



Machine Learning and Data Mining 5DATA001C.2

Course work Report

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Partitioning Clustering Part

Objectives/Deliverables (partitioning clustering)

1st Subtask Objectives:

(a)

Scaling: - Scaling is the process of converting a dataset's numerical features to a standard scale. Due of the sensitivity of many machine learning algorithms to the magnitude of the input features, this is done. Results may be skewed if the features are not weighted equally; when this happens, some features will predominate over others. This bias can be avoided and the machine learning algorithm's performance can be enhanced by scaling the features. Scaling can also hasten the optimization process during training, hastening convergence and improving outcomes.

Outliers Detection: - Data points known as outliers differ significantly from other data points in the dataset. They may appear for a number of reasons, including measurement errors, data corruption, or just the basic nature of the data itself. Outliers can significantly affect how well machine learning algorithm's function since they can distort the findings and produce false models. In order to improve data collecting and processing procedures, outliers can be used to discover data quality problems and abnormalities.

Removal: - To increase the model's accuracy and resilience, outliers can be eliminated. However, it's crucial to remember that eliminating too many outliers can also result in the loss of important data, so a careful balance must be struck. Removing outliers can aid in discovering and fixing problems with data collection and processing, which can assist to improve data quality and reliability. Additionally, eliminating outliers can aid in enhancing the model's interpretability by assisting in the discovery of data patterns and relationships that may have been hidden by the presence of outliers.

(b)

1. NbClust methods.

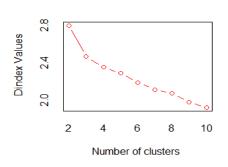
The R package NbClust is used to determine the ideal number of clusters to include in a dataset. It offers a number of indices and techniques, including the Silhouette coefficient, Calinski-Harabasz index, Dunn index, and Gap statistic, to assess clustering solutions. In order to facilitate the selecting process, it also offers graphical representations of the indexes.

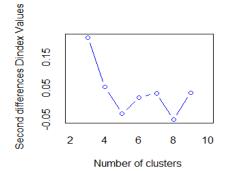
```
#Nbclust method
install.packages("Nbclust")
library(Nbclust)
nb <- Nbclust(vehicles_cleaned, distance = "euclidean", min.nc = 2, max.nc = 10, method = "kmeans")
print(nb)

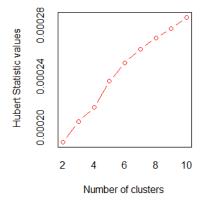
#Nbclust method
install.packages("Nbclust")
print(nb)

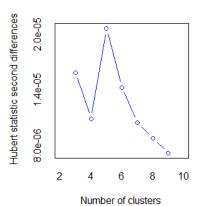
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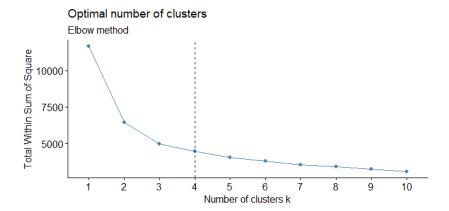


>

2.Elbow methods.

Based on the idea that as the number of clusters increases, the within-cluster sum of squares (WCSS) drops and the between-cluster sum of squares (BCSS) increases, the Elbow approach is a heuristic methodology used to determine the ideal number of clusters in a clustering algorithm. Plotting the WCSS versus the number of clusters and choosing the number of clusters where the rate of WCSS decline begins to level off are both required. The Elbow method can be combined with other techniques to conduct a more thorough study.

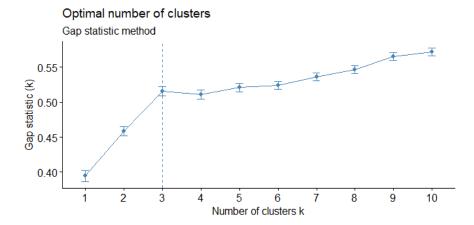
```
#Elbow method
install.packages("factoextra")
library(factoextra)
x11() # creates an X11 graphics device
graphics.off() # reset the graphics device
#plot(x, y) # try plotting again
for iz_nbclust(vehicles_cleaned, kmeans, method = "wss") + geom_vline(xintercept = 4, linetype = 2) + labs(subtitle = "Elbow method")
```



3. Gap statistics methods.

The ideal number of clusters in a dataset can be found using the statistical method known as gap statistics. It compares the overall within-cluster variation with what would be predicted under a null reference distribution for various values of k (number of clusters). The amount of k that maximizes the gap statistic, or the amount of clustering that is most noticeably improved above the null reference distribution, is the optimal number of clusters.

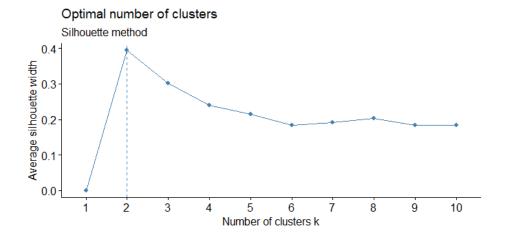
```
4/ # Gap statistic method
48 set.seed(123)
49 fviz_nbclust(vehicles_cleaned, kmeans, nstart = 25, method = "gap_stat", nboot = 50) + labs(subtitle = "Gap statistic method")
50
```



4.silhouette methods.

When comparing an observation to other clusters, the silhouette method, a clustering evaluation technique, determines how similar the observation is to its own cluster. For each observation, it generates a silhouette coefficient that spans from -1 to 1. The overall average is generated for the entire dataset and includes the average silhouette coefficient for each observation in each cluster.

```
# Silhouette method
install.packages("cluster")
library(cluster)
library(factoextra)
fiviz_nbclust(vehicles_cleaned, kmeans, method = "silhouette") + labs(subtitle = "Silhouette method")
```



(c)

Elbow methods are useful for identifying the optimal number of clusters but have limitations, silhouette methods are useful for determining the quality of the clusters, NbClust methods are useful for determining the optimal number of clusters and the best clustering method, and gap statistics methods are useful for identifying the optimal number of clusters but can be computationally intensive. The particular requirements of the investigation and the features of the dataset will determine which clustering validation technique is used.

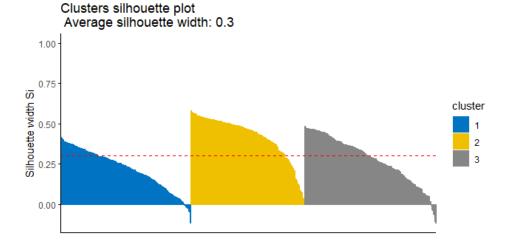
```
58 #compute k-means clustering with k=3
59 set.seed(123)
60 final_stat <- kmeans(vehicles_cleaned,3,nstart = 25)
61 print(final_stat)
62
63 #BSS
64 BSS <- sum(final_stat\size-(colMeans(vehicles_cleaned)-final_stat\centers)^2)
65 cat("BSS:",BSS,"\n")
66
67 #TSS
   TSS <- sum((vehicles_cleaned-colMeans(vehicles_cleaned))^2)
68
69 cat("TSS:",TSS,"\n")
70
71 #WSS
72 WSS <- sum(final_stat$withinss)</pre>
73 cat("WSS:",WSS,"\n")
74
75 #BSS to TSS
76 ratio_BSS_to_TSS <- BSS/TSS
77 cat("ratio_BSS_to_TSS:",ratio_BSS_to_TSS,"\n")
78
79
```

```
> #BSS
> BSS <- sum(final_stat$size-(colMeans(vehicles_cleaned)-final_stat$centers)^2)
> cat("BSS:",BSS,"\n")
BSS: 13579.7
>
> #TSS
> TSS <- sum((vehicles_cleaned-colMeans(vehicles_cleaned))^2)
> cat("TSS:",TSS,"\n")
TSS: 11694.95
>
> #WSS
> WSS <- sum(final_stat$withinss)
> cat("WSS:",WSS,"\n")
WSS: 4975.349
> #BSS to TSS
> ratio_BSS_to_TSS <- BSS/TSS
> cat("ratio_BSS_to_TSS:",ratio_BSS_to_TSS,"\n")
ratio_BSS_to_TSS: 1.161159
> |
```

(d)

ytruyruru

```
80
81 #Silhouette plot
82 pam.res2 <- pam(vehicles_cleaned,3,metric="euclidean",stand = FALSE)
83 fviz_silhouette(pam.res2,palette="jco",ggtheme=theme_classic())
84
```



```
#average Silhouette width score
sil <- silhouette(final_stat$cluster, dist(vehicles_cleaned))
avg_Sil_width <-mean(sil[,3])
cat("average Silhouette width score:",avg_Sil_width,"\n")

> #average Silhouette width score
> sil <- silhouette(final_stat$cluster, dist(vehicles_cleaned))
> avg_Sil_width <-mean(sil[,3])
> cat("average Silhouette width score:",avg_Sil_width,"\n")
average Silhouette width score: 0.3015821
> |
```

2nd Subtask Objectives:

(e) The amount of variance explained by the main components and the complexity of the model must be balanced when deciding how many principal components to keep. In this instance, we decided to keep the eight major components out of the original 18 features that provided a total score of at least 92%. This decision strikes a compromise between the complexity of the modified dataset and the amount of variance that the model can explain.

```
8
9 # Show the eigenvalues and eigenvectors
10 summary(pca_data)
11
```

```
> # Show the eigenvalues and eigenvectors
> summary(pca_data)
    PCA(X = vehicles_cleaned, graph = FALSE)
  Eigenvalues
  Dim.1 Dim.2 Dim.3 Dim.4 Dim.5 Dim.6 Dim.7 Dim.8 Dim.9 Dim.10 Dim.11 Dim.12 Dim.13 Dim.14 Dim.15 Dim.16 Dim.17 Dim.8 Dim.9 Dim.10 Dim.11 Dim.12 Dim.13 Dim.14 Dim.15 Dim.16 Dim.17 Dim.18 Dim.14 Dim.15 Dim.16 Dim.17 Dim.18 Dim.16 Dim.17 Dim.18 Dim.18
   variance
                                                                                          0.000
   % of var.
  Cumulative % of var. 100.000
  Individuals (the 10 first)
                                                                                                                         ctr cos2
0.006 0.070 |
0.030 0.468 |
0.222 0.803 |
                                                                                                                                                                                        Dim.2 ctr cos2
0.630 0.016 0.063 |
0.408 0.007 0.035 |
-0.336 0.005 0.005 |
                                                                                                                                                                                                                                                                                  Dim.3 ctr
0.511 0.029
0.303 0.010
1.199 0.158
                                                                                                                                                                                                                                                                                                                                      cos2
0.041
0.019
0.070
                                                                                       1.497
                                                               2.189
                                                                4.542
                                                               3.648
                                                                                             -1.453
                                                                                                                            0.028
                                                                                                                                                       0.159
                                                                                                                                                                                           3.120
2.318
                                                                                                                                                                                                                    0.388
                                                                                                                                                                                                                                                0.731
                                                                                                                                                                                                                                                                                   0.461
                                                                                                                                                                                                                                                                                                               0.023
                                                                                                                                                                                                                                                                                                                                        0.016
                                                                                                                                                                                                                   0.388
0.214
0.104
0.479
0.179
0.188
0.251
                                                                 3 572
                                                                                             -0 712
                                                                                                                                                                                                                                                                                   1.929
                                                                                                                                                                                                                                                                                                               0.408
                                                                                                                                                                                                                                                                                                                                        0 292
                                                 | 3.572 | -0.712 | 0.007
| 3.173 | -1.924 | 0.050
| 5.710 | -4.357 | 0.254
| 3.185 | 1.617 | 0.035
| 4.185 | -3.300 | 0.146
| 5.615 | -4.437 | 0.264
                                                                                                                                                                                          2.318
1.617
3.466
2.121
2.170
2.511
                                                                                                                                                       0.368
0.582
0.258
                                                                                                                                                                                                                                                0.421
0.260
0.369
0.444
                                                                                                                                                                                                                                                                               1.929
1.079
-0.568
-0.576
-0.279
                                                                                                                                                                                                                                                                                                              0.408
0.128
0.035
0.036
                                                                                                                                                                                                                                                0.269
                                                                                                                                                       0.622 | 0.624 |
  10
  Variables (the 10 first)
  Dim.2 ctr cos2
0.159 0.763 0.025 |
-0.270 2.195 0.073 |
0.073 0.162 0.005 |
                                                                                                                                                                                                                                              Dim. 3
0.031
0.234
-0.076
                                                                                                                                                                                                                                                                          ctr cos2
0.080 0.001
4.527 0.055
0.483 0.006
                                                                                                                                                                                   3.689
                                                                                                                                                                                                             0.122
                                                                                                                                                                                                                                              -0.069
                                                                                                                                                                                                                                                                           0.398
                                                                                                                                                                                                                                                                                                      0.005
                                                                                                                                                                                   6.466
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                                                                                                                                                                                                                                                                           0.569
                                                                                                                                                                                                                                                                                                      0.007
                                                                                                                                                                                  0.667
                                                                                                                                                                                                              0.022
                                                                                                                                                                                                                                                0.170
                                                                                                                                                                                                                                                                          2.405
                                                                                                                                                                                                                                                                                                      0.029
                                                                                                                                                                                  0.640
                                                                                                                                                                                                           0.029 | 0.061 |
                                                                                                                                                                                                                                              -0.124 1.275 0.015
0.257 5.458 0.066
                                                                                                                                                                                1.850
```

cumulative score per principals

```
# Show the cumulative percentage of variance explained
 15
          eig_val <- get_eigenvalue(pca_data)
          eig_val
 17
 18
         cumulative_variances <- cumsum(eig_val/sum(eig_val)*100)
 20 cumulative variances
         barplot(cumulative_variances, main = "Cumulative Percentage of Variance Explained", xlab = "Number of Components", ylab = "Cumulative %")
 23
 # Choose the PCs that provide at least cumulative score > 92%

num_components <- length(cumulative_variances[cumulative_variances > 92])

print(paste("Number of components needed to explain at least 92% of the variance:", num_components))
 28 # Create a transformed data set
 29
         pca_result <- PCA(vehicles_cleaned, ncp = num_components, graph = FALSE)$ind$coord
> # Show the cumulative percentage of variance explained > eig_val <- get_eigenvalue(pca_data) > eig_val
cumulative_variances <- cumsum(eig_val/sum(eig_val)*100)

    mulative_variances

    0.5517772
    0.7372497
    0.8046644
    0.8682667
    0.9191475
    0.9550829
    0.9726719
    0.9852338
    0.9915344
    0.9958870
    0.9992030

    1.0030547
    1.0030547
    1.0031077
    1.0051824
    1.0061332
    1.0061332
    1.001831
    5.1019851
    5.4765112
    5.8295568
    6.1125284

    6.4098861
    6.4796743
    6.5146778
    6.5388585
    6.5572105
    6.5703768
    6.5786795
    6.582105
    6.5900846
    6.938276
    6.997818

    9.6613216
    13.7571535
    18.2275115
    23.0512151
    28.1575903
    33.4636064
    38.8673394
    44.3408605
    49.8493852
    55.3820905
    60.9332184

    72.0699683
    77.6490257
    83.2329572
    88.8206316
    94.4102603
    100.0000000
    88.8673394
    44.3408605
    49.8493852
    55.3820905
    60.9332184

  > barplot(cumulative_variances, main = "Cumulative Percentage of Variance Explained", xlab = "Number of Components", ylab = "Cumulative %")
 >> # Choose the PCs that provide at least cumulative score > 92% 
> num_components <- length(cumulative_variances[cumulative_variances > 92]) 
> print(paste("Number of components needed to explain at least 92% of the variance:", num_components)) 
[1] "Number of components needed to explain at least 92% of the variance: 2"
> # Create a transformed data set
> pca_result <- PCA(vehicles_cleaned, ncp = num_components, graph = FALSE)$ind$coord
```

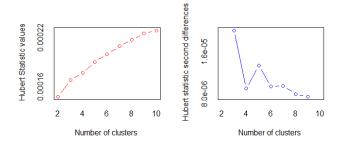
1.NbClust methods.

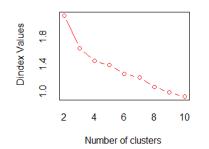
```
34 #Nbclust method
35 library(Nbclust)
36 nb <- Nbclust(pca_result, distance = "euclidean", min.nc = 2, max.nc = 10, method = "kmeans")
37 print(nb)
```

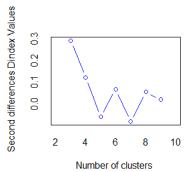
```
> nb <- Nbclust(pca_result, distance = "euclidean", min.nc = 2, max.nc = 10, method = "kmeans")
****: The Hubert index is a graphical method of determining the number of clusters.
In the plot of Hubert index, we seek a significant knee that corresponds to a
significant increase of the value of the measure i.e the significant peak in Hubert
index second differences plot.
     ***: The D index is a graphical method of determining the number of clusters.

In the plot of D index, we seek a significant knee (the significant peak in Dindex second differences plot) that corresponds to a significant increase of the value of the measure.
     **** Conclusion ****
     * According to the majority rule, the best number of clusters is 3
     Warning message:
did not converge in 10 iterations
> print(nb)
| Sprint(nb) | Spr
     SAll.index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          houette Duda Pseudot2 Beale 0.5159 0.5500 428.7110 0.8165 0.4646 1.6603 -159.0779 -0.3958 0.4182 1.5443 -131.4722 -0.3506 0.4017 0.8884 1.3379 -56.5676 -0.2508 0.3608 0.3703 2.9648 -131.8793 -0.6549 0.3677 2.8925 -100.1042 -0.6468 0.3706 2.3506 -103.9981 -0.5678
     $All.CriticalValues
                  CritValue_Duda CritValue_PseudoT2 Fvalue_Beale
0.5713 393.1612 0.4423
0.5134 379.1353 1.0000
                                                                                                                                        393.1612
379.1353
366.3245
     2
                                                                                                                                                                                                                  1.0000
                                                         0.5012
```

5 0.5012 6 0.4760 7 0.4913 8 0.4156 9 0.4186 10 0.4156	233.8476 246.5460 211.2269 279.8445 212.4822 254.5320	0.8602 1.0000 0.7760 1.0000 1.0000				
Value_Index 5.2637 1100	tBiserial Frey 2.0000 1	3.0000 3.0000 5.1869 774.2957	3 4	3.000 10.0000 6 136.971 5.1311 -0		e Duda PseudoT2 Beale Ratkowsky 0 3.0000 3.0000 2.0000 3.0000 9 1.6603 -159.0779 0.8165 0.4722
40 41 42 43 44 45 43 44 45 43 2 1 1 1 3 79 80 81 82 83 84 81 3 1 3 1 2 3 3 1 1 3 1 2 3 3 1 1 2 1 3 1 3 1 1 3 1 1 3 1 1 3 1 1 3 1 1 3 1 1 3 1 1 3 1 1 3 1 1 3 1 1 3 1 3 1 1 3 1 3 1 1 3 1 3 1 1 3 1 3 1 1 3 1 3 1 3 1 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3 1 3 3	3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3	3 3 3 2 2 3 3 3 2 9 3 9 9 9 9 1 92 1 2 1 3 3 2 6 1 4 5 1 5 1 5 1 5 1 5 1 5 1 5 1 5 1 5 1	3	1 1 3 3 2 2 1 3 3 1 1 1 1 3 1 1 1 1 1 3 1 3	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	1 3 3 3 1 1 3 2 2 3 8 149 150 151 152 153 13 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3







2.Elbow methods.

```
#Elbow method

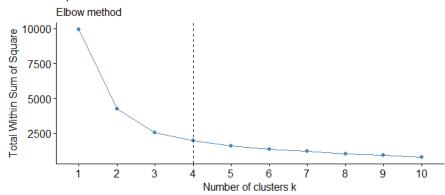
xil() # creates an Xil graphics device

graphics.off() # reset the graphics device

#plot(x, y) # try plotting again

fviz_nbclust(pca_result, kmeans, method = "wss") + geom_vline(xintercept = 4, linetype = 2) + labs(subtitle = "Elbow method")
```

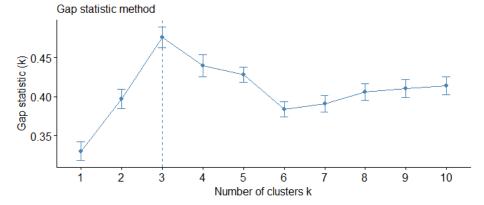
Optimal number of clusters



3.Gap statistics methods

```
45 # Gap statistic method
46 set.seed(123)
47 fviz_nbclust(pca_result, kmeans, nstart = 25, method = "gap_stat", nboot = 50) + labs(subtitle = "Gap statistic method")
48
```

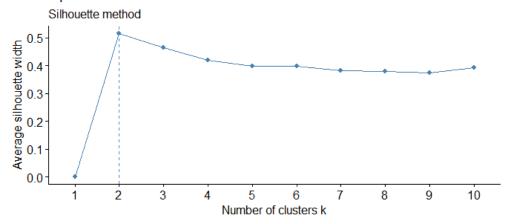
Optimal number of clusters



4.silhouette methods.

```
49
50 # Silhouette method
51 library(cluster)
52 library(factoextra)
53 fviz_nbclust(pca_result, kmeans, method = "silhouette") + labs(subtitle = "Silhouette method")
```

Optimal number of clusters



(g)

```
55 #compute k-means clustering with k=3
56  set.seed(123)
57  final_stat_pca <- kmeans(pca_result,3,nstart = 25)</pre>
58 print(final_stat_pca)
59
60 #BSS
61 BSS_pca <- sum(final_stat_pca\size-(colMeans(pca_result)-final_stat_pca\scenters)\^2)
62 cat("BSS:",BSS_pca,"\n")
63
64
65 TSS_pca <- sum((pca_result-colMeans(pca_result))^2)
66 cat("TSS:",TSS_pca,"\n")
67
68 #WSS
69 WSS_pca <- sum(final_stat_pca$withinss)</pre>
70 cat("WSS:",WSS_pca,"\n")
71
72 #BSS to TSS
73 ratio_BSS_to_TSS_pca <- BSS_pca/TSS_pca
74 cat("ratio_BSS_to_TSS:",ratio_BSS_to_TSS_pca,"\n")
```

```
**Compared & segment Clustering with %-1
**Installation & segment (Clustering with %-1)
**Installation & segment (Clust
```

(h)

> WSS_pca <- sum(final_stat_pca\$withinss)

> ratio_BSS_to_TSS_pca <- BSS_pca/TSS_pca

> cat("ratio_BSS_to_TSS:",ratio_BSS_to_TSS_pca,"\n")

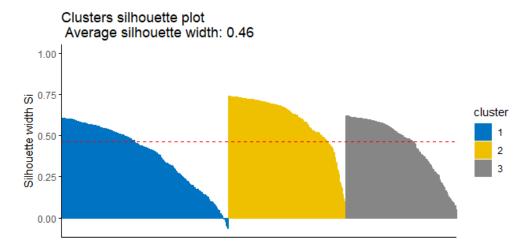
> cat("WSS:",WSS_pca,"\n")

ratio_BSS_to_TSS: 0.148429

WSS: 2530.801

```
86 #Silhouette plot
 87 library(cluster)
88 pam.res2 <- pam(pca_result,3,metric="euclidean",stand = FALSE)
 89 fviz_silhouette(pam.res2,palette="jco",ggtheme=theme_classic())
 90
 91 #average Silhouette width score
 92 sil <- silhouette(final_stat_pca$cluster, dist(pca_result))
 93 avg_Sil_width <-mean(sil[,3])
 94 cat("average Silhouette width score:",avg_Sil_width,"\n")
95
> pam.res2 <- pam(pca_result,3,metric="euclidean",stand = FALSE)
> fviz_silhouette(pam.res2,palette="jco",ggtheme=theme_classic())
 cluster size ave.sil.width
       1 319
2 225
                        0.38
1
2
                        0.60
       3 212
                       0.45
3
```

```
> #average Silhouette width score
> sil <- silhouette(final_stat_pca$cluster, dist(pca_result))
> avg_Sil_width <-mean(sil[,3])
> cat("average Silhouette width score:",avg_Sil_width,"\n")
average Silhouette width score: 0.4645933
```



(i) Calinski-Harabasz Index

```
87 library(fpc)
88 ch_index <- calinhara(pca_result, final_stat_pca$cluster)
89 print(ch_index)
90 barplot(ch_index, main="Calinski-Harabasz Index for K-Means Clustering", xlab="Number of Clusters", ylab="Calinski-Harabasz Index")
91 plot(ch_index, type="b", xlab="Number of Clusters", ylab="Calinski-Harabasz:Index")
```

```
> library(fpc)
> ch_index <- calinhara(pca_result, final_stat_pca$cluster)
> print(ch_index)
[1] 1106.878
```

Energy Forecasting Part (part of Work Based Learning Activity)

Objectives/Deliverables (Multi-layers Neural Network)

1st Subtask Objectives:

(a)

Time-based Approach: - Time-based characteristics can be an effective tool for defining the input vector when dealing with predicting issues for electricity load. By incorporating elements like the day of the week, the hour of the day, and the month of the year, the neural network can detect seasonality and temporal trends in the data. If the dataset contains hourly electricity consumption data for several years, the neural network, for example, can use time-based characteristics to distinguish between weekday and weekend patterns, morning and evening peaks, and seasonal swings in demand.

Calendar-based Approach: - In this method, the input variables are calendar elements like the day of the week, the month of the year, and the holiday indicators. This strategy acknowledges that the amount of electricity consumed has a seasonal rhythm and is influenced by occasions on the calendar, such as vacations. This strategy has been demonstrated to increase the precision of load forecasting models.

Weather-based Approach: - This method makes use of weather input variables like temperature, humidity, and wind speed. This strategy acknowledges how the demand for electricity, particularly for heating and cooling, is influenced by the weather. This strategy has also been demonstrated to increase the precision of load forecasting models.

Economic-based Approach: - Economic input factors used in this strategy include GDP, employment, and industrial production. This strategy acknowledges how the demand for electricity is influenced by the economy, particularly in the industrial and commercial sectors. This strategy has also been demonstrated to increase the precision of load forecasting models.

(b)

```
| library(dplyr) |
| 1ibrary(neuralnet) |
| 10 library(ggplot2) |
| 11 library(readxl) |
| 12 setwd("D:/2nd Year/ML_CourseWork") |
| 13 UOW_data <- read_excel("uow_consumption.xlsx") |
| 14 |
| 15 # Rename columns |
| 16 colnames(UOW_data) <- c("date", "time_eighteen", "time_nineteen", "time_twenty") |
| 17 head(UOW_data) |
| 18 |
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```

```
> # Rename columns
> colnames(UOW_data) <- c("date", "time_eighteen", "time_nineteen", "time_twenty")
> head(UOW_data)
# A tibble: 6 × 4
                   time_eighteen time_nineteen time_twenty
  date
                                        <db1>
  <dttm>
                            <db1>
1 2018-01-01 00:00:00
                            38.9
                                         38.9
                                                    38.9
2 2018-01-02 00:00:00
                            42.3
                                         41.9
                                                     41.9
                           40.8
3 2018-01-03 00:00:00
                                         40.5
                                                    40.7
4 2018-01-04 00:00:00
                                         41.9
                                                    41.9
                           44
5 2018-01-05 00:00:00
                                         44.1
                                                    44
6 2018-01-06 00:00:00
                           45.6
                                                    44.3
                                         44.5
>
```

```
#apply lag method
UOW_data$lag_1 <- lag(UOW_data$time_twenty, 1)
UOW_data$lag_2 <- lag(UOW_data$time_twenty, 2)
UOW_data$lag_3 <- lag(UOW_data$time_twenty, 3)
UOW_data$lag_4 <- lag(UOW_data$time_twenty, 4)
UOW_data$lag_7 <- lag(UOW_data$time_twenty, 7)
UOW_data <- na.omit(UOW_data)

#TASK 3
#dividing data to testing and training
UOW_train <- UOW_data[1:380,]
UOW_test <- UOW_data[381:nrow(UOW_data),]</pre>
```

(c)

Normalization

normalizing data before using them in an MLP structure can help improve the convergence, avoid bias, and improve the performance of the MLP. Normalization is a standard pre-processing step in machine learning, and it is important to carefully choose the normalization method that is appropriate for the specific problem and data set.

```
#TASK 4

#normalization

as a mormalize <- function(x) {

return((x - min(x)) / (max(x) - min(x)))

as a mormalize <- as.data.frame(lapply(Uow_train[-1], normalize))

Uow_train_normalized <- as.data.frame(lapply(Uow_test[-1], normalize))

Uow_test_normalized <- as.data.frame(lapply(Uow_test[-1], normalize))

# Set the column names of the test_normalized data frame

colnames(Uow_test_normalized) <- colnames(Uow_train_normalized)
```

(d)

```
48 input_vectors <- list(
          49
51
52
53
54
56
57
59     build_mlp_model <- function(train_data, test_data, input_vars, hidden_structure) {
60         formula <- paste("time_twenty ~", paste(input_vars, collapse = " + "))
61         nn <- neuralnet(as.formula(formula), train_data, hidden = hidden_structure)
62         test_matrix <- as.matrix(test_data[, input_vars, drop = FALSE])
63         colnames(test_matrix) <- colnames(train_data[, input_vars, drop = FALSE])
64         redistrient</pre>
           predictions <- predict(nn, test_matrix)
return(list(model = nn, predictions = predictions))</pre>
64
65
67
68 models <- list()
69 - for (i in 1:length(input_vectors)) {
          models[[i]] <- build_mlp_model(Uow_train_normalized, UOW_test_normalized, input_vectors[[i]], c(5))
70
```

(e)

RMSE: - The average difference between predicted values and actual values is measured by the term "RMSE," which stands for root mean squared error. The square root of the average of the squared differences between the predicted and actual values is used to calculate RMSE. To assess the precision of predictions, regression analysis frequently uses RMSE.

MAE: - MAE, or mean absolute error, is another way to quantify the average difference between projected and actual data. The average of the absolute discrepancies between projected and actual values is used to determine MAE. To assess the precision of predictions in regression analysis, MAE is frequently utilized.

MAPE: - Mean Absolute Percentage Error, or MAPE, is a measurement of the typical percentage difference between predicted values and actual values. By averaging the absolute percentage deviations between projected and actual values, MAPE is calculated. When the scale of the data varies significantly from prediction to prediction, MAPE is frequently used to assess the accuracy of forecasts.

sMAPE: - A modification of MAPE that overcomes some of its drawbacks is called sMAPE, or symmetric MAPE. sMAPE accounts for the magnitude of the projected and actual values when calculating the average percentage difference between predicted and actual values. The problem of MAPE creating infinite or undefined values when the actual values are zero is helped by this.

```
76 calculate_metrics <- function(actual, predicted) {
77    rmse <- sqrt(mean((actual - predicted)^2))
78    mae <- mean(abs(actual - predicted))
79    mape <- mean(abs(actual - predicted) / predicted)
80    smape <- mean(abs(actual - predicted) / (abs(actual) + abs(predicted)) * 2) * 100
81    return(list(RMSE = rmse, MAE = mae, MAPE = mape, sMAPE = smape))
82    }
83    evaluation_metrics <- list()
85    for (i in 1:length(models)) {
86        evaluation_metrics[[i]] <- calculate_metrics(UOW_test_normalized$time_twenty, models[[i]]$predictions)
87    }
88</pre>
```

(d)

```
#Create a comparison table of their testing performances comparison_table <- data.frame(

Model_Description = c("AR(1)", "AR(2)", "AR(3)", "AR(4)", "AR(1,7)", "AR(2,7)", "AR(3,7)", "AR(4,7)"),

MAE = sapply(evaluation_metrics, function(x) x$MAE),

MAE = sapply(evaluation_metrics, function(x) x$MAE),
       MAPE = sapply(evaluation_metrics, function(x) x$MAPE)
       sMAPE = sapply(evaluation_metrics, function(x) x$sMAPE)
 98
 99
100 print(comparison_table)
> print(comparison_table)
  Model_Description
                                  RMSE
                                                              MAPE
                                                  MAE
                                                                          SMAPE
                   AR(1) 0.1951943 0.1560261 0.3217749 34.11284
2
                   AR(2) 0.1973886 0.1587065 0.3223764 34.44747
3
                   AR(3) 0.2077762 0.1685484 0.3324528 35.63873
                   AR(4) 0.2087940 0.1692384 0.3517057 36.37538
5
                AR(1,7) 0.1627568 0.1313272 0.2663793 28.67405
6
                AR(2,7) 0.1635911 0.1306763 0.2638402 28.61209
7
               AR(3,7) 0.1602318 0.1300400 0.2664877 28.21815
8
               AR(4,7) 0.1677774 0.1342914 0.2765529 29.36676
```

(g)

The one-hidden layer network in this situation has fewer weight parameters (2 vs. 3) than the two-hidden layer network. As a result, the one-hidden layer network uses fewer weight parameters than other networks.

```
# Efficiency comparison between one-hidden layer and two-hidden layer networks

105

106

model_l_hidden <- build_mlp_model(uow_train_normalized, uow_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(5))

107

model_2_hidden <- build_mlp_model(uow_train_normalized, uow_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(3, 2))

108

109

# Check the total number of weight parameters per network

110

111

112

113

cat("Total number of weight parameters for the one-hidden layer network:", num_weights_1_hidden, "\n")

114

cat("Total number of weight parameters for the two-hidden layer network:", num_weights_2_hidden, "\n")

> model_1_hidden <- build_mlp_model(uow_train_normalized, uow_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(5))

> model_2_hidden <- build_mlp_model(uow_train_normalized, uow_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(3))

> model_1_hidden <- build_mlp_model(uow_train_normalized, uow_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(3))

> model_1_hidden <- build_mlp_model(uow_train_normalized, uow_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(3))

> model_2_hidden <- build_mlp_model(uow_train_normalized, uow_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(3))

> mum_weights_1_hidden <- sum(sapply(model_1_hiddenSmodelSweights, length))

> num_weights_2_hidden <- sum(sapply(model_2_hiddenSmodelSweights, length))

> num_weights_2_hidden <- sum(sapply(model_2_hiddenSmodelSweights, length))

> cat("Total number of weight parameters for the one-hidden layer network:", num_weights_1_hidden, "\n")

Total number of weight parameters for the two-hidden layer network:", num_weights_2_hidden, "\n")

Total number of weight parameters for the two-hidden layer network:", num_weights_2_hidden, "\n")
```

2nd Subtask Objectives:

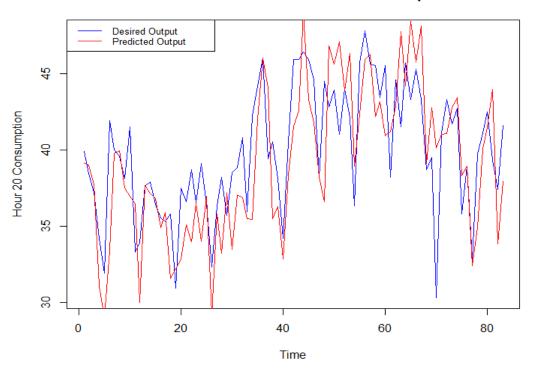
(h)

```
#Task1
# Add the 18th and 19th hour attributes to the input vectors
narx_input_vectors < - list(
c("lag_1", "time_eighteen", "time_nineteen"),
c("lag_1", "lag_2", "lag_3", "lag_3", "lag_1", "time_eighteen", "time_nineteen"),
c("lag_1", "lag_2", "lag_3", "lag_3", "lag_3", "lag_1", "time_eighteen", "time_nineteen"),
c("lag_1", "lag_2", "lag_3", "lag_3", "lag_7", "time_eighteen", "time_nineteen"),
c("lag_1", "lag_2", "lag_3", "lag_3", "lag_7", "time_eighteen", "time_nineteen")
c("lag_1", "lag_2", "lag_3", "lag_7", "time_eighteen", "time_nineteen")

## Build NARX models
## Build NARX models
## Build NARX models
## Build NARX models
## Evaluate NARX models
## Evaluation_metrics <- list()
## Evaluate NARX models
## Evaluation_metrics <- calculate_metrics(UOW_test_normalizedstime_twenty, narx_models[[i]]$predictions)
## ## ## Create a comparison table for NARX models
## Create a comparison table for NARX models
## Model_Description = c("NARX(1,18,19)", "NARX(2,18,19)", "NARX(3,18,19)", "NARX(3,7,18,19)", "NARX(4,7,18,19)"),
## RNSE = sapply(narx_evaluation_metrics, function(x) x$MAPE,
## MAPE = sapply(narx_evaluation_metrics, function(x) x$MAPE,
## Point(narx_comparison_table)
## Point(narx_comparison_
```

(i)

Line Chart of Desired vs. Predicted Output



Appendix

Part 1 sub task 1,

```
1 #lord the data set
   2 install.packages("readxl")
  3 library(readxl)
   4 setwd("D:/2nd Year/ML_CourseWork")
   5 vehicles <- read_excel("vehicles.xlsx")</pre>
  7 #define 18 attributes
  8 install.packages("dplyr")
  9 library(dplyr)
 10 vehicles_subset <- select(vehicles,Comp:Holl.Ra)</pre>
  11
 12 #Scaling.
 13 vehicles_scaled<-scale(vehicles_subset)</pre>
 14
 15 #set outlines using IQR method.
 16 out1<- apply(vehicles_scaled,2,quantile,probs=0.25,na.rm = TRUE)</pre>
 17 out3<- apply(vehicles_scaled,2,quantile,probs=0.75,na.rm =TRUE)
  18 IQR<- out3 - out1
 19
  20 #identifies outliers.
  21 - outliers<-apply(vehicles_scaled,2,function(x){
  22
      lower <- out1-1.5*IQR
  23
      upper <- out3+1.5*IQR
  24
       x < 1ower | x > upper
  25 4 })
  26
  27 #remove outliers.
  28 vehicles_cleaned <- vehicles_scaled
  29 vehicles_cleaned[outliers] <- NA
  30 vehicles_cleaned <- na.omit(vehicles_cleaned)
  31
  32 #determine the number of cluster centers
  33 #NbClust method
  34 install.packages("NbClust")
  35 library(NbClust)
  36 nb <- NbClust(vehicles_cleaned, distance = "euclidean", min.nc = 2, max.nc = 10, method = "kmeans")
  37 print(nb)
  38
  39 #Elbow method
 40 install.packages("factoextra")
41 library(factoextra)
 42 x11() # creates an X11 graphics device
 43 graphics.off() # reset the graphics device
      #plot(x, y) # try plotting again
 45 fviz_nbclust(vehicles_cleaned, kmeans, method = "wss") + geom_vline(xintercept = 4, linetype = 2) + labs(subtitle = "Elbow method")
 46
  47 # Gap statistic method
 48 set.seed(123)
  49 fviz_nbclust(vehicles_cleaned, kmeans, nstart = 25, method = "gap_stat", nboot = 50) + labs(subtitle = "Gap statistic method")
 50
```

```
51
52 # Silhouette method
53 install.packages("cluster")
54 library(cluster)
55 library(factoextra)
56 fviz_nbclust(vehicles_cleaned, kmeans, method = "silhouette") + labs(subtitle = "Silhouette method")
57
58 #compute k-means clustering with k=3
59 set.seed(123)
60 final_stat <- kmeans(vehicles_cleaned,3,nstart = 25)
61 print(final_stat)
62
63 #BSS
64 BSS <- sum(final_stat$size-(colMeans(vehicles_cleaned)-final_stat$centers)^2)
65 cat("BSS:",BSS,"\n")
66
67
   #TSS
68 TSS <- sum((vehicles_cleaned-colMeans(vehicles_cleaned))^2)
69 cat("TSS:",TSS,"\n")
70
71 #WSS
72
   WSS <- sum(final_stat$withinss)
73 cat("WSS:",WSS,"\n")
74
75 #BSS to TSS
76 ratio_BSS_to_TSS <- BSS/TSS</pre>
77
   cat("ratio_BSS_to_TSS:",ratio_BSS_to_TSS,"\n")
78
79
80
81 #Silhouette plot
82 pam.res2 <- pam(vehicles_cleaned,3,metric="euclidean",stand = FALSE)</pre>
83 fviz_silhouette(pam.res2,palette="jco",ggtheme=theme_classic())
84
85 #average Silhouette width score
86 sil <- silhouette(final_stat$cluster, dist(vehicles_cleaned))</pre>
87 avg_Sil_width <-mean(sil[,3])</pre>
88 cat("average Silhouette width score:",avg_Sil_width,"\n")
89
90
```

Part 1 sub task 2,

```
↓□ ♦ I ☐ Source on Save | Q / I | E

                                                                                                                                            Run 💝
   1 install.packages("FactoMineR")
   2 install.packages("factoextra")
  3 library(factoextra)
4 library(FactoMineR)
   6 #apply PCA
   7 pca_data <- PCA(vehicles_cleaned, graph = FALSE)</pre>
   9 # Show the eigenvalues and eigenvectors
  10 summary(pca_data)
  11
 12 # Show the scree plot
 13 fviz_eig(pca_data, addlabels = TRUE)
  14
  15 # Show the cumulative percentage of variance explained
  16 eig_val <- get_eigenvalue(pca_data)</pre>
  17 eig_val
  18
  19 cumulative_variances <- cumsum(eig_val/sum(eig_val)*100)</pre>
  20 cumulative_variances
  21
  22 barplot(cumulative_variances, main = "Cumulative Percentage of Variance Explained", xlab = "Number of Components", ylab = "Cumulative %")
 23
  24 # Choose the PCs that provide at least cumulative score > 92%
  25 num_components <- length(cumulative_variances[cumulative_variances > 92])
     print(paste("Number of components needed to explain at least 92% of the variance:", num_components))
  27
  28 # Create a transformed data set
  29 pca_result <- PCA(vehicles_cleaned, ncp = num_components, graph = FALSE)$ind$coord
  30
  31
  32 #determine the number of cluster centers
  33
  34 #NbClust method
  35 library(NbClust)
  36 nb <- NbClust(pca_result, distance = "euclidean", min.nc = 2, max.nc = 10, method = "kmeans")
  37
     print(nb)
  38
  39 #Elbow method
  40 x11() # creates an X11 graphics device
  41 graphics.off() # reset the graphics device
 42 #plot(x, y) # try plotting again
 43 fviz_nbclust(pca_result, kmeans, method = "wss") + geom_vline(xintercept = 4, linetype = 2) + labs(subtitle = "Elbow method")
  44
  45 # Gap statistic method
  46 set.seed(123)
  47 fviz_nbclust(pca_result, kmeans, nstart = 25, method = "gap_stat", nboot = 50) + labs(subtitle = "Gap statistic method")
  48
 49
```

```
49
 50 # Silhouette method
 51 library(cluster)
 52 library(factoextra)
 53 fviz_nbclust(pca_result, kmeans, method = "silhouette") + labs(subtitle = "Silhouette method")
 54
 55 #compute k-means clustering with k=3
 56 set.seed(123)
 57 final_stat_pca <- kmeans(pca_result,3,nstart = 25)</pre>
 58 print(final_stat_pca)
 59
 60 #BSS
 61 BSS_pca <- sum(final_stat_pca\size-(colMeans(pca_result)-final_stat_pca\centers)\^2)
 62 cat("BSS:",BSS_pca,"\n")
 63
 64 #TSS
 65 TSS_pca <- sum((pca_result-colMeans(pca_result))^2)
66 cat("TSS:",TSS_pca,"\n")
 67
 68 #WSS
 69 WSS_pca <- sum(final_stat_pca$withinss)
 70 cat("WSS:",WSS_pca,"\n")
 71
 72 #BSS to TSS
73 ratio_BSS_to_TSS_pca <- BSS_pca/TSS_pca
74 cat("ratio_BSS_to_TSS:",ratio_BSS_to_TSS_pca,"\n")
 75
 76 #Silhouette plot
 77 pam.res2 <- pam(pca_result,3,metric="euclidean",stand = FALSE)</pre>
 78 fviz_silhouette(pam.res2,palette="jco",ggtheme=theme_classic())
 79
 80 #average Silhouette width score
 81 sil <- silhouette(final_stat_pca$cluster, dist(pca_result))
 82 avg_Sil_width_pca <-mean(sil[,3])</pre>
 83 cat("average Silhouette width score:",avg_Sil_width_pca,"\n")
 84
 85 #Calinski-Harabasz Index
 86 install.packages("fpc")
    library(fpc)
 87
 88 ch_index <- calinhara(pca_result, final_stat_pca$cluster)
 89
     print(ch_index)
 90 barplot(ch_index, main="Calinski-Harabasz Index for K-Means Clustering", xlab="Number of Clusters", ylab="Calinski-Harabasz Index")
 91 plot(ch_index, type="b", xlab="Number of Clusters", ylab="Calinski-Harabasz_Index")
 92
 93
 94 #Silhouette plot
 95 library(cluster)
 96 pam.res2 <- pam(pca_result,3,metric="euclidean",stand = FALSE)
 97
    fviz_silhouette(pam.res2,palette="jco",ggtheme=theme_classic())
 99 #average Silhouette width score
100 sil <- silhouette(final_stat_pca$cluster, dist(pca_result))</pre>
101 avg_Sil_width <-mean(sil[,3])</pre>
102 cat("average Silhouette width score:",avg_Sil_width,"\n")
```

Part 2

```
1 #1A5K 1
       install.packages("readxl")
       install.packages("neuralnet")
install.packages("ggplot2")
install.packages("dplyr")
   /
| brary(dplyr) |
| bibrary(neuralnet) |
| bibrary(ggplot2) |
| bibrary(readx1) |
| setwd("b:/2nd Year/ML_CourseWork")
  12
  13
        UOW_data <- read_excel("uow_consumption.xlsx")
  15
       # Rename columns
  16
       colnames(UOW_data) <- c("date", "time_eighteen", "time_nineteen", "time_twenty")
  17
        head(UOW_data)
  19
       #apply lag method
  20 UOW_data$lag_1 <- lag(UOW_data$time_twenty, 1)
21 UOW_data$lag_2 <- lag(UOW_data$time_twenty, 2)
       UOW_data$lag_3 <- lag(UOW_data$time_twenty, 3)
UOW_data$lag_3 <- lag(UOW_data$time_twenty, 3)
UOW_data$lag_7 <- lag(UOW_data$time_twenty, 4)
UOW_data$lag_7 <- lag(UOW_data$time_twenty, 7)
UOW_data <- na.omit(UOW_data)
  22
23
  24
25
  26
  27
28
  # #dividing data to testing and training
UOW_train <- UOW_data[1:380,]
UOW_test <- UOW_data[381:nrow(UOW_data),]</pre>
  31
  33
       #normalization
  35 normalize <- function(x) {
36    return((x - min(x)) / (max(x) - min(x)))
  39 # Exclude the date column from normalization
  40 UOW_train_normalized <- as.data.frame(lapply(UOW_train[-1], normalize))
41 UOW_test_normalized <- as.data.frame(lapply(UOW_test[-1], normalize))
  42
       # Set the column names of the test_normalized data frame colnames(UOW test normalized) <- colnames(UOW train normalized)
 47
      input_vectors <- list(
    c("lag_1"),
    c("lag_1", "lag_2"),
    c("lag_1", "lag_2", "lag_3"),
    c("lag_1", "lag_2", "lag_3", "lag_4"),
    c("lag_1", "lag_7"),
    c("lag_1", "lag_2", "lag_7"),
    c("lag_1", "lag_2", "lag_3", "lag_7"),
    c("lag_1", "lag_2", "lag_3", "lag_4", "lag_7"))
)</pre>
  48
  49
  50
  51
  52
  53
  54
  55
  56
  57
        )
  58
  59 build_mlp_model <- function(train_data, test_data, input_vars, hidden_structure) {
           formula <- paste("time_twenty ~", paste(input_vars, collapse = " + "))
nn <- neuralnet(as.formula(formula), train_data, hidden = hidden_structure)
  60
  61
            rm <- Retrainet(as.formula(formula), train_data, finder - finder_stracture
test_matrix <- as.matrix(test_data[, input_vars, drop = FALSE])
colnames(test_matrix) <- colnames(train_data[, input_vars, drop = FALSE])
  62
  63
  64
             predictions <- predict(nn, test_matrix)</pre>
  65
            return(list(model = nn, predictions = predictions))
  66 ^ }
  67
  68
       models <- list()
  69 for (i in 1:length(input_vectors)) {
  70
           models[[i]] <- build_mlp_model(UOW_train_normalized, UOW_test_normalized, input_vectors[[i]], c(5))
  71 ^ }
72
73
  74
         #calculated using the standard statistical indices (RMSE, MAE, MAPE and SMAPE - symmetric MAPE)
  76 - calculate_metrics <- function(actual, predicted) {
  77
78
            rmse <- sqrt(mean((actual - predicted)^2))</pre>
            mae <- mean(abs(actual - predicted))
mape <- mean(abs(actual - predicted) / predicted)
smape <- mean(abs(actual - predicted) / (abs(actual) + abs(predicted)) * 2) * 100</pre>
  79
  80
  81
            return(list(RMSE = rmse, MAE = mae, MAPE = mape, sMAPE = smape))
  82 ^ }
  83
  84 evaluation_metrics <- list()
  85 - for (i in 1:length(models)) {
  86
             evaluation_metrics[[i]] <- calculate_metrics(UOW_test_normalized$time_twenty, models[[i]]$predictions)
  87 ^ }
  88
 89
```

```
89
   90
           #TASK 7
   91
             #Create a comparison table of their testing performances
   92
           comparison_table <- data.frame(
Model_Description = c("AR(1)", "AR(2)", "AR(3)", "AR(4)", "AR(1,7)", "AR(2,7)", "AR(3,7)", "AR(4,7)"),
                  MAKE = sapply(evaluation_metrics, function(x) x$MMSE),
MAE = sapply(evaluation_metrics, function(x) x$MAE),
MAPE = sapply(evaluation_metrics, function(x) x$MAPE),
   94
   95
   96
   97
                  sMAPE = sapply(evaluation_metrics, function(x) x$sMAPE)
   99
100
          print(comparison_table)
101
102
           # Add more models with different hidden layer structures and input vectors to create 12-15 models in total
103
104
          # Efficiency comparison between one-hidden layer and two-hidden layer networks
105
106
          model_1_hidden <- build_mlp_model(Uow_train_normalized, UOW_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(5))
model_2_hidden <- build_mlp_model(UoW_train_normalized, UOW_test_normalized, c("lag_1", "lag_2", "lag_3", "lag_7"), c(3, 2))
107
108
          # Check the total number of weight parameters per network num_weights_1_hidden <- sum(sapply(model_1_hidden$model$weights, length))
109
110
111
          num_weights_2_hidden <- sum(sapply(model_2_hidden$model$weights, length))
112
cat("Total number of weight parameters for the one-hidden layer network:", num_weights_1_hidden, "\n")

114 cat("Total number of weight parameters for the two-hidden layer network:", num_weights_2_hidden, "\n")
115
  139 narx_evaluation_metrics <- list()
136 of for (i in 1:length(narx_models)) {
137 narx_evaluation_metrics[[i]] <- calculate_metrics(UOW_test_normalized$time_twenty, narx_models[[i]]$predictions)
138 of the product 
   139
  139
# Create a comparison table for NARX models
141 narx_comparison_table <- data_frame(
142 model_Description = c("NARX(1,18,19)", "NARX(2,18,19)", "NARX(3,18,19)", "NARX(3,7,18,19)", "NARX(4,7,18,19)"),
143 RMSE = sapply(narx_evaluation_metrics, function(x) x$RMSE),
144 MAE = sapply(narx_evaluation_metrics, function(x) x$MAE),
145 MAPE = sapply(narx_evaluation_metrics, function(x) x$MAPE,
146 sMAPE = sapply(narx_evaluation_metrics, function(x) x$MAPE),
147 sMAPE = sapply(narx_evaluation_metrics, function(x) x$MAPE)
   147
   148
   149 print(narx_comparison_table)
   151 #Task 2
   152
   # Denormalize the predictions

154 - denormalize <- function(x, min_value, max_value) +

return(x * (max_value - min_value) + min_value)
   156 ^ }
   157
   158 best_model_index <- which.min(sapply(evaluation_metrics, function(x) x$RMSE))
159 best_model <- models[[best_model_index]]
   160 best_model_predictions <- best_model$predictions
   161
  162 min_value <- min(UOW_train$time_twenty)
163 max_value <- max(UOW_train$time_twenty)
   165 denormalized_predictions <- denormalize(best_model_predictions, min_value, max_value)
   166
  # Plot the predicted output vs. desired output using a line chart

168 plot(UoW_ttest$time_twenty, type = "l", col = "blue", xlab = "rime", ylab = "Hour |

169 lines (denormalized_predictions, col = "red")

170 legend("topleft", legend = c("Desired Output", "Predicted Output"), col = c("blue", "red"), lty = 1, cex = 0.8)
  173
```

References

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