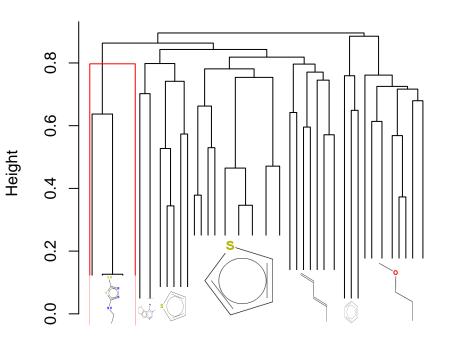
## **Cluster Dendrogram**



as.dist(fp.dist) hclust (\*, "complete")