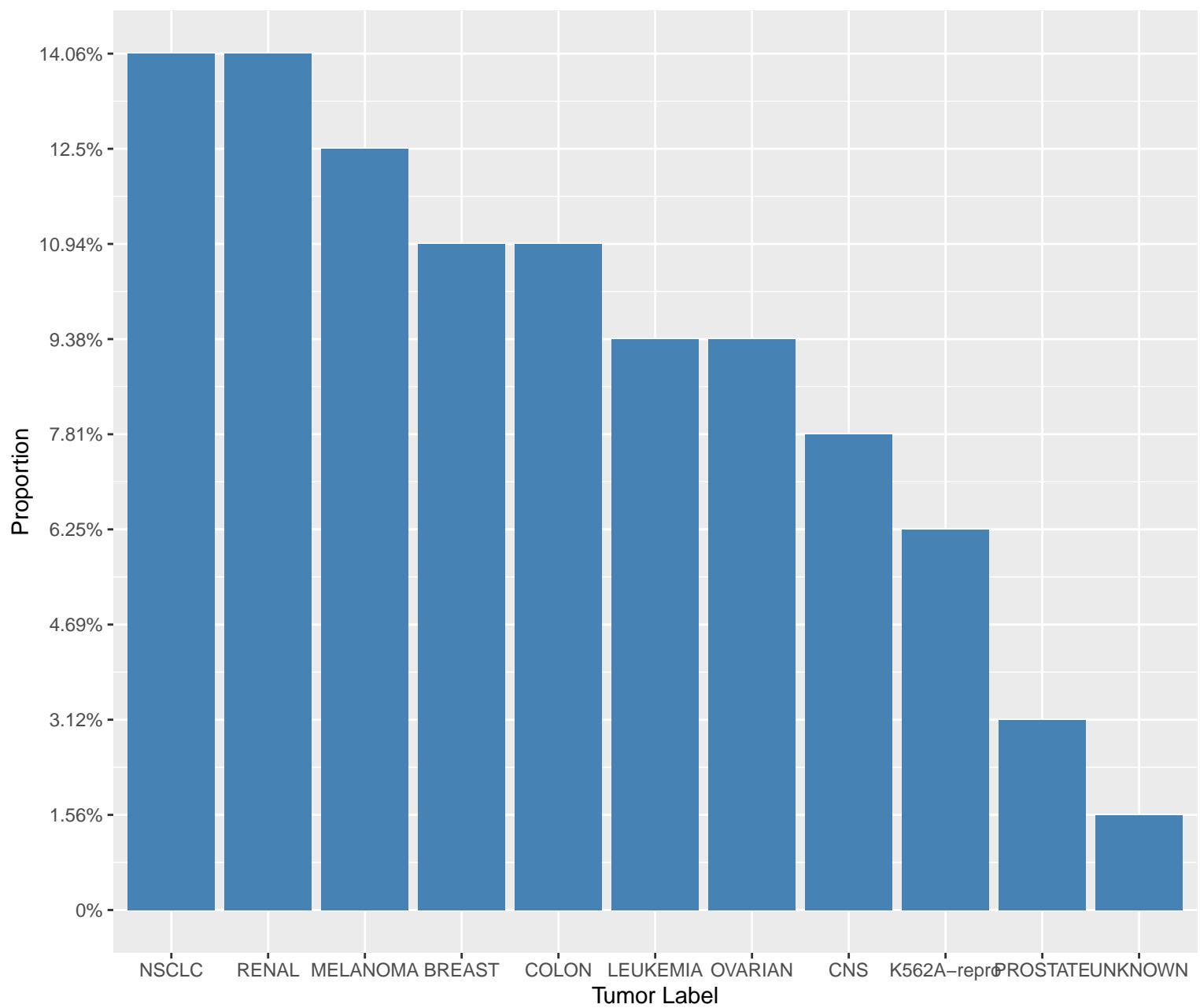
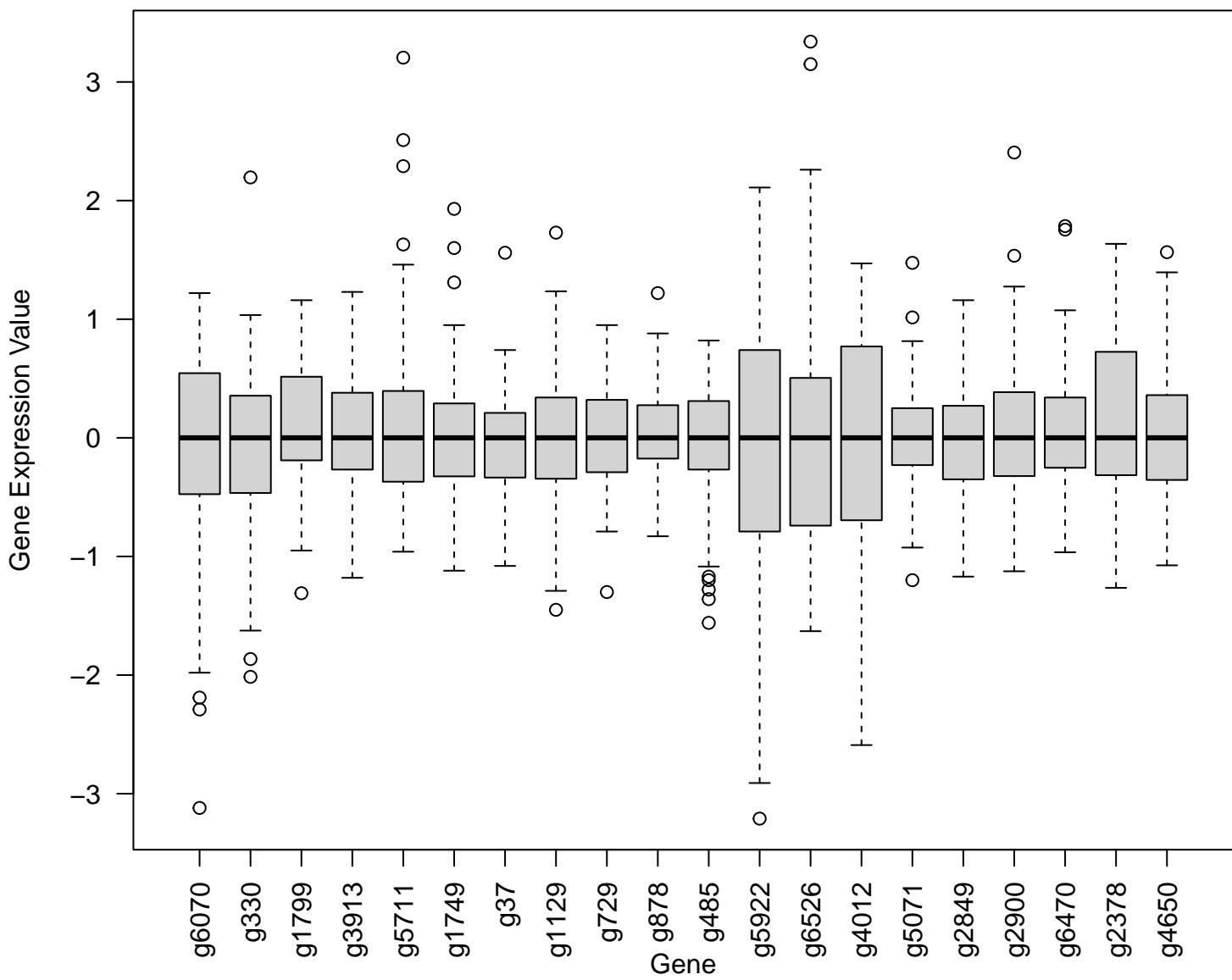


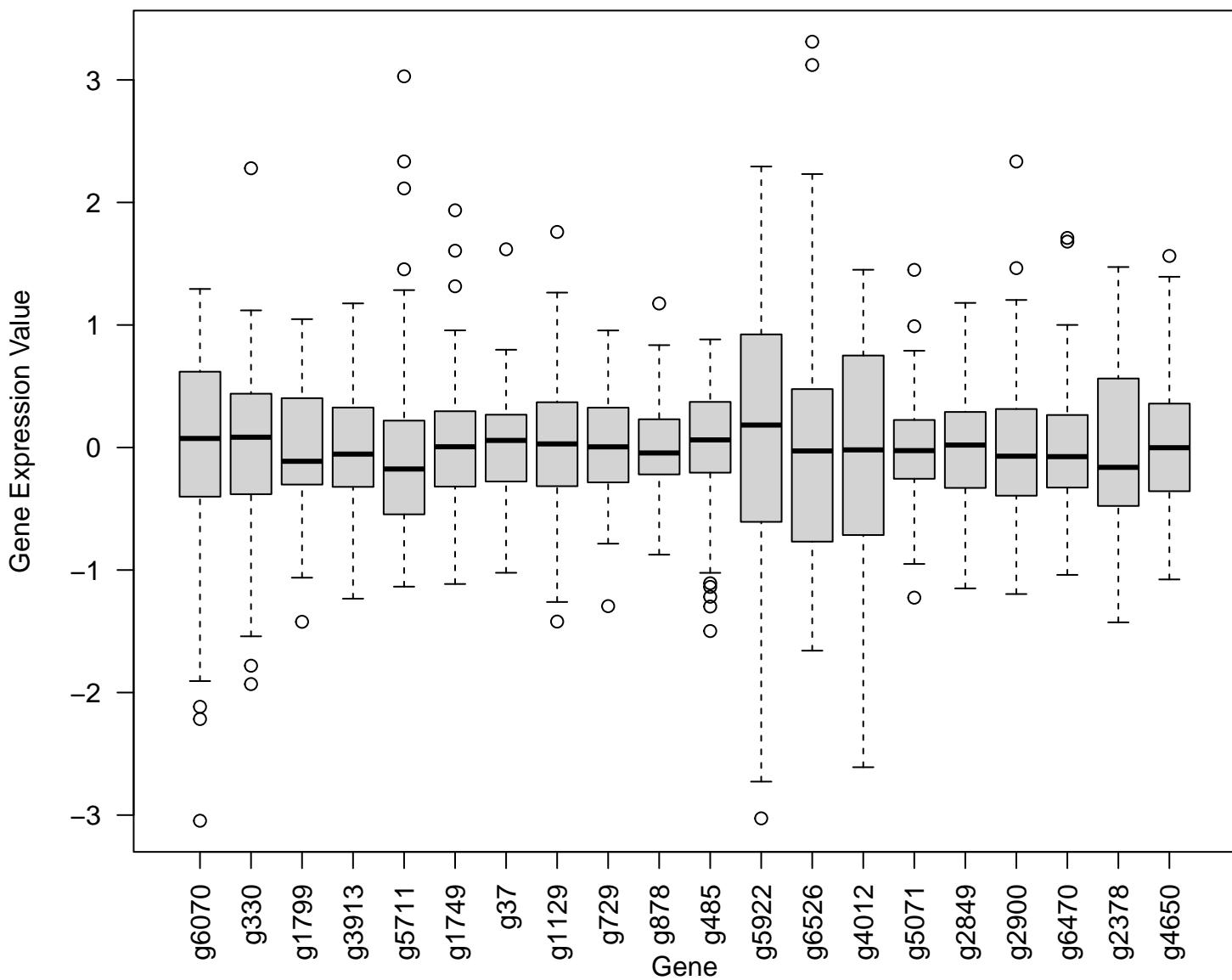
Tumor Label Distribution



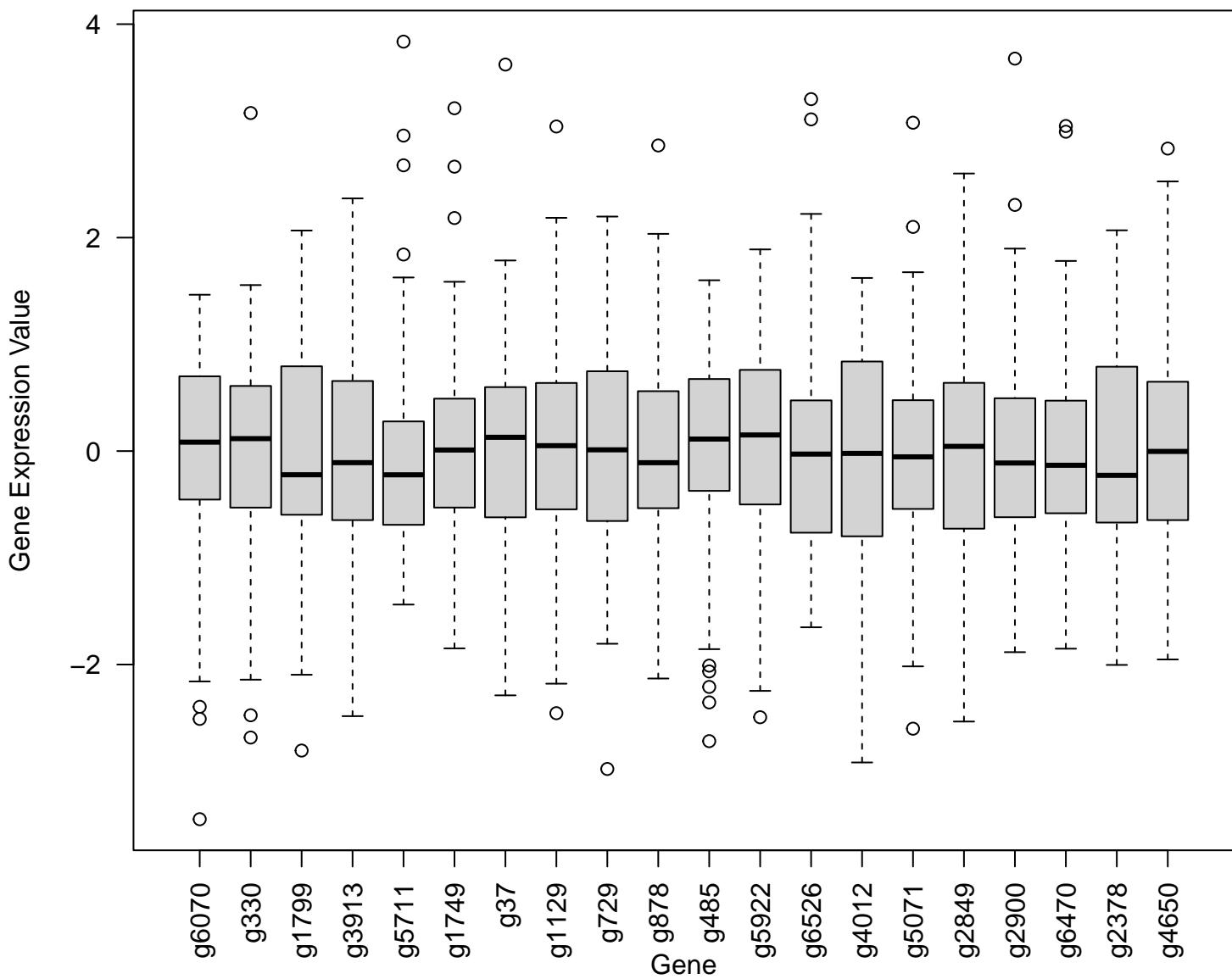
Subset Gene Expression marginalised over patients

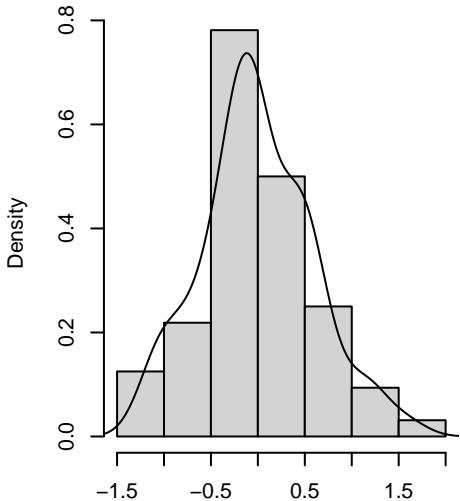
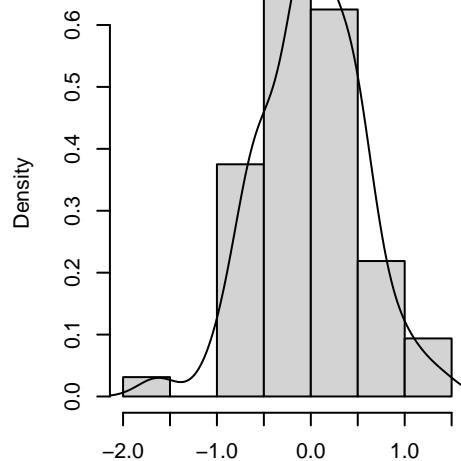
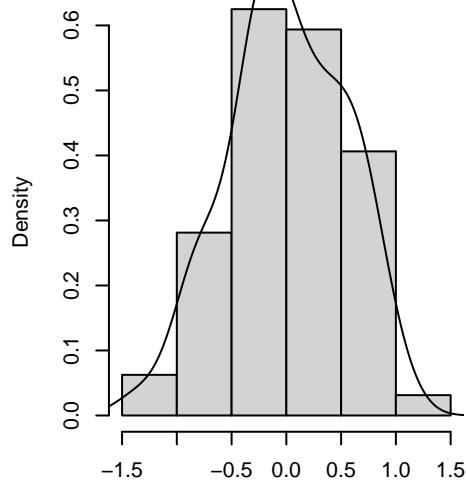
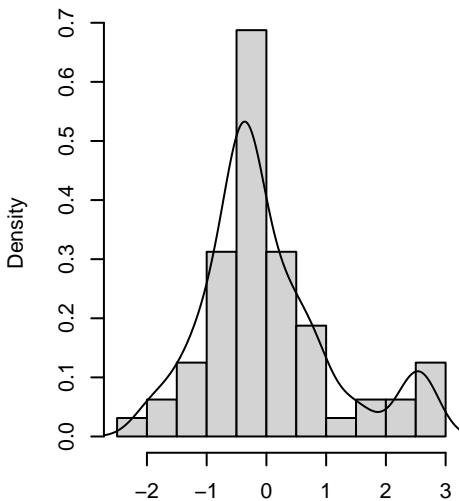
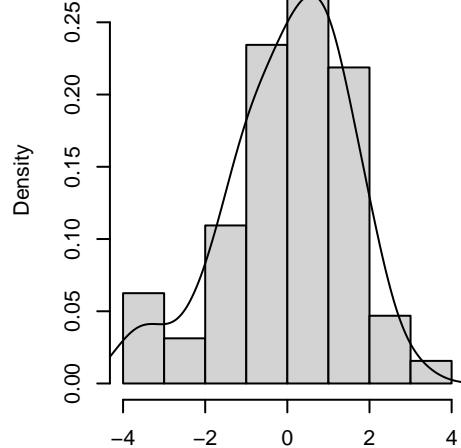
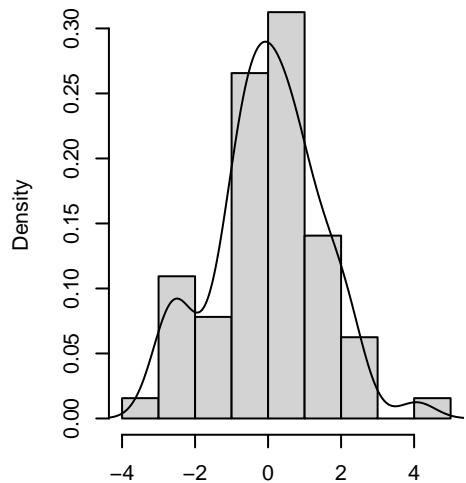


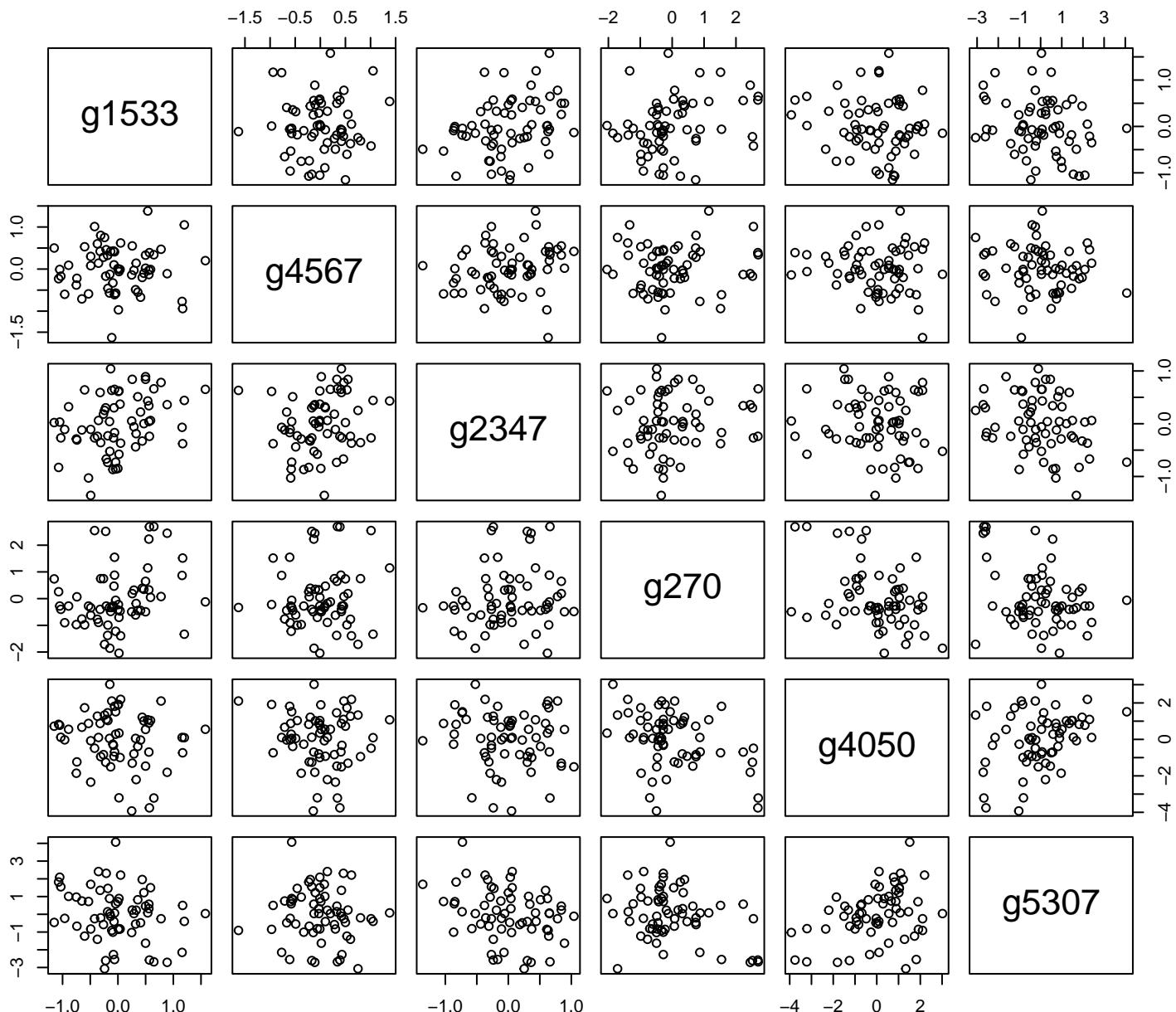
Subset Gene Expression marginalised over patients (centered)

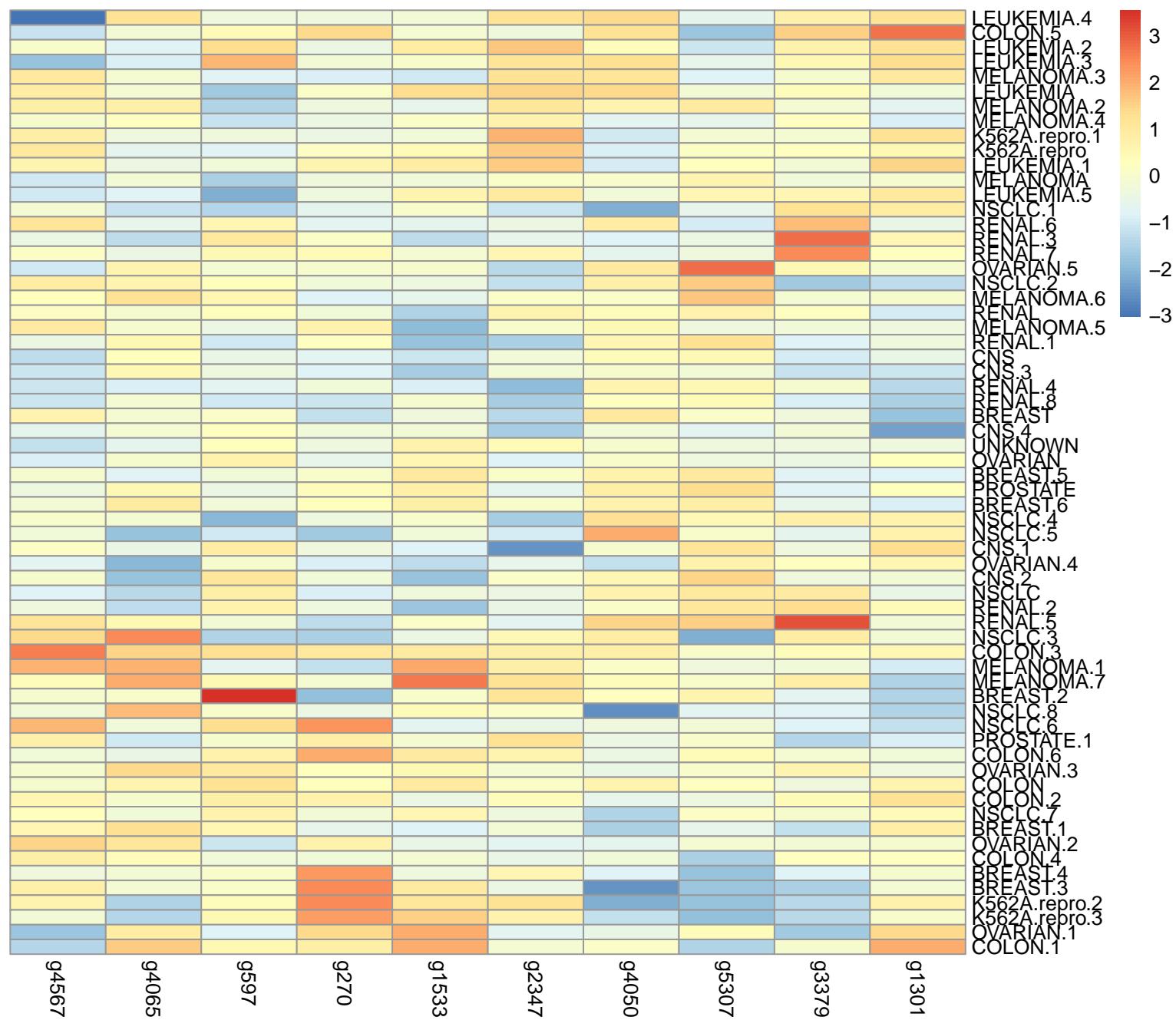


Subset Gene Expression marginalised over patients (centered and scaled)

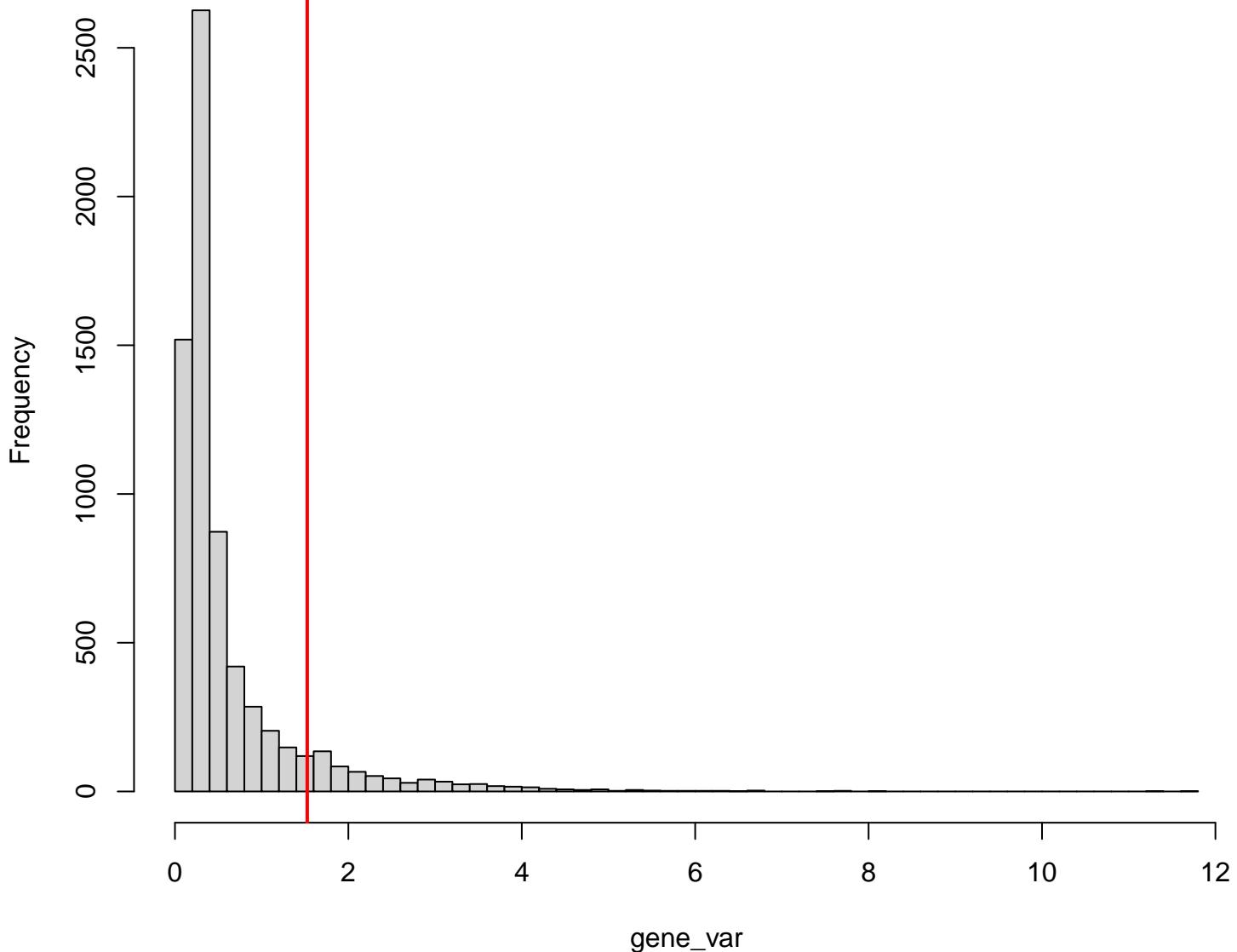


g1533 Distribution**g4567 Distribution****g2347 Distribution****g270 Distribution****g4050 Distribution****g5307 Distribution**

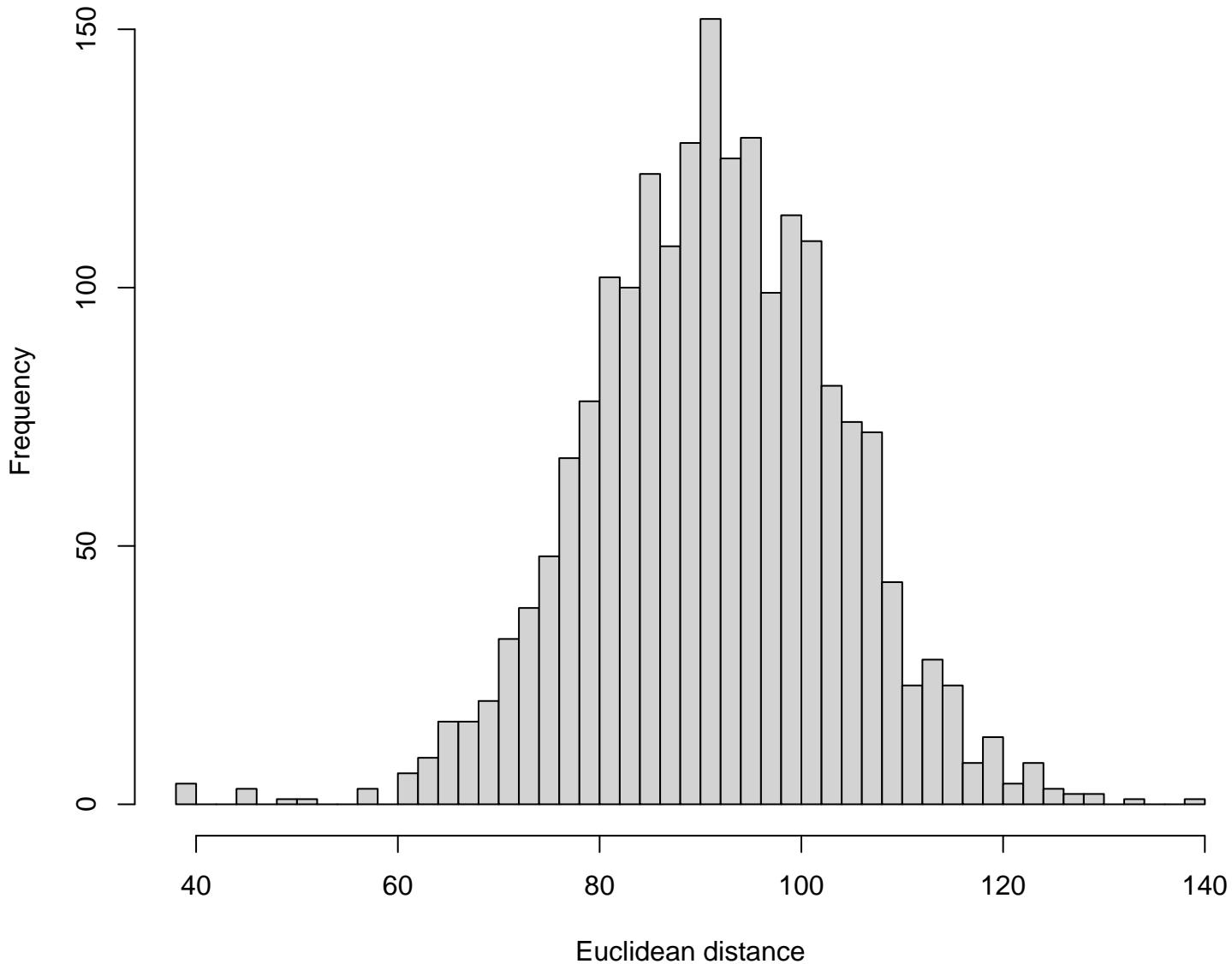




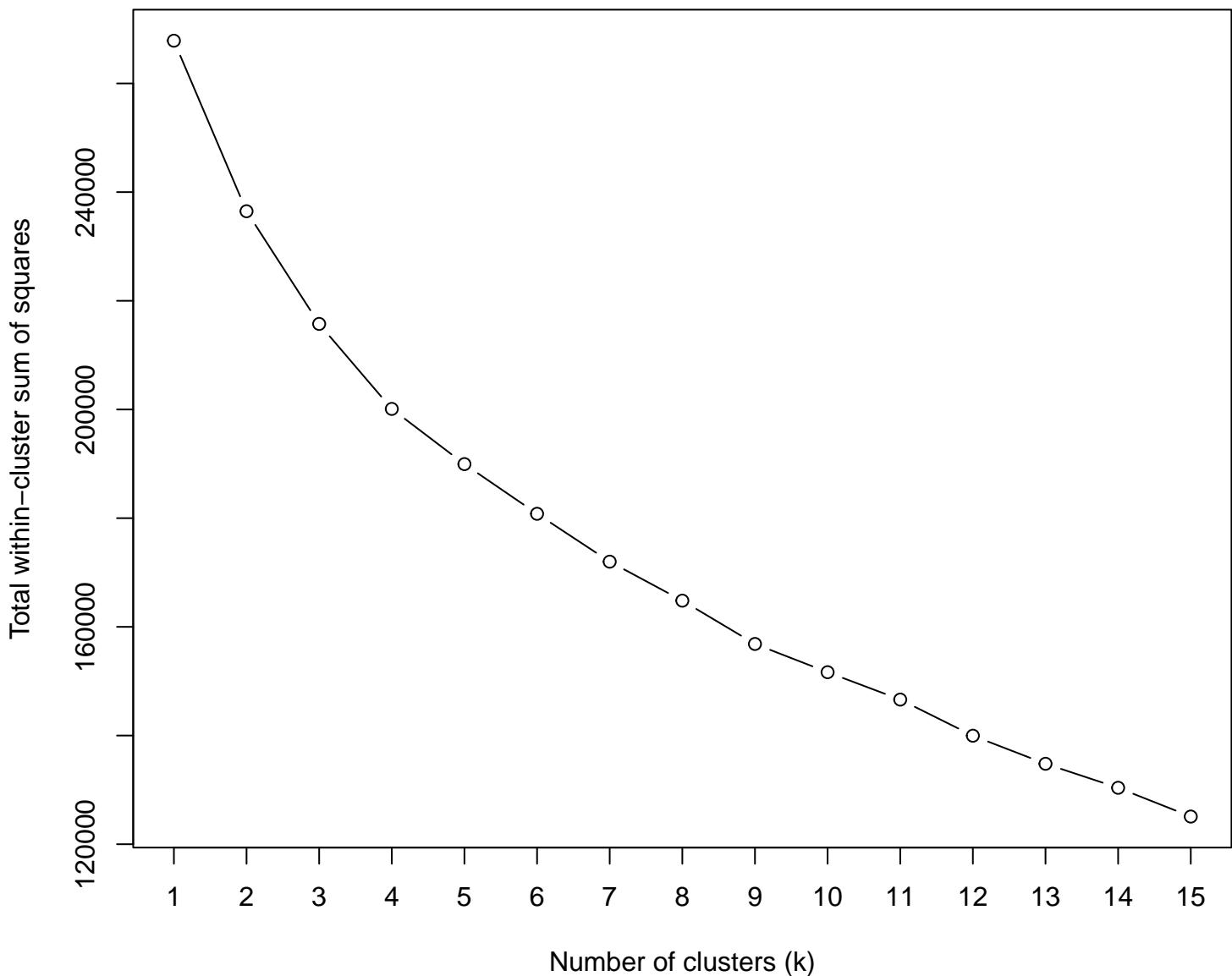
Gene-wise variance distribution



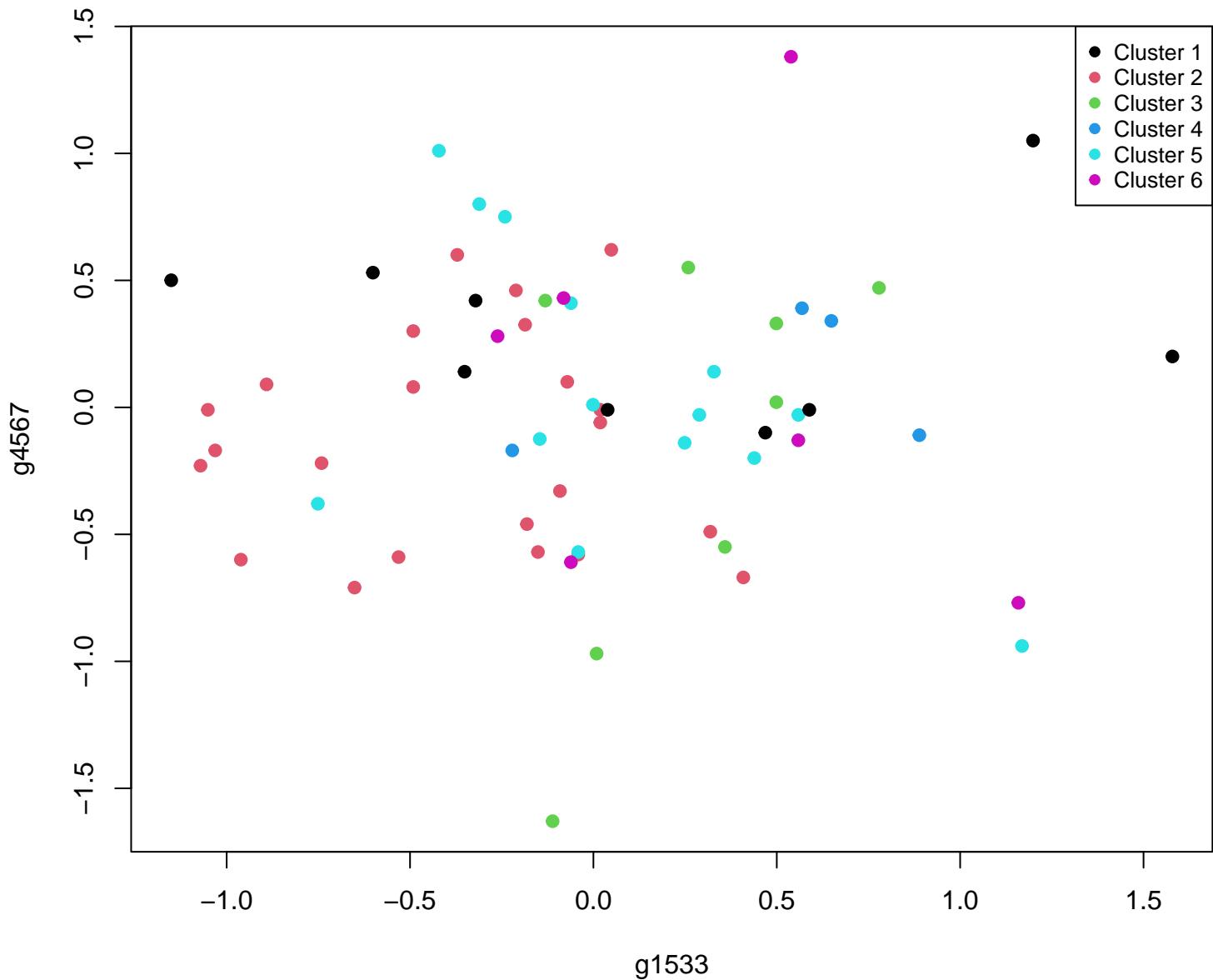
Pairwise distances between samples



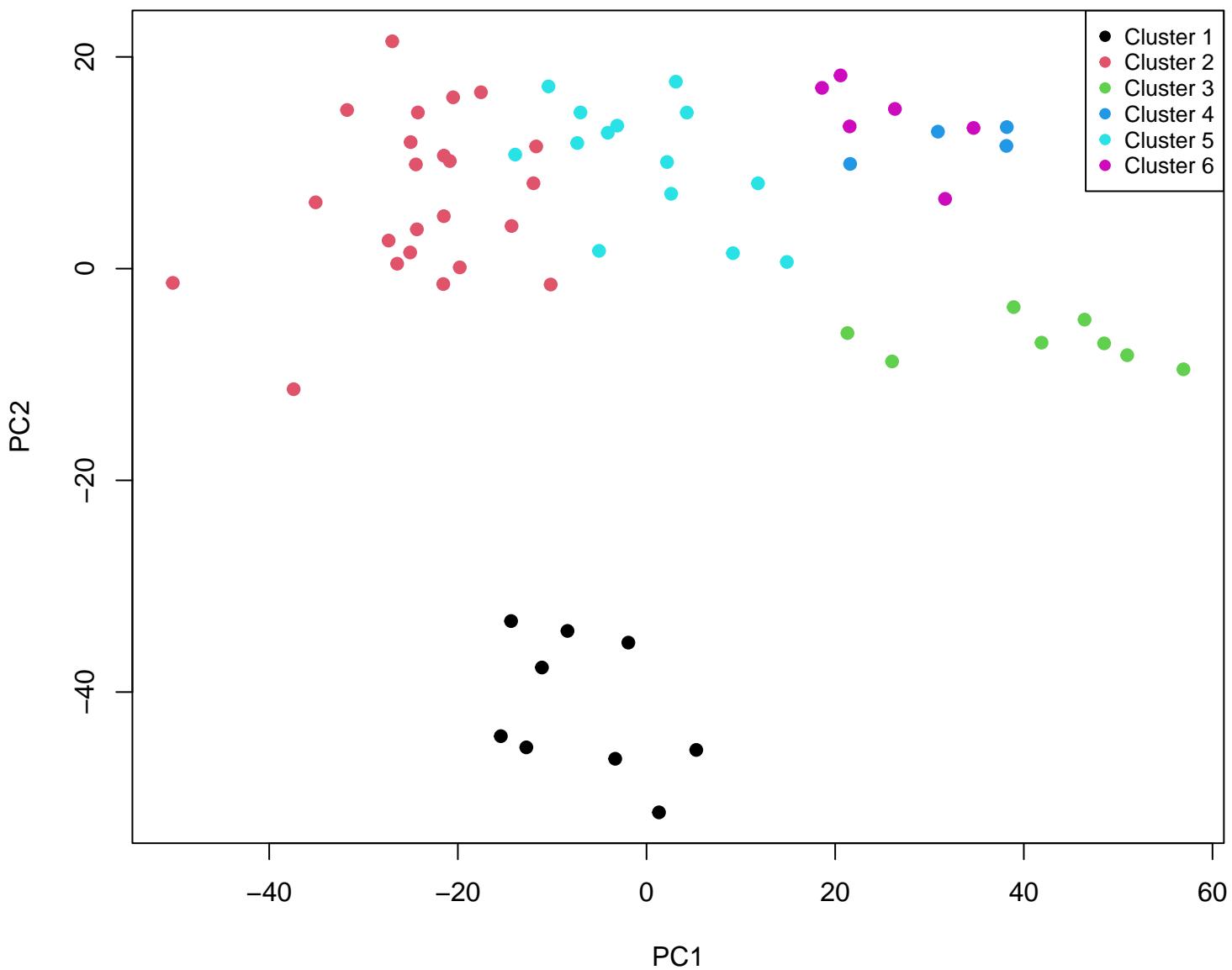
Elbow plot for k-means clustering



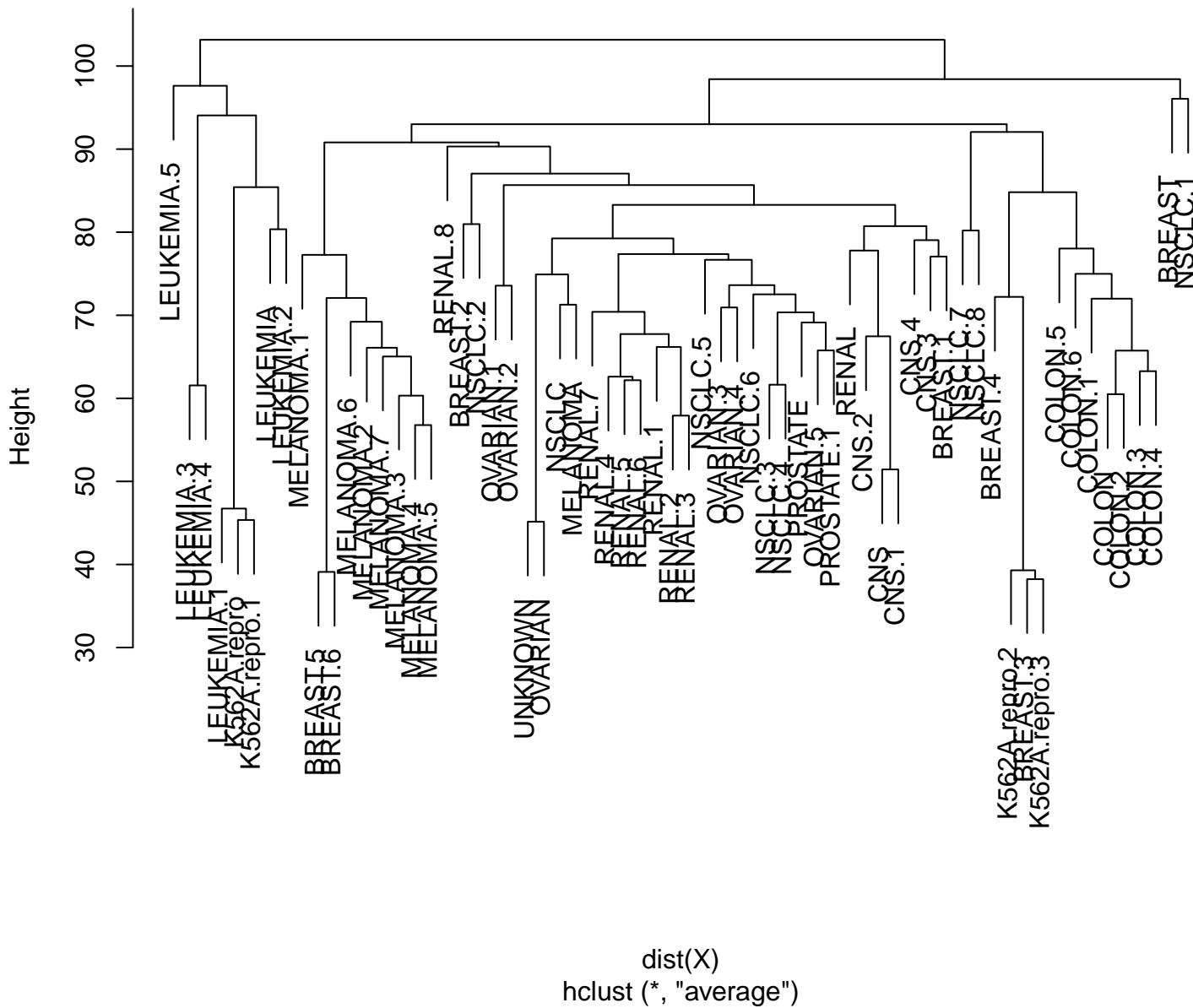
k-means clusters with 6 clusters (using 2 randomly sampled genes)



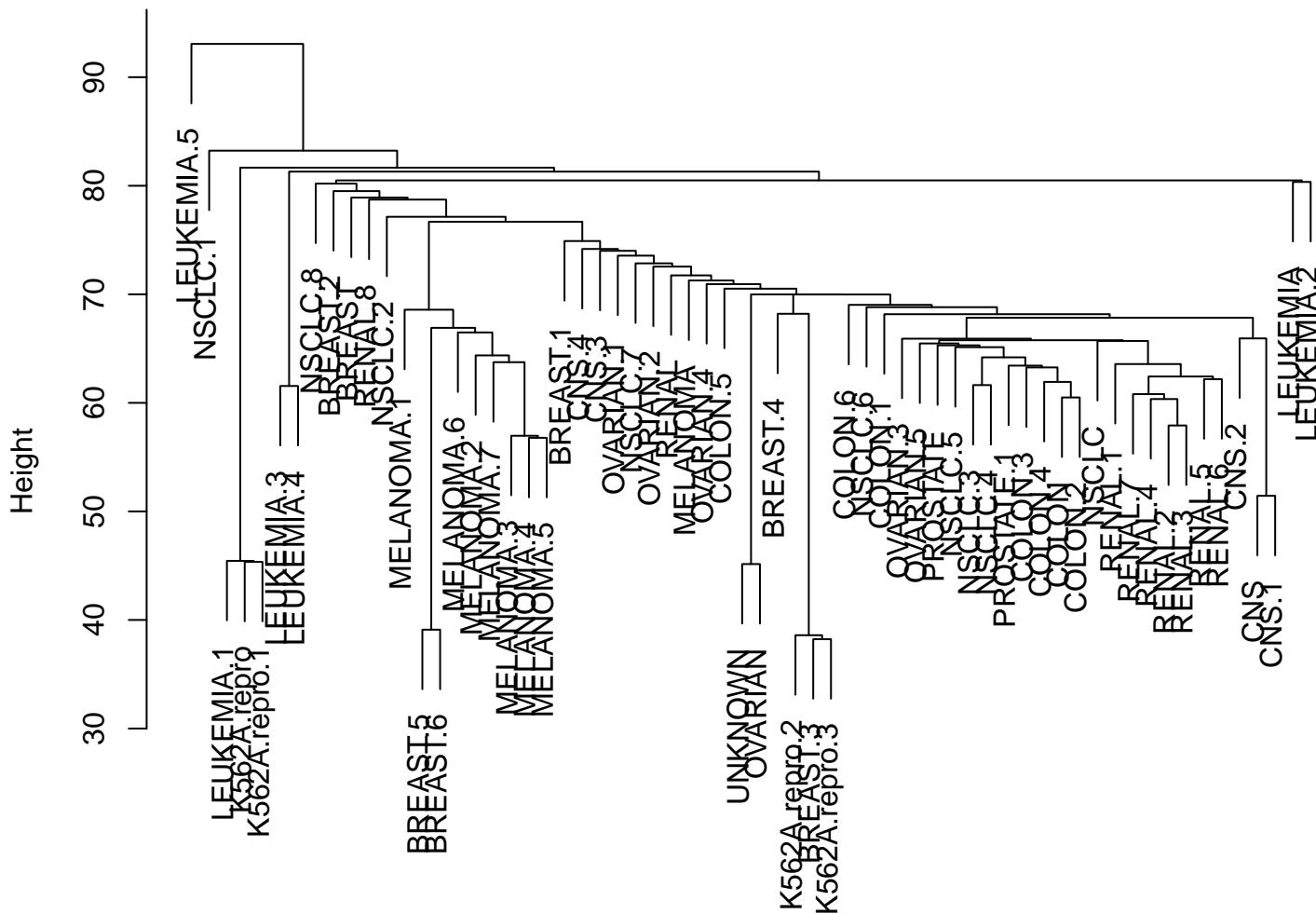
K-means clusters (first 2 PCs, raw clustering)



Average



Single



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

Complete

