

Figure 1: Cross-validation estimated classification error rates versus number of principal components. Solid line: $M = 25$; dashed line, $M = 1500$; dotted line, $M = 3226$. Variable selection was not performed in the cross-validation procedure.

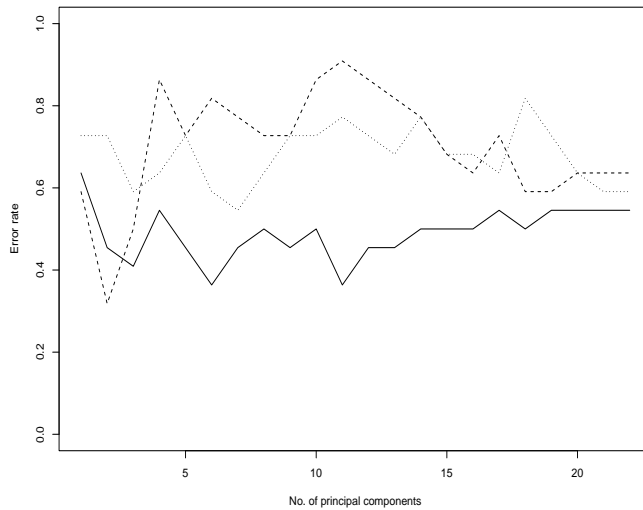


Figure 2: Cross-validation estimated classification error rates versus number of principal components. Solid line: $M = 25$; dashed line, $M = 1500$; dotted line, $M = 3226$. Variable selection was performed in the cross-validation procedure.

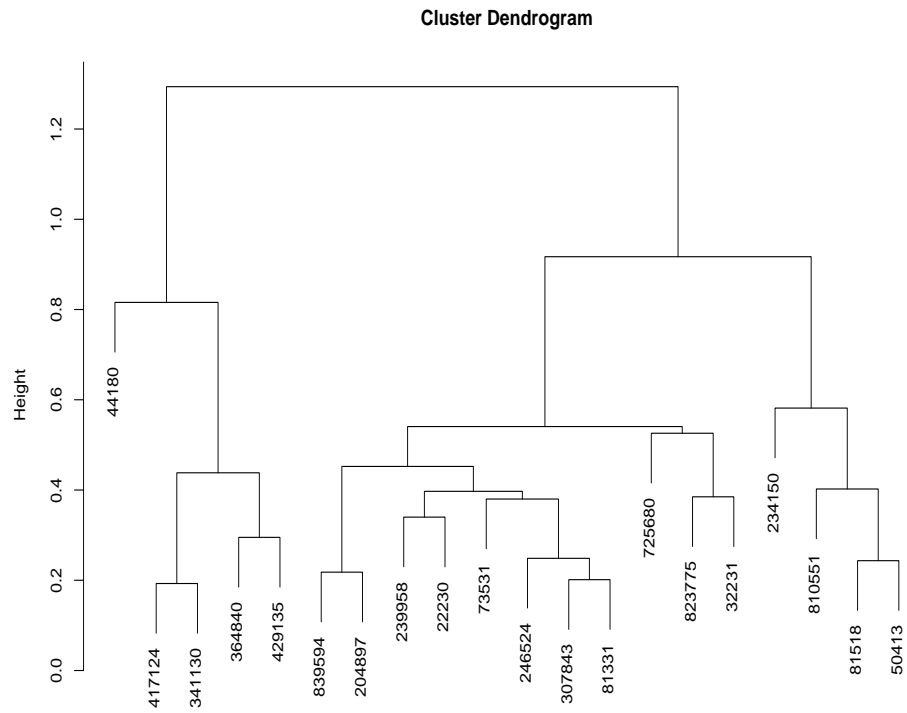


Figure 3: Dendrogram based on average linkage hierarchical clustering of genes in Table 1. Dissimilarity matrix based on correlation between gene expression measurements.

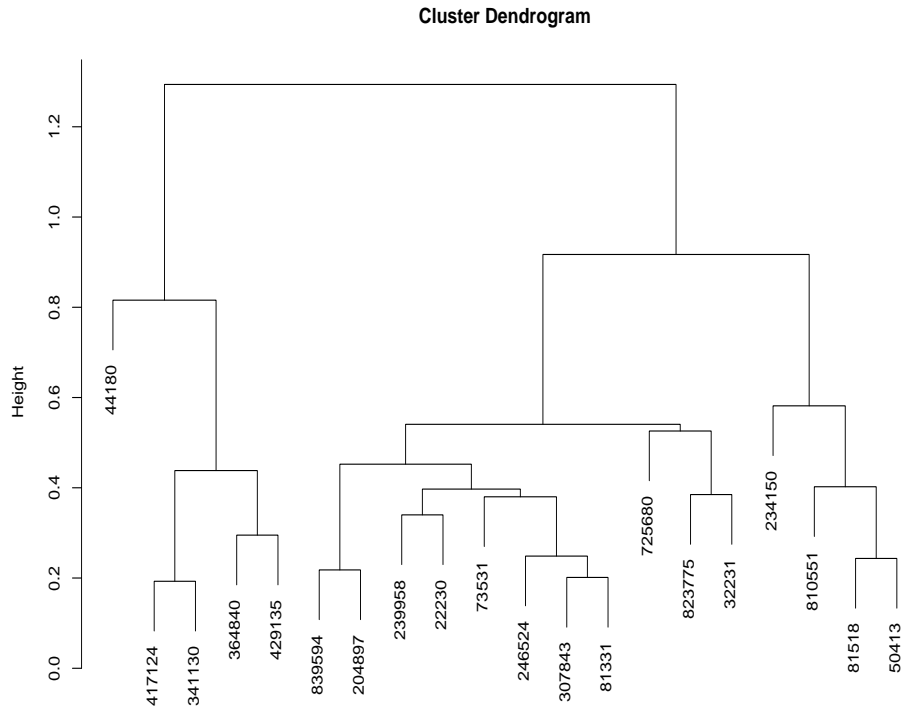


Figure 4: Dendrogram based on average linkage hierarchical clustering of genes in Table 1. Dissimilarity matrix based on correlation between estimated regression coefficients from singular value decomposition regression model.