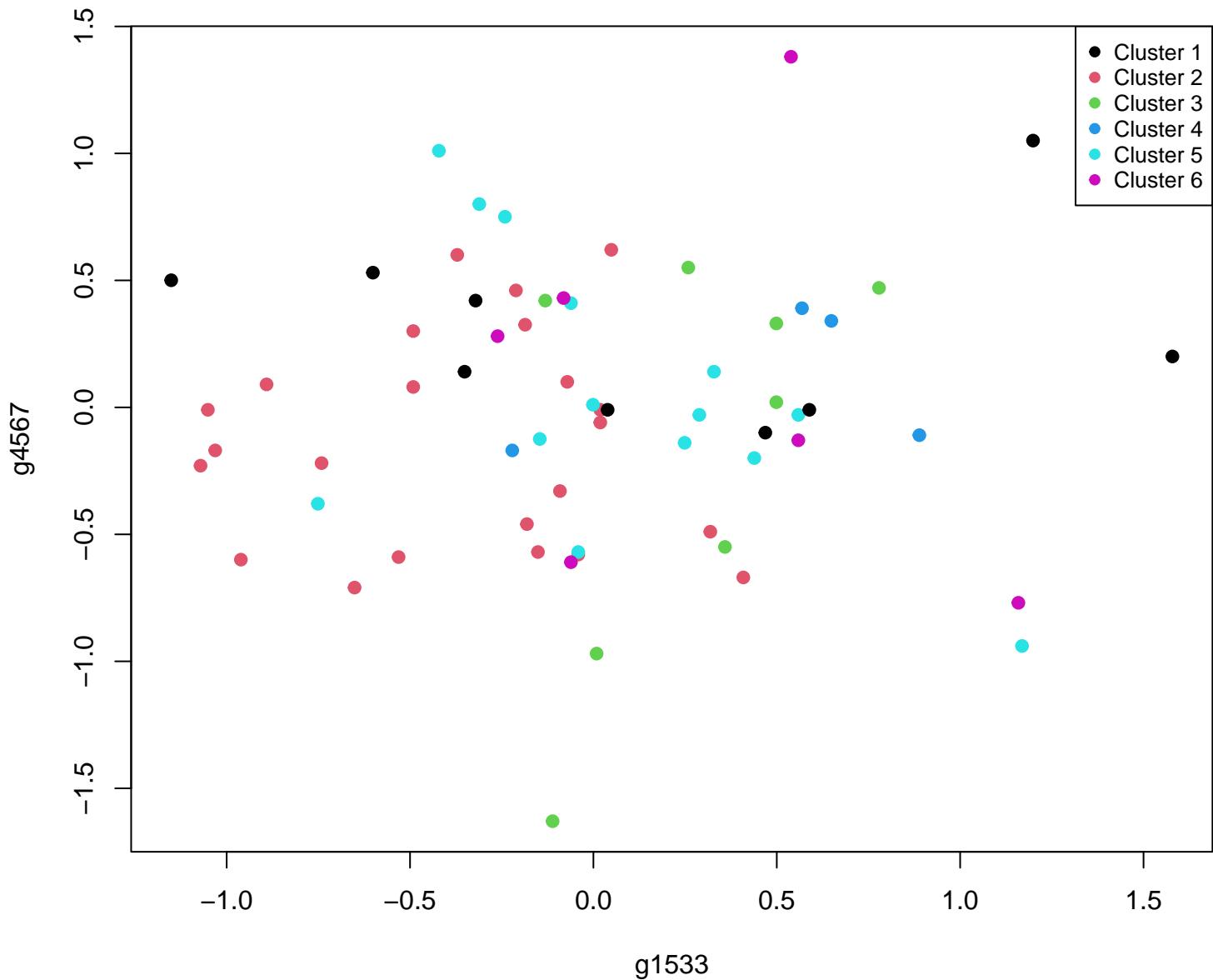
Pairwise distances between samples



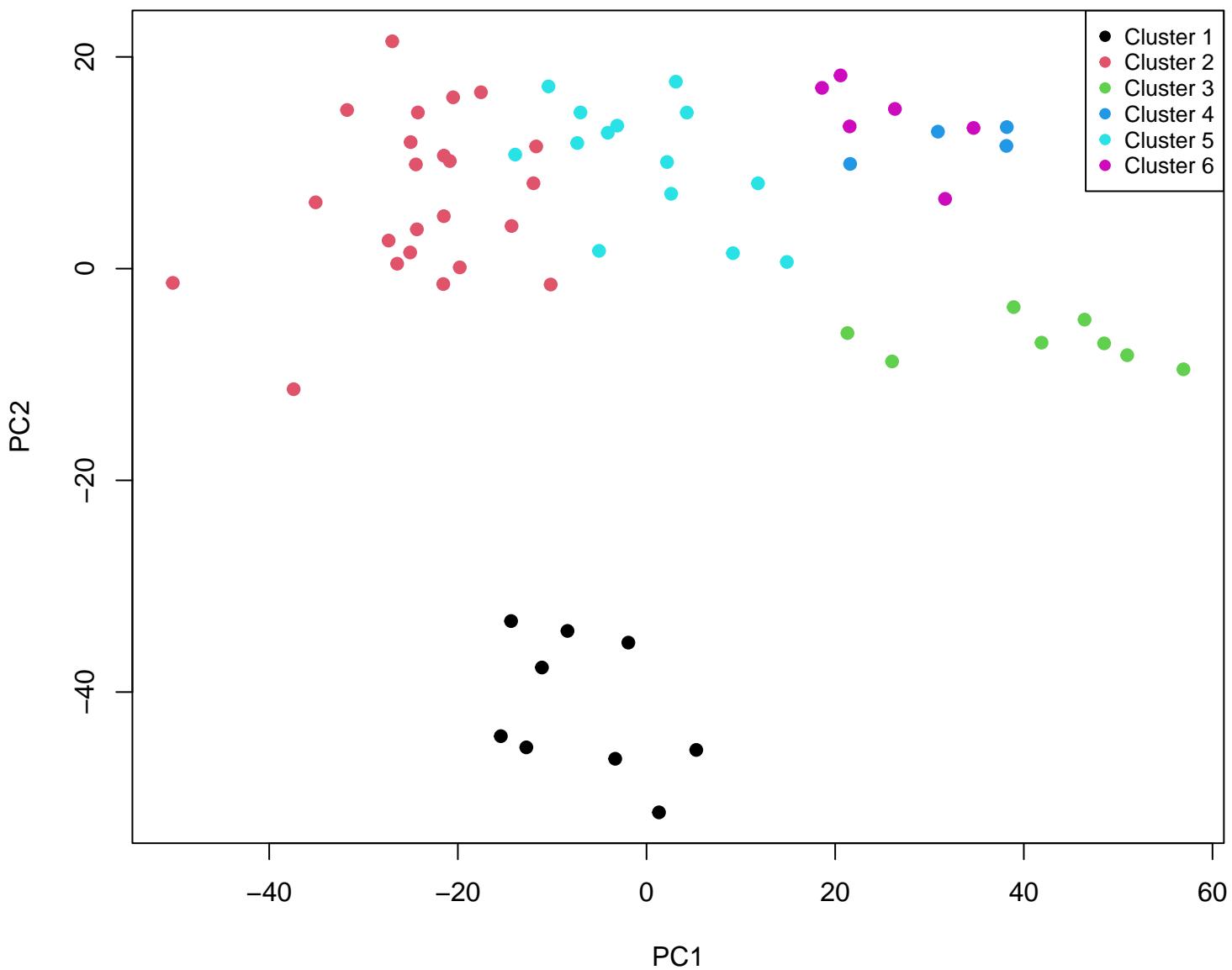
Elbow plot for k-means clustering



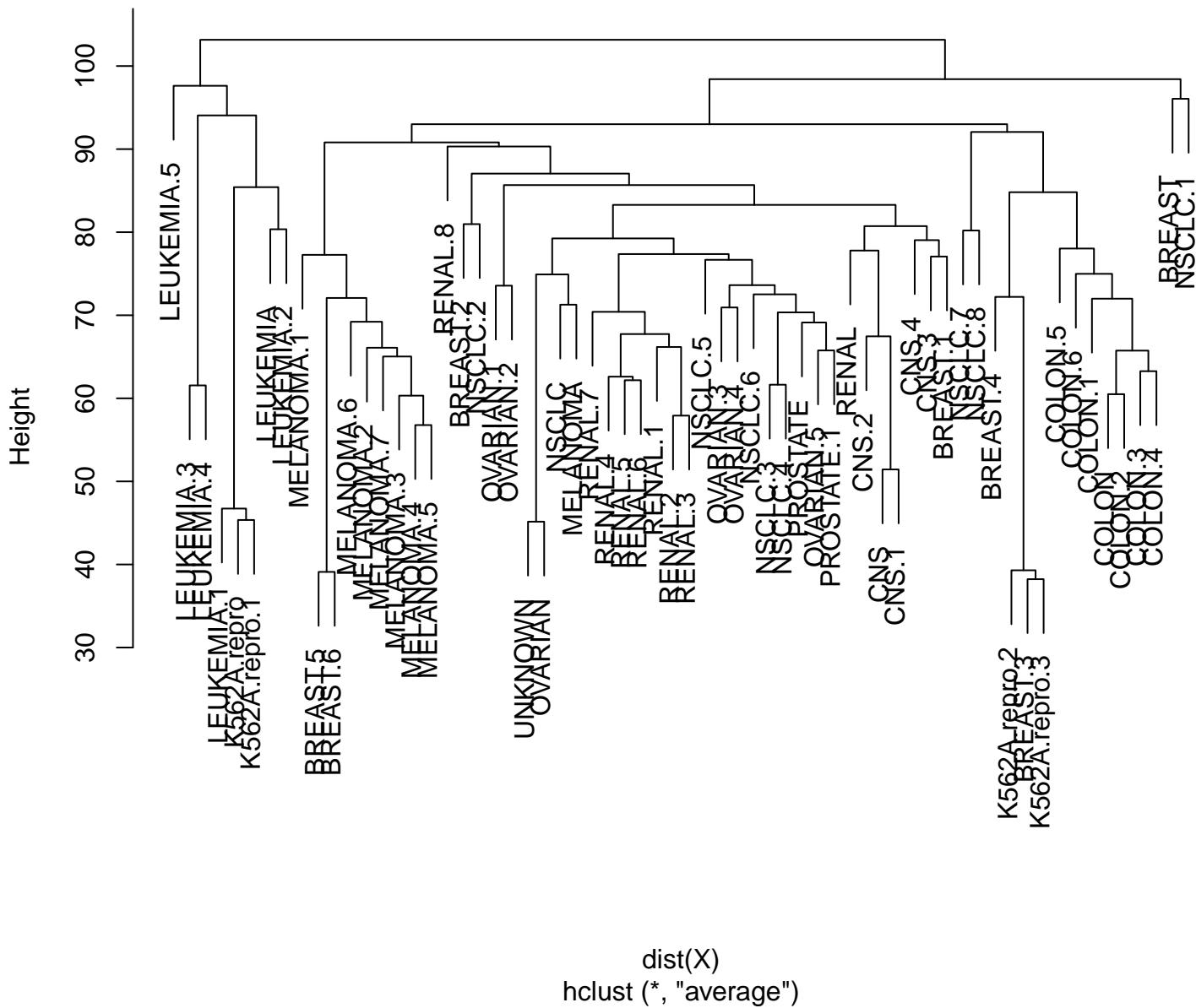
k-means clusters with 6 clusters (using 2 randomly sampled genes)



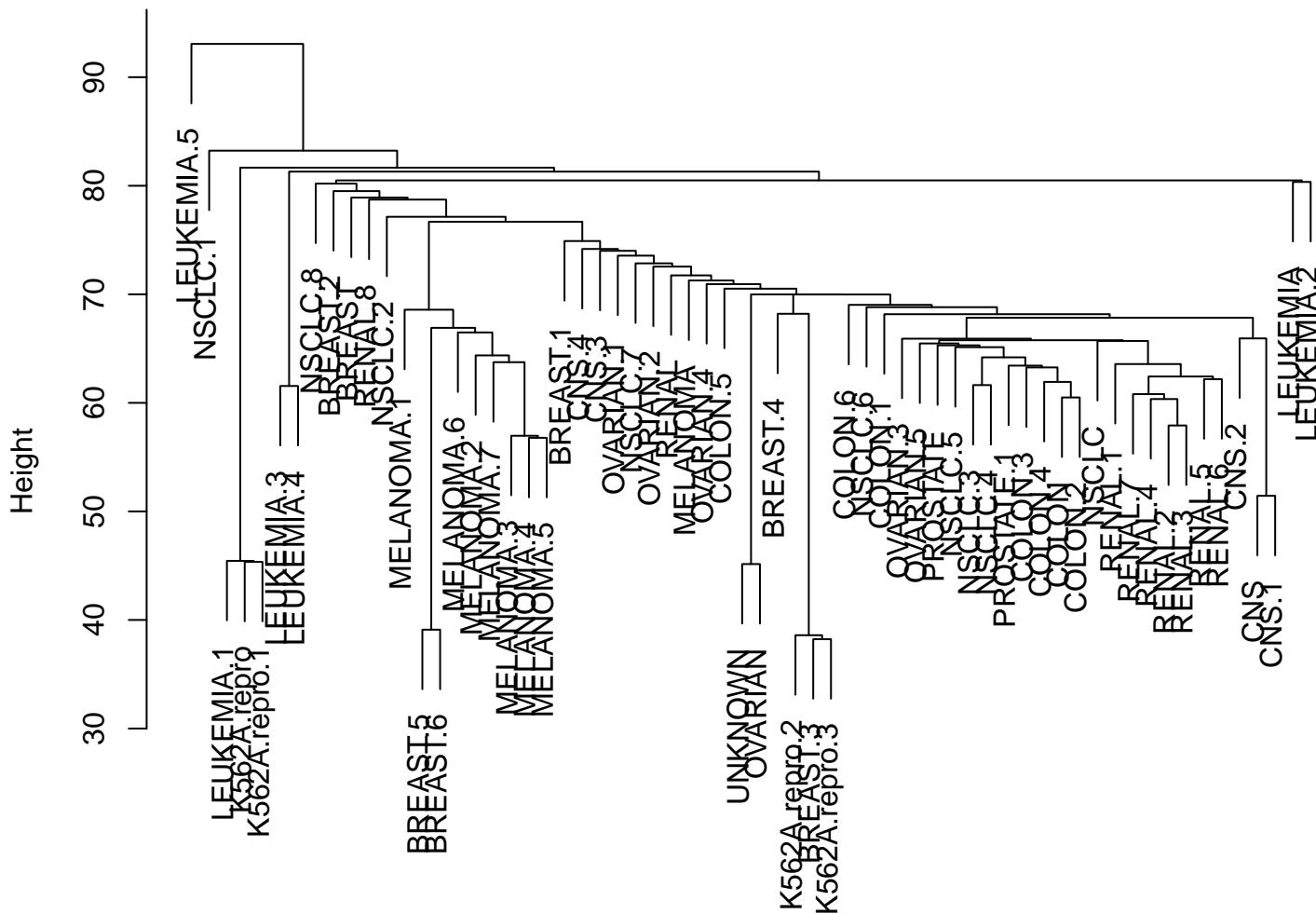
K-means clusters (first 2 PCs, raw clustering)



Average



Single



$\text{dist}(X)$
hclust (*, "single")

Complete

