2.

No linear boundry can separate label 1 and -1. [10] . not linearly separable.

D1 = -1, -2, 3, 3, 6, 7  
D2 = -3, -2, 3, 5, 8  
Wildow width h = 2  
Window function 
$$\phi(u) = \begin{cases} -3_4 u^2 + 3_1 4 & \text{if } -1 \le u \le 1 \end{cases}$$
  
 $x = 4 \text{ (what class?)}$   
 $P(n) = \frac{k/n}{v}$   
 $v = h^1 = 2$   
For class 1

$$P(n) = P(4) = \frac{1}{h} \underbrace{\begin{cases} \frac{1}{5} & \frac{1}{4} & \frac{1}{3} & \frac{1}{4} \\ \frac{1}{5} & \frac{1}{12} & \frac{1}{5} & \frac{1}{4} \\ \frac{1}{5} & \frac{1}{5} & \frac{1}{5} & \frac{1}{4} & \frac{1}{3} & \frac{1}{4} \\ \frac{1}{5} & \frac{1}{5} \\ \frac{1}{5} & \frac{1}{5}$$

$$\frac{1}{5} \sum_{j=1}^{5} \frac{1}{2} \varrho \left( \frac{4-nj}{2} \right) = \frac{1}{10} (\varrho(\frac{1}{2}) + \varrho(-\frac{1}{2}) + \varrho(-1) + \varrho(-\frac{1}{2}) + \varrho(-1)$$

$$= \frac{\varrho(\frac{1}{2})}{5} = \frac{\varrho}{16 \times 5} = 0.1125 \quad ((lass 2))$$

Class 2 > Class 1

We are comparing Published densities directly, as no information about prior is given.

(las) = 0.09375

Using Single-link algorithm

+ This algorithm uses smin as the similarity measure + We will combine clusters that have high similarity measure

From the matrix Atcan be seen. P2 and P3 have high similarity.

Step! (P1 (P2, P3), P4, P5, P6)

Step 2 P5 and P6 have high similarity sim (PS,P6) = (0.85) {P1 {P2, P3}, P4, {P5, P63}}

Step 3
P(5,6)-74 = mln (ds,4, 16,4) = max (similarity 5,4, simularity 6,4) Company with all clusters P4 is similar to P5, P6

 $Sim_{(5,6)} \Rightarrow 4 = max(0.80, 0.65)$ {P1, {P2, 833, {P4, {P5, P6}}}

Shm(1,3) 74 = max (0.70,0.75)

 $Sun(P_2, P_3) = 0.95$ 

By comparing all clusters Elustors LP2, P3 3 and & P4, {P5, P64} are more similar.

=> {P1, {P2, P33, 1P4, 1P5, P633}}

Step 5 and the last cluster.

= {{ P,7, { { P2, P37, { P4, { P5, P6} } } }}

The Dandrogram D' cost depicts the Clusters.

(emplote linkage Algorithm.)

+ In Single linkage we took max (similarity duto, similarity that )

+ Here we will highest (forthest) neighbour bictures clusters

unlike in single linkage.

Step 1 Clusters P2 and P3 are more stenclar.

= { P1, {P2,P3}, P4,P5, P6}

shm (P2, P3) =0.95

Cluster P5 and P6 has the highest similarity

= { Pi, 182, P3}, P4, {P5, P63}

SLM (PS, P6)=0.85

Step 3

we compare all clusters with all clusters the difference here is we compare with farthest heighbour.

 $d_{5,6}\rightarrow 4 = min(0.80,0.65) = 0.65$   $d_{2,3}\rightarrow 4 = min(0.70,0.78) = 0.75$ hence cluster 4 is smaller to cluster 2,3

=> { 1, 1{2,33,43, {5,63}}

Step 4

like in step 3 we compare all, cluster, is sunclar to cluster \$2,3,44 than \$5,63

= { {1, {{2,33,43}, 55,63}}

Step 5

Add all cluster together (the last cluster)

= { {1, { {2,33,4333,4333, {5,633

:. The dendrogram B'ss correct for mutilinkage Algorithm.