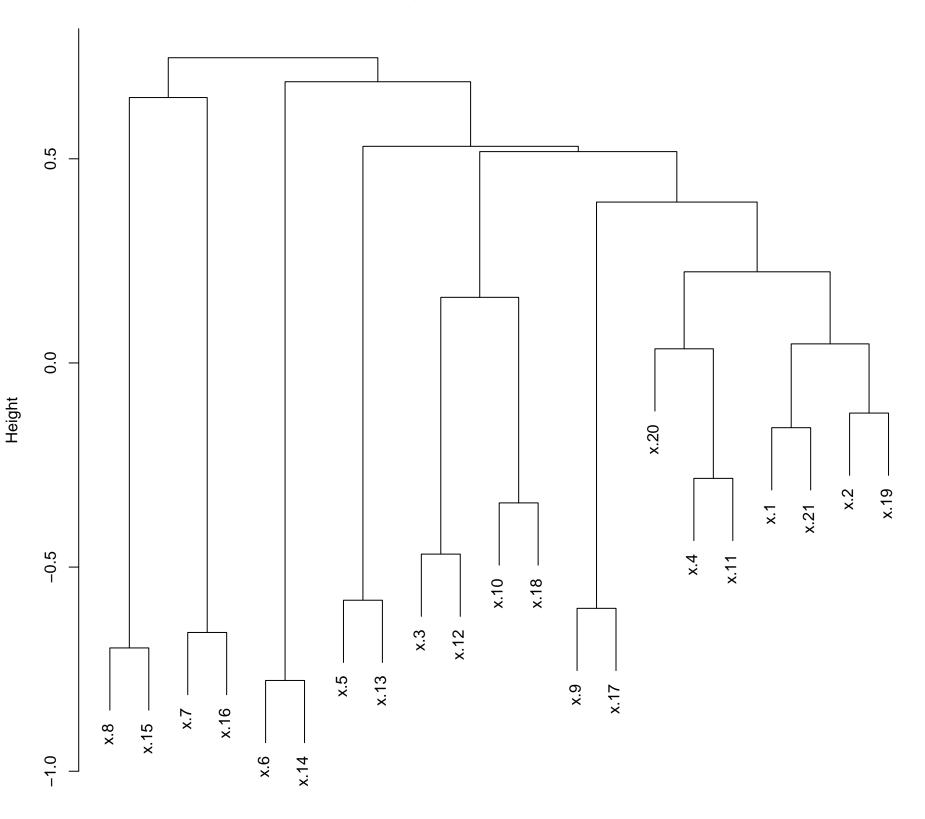
complete linkage and correlation-based distance



as.dist(cor) hclust (*, "complete")