

8281_Application

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```
library("dtwclust")
data("uciCT")
```

Character Trajectories Data Set

20 characters: A B C D E G H L M N O P Q R S U V W Y Z

5 samples for each

```
length(CharTraj)
```

```
## [1] 100
```

```
names(CharTraj)
```

```
## [1] "A. V1" "A. V2" "A. V3" "A. V4" "A. V5" "B. V1" "B. V2" "B. V3" "B. V4" "B. V5"
## [11] "C. V1" "C. V2" "C. V3" "C. V4" "C. V5" "D. V1" "D. V2" "D. V3" "D. V4" "D. V5"
## [21] "E. V1" "E. V2" "E. V3" "E. V4" "E. V5" "G. V1" "G. V2" "G. V3" "G. V4" "G. V5"
## [31] "H. V1" "H. V2" "H. V3" "H. V4" "H. V5" "L. V1" "L. V2" "L. V3" "L. V4" "L. V5"
## [41] "M. V1" "M. V2" "M. V3" "M. V4" "M. V5" "N. V1" "N. V2" "N. V3" "N. V4" "N. V5"
## [51] "O. V1" "O. V2" "O. V3" "O. V4" "O. V5" "P. V1" "P. V2" "P. V3" "P. V4" "P. V5"
## [61] "Q. V1" "Q. V2" "Q. V3" "Q. V4" "Q. V5" "R. V1" "R. V2" "R. V3" "R. V4" "R. V5"
## [71] "S. V1" "S. V2" "S. V3" "S. V4" "S. V5" "U. V1" "U. V2" "U. V3" "U. V4" "U. V5"
## [81] "V. V1" "V. V2" "V. V3" "V. V4" "V. V5" "W. V1" "W. V2" "W. V3" "W. V4" "W. V5"
## [91] "Y. V1" "Y. V2" "Y. V3" "Y. V4" "Y. V5" "Z. V1" "Z. V2" "Z. V3" "Z. V4" "Z. V5"
```

1-nearest-neighbor classifier

```
train <- CharTraj[-66L]
test <- CharTraj[66L]
d <- proxy::dist(train, test, method = "dtw", window.size = 10L)
cbind(test_data = names(CharTraj)[66], nearest_neighbor = rownames(d)[which.min(d)])
```

```
## test_data nearest_neighbor
## [1,] "R. V1" "R. V2"
```

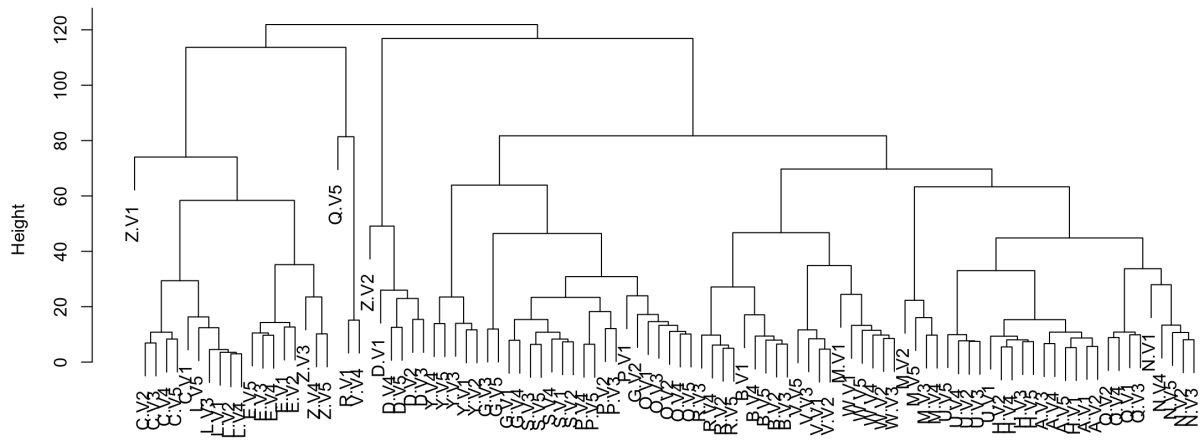
Hierarchical Clustering

```
hc_dtw <- tsclust(CharTraj, type = 'hierarchical', k = 20L, distance = 'dtw',
  control = hierarchical_control(method = "average"),
  args = tsclust_args(dist = list(window.size = 18L)))
```

```
## Warning in tsclust(CharTraj, type = "hierarchical", k = 20L, distance =
## "dtw", : Distance matrix is not symmetric, and hierarchical clustering
## assumes it is (it ignores the upper triangular).
```

```
##plot the dendrogram
plot(hc_dtw, type = "dendrogram")
```

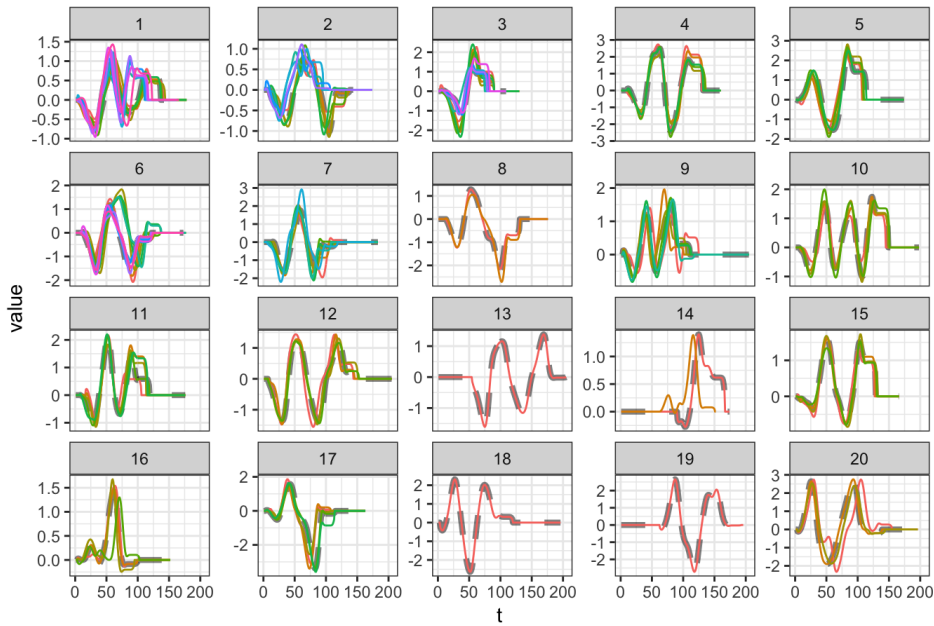
Cluster Dendrogram



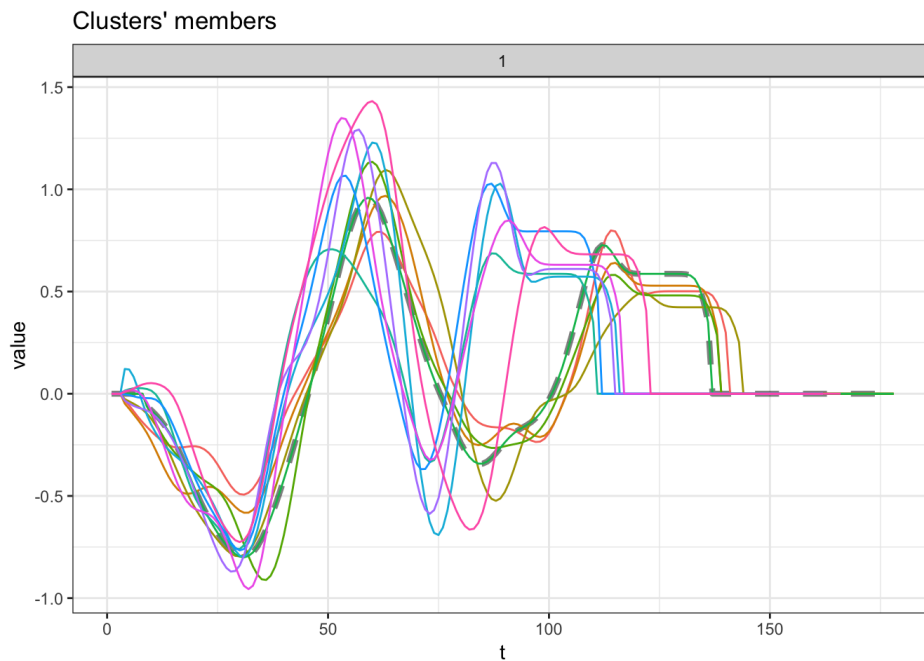
```
stats::as.dist(distmat)
stats::hclust (*, "average")
```

```
##plot the series and centroids
plot(hc_dtw, type = "sc")
```

Clusters' members



```
##zoom in the first cluster
plot(hc_dtw, type = "sc", clus = 1L)
```



```
##check the ground labels and the resulted clusters
matrix(hc_dtw@cluster, ncol = 5, byrow = T)
```

```
##      [,1] [,2] [,3] [,4] [,5]
## [1,]  1   1   1   1   1
## [2,]  2   2   2   2   2
## [3,]  3   3   3   3   3
## [4,]  4   4   4   4   4
## [5,]  5   5   5   5   5
## [6,]  6   7   8   6   8
## [7,]  1   1   1   1   1
## [8,]  3   3   3   3   3
## [9,]  9  10  10  10  10
## [10,] 11  11  11  11  11
## [11,]  7   7   7   7   7
## [12,]  7   6   6   6   6
## [13,] 12  12  12  12  13
## [14,] 14   2   2   2   2
## [15,]  6   6   6   6   6
## [16,]  1  15  15  15  15
## [17,] 16  16  16  14  16
## [18,]  9   9   9   9   9
## [19,] 17  17  17  17  17
## [20,] 18  19  20  20  20
```

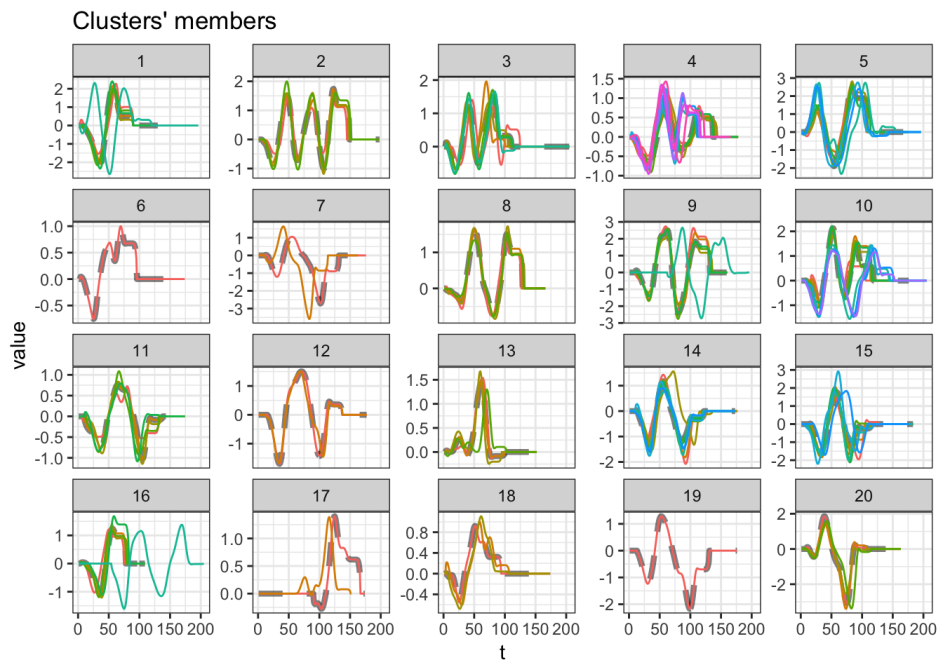
The 1st character (A) and the 7th character (H) are clustered into one. Let's apply the nearest-neighbor classifier again to see which has the closest DTW distance with the 7th character

```
d2 <- proxy::dist(CharTraj[-c(31:35)], CharTraj[31:35], method = "dtw", window.size = 10L)
which.min(apply(matrix(apply(d2, 1, mean), ncol = 5, byrow = T), 1, mean))
```

```
## [1] 1
```

Partitional Clustering

```
pc_dtw <- tsclust(CharTraj, type = 'partitional', k = 20L, distance = 'dtw', centroid = "pam",
  seed = 8,
  args = tsclust_args(dist = list(window.size = 18L)))
##plot the series and centroids
plot(pc_dtw, type = "sc")
```



```
##check the ground labels and the resulted clusters
matrix(pc_dtw@cluster, ncol = 5, byrow = T)
```

```
##      [,1] [,2] [,3] [,4] [,5]
## [1,]   4   4   4   4   4
## [2,]  11  11  11  11  11
## [3,]   1   1   1   1   1
## [4,]   9   9   9   9   9
## [5,]   5   5   5   5   5
## [6,]  14  15  19  14   7
## [7,]   4   4   4   4   4
## [8,]  16  16  16  16  16
## [9,]   3   2   2   2   2
## [10,]  10  10  10  10  10
## [11,]  15  15  15  15  15
## [12,]  15  15  14  12  12
## [13,]  10  10  10  10  16
## [14,]  17  18   6  18  18
## [15,]  14  14  14  14  14
## [16,]   4   8   8   8   8
## [17,]  13  13  13  17  13
## [18,]   3   3   3   3   3
## [19,]  20  20  20  20   7
## [20,]   1   9   5   5   5
```

```
##evaluate the two algorithms using "VI" values
sapply(list(DTW = pc_dtw, hc_dtw),
       cvi, b = CharTrajLabels, type = "VI")
```

```
##      DTW.VI      VI
## 0.3634873 0.3018201
```