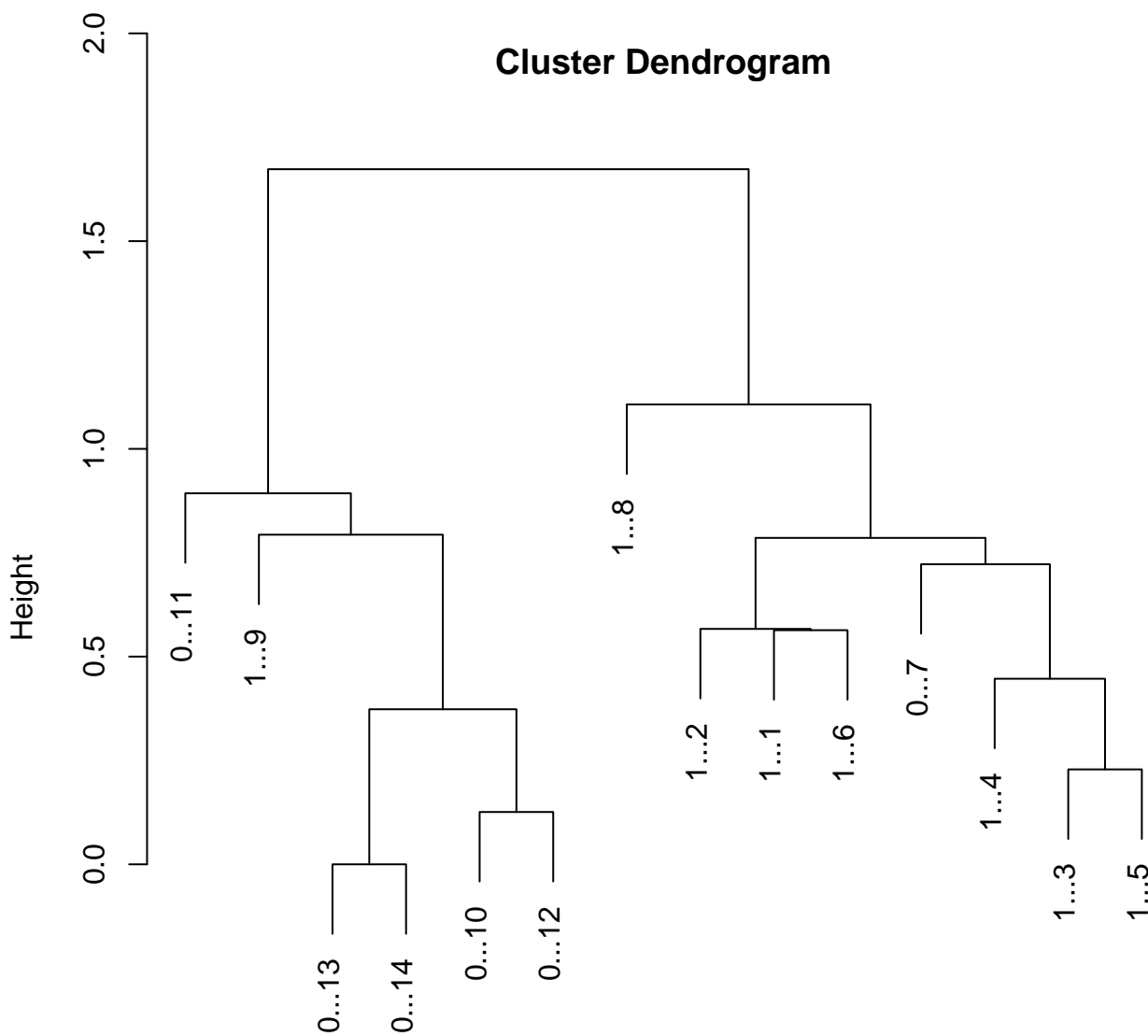


Cluster Dendrogram



as.dist(1 - cor(swi))
hclust (*, "complete")