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Essentials of Data Analytics

Tasks for Week-8: Hierarchical Clustering

Understand the following operations/functions on 'USArrests' data and perform similar operations on 'iris' dataset based on given instructions.

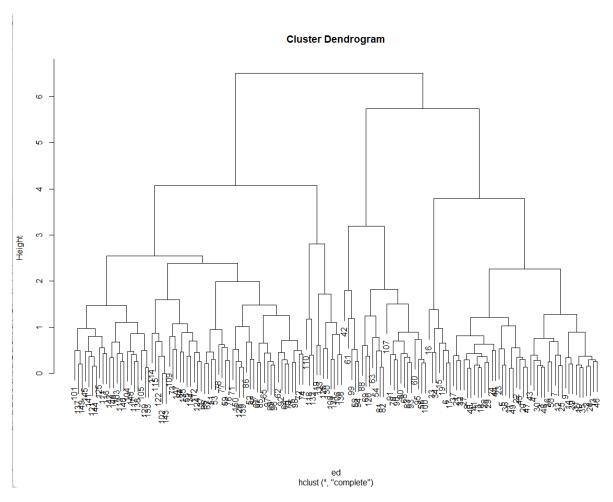
AIM:

To develop a hierarchical clustering model - a dendogram for the given data using R programming and perform operations on the data "iris".

Algorithm:

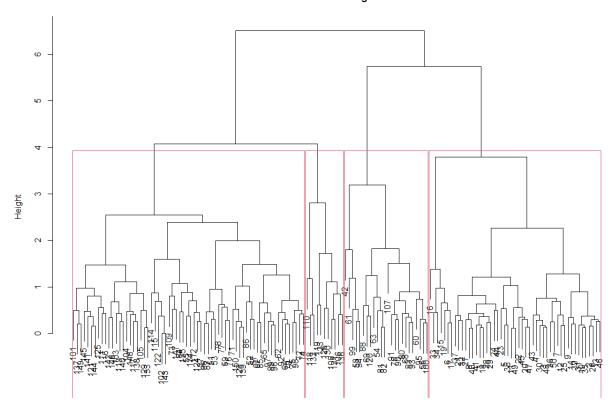
- Set the working directory to the dataset where we store by using setwd() and read data.
- By using scale function. We scale the data and store it in another variable.
- Find the Euclidean distance using the dist() function and method="euclidean".
- Find the hierarchical clustering using hclust and method = "complete.
- For plotting the dendrogram we use plot function.
- Grouping the dataset into number of cluster using cutree function and plotting them.
- Use rec.hclust to find the cluster.

RESULTS:



> c	> cluster																												
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60
1	1	1	1	1	1	1	1	1	1	1	2	1	1	1	1	1	1	1	1	3	3	3	2	3	2	3	2	3	2
61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90
2	3	2	3	3	3	3	2	2	2	3	3	3	3	3	3	3	3	3	2	2	2	2	3	3	3	3	2	3	2
91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	113	114	115	116	117	118	119	120
2	3	2	2	2	3	3	3	2	2	3	3	3	3	3	4	2	4	3	4	3	3	3	3	3	3	3	4	4	2
121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150
3	3	4	3	3	4	3	3	3	4	4	4	3	3	3	4	3	3	3	3	3	3	3	3	3	3	3	3	3	3
> re	ect.h	ic1us	st (h	erclu	ustei	r,k=4	4)																						





CODE:

```
rm(list=ls())
setwd("C:/Users/VIKRAM SURYA/Desktop/EDA_LAB")
data <- read.csv("iris.csv",row.names=1)
View(data)
df<-scale(data)
ed<-dist(df,method="euclidean")
hercluster<-hclust(ed,method='complete')</pre>
```

```
plot(hercluster)
cluster<-cutree(hercluster,k=4)
cluster
rect.hclust(hercluster,k=4)</pre>
```