Point	X. Coordinate	Y-coordinate
PI	0.4005	0.5306
P2	0.2148	0.3854
P3	0.3457	0.3156
P4	0.2652	0.1875
P5	0.0789	0.4139
P6	0,4548	0-3022

e) Single link; - We need to Consider the minimum distance blue the points and Choose the minimum distance points as a group on cluster.

Given Distance Matrix 1-

	PI	P2	P3	P4	P5	PG
PI	0.00	0.2357	0.2218	0.3688	0.3421	0.2347
P2	0.2357	0-00	0.1483	0.2042	0.1388	0.2540
P3	0.2218	0.1483	0.00	0.1513	0.2843	0.1100
РЧ	o·3688	0.2042	0.1513	0.00	0.2932	0.9316
P5	0.3421	0.1388	0.2843	० • २१३२	0,00	0.3921
P6	0.2347	0.2540	0.1100	0.2216	०. अग	0.00

Broallest distance b/w the points is 0.1100 b/w (P3, P6) so we need to group/cluster P3, P6 as a single Unit.

Dendo gram

3 6

3.6

```
Repeating the process by updating the Distance matrix
  min( dist (P_3, P_6), P_1) = min (dist (P_1, P_3), dist (P_1, P_6))
                     = min (0.2218, 0.2347)
                        = 0.2218
min (dist (P3,P6), P2) = min (dist (P2P3), dist (P2,P6))
                    = min(0.1483, 0.2540)
                      = 0.1483
  min (dist (P3, P6), P4) = min (dist (P3, P4), dist (P4, P6))
                    = min (0.1513, 0.2216)
                    = 0.1513.
   min(dist (P3, P6), P5) = min(dist (P3, P5), dist (P5, P6))
                   = min (0.2843, 6.3921)
                   2 0.2843.
Updating the distance matrix
                       B, Pc Py P5
                  P2
         P,
   P,
         0.00
   P2 0.2357 0.00
   P3, P7 0.2218 0.1483 0.00
      0.3688 0.2043 0.1513
   Py
    Ps 0-3421 0-1388 6-2843 0-2932 0.00
  smallest distance is .0.1388 blu (P2, P5)
  After forming the Cluster.
 Dendo gram
```

min (dist (P2, P5), P1) = min (dist (P1, P2), dist (P1, P3)) min (0.2357) 0.3421) > 0.2357 min (dist (P2P5), dist (B3P8)) > min (dist (P2P6), P), dist (P3, P6), P5) min (0.1483, 0.2843) > 0.1483 min (dist (P2, P5), P4) = min (dist (P2, P4), dist (P4, P5)) = min (0.2043, 0.2932) = 0.2043 Updating distance matrix Pr. Ps B. P. 0,00 P2,P5 0.2357 0.00 P3,P6 0.2218 0.1483 0000 Py 0.3688 0.2043 0.1513 0,00 The smallest distance is 0.1483 blu the points (P215) x (P3,P8) group these two into a Single Cluster

Dendo 8 (2 5)

Represting the steps above

min (dist (C(P2Ps), (P3,P6))), P1) = min (dist (P2Ps), P1), dist (P3,P6), P1) = min(0.2357, 0.2218) = 0-2218 min(dist (C (P2/P5), P3/P2)), P4) = min ((P2/P5), P4), ((P3/P6), P4)) = min (0.2043, 0.1513) = 0.1513. updating the distance moutrin (P2P5) (P3,P6) P, 0 (P2Pg), (P3,P6) 0.2218 0.1513 0.3688 Py The smallest distance is 0.150 blu P4 & (Page) (Bips) Dendo gram Alpert Min (dia (P3, PD) (P2, P5), P4), P1) = min ((P3, P6), P1, (P2, P5), (P4, P)) = min (0.2218, 0.3688) - 0.2218realist Singe this is the last point we need to group into the Cluster. Dendo gram

## Complete link Proximity function:

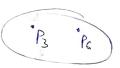
complete link considers the trace distance while update Collaborating the distance mouther

from the given distance matrix the smallest dustance is 0.1100 b/w P2 4-R

Derdogram

Christer





Here we need to find the movimum distance blu the cluster and the rest of the points.

Marx (dist (B, 187, P) = Marx (dist (P, 183), dist (P, 18)) = Man (0.2218, 0.2347) - 0.2347

Man (dist (P3, P6), P2) = Meur (dist (P2, P3), dist (P2, P3)

= max(0.1403, 0.25 40) = 0.2540

max (dist(fs, Pa), fy) = may (dist (P3, P4), dist (P4, Pa))

= man (0.1513,00000) 0.2216)

= 0.2716

man (dist (P3,P6), P5) = man (dist (P3, P5) dist (P5,P6))

= max (0.2843, 0.3921)

= 0.3921

Distance Matrix

P (P3Rc) PZ P5 P. 0 0.2540 0.2347 P3,P6 0.2042 0.2216 0 0.3688 Py 0.3921 0.2932 0.1388 P= 0.3921 The Smallet distance is \$ 0.1388 5/w Prefs. forming the denderson of Cluster. Distance new from (P2, P5) Man (dist (P2, P5), P1) = max (dist (P1, P2), dist (P1, P5)) = Mar ( 6.2357, 0.3921) 5 0.3921 Man (dist (P2, P5), (P3, P6)) = max (dist (P3, P6), P3), dist ((P3, P6), P5)) = man (0.2540,0.3921) = 0.3921 man (dist (P2, P5 2 Ry) = move (dist (P2, Ry), dist (Ry, P5)) = man (0.2042, 0.2932)

= 0.2932

Distance Mouth !

PI P2, P5 P3, P1 Py. PI 0 P2,75 0.3921 0 0 0.3921 P3, P6 0.2216 012932 0.388 Ry to By, Ps Chysten smalles + distances 0.2216 from Dendogram & Clyster. 2 5 Distance from Chykr to all other points marx (dist ((B3,P6),Pr), (12.P5)) = marx (dist (P3,P6), (P2.P5)) (R2, P5) P4)) = mar (0.3921, 0.2932) = 0.3921 man (dist ((P3,P6),P4),P,1)) = man (dist((P3,P6),P,), (P4,P,)) = man (0.2347) 0.3688) = 0.3(8% updated distance manna y P PZPS (P3,P6), R P. 0 0.3421 P2,85 6.3688 0.3921 (P3,P6),P4 0

distance is 0.3421 b/w P, f(Pz, ts) smallest pender gram Cluster-Distance from Chyter is. Marc (dist (Pz, Ps), P, ), (Pz, Ps), Py) = Man (dist (P2P5), (B,POP4), dift P, (B,POP4)) man (0.3921, 0.3688) 2 0:394 Updated distance reating v. P3, P4, Px Pr. P2, P5 P1, P2, P5 0 0.3924 P3, Ry, Pc

Average link; -Average link proximity function Considers the average while collulating the distance motion from the given distance matrix the Snallest distance is 0.1100 b/w P3 & P6 Dendagram Chuter The distance from the Cluster (P3, P6) & Ag ( dist (P3 (P8), P1) = dist ((P, P1)+ (P, 1P8))/2 = (0.2218+0.2347)2 - 0.22825 Ang (dis+(P3 R6), P2) = dis ((P2/2) + (P2/2))/2 = (0,1483 +0,2540)/2. Avg (disa (P3,P6),P4) = Aisa (B,P4) + (P4,P0)/2 = 0.20115

= (0.1513 + 0.2216)/2 = 0.18645 ANG (dist (P3, Pr), P5) = (0.2843+0.3921)/2 = 0.3382. Updated distance matrix b. PI PZ PI P2 0:2357 0

P3, P6

PY

Ps

Bip 0-21825 0:20115 0

Pu 0.3688 0.2042 0.18645 0

P5 0.3421 0.1388 0.3382 0.2932 Dandegroum

6 2 5

Distance from the Chusta is

Arg (dist (P2, P5), P,) = distal (0.2357 + 0.3421)/2

- 0.2889

(25)

Avg (dist (P2, P5), (P2, P6)) = (0.20115 + 0.3382)/2

- 0.269675

Avg (dist (P2, P3), P4) = 10.204240:2922)/2

= 0.2487

updated matrix i-PI P2, P5 P3, Ps P4 PI P2, P5 0.2689 0 0.22825 0.269675 0 P3, P 0.5487 0.18645 0.3421 PL clustry Dendegram Distance from chuster: Ag (dist (P3,P6), P4, P1)) = (0.22825 + 0.3421)/2 - 0.285175 Avg (dist (P2, P0, P4), (P2, P5)) > (0.289675+0.2487)/2 = 0.2591875 U Padated distance moutrix (P3, P6) P4 P, P2 P5 P.  $\Diamond$ P2P5 0.2869.  $\circ$ (P3, P6), P4 0.285175 0.2591875 O

Den do gram Cluster The distance from Clusters is. AVS ( B, P(), A), (R, PS), P) (0.285175+0.2889)/2 0.5879375 Dandogram Clusters B