

## Chapter 6

1. Commands in R to know.

prcomp, drawst, dist, plot, text, as.matrix, hclust, cutree, cmdscale, isoMDS, round, scale, cov, rownames, colnames, Stir2, pairsbg, mahalanobis, kmeans, scatterplot3d, cclust, clusters, barplot, clusterSim, dist2, parameters, head, info, shadow, shadowStars, stepFlexclust, polygon, chull, points, stripes.

2. Be able to implement by hand for very small data sets (and with R for larger ones):

- perform agglomerative hierarchical clustering (using single, complete and average linkage); plot cluster dendograms, obtain clusters.
- plot clusters on principal components, or other lower dimensional representations.
- obtain approximate K-means clusters.
- display cluster solutions graphically (using flexclust ideas such as shadows, barplots, shadow stars, stripes).
- determine reasonable numbers of clusters.
- determine the sum of within cluster distances.

### Problems

Show your work. Check your answers with R. Feel free to liberally round solutions to, say, three decimal places for display purposes.

1. (Use R only for Part (a)) Consider the Euclidean distance matrix.

	1	2	3	4	5	6
1	0	8	9	13	18	16
2	8	0	1	5	10	8
3	9	1	0	4	9	7
4	13	5	4	0	5	3
5	18	10	9	5	0	2
6	16	8	7	3	2	0

- Find a data set with (integer coordinates and) the above Euclidean distances (you can use R for this part).
- Obtain cluster dendograms using the single and complete linkage methods.
- State any differences in the dendograms found in (b).
- Obtain the three-cluster solution for the complete method in (b). State the corresponding sum of within cluster distances.

- (e) Find the centroids for the three clusters obtained in (d), and the matrix of shadow values between centroids, using the formula

$$S_{i,j} = \frac{1}{n_i} \sum_{x \in A_{i,j}} \frac{2d(x, c_i)}{d(x, c_i) + d(x, c_j)}, \quad 1 \leq i \neq j, \leq 2, \quad (1)$$

where  $A_{i,j}$  is the set of points whose closest centroid is  $c_i$  and second closest centroid is  $c_j$ , and  $n_i$  is the number of points closest to the centroid  $c_i$ .

- (f) Draw a rough shadow stars plot for the clusters in (b).

2. (Use R, throughout) Consider the data matrix given in the file that was E-mailed to you. The matrix begins

```
> head(Q2)
      V1      V2      V3
1 -0.09580978 0.06216742 -0.07684248
2  0.79640420 0.08455216  1.43648000
3  1.16606100 0.84623770 -0.77053320
4 -0.68268140 0.49050940 -0.12621180
5  0.96462750 0.94635740  1.13541900
6  0.08437982 0.05538653  1.66976800

> dim(Q2)
[1] 600  3
```

You can read the data into the matrix Q2, (check the “Change dir” tab under “File”, first) by typing

```
> Q2<-read.table(file="Q2data.txt")
```

Note that matrix was written out with the command

```
> write(t(Q2), file="Q2data.txt",ncol=3)
```

- Plot the points in three dimensional space.
- Use `cclust`/flexclust a few times to obtain some insight on appropriate number of clusters.
- Obtain a `cclust` run that gives a sum of within cluster distances that is less than 312.03. Plot the `cclust` run (using `prcomp`). You may need to run a few iterations of `cclust` to obtain a low enough sum of within cluster distances. If you don't get a fairly nice looking (perhaps even “cool”) plot, you may have an error in your coding. :)

- (d) Draw a new three dimensional scatterplot, this time with colored cluster numbers at each of the points (with colors distinguishing clusters).
- (e) The data arose by adding some noise to some nice points in three dimensional space. Come up with a theory on what those points were. Use any information from the package `flexclust` that may be helpful.
- (f) Obtain a barplot for the `cclust` result in (c).
- (g) For the `cclust` result from (c), use `clusterSim` to obtain the matrix of shadow values. Is the matrix symmetric?
- (h) For the `cclust` result from (c), determine the centroids,  $c_a$ , closest to the point  $(0,0,0)$  and the centroid,  $c_b$ , closest to the point  $(1,1,0)$ . Find the set of points  $A_{a,b}$  (as in Equation (1), above), whose closest centroid is  $c_a$  and second closest centroid is  $c_b$ . Hint: Use the function `getshads`, below.
- (i) (Show some manipulations.) Verify the value,  $S_{a,b}$  in the matrix of shadow values corresponding to the centroids  $c_a$  and  $c_b$  (as found in Part (h)). Hint: Use the interim values outputted by the function `getshads`, below, as a check.
- (j) Locate the lines for the corresponding values,  $2d(x, c_a)/(d(x, c_a) + d(x, c_b))$  (as in (1)), for  $x \in A_{a,b}$ , in a shadow stars plot. Hint: The corresponding values comprise the `[[5]]` component in the list of values outputted by `getshads`.

```
> getshads<-
function (M,clM,i,j)
{
  L<-1:dim(M)[1]
  DM<-dist2(M,parameters(clM))
  DMo<-t(apply(DM,1,order))
  DMoi<-DMo[clusters(clM)==i,]
  ni<-dim(DMoi)[1]
  v<-L[(DMo[,1]==i)&(DMo[,2]==j)]
  dii<-DM[(DMo[,1]==i)&(DMo[,2]==j),i]
  dij<-DM[(DMo[,1]==i)&(DMo[,2]==j),j]
  d<-2*dii/(dii+dij)
  list(v,ni,dii,dij,2*dii/(dii+dij),sum(d)/ni)
}
```

3. (Use R, throughout) Consider the crime data for states. You may need to run

```
> demo("Ch-CA")
```

to load the data into R.

- (a) Restrict to the data on the variables rape, burglary and theft. Scale the variables accordingly.
- (b) Find the mahalanobis distances for the states in the new data set.
- (c) Use `pairsbg` to obtain bagplots for the data.
- (d) Remove the two states with the highest mahalanobis distances, and rescale the data. Note that the following will scale the data and remove the attributes (“scaled:scale” and “scaled:scale”).

```
> MQ4s<-matrix(scale(MQ4,scale=TRUE,center=TRUE),ncol=3)
```

The data should now begin as follows.

```
> head(MQ4s)
      Rape   Burglary   Theft
ME -1.3711582 -0.95946793 -0.7384367
NH -0.8828718 -1.07153371 -0.9243492
VT -0.8610082 -0.61860119 -0.2703116
MA -0.2852675 -0.33376734 -0.9497617
RI -0.8901596  0.18687159 -0.4428491
CT -0.7152511 -0.03725997 -0.1887240
```

- (e) Perform a principal components analysis on the new data set in (d). Interpret the coefficients.
- (f) Plot the states, using the first two principal components.
- (g) Cluster the states using `hclust` (and complete linkage). Plot the cluster dendrogram and use the result to form four clusters. State the states in each cluster. Give the sum of the within cluster distances; you can use the following program.

```
> sumclldist<-
function (M,cl)
{
  mv<-NULL
  Dt<-0
  t<-unique(cl)
  for (i in t)
  {
    U<-cbind(M[cl==i,])
    m<-colMeans(U)
    mv<-rbind(mv,m)
    Dm<-dist2(U,m)
    print(Dm)
    Dt<-Dt+sum(Dm)
  }
}
```

```

}
rownames(mv)<-t
list(mv[order(t),],Dt)
}

```

- (h) Obtain a four-clustering of the data using `cclust`. Ensure that your clustering has a sum of within cluster distances that is less than 34.30 and that exactly six  $S_{i,j}$  (produced by `clusterSim`) are zero ( $1 \leq i \leq j \leq 4$ ). List the states in each cluster.
- (i) Plot the result of `cclust`, from Part (h) (without projecting onto the principal components). Label the states using `text` (and a reasonable value of `cex`). Plot the result of `cclust` (projecting onto the principal components). Label the states using `text`.
- (j) Obtain a second four-clustering of the data using `cclust`, this time ensuring that your clustering has a sum of within cluster distances that is greater than 39.50. Plot the results, as in Part (i) (with and without projecting onto the principal components).
- (k) For the result of `cclust` in Part (h) (with sum of within cluster distances less than 34.30), draw a barplot, and give the cluster centroids.
- (l) Give the distances of the data point corresponding to NC, from all four centroids (for the clustering in Part (h)). Locate (approximately) the stripes corresponding to these distances on a `stripes` (type="all") plot.