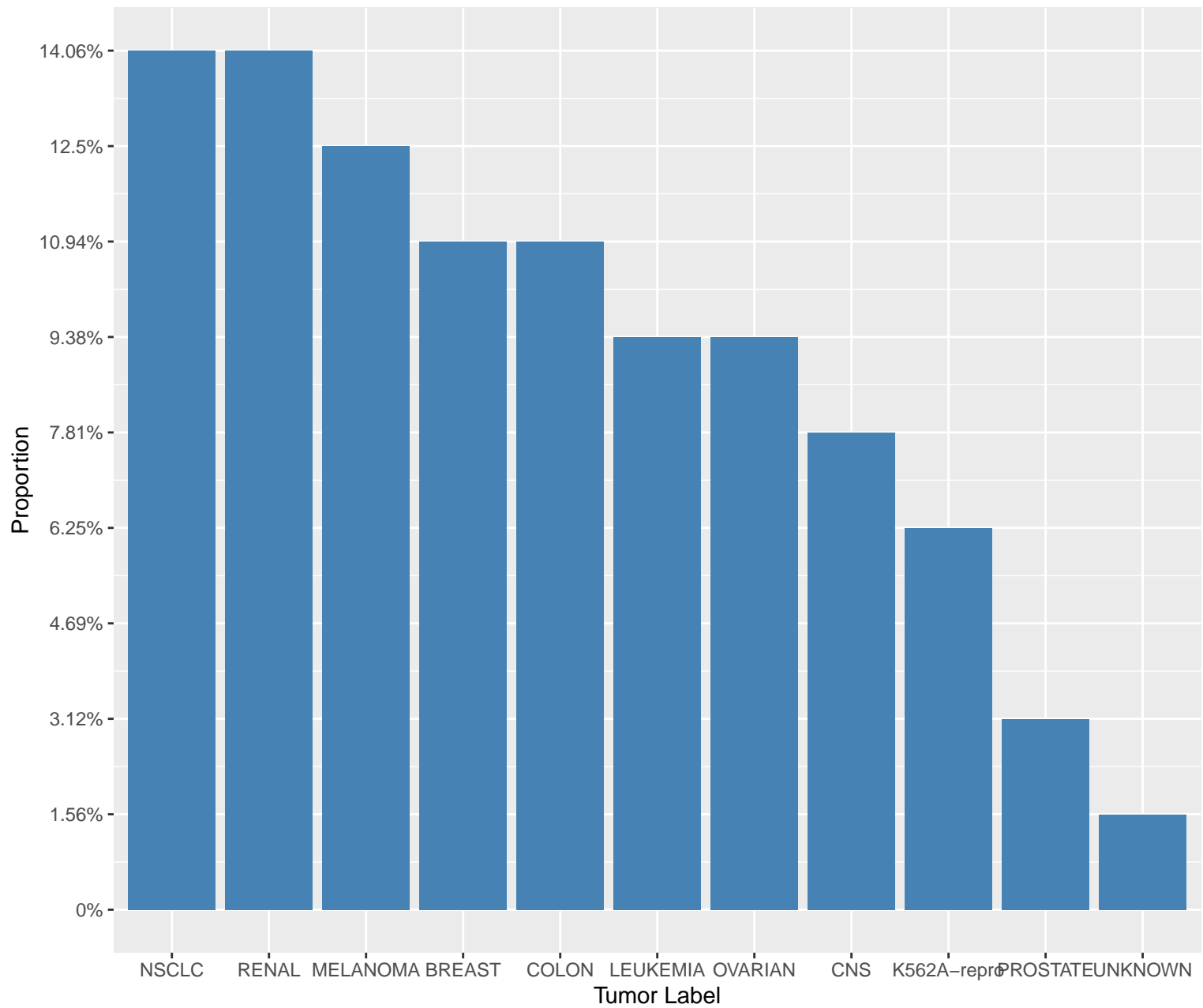
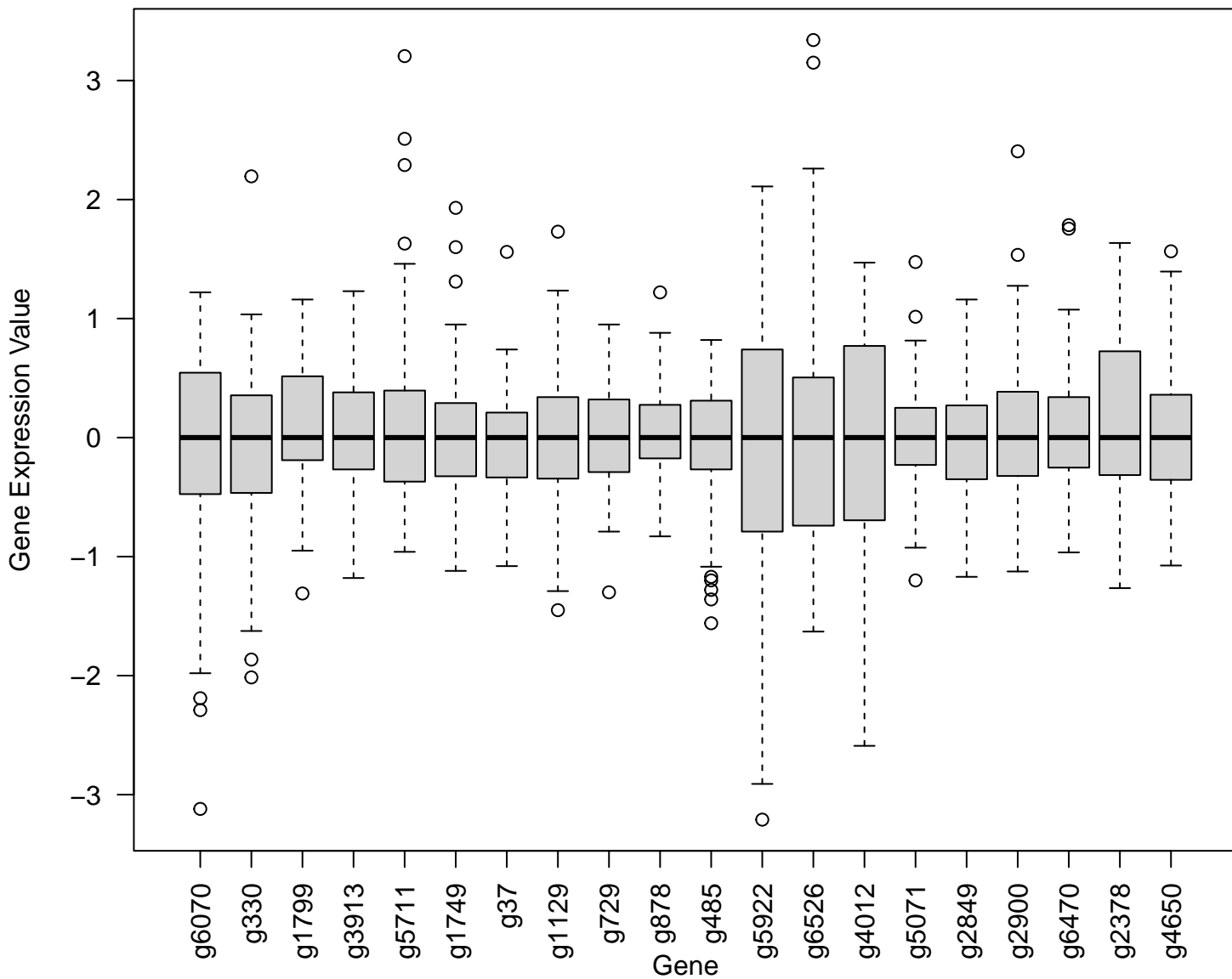


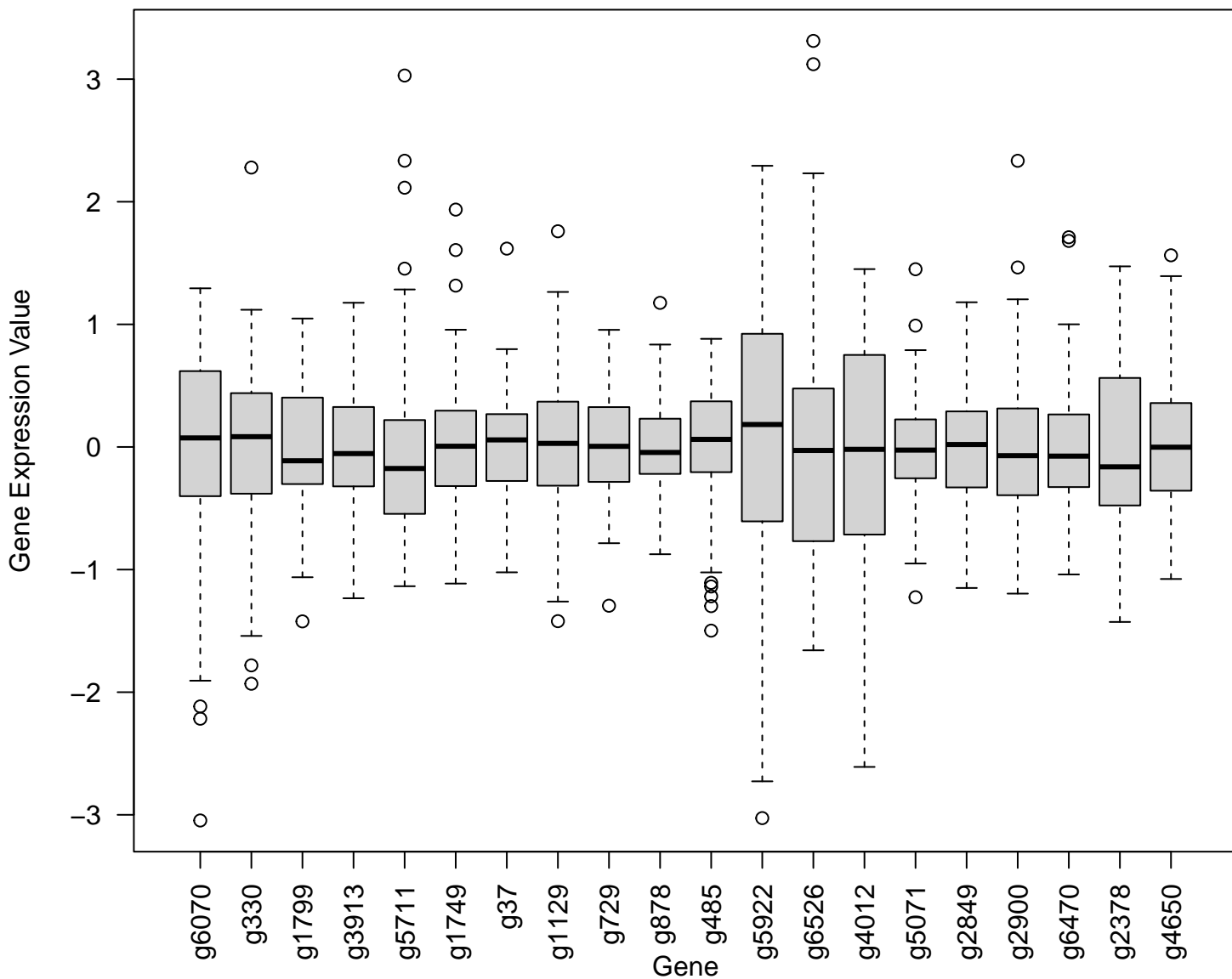
Tumor Label Distribution



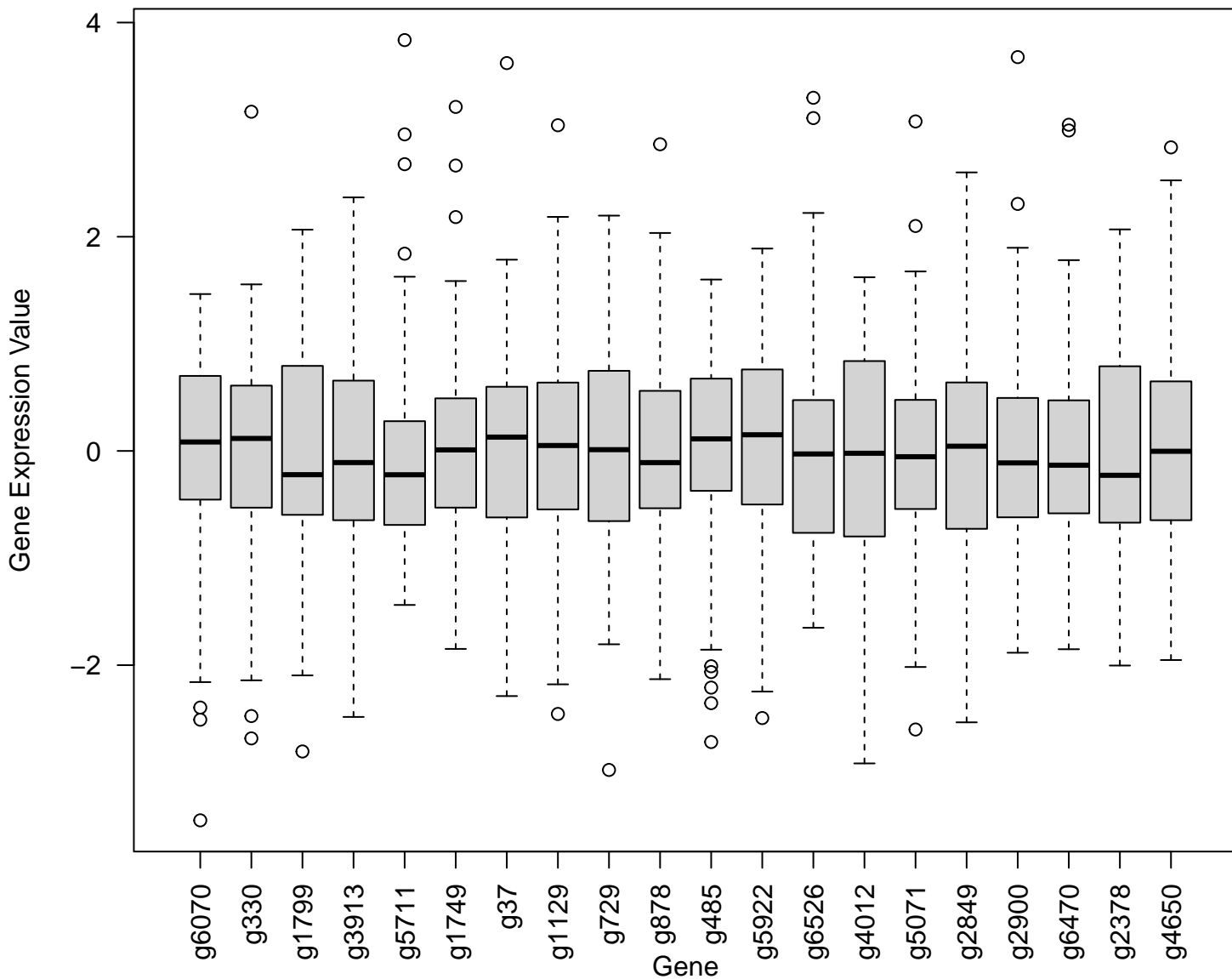
Subset Gene Expression marginalised over patients



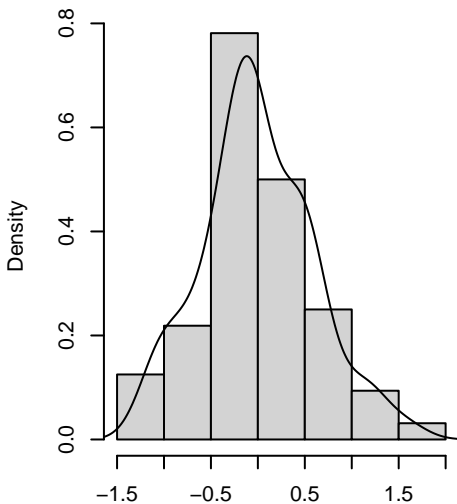
Subset Gene Expression marginalised over patients (centered)



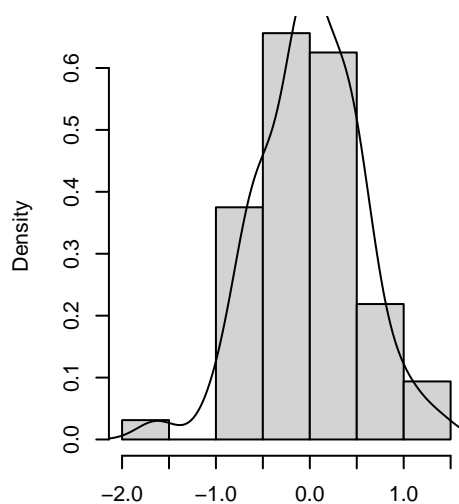
Subset Gene Expression marginalised over patients (centered and scaled)



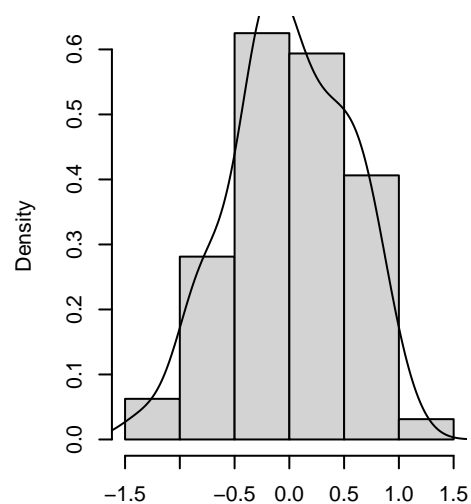
g1533 Distribution



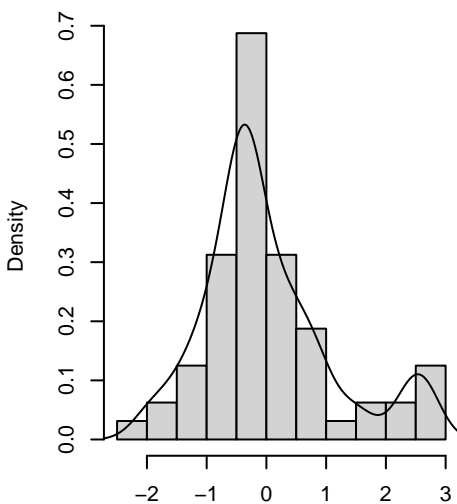
g4567 Distribution



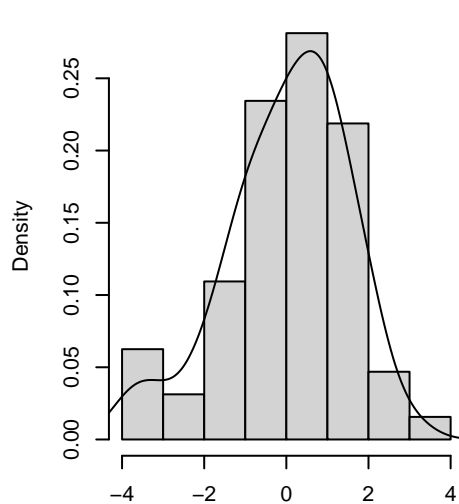
g2347 Distribution



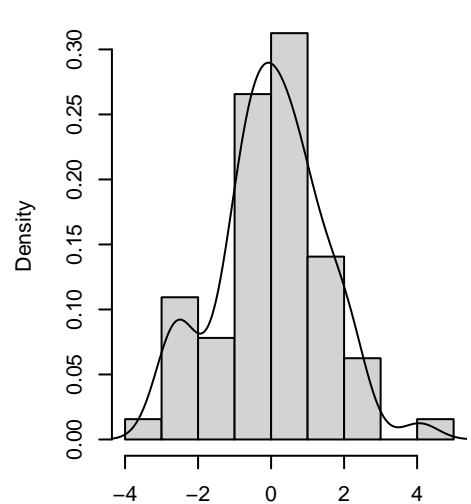
g270 Distribution

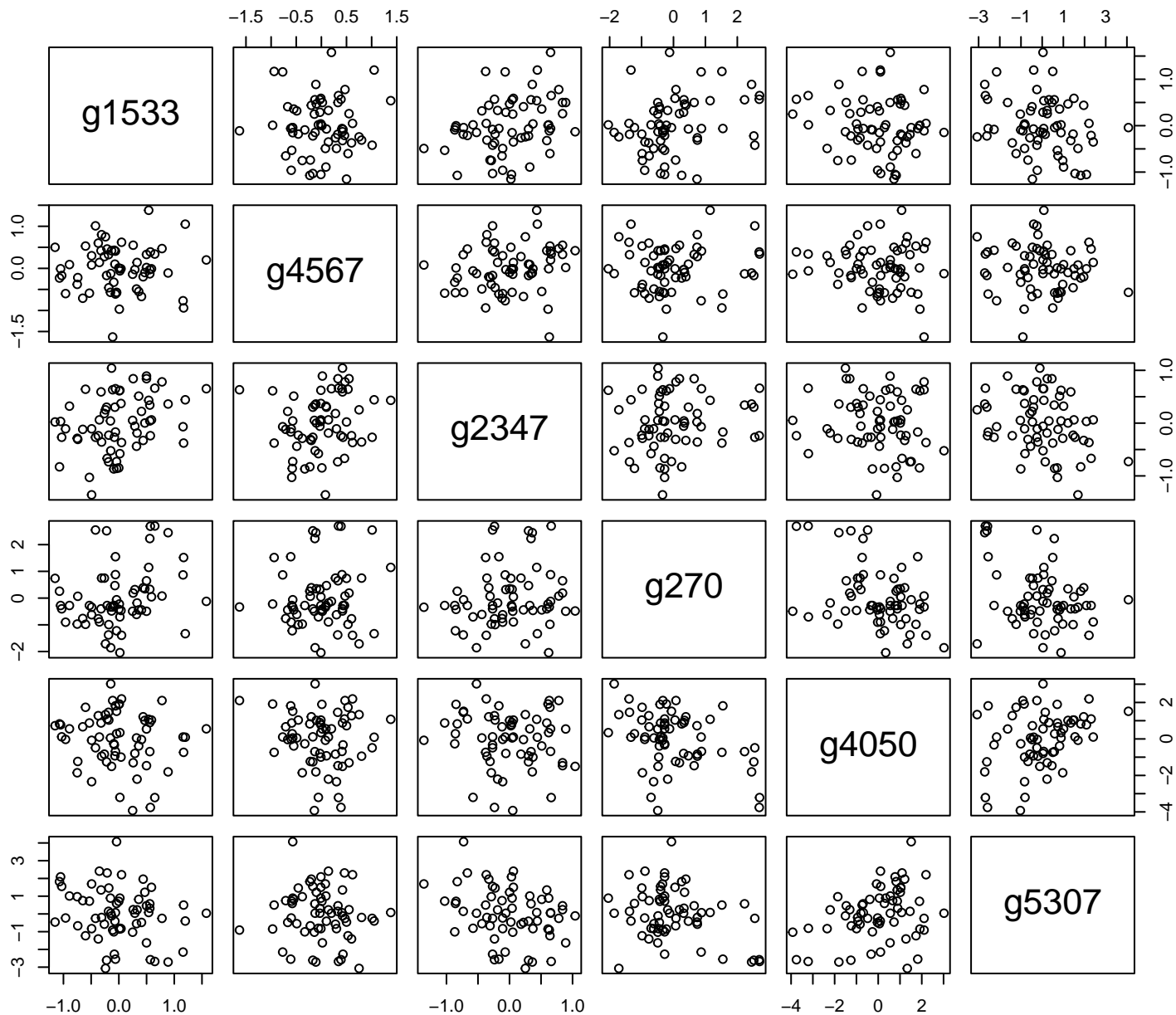


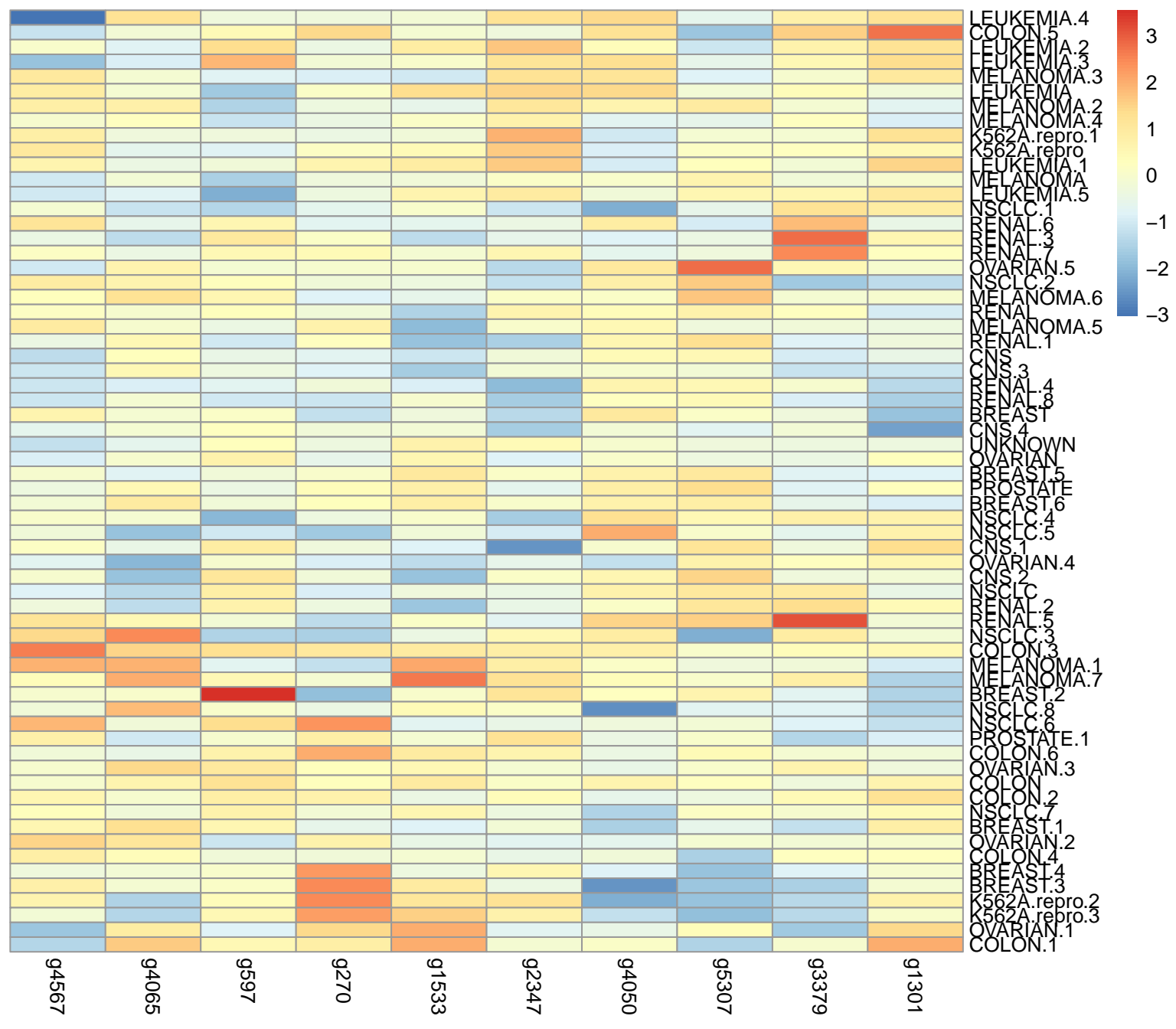
g4050 Distribution



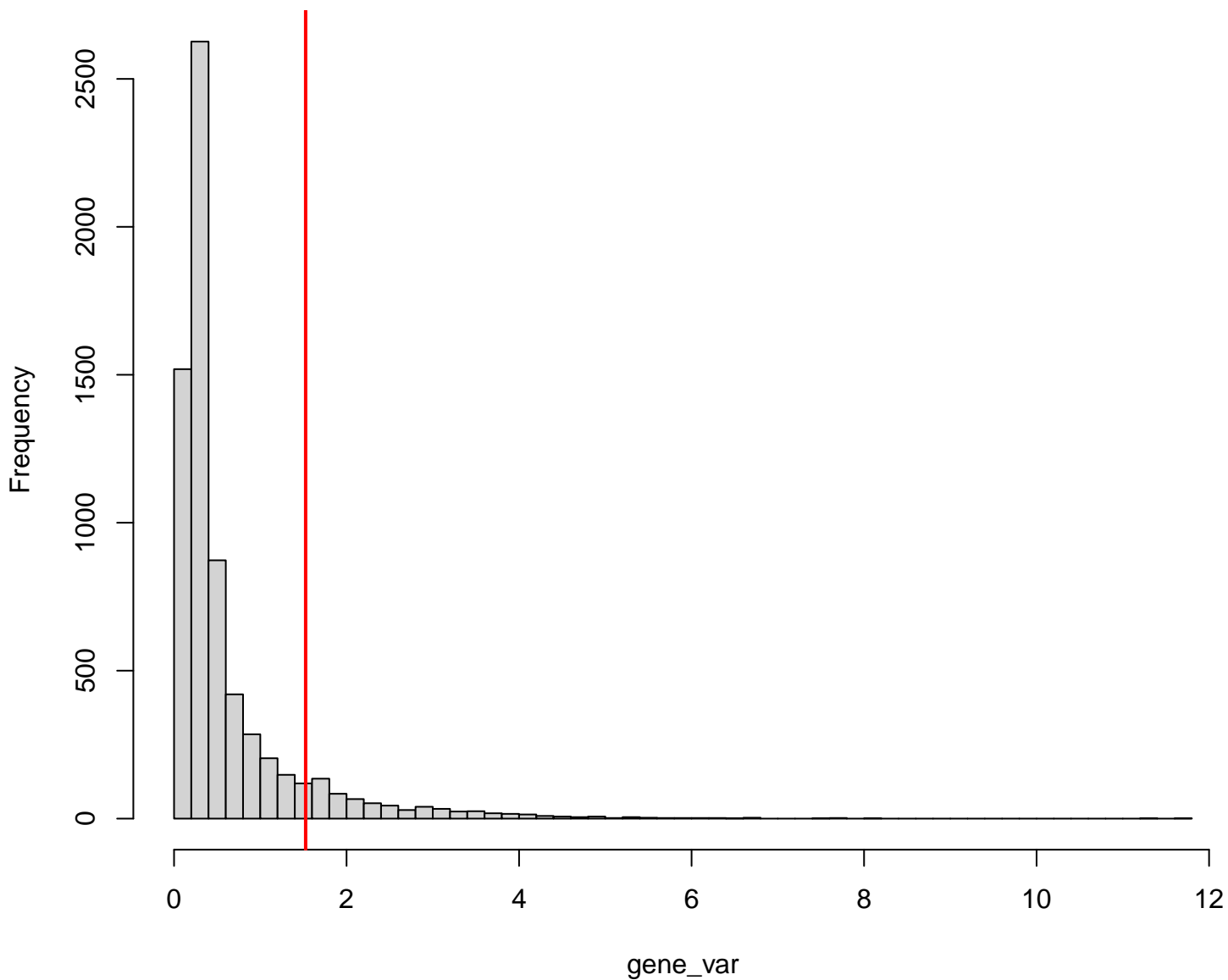
g5307 Distribution



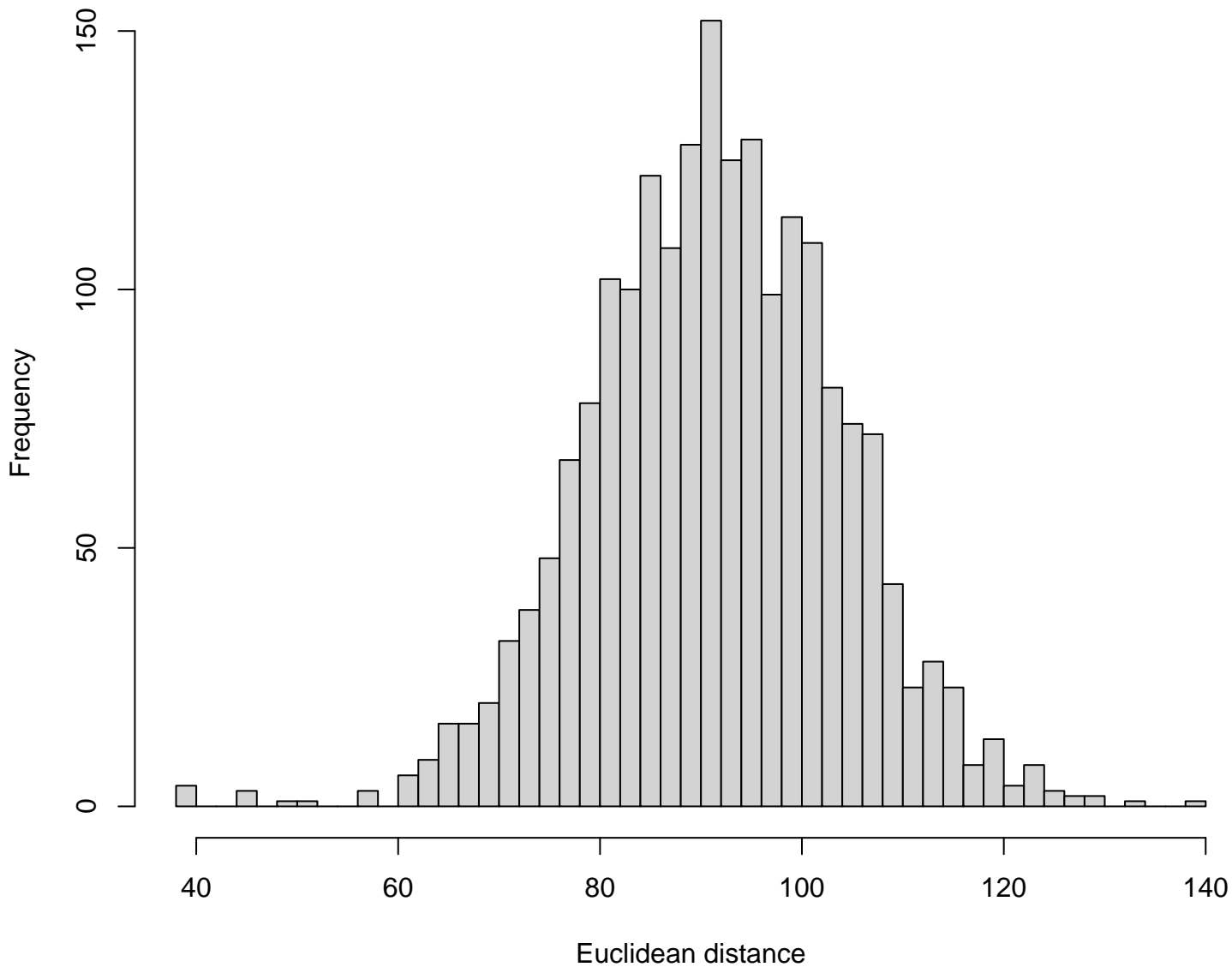




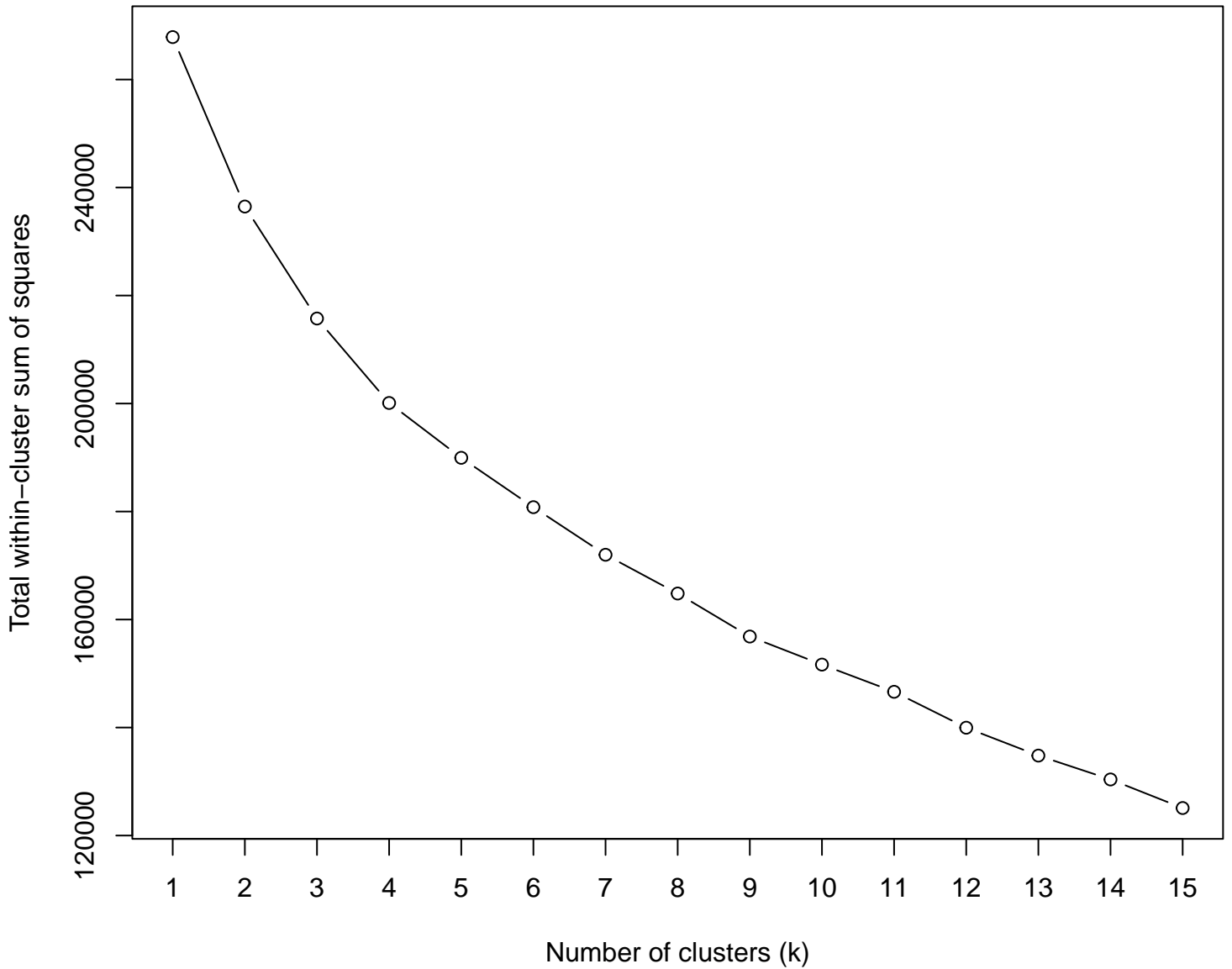
Gene-wise variance distribution



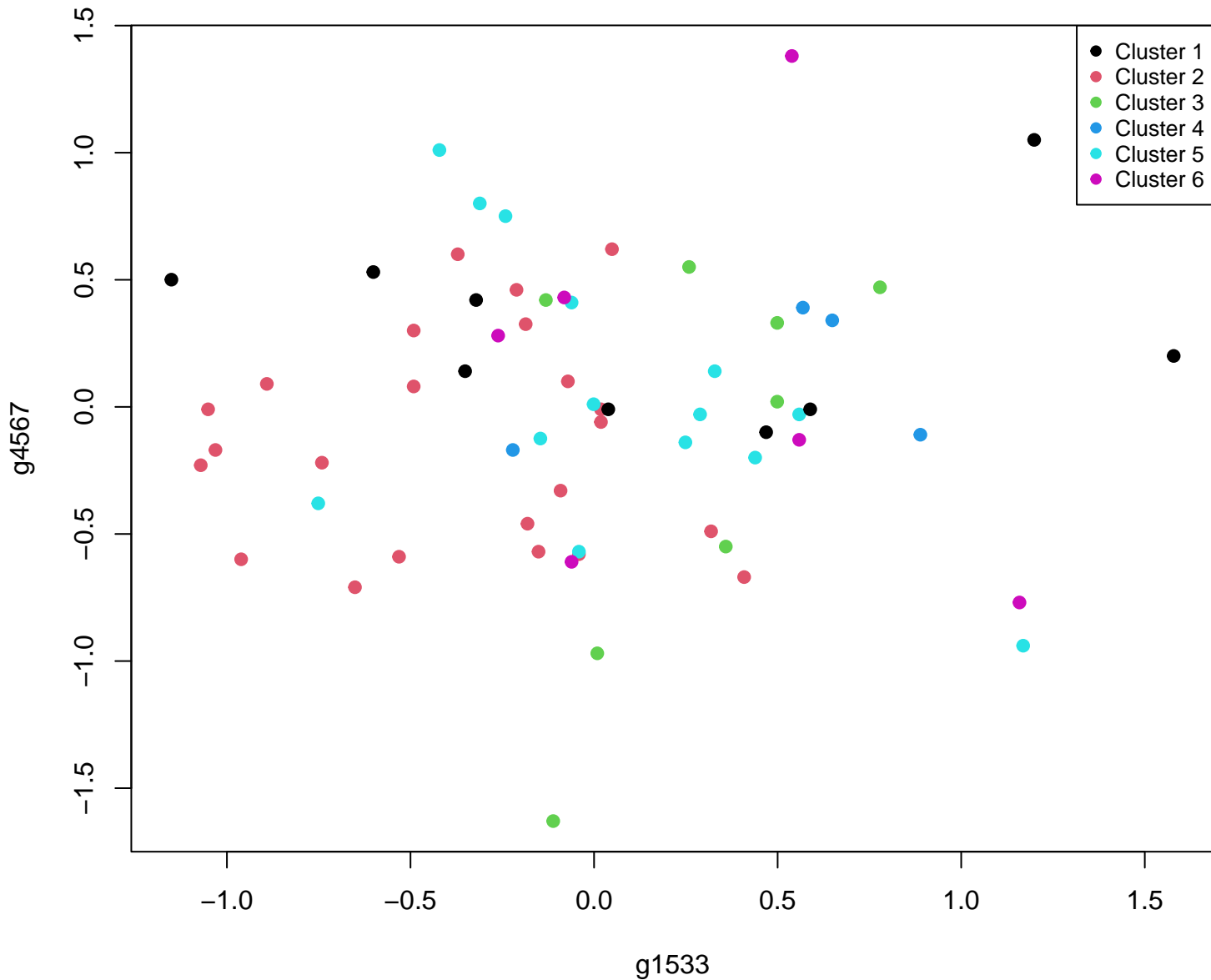
Pairwise distances between samples



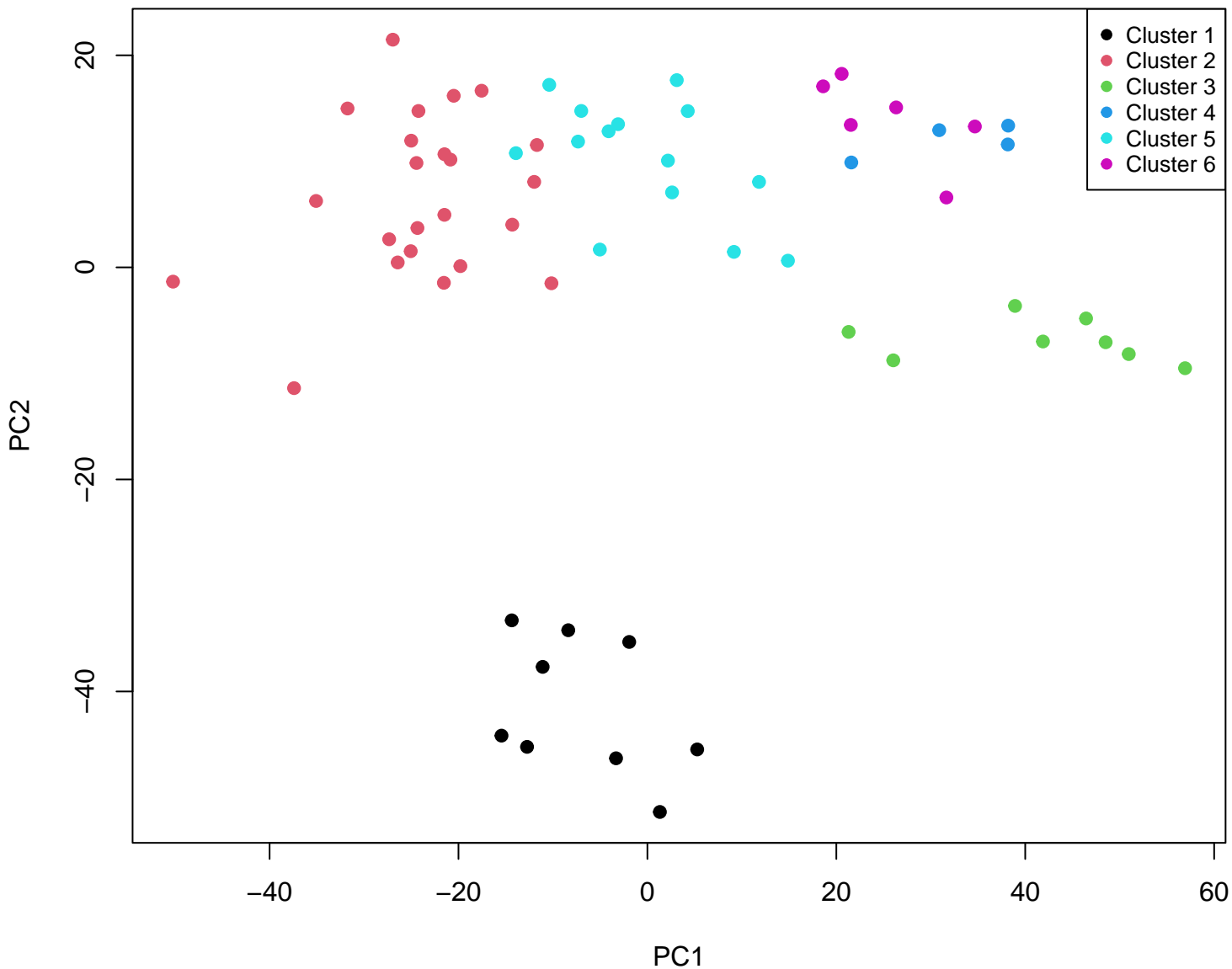
Elbow plot for k-means clustering



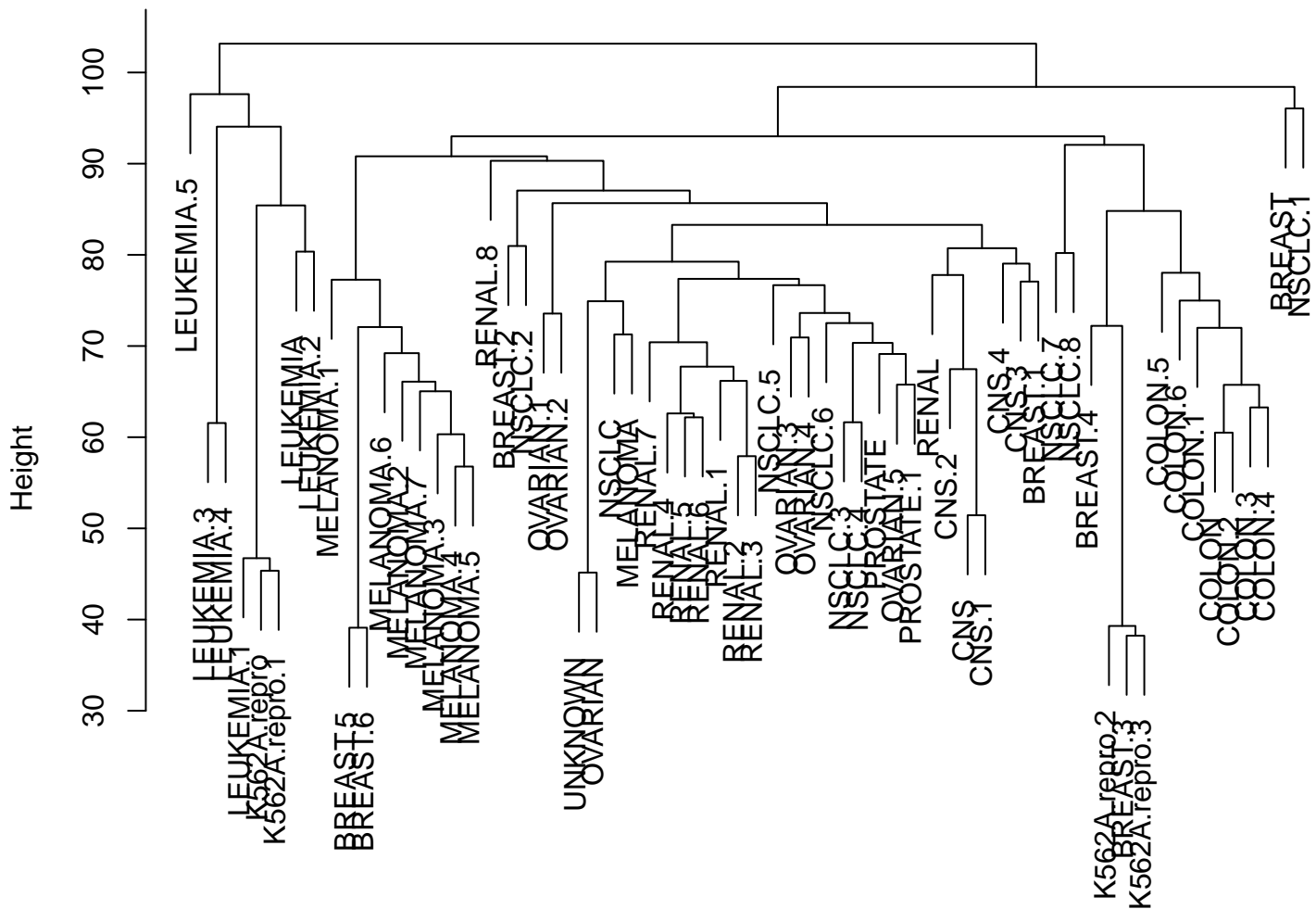
k-means clusters with 6 clusters (using 2 randomly sampled genes)



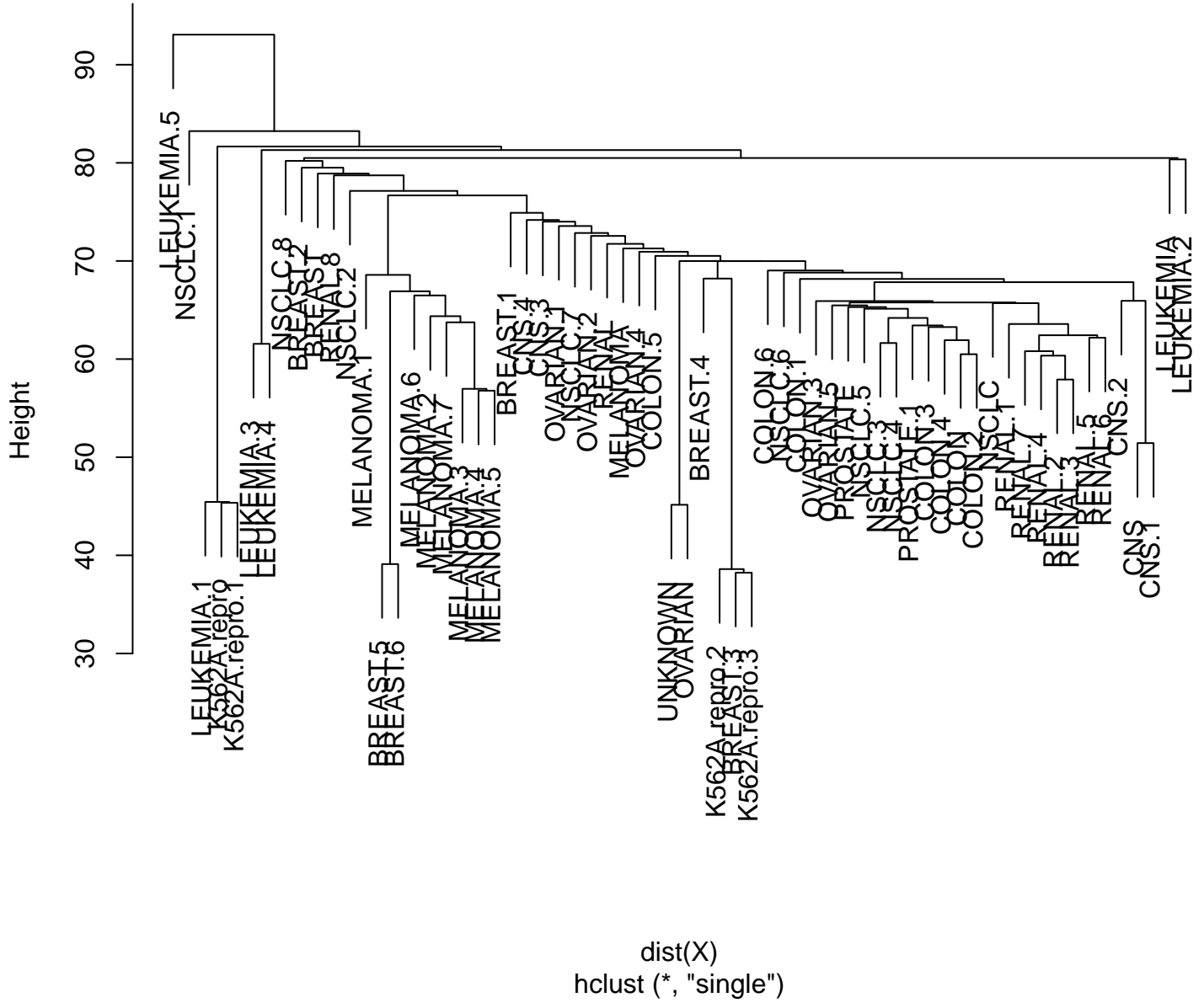
K-means clusters (first 2 PCs, raw clustering)



Average



Single



Complete

