

EJEMPLOS DE CLASIFICACION

Funciones que se usan para hacer cluster

Función	library	Que hace	Observaciones
daisy	cluster	Calcula Matriz disim/dist	
CLASIF. JERARQUICA			
agnes	cluster	Jerárquico Agregativo	Trabaja con matriz de datos y disimilaridades
hcluster	stats	Jerárquico Agregativo	Parte de matriz disimilaridades
diana	cluster	Jerárquico Divisivo	
CLASIF. NO JERARQUICA			
kmeans	stats	No Jerárquico k medias	Fija centros de cada K grupo
pam	cluster	No Jerárquico K medoides	Elige individuos representativos
clara	cluster	No Jerárquico.	Igual que <i>pam</i> pero set de datos grandes.
fanny	cluster	No Jerárquico. Difuso	Para cada observación se obtiene coeficiente de pertenencia a cada grupo
CLASIF. BASADA EN MODELOS			
Mclust	mclust		Trabaja con variables cuantitativas

DATOS

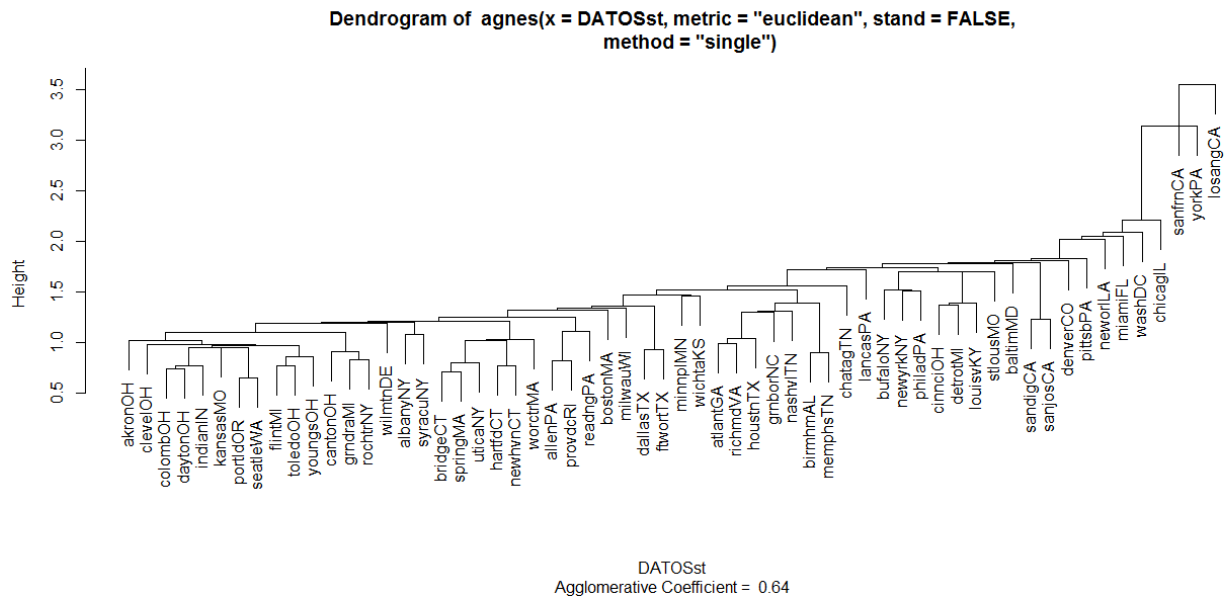
Observaciones: ciudades de USA.

Las variables:

- Promedio anual de precipitaciones (Rainfall).
- Mediana de años de educación completa en aquellos mayores de 25 años, en 1960 (Education).
- Densidad población en áreas urbanas. Población /millas² (Popden).
- Porcentaje de la población no blanca en áreas urbanas. (Nonwhite).
- Polución potencial de óxidos de nitrógeno (NOX).
- Polución potencial de sulfuro (SO2)
- Tasa de mortalidad. Expresada en muertes cada 100.000 (Mortality)

CLASIFICACION JERARQUICA

1. VECINO MAS CERCANO

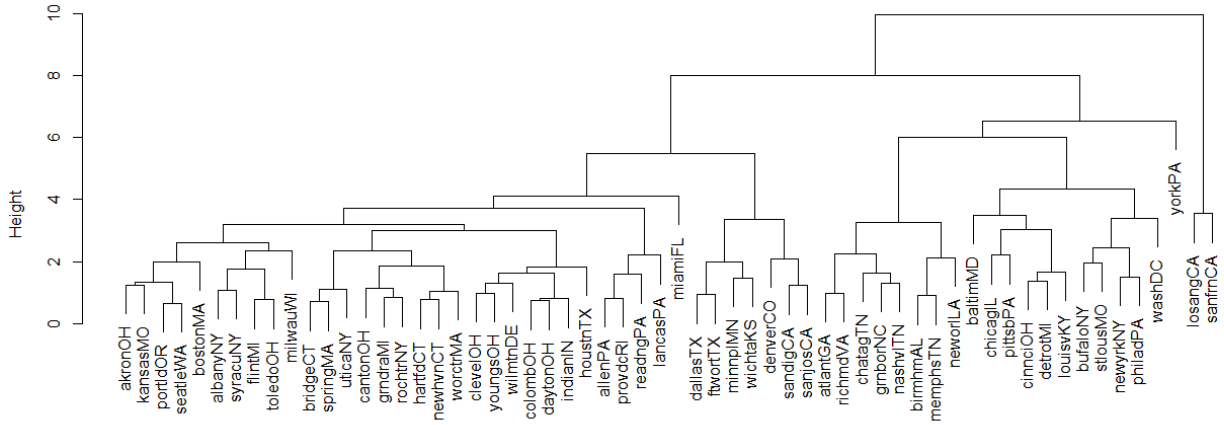


Indicadores

	. history	Freq	Rcuad	psF	psT
39	-34	-56	2	0.8163700	8.669181
40	35	39	32	0.7940983	8.119338
41	-38	-39	2	0.7913895	8.641029
42	40	38	39	0.6814305	5.284669
43	-9	41	3	0.6759521	5.606027
44	42	-11	40	0.6554760	5.580830
45	37	-46	4	0.6501080	5.972223
46	43	45	7	0.6364711	6.195185
47	44	-28	41	0.6207462	6.410631
48	47	46	48	0.5349694	5.019910
49	48	-5	49	0.4883246	4.676384
50	49	29	51	0.4138068	3.921790
51	50	-18	52	0.3901317	4.078076
52	51	-40	53	0.3570034	4.124478
53	52	-37	54	0.3129311	4.023214
54	53	-32	55	0.2933039	4.482382
55	54	-55	56	0.2754953	5.228484
56	55	-12	57	0.2294885	5.559665
57	56	-48	58	0.1862005	6.520912
58	57	-59	59	0.1310792	8.749468
59	58	-29	60	0.0000000	NaN

2. VECINO MAS LEJANO

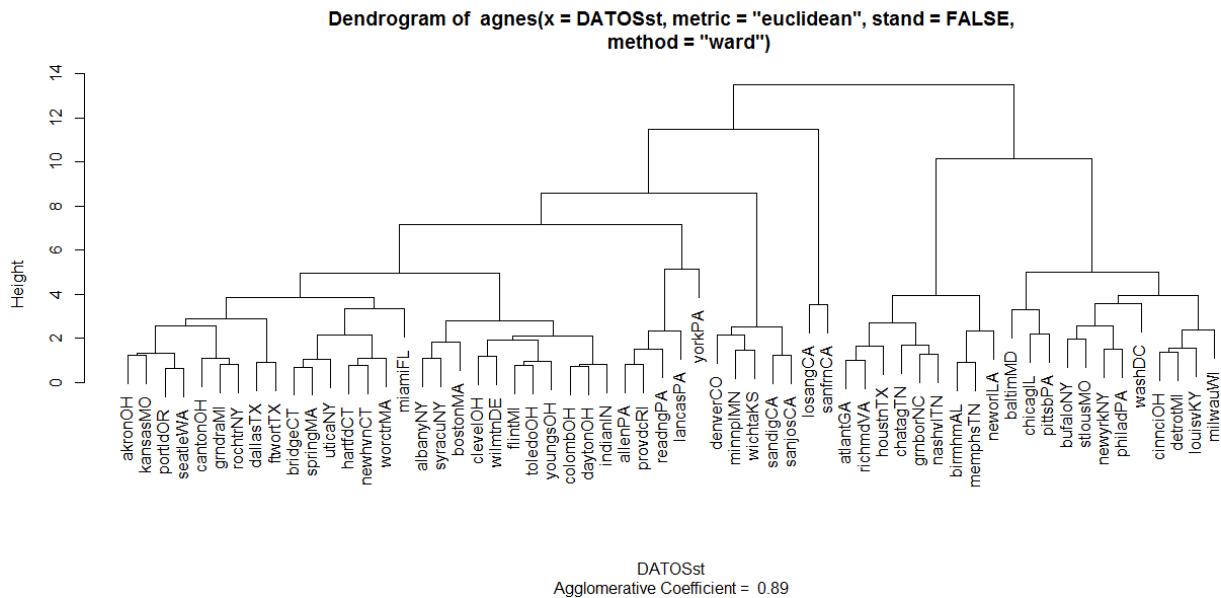
Dendrogram of agnes(x = DATOSst, metric = "euclidean", stand = FALSE,
method = "complete")



DATOSst
Agglomerative Coefficient = 0.85

. history	Freq		Rcuad	psF	psT
39	12	27	5 0.8888508	15.59399	3.601249
40	15	31	9 0.8795355	15.37096	5.001720
41	29	-33	5 0.8723602	15.56758	2.993880
42	32	24	4 0.8645244	15.76582	2.155591
43	33	41	10 0.8525083	15.53386	4.170937
44	40	30	16 0.8266365	13.98680	9.910457
45	37	28	5 0.8137771	14.04614	3.505965
46	43	44	26 0.8029367	14.41751	2.668861
47	39	36	8 0.7880284	14.56065	4.064596
48	34	35	7 0.7685559	14.49032	5.457105
49	42	-55	5 0.7531373	14.94909	3.062090
50	-5	45	6 0.7392332	15.74913	2.330646
51	-29	-48	2 0.7242642	16.74496	NaN
52	46	38	30 0.6905630	16.57816	7.980273
53	52	-32	31 0.6732874	18.20368	3.297157
54	50	49	11 0.6460511	19.71288	3.589412
55	53	48	38 0.5620634	17.64724	14.639348
56	47	54	19 0.4487692	15.19695	14.542043
57	56	-59	20 0.3975090	18.80361	3.754748
58	55	57	58 0.1596907	11.02220	22.667795
59	58	51	60 0.0000000	NaN	11.022205

3. METODO WARD

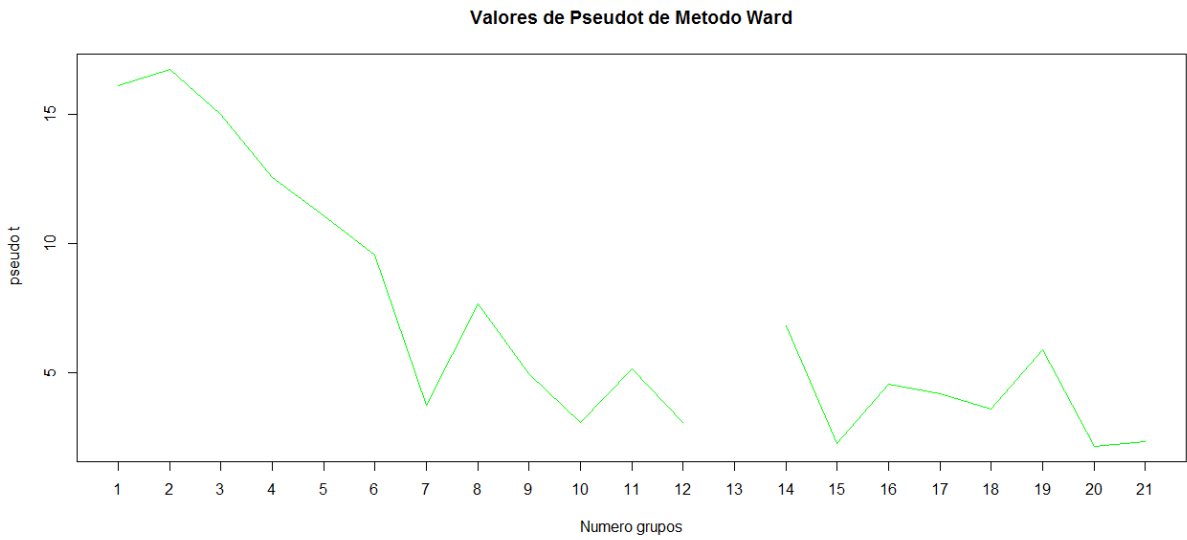
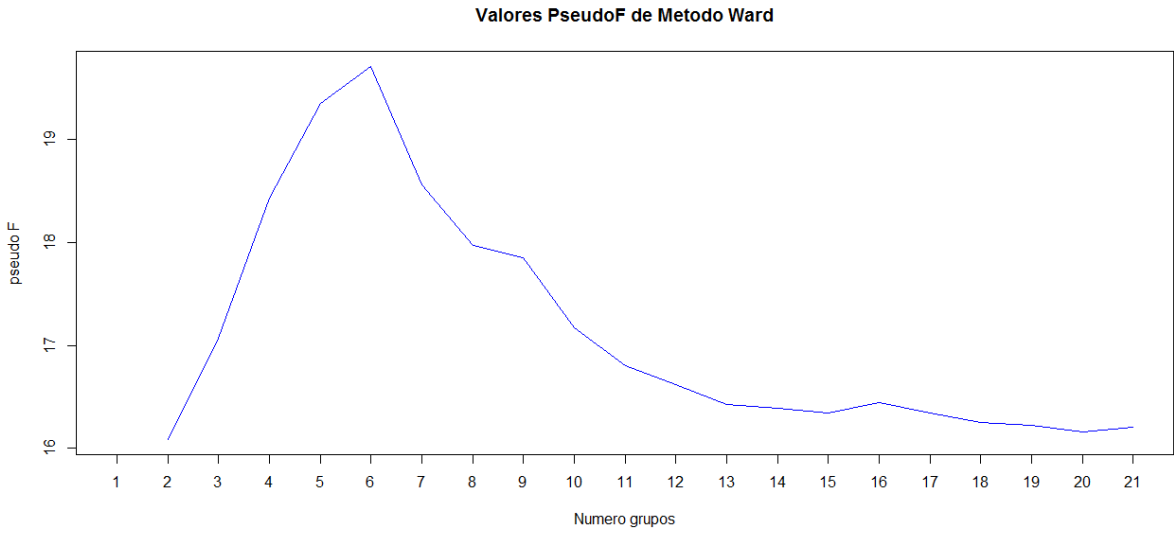
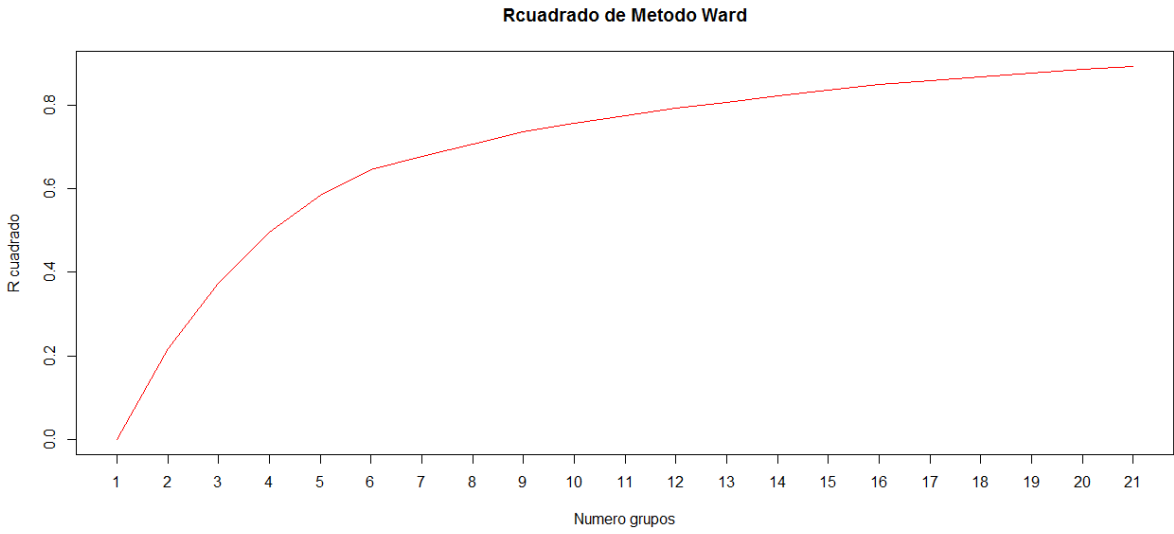


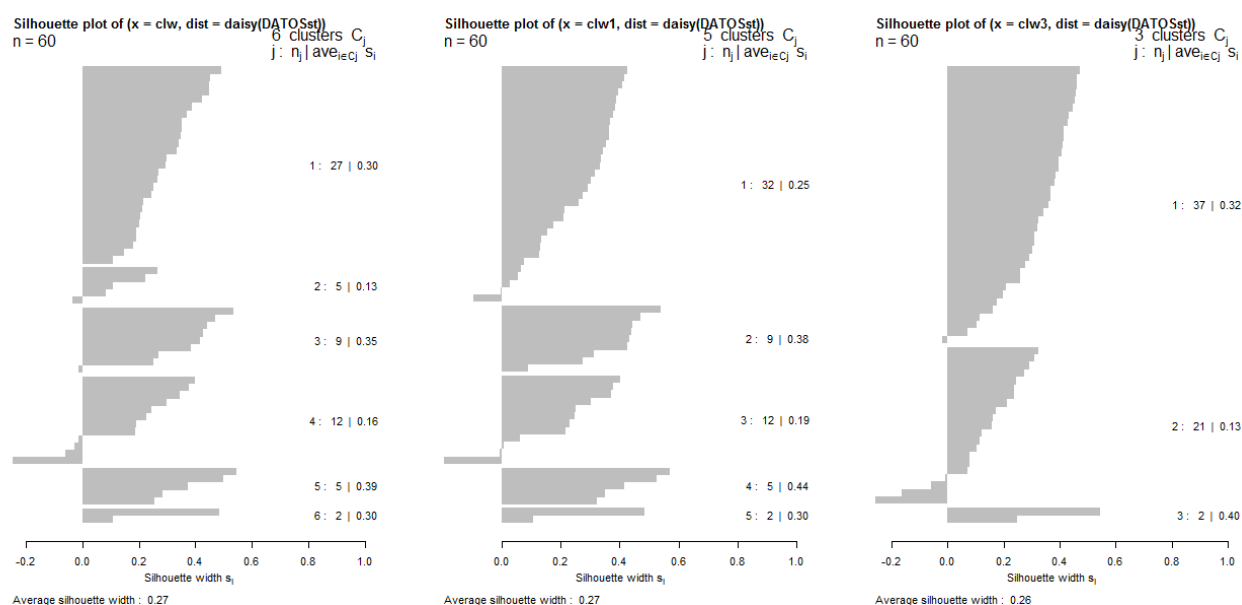
	.	history	Freq		Rcuad	psF	psT
39	33	18	5	0.8925979	16.20607	2.318250	
40	31	24	4	0.8847621	16.16358	2.155591	
41	21	15	7	0.8769154	16.22802	5.880668	
42	27	28	6	0.8680596	16.25444	3.587898	
43	29	32	11	0.8587627	16.34076	4.170042	
44	41	10	9	0.8486633	16.44949	4.545395	
45	-5	35	3	0.8356928	16.34837	2.241636	
46	34	-32	7	0.8224712	16.39330	6.796644	
47	-29	-48	2	0.8075022	16.42989	NaN	
48	40	-55	5	0.7920836	16.62382	3.062090	
49	44	46	16	0.7742029	16.80090	5.150735	
50	48	38	9	0.7555569	17.17184	3.071102	
51	42	37	9	0.7368719	17.85274	4.966924	
52	49	43	27	0.7075625	17.97368	7.645121	
53	45	50	12	0.6775404	18.56028	3.757331	
54	36	-59	5	0.6459748	19.70630	9.543820	
55	52	54	32	0.5845989	19.35054	11.049396	
56	55	39	37	0.4967619	18.42645	12.523144	
57	51	53	21	0.3745617	17.06805	14.984870	
58	56	47	39	0.2171539	16.08863	16.721725	
59	58	57	60	0.0000000	NaN	16.088634	

En los 3 algoritmos se observa que existen ciudades que son "atípicas". En el caso de vecino más cercano, Los Ángeles se une a último momento conformando dos grupos 1 de 59 y uno de 1. En el caso del vecino más lejano, se conforman 2 grupos uno con 58 observaciones y uno con 2 (San Francisco y Los Ángeles).

En el caso del método de Ward si bien se pueden conformar 6 grupos se observa que San Francisco y Los Ángeles conforman un grupo en si mismo, mientras que York forma parte de un grupo de de 5 observaciones pero es la última observación en unirse.

Análisis de Método de Ward





Frecuencia. Cantidad de observaciones en cada grupo.

1	2	3	4	5	6
27	5	9	12	5	2

GRUPO 1

```
[1] "akronOH" "albanyNY" "bostonMA" "bridgeCT" "cantonOH" "clevelOH"
"colombOH" "dallasTX" "daytonOH" "flintMI" "ftwortTX" "grndraMI"
[13] "hartfdCT" "indianIN" "kansasMO" "miamiFL" "newhvnCT"
"portldOR" "rochtrNY" "seattleWA" "springMA" "syracuNY" "toledoOH"
"uticaNY"
[25] "wilmtnde" "worctrMA" "youngsoH"
```

GRUPO 2

```
[1] "allenPA" "lancasPA" "provdCRI" "readngPA" "yorkPA"
```

GRUPO 3

```
[1] "atlantGA" "birmhmAL" "chatagTN" "grnborNC" "houstnTX" "memphsTN"
"nashvltN" "neworlLA" "richmdVA"
```

GRUPO 4

```
[1] "baltimMD" "bufaloNY" "chicagIL" "cinnciOH" "detrotMI"
"louisvKY" "milwauWI" "newyrkNY" "philadPA" "pittsbPA" "stlousMO"
"washDC"
```

GRUPO 5

```
[1] "denverCO" "minnplMN" "sandigCA" "sanjosCA" "wichtaKS"
```

GRUPO 6

```
[1] "losangCA" "sanfrnCA"
```

[1] "Variable 1 : **Rainfall**"

GRUPOS[, 8]: 1

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
30.00	35.00	37.00	38.48	43.00	60.00

GRUPOS[, 8]: 2

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
41.0	42.0	42.0	42.4	43.0	44.0

GRUPOS[, 8]: 3

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
42.00	45.00	47.00	48.11	52.00	54.00

GRUPOS[, 8]: 4

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
30.00	32.50	36.00	36.50	41.25	43.00

GRUPOS[, 8]: 5

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
10.0	13.0	15.0	18.2	25.0	28.0

GRUPOS[, 8]: 6

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
11.00	12.75	14.50	14.50	16.25	18.00

[1] "Variable 2 : **Education**"

GRUPOS[, 8]: 1

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
10.30	10.95	11.30	11.29	11.50	12.20

GRUPOS[, 8]: 2

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
9.0	9.5	9.6	9.6	9.8	10.1

GRUPOS[, 8]: 3

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
9.60	10.10	10.40	10.43	11.00	11.40

GRUPOS[, 8]: 4

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
9.60	10.12	10.55	10.57	10.82	12.30

GRUPOS[, 8]: 5

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
12.10	12.10	12.10	12.14	12.20	12.20

GRUPOS[, 8]: 6

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
12.10	12.12	12.15	12.15	12.18	12.20

[1] "Variable 3 : **Popden**"

GRUPOS[, 8]: 1

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1441	3097	3387	3409	4236	4923

GRUPOS[, 8]: 2

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
3214	3508	4260	5105	4843	9699

GRUPOS[, 8]: 3

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
2082	2302	3125	2910	3325	3768

GRUPOS[, 8]: 4

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
2934	4381	5234	5246	6202	7462

GRUPOS[, 8]: 5

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
2095	2702	3033	3264	3665	4824

GRUPOS[, 8]: 6

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
4253	4365	4476	4476	4588	4700

[1] "Variable 4 : **Nonwhite**"

GRUPOS[, 8]: 1

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1.000	4.400	8.800	8.463	12.500	15.600

GRUPOS[, 8]: 2

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.80	2.20	2.70	2.68	2.90	4.80

GRUPOS[, 8]: 3

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
21.00	22.20	27.10	27.69	31.40	38.50

GRUPOS[, 8]: 4

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
5.80	10.50	14.45	14.71	17.27	25.90

GRUPOS[, 8]: 5

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
2.00	3.00	4.70	4.62	5.90	7.50

GRUPOS[, 8]: 6

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
7.800	9.275	10.750	10.750	12.220	13.700

[1] "Variable 5 : **NOX**"

GRUPOS[, 8]: 1

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1.00	3.00	4.00	7.63	9.50	32.00

GRUPOS[, 8]: 2

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
4.0	6.0	7.0	7.2	8.0	11.0

GRUPOS[, 8]: 3

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
3.00	8.00	9.00	12.67	17.00	32.00

GRUPOS[, 8]: 4

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
12.00	25.25	30.00	32.83	37.25	63.00

GRUPOS[, 8]: 5

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
2.0	8.0	11.0	23.8	32.0	66.0

GRUPOS[, 8]: 6

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
171	208	245	245	282	319

[1] "Variable 6 : **SO2**"

GRUPOS[, 8]: 1

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1.00	9.00	18.00	22.59	36.00	64.00

GRUPOS[, 8]: 2

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
18.0	32.0	33.0	44.2	49.0	89.0

GRUPOS[, 8]: 3

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1.00	5.00	27.00	32.22	48.00	78.00

GRUPOS[, 8]: 4

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
37.0	106.5	135.5	150.9	196.2	278.0

GRUPOS[, 8]: 5

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1.0	3.0	20.0	15.6	26.0	28.0

GRUPOS[, 8]: 6

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
86	97	108	108	119	130

[1] "Variable 7 : **Mortality**"

GRUPOS[, 8]: 1

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
860.1	894.8	919.7	923.5	952.6	1004.0

GRUPOS[, 8]: 2

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
844.1	911.8	938.5	920.6	946.2	962.4

GRUPOS[, 8]: 3

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
952.5	971.1	1006.0	1007.0	1026.0	1113.0

GRUPOS[, 8]: 4

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
929.2	966.4	990.3	989.1	1005.0	1071.0

GRUPOS[, 8]: 5

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
790.7	823.8	839.7	836.7	857.6	871.8

GRUPOS[, 8]: 6

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
861.8	874.3	886.8	886.8	899.2	911.7

>

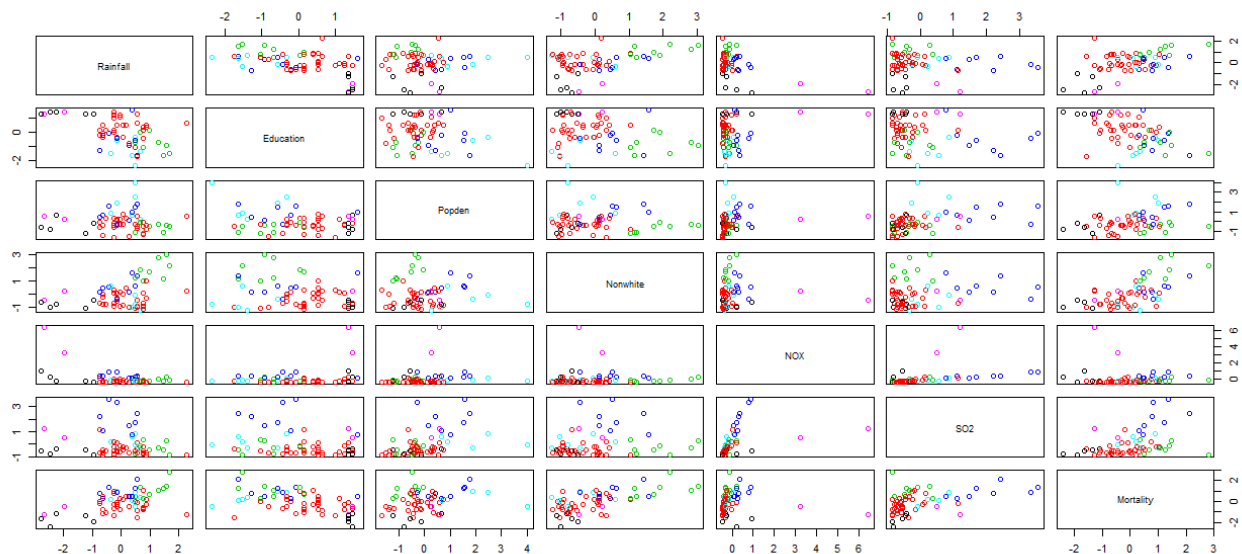
CLASIFICACION NO JERARQUICA

KMEANS

\$size

```
[1] 11 8 11 2 12 16 # con un inicio
```

```
5 31 8 8 6 2 # con 4 inicios
```



\$centers

	Rainfall	Education	Popden	Nonwhite	NOX	SO2	Mortality
1	-1.93580749	1.3918289	-0.4147088	-0.8195338	0.02502962	-0.6071694	-1.6803149
2	0.13564228	0.2550190	-0.3445561	-0.3922458	-0.32096192	-0.4519858	-0.2897502
3	1.11182682	-0.7883717	-0.6359556	1.8826670	-0.19642814	-0.2806501	1.1840037
4	-0.03703284	-0.4453852	0.8504714	0.5530440	0.37217962	2.0737885	0.9463304
5	0.24913001	-1.3003659	1.6996575	-0.4958651	-0.21003119	0.1627956	0.3466424
6	-2.30950250	1.4037589	0.4203553	-0.1266038	4.83942337	0.8627639	-0.8693482

\$withinss

```
[1] 7.338411 68.539602 15.504540 25.780220 18.360161 6.286985
```

De ayuda de R

The data given by `x` are clustered by the *k*-means method, which aims to partition the points into *k* groups such that the sum of squares from points to the assigned cluster centres is minimized. At the minimum, all cluster centres are at the mean of their Voronoi sets (the set of data points which are nearest to the cluster centre).

The algorithm of Hartigan and Wong (1979) is used by default. Note that some authors use *k*-means to refer to a specific algorithm rather than the general method: most commonly the algorithm given by MacQueen (1967) but sometimes that given by Lloyd (1957) and Forgy (1965). The Hartigan–Wong algorithm generally does a better job than either of those, but trying several random starts (`nstart > 1`) is often recommended. In rare

cases, when some of the points (rows of x) are extremely close, the algorithm may not converge in the “Quick-Transfer” stage, signalling a warning (and returning `ifault = 4`). Slight rounding of the data may be advisable in that case.

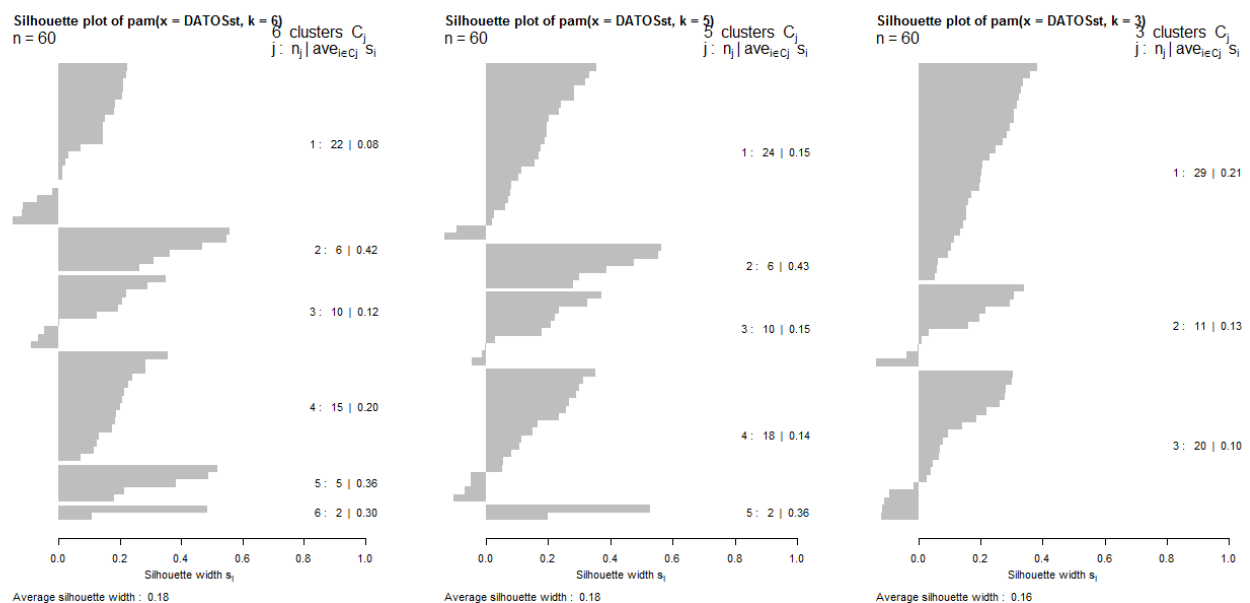
For ease of programmatic exploration, $k=1$ is allowed, notably returning the center and withinss.

Except for the Lloyd–Forgy method, k clusters will always be returned if a number is specified. If an initial matrix of centres is supplied, it is possible that no point will be closest to one or more centres, which is currently an error for the Hartigan–Wong method.

K-MEDOIDES

Medoids:

	ID	Rainfall	Education	Popden	Nonwhite	NOX	SO2	Mortality
youngsOH	60	0.06396581	-0.3260856	-0.2858030	-0.01921666	-0.2100312	-0.2349136	0.2272317
memphsTN	31	1.27594963	-0.6839845	-0.2541275	2.80676205	-0.1012067	-0.3144554	1.0636474
philadPA	39	0.46796042	-0.5646849	1.5327870	0.63641040	0.2035017	1.7059075	1.2095339
hartfdCT	24	0.56895907	0.6283113	-0.6590237	-0.52789282	-0.4276801	-0.6962563	-0.8571910
sanjosCA	49	-2.46100048	1.4634087	-0.8015637	-1.00265724	0.2035017	-0.8076149	-2.4262810
sanfrnCA	48	-1.95600722	1.4634087	0.2664534	0.20686164	3.2288215	0.5127798	-0.4649185



De ayuda de R

The basic `pam` algorithm is fully described in chapter 2 of Kaufman and Rousseeuw(1990). Compared to the k -means approach in `kmeans`, the function `pam` has the following features: (a) it also accepts a dissimilarity matrix; (b) it is more robust because it minimizes a sum of dissimilarities instead of a sum of squared euclidean distances; (c) it provides a novel graphical display, the silhouette plot (see `plot.partition`) (d) it allows to select the number of clusters using `mean(silhouette(pr))` on the result `pr <- pam(..)`, or directly its component `pr$silinfo$avg.width`, see also [pam.object](#).

When `cluster.only` is true, the result is simply a (possibly named) integer vector specifying the clustering, i.e., `pam(x,k, cluster.only=TRUE)` is the same as `pam(x,k)$clustering` but computed more efficiently.

The `pam`-algorithm is based on the search for k representative objects or medoids among the observations of the dataset. These observations should represent the structure of the data. After finding a set of k medoids, k clusters are constructed by assigning each observation to the nearest medoid. The goal is to find k representative objects which minimize the sum of the dissimilarities of the observations to their closest representative object.

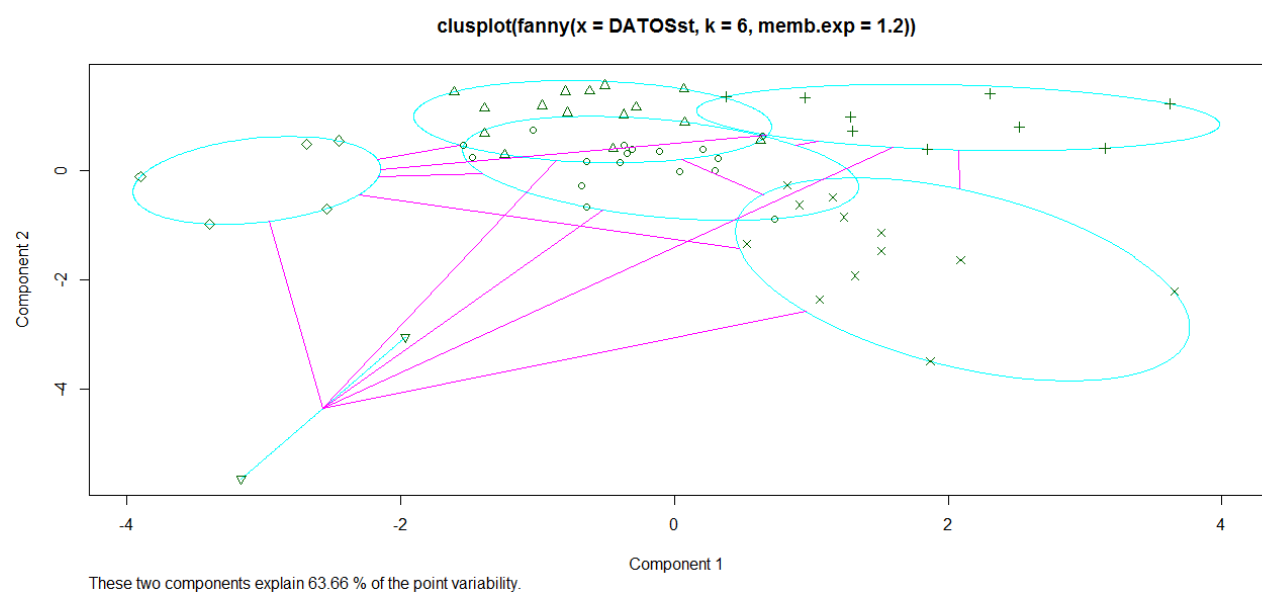
By default, when medoids are not specified, the algorithm first looks for a good initial set of medoids (this is called the **build** phase). Then it finds a local minimum for the objective function, that is, a solution such that there is no single switch of an observation with a medoid that will decrease the objective (this is called the **swap** phase).

When the medoids are specified, their order does *not* matter; in general, the algorithms have been designed to not depend on the order of the observations.

The `pamonce` option, new in cluster 1.14.2 (Jan. 2012), has been proposed by Matthias Studer, University of Geneva, based on the findings by Reynolds et al. (2006).

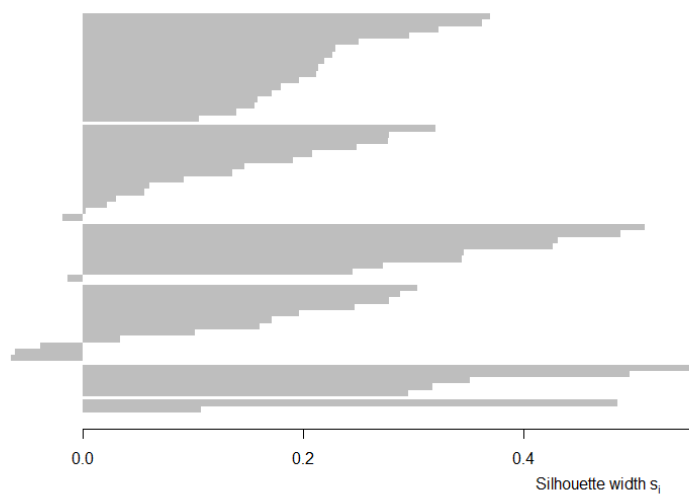
The default FALSE (or integer 0) corresponds to the original “swap” algorithm, whereas `pamonce = 1` (or TRUE), corresponds to the first proposal and `pamonce = 2` additionally implements the second proposal as well.

FUZZY



Silhouette plot of fanny(x = DATOSst, k = 6, memb.exp = 1.2)

n = 60



6 clusters C_j
 $j: n_j | \text{ave}_{i \in C_j} s_i$

1: 17 | 0.22

2: 15 | 0.14

3: 9 | 0.34

4: 12 | 0.13

5: 5 | 0.40

6: 2 | 0.30

Average silhouette width: 0.22

Coefficientes de pertenencia

```
round(A$membership, 2)
      [,1] [,2] [,3] [,4] [,5] [,6]
akronOH  0.93 0.07 0.00 0.00 0.00 0.00
albanyNY  0.88 0.09 0.00 0.02 0.00 0.00
allenPA   0.14 0.78 0.02 0.06 0.00 0.00
atlantGA  0.01 0.00 0.99 0.00 0.00 0.00
baltimMD  0.01 0.00 0.02 0.96 0.00 0.00
birmhmAL  0.00 0.00 0.99 0.00 0.00 0.00
bostonMA  0.83 0.13 0.00 0.02 0.02 0.00
bridgeCT  0.01 0.99 0.00 0.00 0.00 0.00
bufaloNY  0.21 0.06 0.02 0.69 0.01 0.00
cantonOH  0.36 0.63 0.00 0.00 0.00 0.00
chatagTN  0.00 0.01 0.99 0.00 0.00 0.00
chicagIL  0.01 0.00 0.01 0.97 0.00 0.00
cinnciOH  0.02 0.01 0.01 0.97 0.00 0.00
clevelOH  0.92 0.03 0.02 0.02 0.00 0.00
colombOH  0.98 0.01 0.00 0.00 0.00 0.00
dallasTX  0.27 0.41 0.02 0.01 0.29 0.00
daytonOH  0.99 0.01 0.00 0.00 0.00 0.00
denverCO  0.01 0.00 0.00 0.00 0.99 0.00
detroitMI 0.06 0.01 0.01 0.92 0.00 0.00
flintMI   0.90 0.09 0.00 0.00 0.00 0.00
ftwortTX  0.35 0.48 0.01 0.00 0.16 0.00
```

De la ayuda del R

Fanny aims to minimize the objective function

$$\sum_{v=1..k} (\sum_{i,j} u(i,v)^r u(j,v)^r d(i,j)) / (2 \sum_j u(j,v)^r)$$

where n is the number of observations, k is the number of clusters, r is the membership exponent `memb.exp` and $d(i,j)$ is the dissimilarity between observations i and j .

Note that $r \rightarrow 1$ gives increasingly crisper clusterings whereas $r \rightarrow \infty$ leads to complete fuzzyness. K&R(1990), p.191 note that values too close to 1 can lead to slow convergence. Further note that even the default, $r = 2$ can lead to complete fuzzyness, i.e., memberships $u(i,v) == 1/k$. In that case a warning is signalled and the user is advised to chose a smaller `memb.exp` ($=r$).

Compared to other fuzzy clustering methods, fanny has the following features: (a) it also accepts a dissimilarity matrix; (b) it is more robust to the spherical cluster assumption; (c) it provides a novel graphical display, the silhouette plot (see [plot.partition](#)).

SCRIPT R

#lee datos de R

```
pol<-source("chap2airpoll.dat")$value
```

```
summary(pol)
```

```
#####  
###CLUSTER JERARQUICOS###  
#####
```

##cargar la biblioteca 'cluster' es necesario porque se necesita la funcion 'agnes'

```
library(cluster)
```

##carga las funciones necesarias para TIPIFICAR y OBTENER LOS PSEUDO-INDICADORES

```
source('cluster/standard.R') #ojo con la ubicacion  
source('cluster/indicadores.R') #ojo con la ubicacion
```

1

LECTURA DE DATOS carga en DATOS

DATOS.txt _ nombre del archivo que ud quiere importar

header=T _ si las columnas(variables) traen sus nombres en la primera fila

sep = '\t' _ si el archivo de texto viene separado por TABULADORES, las otras opciones son ESPACIO o COMA

quote = ' ' _ que identifica texto o caracter en el archivo a importar

dec= '.' _ que separa decimales en el archivo.

```
DATOS = pol
```

2

##Tipifica los datos de DATOS y los carga en DATOSst

```
DATOSst <- standard(DATOS)
```

3

Ejecuta la funcion agnes sobre los datos tipificados y los carga en AGRUPO

en general aca los unico que se modificara sera: method = 'ward', 'single', 'complete', 'average'.

#Metodo Vecino mas cercano

```
AGRUPO <- agnes(DATOSst, metric = "euclidean", stand = FALSE, method = "single")
```


4

Ejecuta la funcion indicadores y guarda en IND

aca se pueden tocar todos los parametros:

AGRUPO[4] _ es el elemento 4 o \$merge del objeto generado en el paso anterior a traves de agnes

DATOSst _ es el conjunto de datos sobre los que ejecute agnes

imprime _ es el numero de pasos de la jerarquia que queremos ver impresos

IND <- indicadores(AGRUPPO[4],DATOSst,imprime=20)

grafica un dendograma a partir de AGRUPO (objeto de tipo agnes)

plot(AGRUPPO,which=2)

#Metodo Vecino mas Lejano

AGRUPPO1 <- agnes(DATOSst, metric = "euclidean", stand = FALSE, method = "complete")

IND <- indicadores(AGRUPPO1[4],DATOSst,imprime=20)

grafica un dendograma a partir de AGRUPO (objeto de tipo agnes)

plot(AGRUPPO1,which=2)

#Metodo Ward

AGRUPPO2 <- agnes(DATOSst, metric = "euclidean", stand = FALSE, method = "ward")

IND <- indicadores(AGRUPPO2[4],DATOSst,imprime=20)

grafica un dendograma a partir de AGRUPO (objeto de tipo agnes)

plot(AGRUPPO2,which=2)

Grafica R2, pseudo F, pseudot

a=c(21:1)

IND1=cbind(IND[,4:6],a)

x=seq(1,21)

plot(a,IND1[,1], type='l', col='red', xaxt="n", xlab='Numero grupos', ylab='R cuadrado',main='Rcuadrado de Metodo Ward')

```
axis(1, at=x, labels=x, las=0)
```

```
plot(a, IND1[,2], type='l', col='blue', xaxt="n", xlab='Numero grupos', ylab='pseudo F', main='Valores PseudoF de Metodo Ward')
```

```
axis(1, at=x, labels=x, las=0)
```

```
plot(a, IND1[,3], type='l', col='green', xaxt="n", xlab='Numero grupos', ylab='pseudo t', main='Valores de Pseudot de Metodo Ward')
```

```
axis(1, at=x, labels=x, las=0)
```

#grafica silhouette plot

```
par(mfrow=c(1,3))
```

```
k <- 6
```

```
clw <- cutree(AGRUP02[4], k)
```

```
ss = silhouette(clw, daisy(DATOSst))
```

```
plot(ss)
```

```
k1 <- 5
```

```
clw1 <- cutree(AGRUP02[4], k1)
```

```
ss1 = silhouette(clw1, daisy(DATOSst))
```

```
plot(ss1)
```

```
k3 <- 3
```

```
clw3 <- cutree(AGRUP02[4], k3)
```

```
ss3 = silhouette(clw3, daisy(DATOSst))
```

```
plot(ss3)
```

#Numero grupos

```
k <- 6
```

```
clw <- cutree(AGRUP02[4], k)
```

guarda en GRUPOS la union de los DATOS y las MEMBRESIAS

```
GRUPOS <- cbind(DATOS, clw)
```

```
##### 4 #####
```

```
table(GRUPOS[,8])
```

```
row.names(GRUPOS[GRUPOS[,8]==1,])
```

```
row.names(GRUPOS[GRUPOS[,8]==2,])
```

```
row.names(GRUPOS[GRUPOS[,8]==3,])
```

```
row.names(GRUPOS[GRUPOS[,8]==4,])
```

```

row.names(GRUPOS[GRUPOS[,8]==5,])
row.names(GRUPOS[GRUPOS[,8]==6,])
by(GRUPOS[,1:7],GRUPOS[,8],summary)

for (j in 1:(ncol(GRUPOS)-1))
{
  cat(' ', '\n')
  print(paste('Variable',j,':',names(GRUPOS)[j]))
  print(by(GRUPOS[,j],GRUPOS[,8],summary))
  cat(' ', '\n')
  cat('*****')
  cat(' ', '\n')
}

```

#CLASIFICACION NO JERARQUICA

#kmeans

```

nojer=kmeans(DATOSst,6,nstart=4)
plot(DATOSst, col = nojer$cluster)
points(nojer$centers, col = 1:7, pch = 8, cex = 2)

```

```

par(mfrow=c(1,3))
pp=pam(DATOSst,6)
si=silhouette(pp)
plot(si)

```

#kmedoides

```

pp1=pam(DATOSst,5)
si1=silhouette(pp1)
plot(si1)

```

```

pp2=pam(DATOSst,3)
si2=silhouette(pp2)
plot(si2)

```

Fuzzy

```

aa=fanny(DATOSst, 6, memb.exp=1.2)
summary(A)
plot(A)

```