Cluster Dendrogram

Height
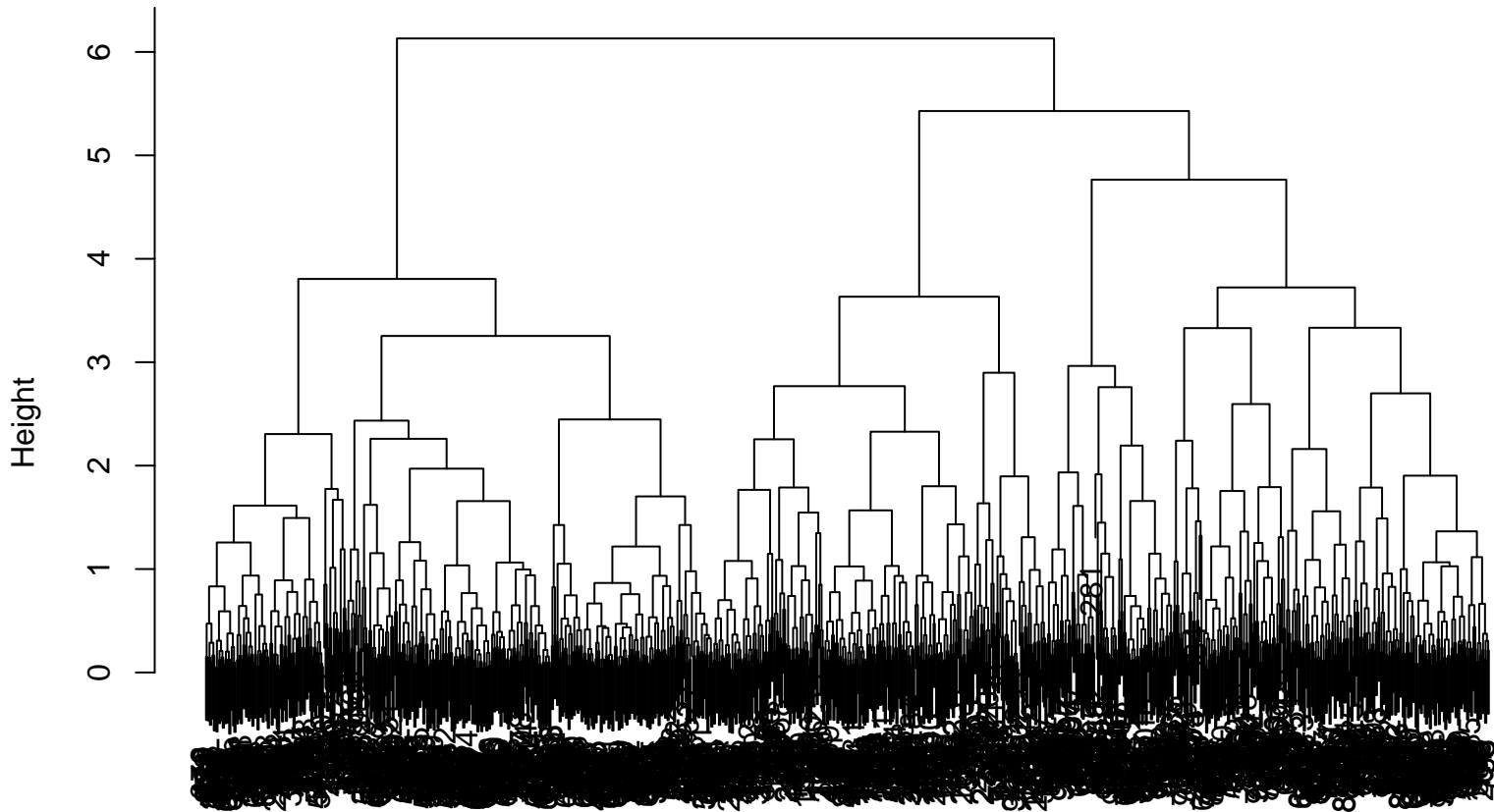
dist(patients_scaled)
hclust (*, "complete")