Code ▼

## R Notebook

This is an R Markdown (http://rmarkdown.rstudio.com) Notebook. When you execute code within the notebook, the results appear beneath the code.

Hide

install.packages("readr")

WARNING: Rtools is required to build R packages but is not currently installed. Please downlo ad and install the appropriate version of Rtools before proceeding:

https://cran.rstudio.com/bin/windows/Rtools/ 'C:/Users/leewq/AppData/Local/R/win-library/4.2'의 위치에 패키지(들)을 설치합니다. (왜냐하면 'lib'가 지정되지 않았기 때문입니다) trying URL 'https://cran.rstudio.com/bin/windows/contrib/4.2/readr 2.1.4.zip'

Content type 'application/zip' length 1192968 bytes (1.1 MB) downloaded 1.1 MB

패키지 'readr'를 성공적으로 압축해제하였고 MD5 sums 이 확인되었습니다

다운로드된 바이너리 패키지들은 다음의 위치에 있습니다

C:\Users\leewq\AppData\Local\Temp\Rtmp4K1qIn\downloaded\_packages

Hide

install.packages("FactoMineR")

Error in install.packages : Updating loaded packages

Hide

install.packages("FNN")

WARNING: Rtools is required to build R packages but is not currently installed. Please downlo ad and install the appropriate version of Rtools before proceeding:

https://cran.rstudio.com/bin/windows/Rtools/

'C:/Users/leewq/AppData/Local/R/win-library/4.2'의 위치에 패키지(들)을 설치합니다.

(왜냐하면 'lib'가 지정되지 않았기 때문입니다)

trying URL 'https://cran.rstudio.com/bin/windows/contrib/4.2/FNN\_1.1.3.2.zip'

Content type 'application/zip' length 450109 bytes (439 KB)

downloaded 439 KB

패키지 'FNN'를 성공적으로 압축해제하였고 MD5 sums 이 확인되었습니다

다운로드된 바이너리 패키지들은 다음의 위치에 있습니다

C:\Users\leewq\AppData\Local\Temp\Rtmp4K1qIn\downloaded\_packages

install.packages("FactoMineR")

WARNING: Rtools is required to build R packages but is not currently installed. Please downlo ad and install the appropriate version of Rtools before proceeding:

https://cran.rstudio.com/bin/windows/Rtools/

'C:/Users/leewq/AppData/Local/R/win-library/4.2'의 위치에 패키지(들)을 설치합니다.

(왜냐하면 'lib'가 지정되지 않았기 때문입니다)

trying URL 'https://cran.rstudio.com/bin/windows/contrib/4.2/FactoMineR\_2.7.zip'

Content type 'application/zip' length 3804028 bytes (3.6 MB)

downloaded 3.6 MB

패키지 'FactoMineR'를 성공적으로 압축해제하였고 MD5 sums 이 확인되었습니다

Warning in install.packages :

패키지 'FactoMineR'의 이전설치를 삭제할 수 없습니다

Warning in install.packages :

problem copying C:\Users\leewq\AppData\Local\R\win-library\4.2\00LOCK\FactoMineR\libs\x64\FactoMineR.dll to C:\Users\leewq\AppData\Local\R\win-library\4.2\FactoMineR\libs\x64\FactoMineR\

R.dll: Permission denied

Warning in install.packages :

'FactoMineR'를 복구하였습니다

다운로드된 바이너리 패키지들은 다음의 위치에 있습니다

C:\Users\leewq\AppData\Local\Temp\Rtmp4K1qIn\downloaded\_packages

Try executing this chunk by clicking the *Run* button within the chunk or by placing your cursor inside it and pressing *Ctrl+Shift+Enter*.

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library(readr)

경고: 패키지 'readr'는 R 버전 4.2.3에서 작성되었습니다

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# Load the dataset from the CSV file
data <- read csv("perth.csv")</pre>

Rows: 33656 Columns: 19— Column specification

Delimiter: ","

chr (7): ADDRESS, SUBURB, GARAGE, BUILD\_YEAR, NEAREST\_STN, DATE\_SOLD, NEAREST\_SCH

dbl (12): PRICE, BEDROOMS, BATHROOMS, LAND\_AREA, FLOOR\_AREA, CBD\_DIST, NEAREST\_STN\_DIST, POST CODE, LATITUDE...

i Use `spec()` to retrieve the full column specification for this data.

 ${f i}$  Specify the column types or set `show\_col\_types = FALSE` to quiet this message.

Add a new chunk by clicking the *Insert Chunk* button on the toolbar or by pressing *Ctrl+Alt+I*.

```
# Convert all non-numeric columns to numeric
# Identify non-numeric columns
non_numeric_columns <- sapply(data, function(x) !is.numeric(x))</pre>
# Remove non-numeric columns
data numeric <- data[, !non numeric columns]</pre>
# Check if all columns are now numeric
str(data_numeric)
tibble [33,656 × 12] (S3: tbl df/tbl/data.frame)
                  : num [1:33656] 565000 365000 287000 255000 325000 409000 400000 370000 56
$ PRICE
5000 685000 ...
                : num [1:33656] 4 3 3 2 4 4 3 4 4 3 ...
 $ BEDROOMS
$ BATHROOMS
                 : num [1:33656] 2 2 1 1 1 2 2 2 2 2 ...
                 : num [1:33656] 600 351 719 651 466 759 386 468 875 552 ...
 $ LAND_AREA
                 : num [1:33656] 160 139 86 59 131 118 132 158 168 126 ...
$ FLOOR AREA
 $ CBD DIST
                 : num [1:33656] 18300 26900 22600 17900 11200 27300 28200 41700 12100 5900
 $ NEAREST_STN_DIST: num [1:33656] 1800 4900 1900 3600 2000 1000 3700 1100 2500 508 ...
              : num [1:33656] 6164 6167 6111 6056 6054 ...
$ POSTCODE
 $ LATITUDE
                  : num [1:33656] -32.1 -32.2 -32.1 -31.9 -31.9 ...
                  : num [1:33656] 116 116 116 116 116 ...
 $ LONGITUDE
 $ NEAREST_SCH_DIST: num [1:33656] 0.828 5.524 1.649 1.571 1.515 ...
```

library(FactoMineR)

경고: 패키지 'FactoMineR'는 R 버전 4.2.3에서 작성되었습니다

Hide

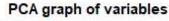
```
# Perform PCA
res.pca <- PCA(data_numeric, scale.unit = TRUE, graph = FALSE)</pre>
```

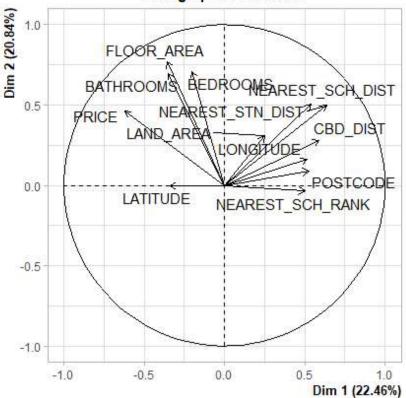
\$ NEAREST SCH RANK: num [1:33656] NA 129 113 NA NA NA NA NA NA 29 ...

경고: Missing values are imputed by the mean of the variable: you should use the imputePCA function of the missMDA package

Hide

# Determine the optimal number of components by plotting the explained variance ratio
plot(res.pca, choix = "var")





NA NA

Hide

```
# Set the target variable as 'price'
target <- "PRICE"

# Remove rows with NA values
data_numeric_clean <- na.omit(data_numeric)

# Split the cleaned dataset into features and target variable(s)
X_clean <- data_numeric_clean[, !names(data_numeric_clean) %in% target]
y_clean <- data_numeric_clean[, target]

# Convert y_clean to a vector
y_clean_vector <- as.vector(unlist(y_clean))

# Check the dimensions of X_clean and y_clean
cat("Dimensions of X_clean:", dim(X_clean), "\n")</pre>
```

Dimensions of X\_clean: 22704 11

Hide

```
cat("Length of y_clean_vector:", length(y_clean_vector), "\n")
```

Length of y\_clean\_vector: 22704

```
# Perform LDA
library(MASS)
res.lda <- lda(X_clean, y_clean_vector)</pre>
```

Hide

- # Load the necessary libraries
- # Load the necessary libraries
  library(caret)

```
필요한 패키지를 로딩중입니다: ggplot2
필요한 패키지를 로딩중입니다: lattice
```

Hide

library(FNN)

경고: 패키지 'FNN'는 R 버전 4.2.3에서 작성되었습니다

Hide

```
# Split the data into train and test sets
```

set.seed(42)

trainIndex <- createDataPartition(y\_vector, p = 0.8, list = FALSE)</pre>

 $\label{lem:cond} $$\operatorname{train\_pca} \leftarrow \operatorname{res.pca} \quad $\inf \circ \operatorname{components} \ chosen$ 

test\_pca <- res.pca\$ind\$coord[-trainIndex, 1:2] # Replace 2 with the number of components cho
sen</pre>

y\_train <- y\_vector[trainIndex]</pre>

y\_test <- y\_vector[-trainIndex]</pre>

# Train and evaluate the kNN regression on PCA-reduced data

knn\_pca <- knn.reg(train = train\_pca, test = test\_pca, y = y\_train)</pre>

# Calculate the mean squared error (MSE) and root mean squared error (RMSE)

mse\_pca <- mean((knn\_pca\$pred - y\_test)^2)</pre>

경고: longer object length is not a multiple of shorter object length

Hide

```
rmse_pca <- sqrt(mse_pca)</pre>
```

# Print the results

cat("MSE for kNN regression on PCA-reduced data:", mse\_pca, "\n")

MSE for kNN regression on PCA-reduced data: 201988363475

```
cat("RMSE for kNN regression on PCA-reduced data:", rmse_pca, "\n")
```

RMSE for kNN regression on PCA-reduced data: 449431.2

Hide

```
library(MASS)
# Perform LDA
res_lda <- lda(X_clean, y_clean_vector)</pre>
# Project data using LDA
X_lda <- predict(res_lda, X_clean)$x</pre>
# Split the LDA-reduced data into train and test sets
trainIndex <- createDataPartition(y_clean_vector, p = 0.8, list = FALSE)</pre>
train_lda <- X_lda[trainIndex,]</pre>
test lda <- X lda[-trainIndex,]</pre>
y_train <- y_clean_vector[trainIndex]</pre>
y_test <- y_clean_vector[-trainIndex]</pre>
# Train and evaluate the kNN regression on LDA-reduced data
knn_lda <- knn.reg(train_lda, test_lda, y_train)</pre>
mse_lda <- mean((knn_lda$pred - y_test)^2)</pre>
rmse_lda <- sqrt(mse_lda)</pre>
cat("MSE for kNN regression on LDA-reduced data:", mse_lda, "\n")
```

MSE for kNN regression on LDA-reduced data: 41061045892

Hide

cat("RMSE for kNN regression on LDA-reduced data:", rmse lda, "\n")

RMSE for kNN regression on LDA-reduced data: 202635.3

Hide

```
library(caret)
library(FNN)
library(MASS)
# Replace X and y_vector with your dataset and target variable
X <- X_clean
y_vector <- y_clean_vector</pre>
# Split the data into train and test sets
set.seed(42)
trainIndex <- createDataPartition(y_vector, p = 0.8, list = FALSE)</pre>
train <- X[trainIndex,]</pre>
test <- X[-trainIndex,]</pre>
y train <- y vector[trainIndex]</pre>
y test <- y vector[-trainIndex]</pre>
# Impute missing values in the train and test datasets
# Convert data frames to matrices
train imputed <- as.matrix(train imputed)</pre>
test_imputed <- as.matrix(test_imputed)</pre>
# Train and evaluate the kNN regression on the imputed original data
knn_original <- knn.reg(train = train_imputed, test = test_imputed, y = y_train)</pre>
# Make predictions on test set
pred_original <- knn.reg(train = train_imputed, test = test_imputed, y = y_train)$pred</pre>
# Perform PCA
res.pca <- prcomp(X, scale. = TRUE)</pre>
# Project data using PCA
train_pca <- res.pca$x[trainIndex, 1:2] # Replace 2 with the number of components chosen
test_pca <- res.pca$x[-trainIndex, 1:2] # Replace 2 with the number of components chosen</pre>
# Train and evaluate the kNN regression on PCA-reduced data
knn_pca <- knn.reg(train_pca, test_pca, y_train)</pre>
mse_pca <- mean((knn_pca$pred - y_test)^2)</pre>
rmse_pca <- sqrt(mse_pca)</pre>
cat("MSE for kNN regression on PCA-reduced data:", mse pca, "\n")
```

MSE for kNN regression on PCA-reduced data: 95613532827

Hide

```
cat("RMSE for kNN regression on PCA-reduced data:", rmse_pca, "\n")
```

```
RMSE for kNN regression on PCA-reduced data: 309214.4
```

```
# Perform LDA
res_lda <- lda(X, y_vector)</pre>
# Project data using LDA
X_lda <- predict(res_lda, X)$x</pre>
train_lda <- X_lda[trainIndex,]</pre>
test_lda <- X_lda[-trainIndex,]</pre>
# Train and evaluate the kNN regression on LDA-reduced data
knn_lda <- knn.reg(train_lda, test_lda, y_train)</pre>
mse_lda <- mean((knn_lda$pred - y_test)^2)</pre>
rmse_lda <- sqrt(mse_lda)</pre>
cat("MSE for kNN regression on LDA-reduced data:", mse lda, "\n")
MSE for kNN regression on LDA-reduced data: 42561456705
                                                                                                Hide
cat("RMSE for kNN regression on LDA-reduced data:", rmse lda, "\n")
RMSE for kNN regression on LDA-reduced data: 206304.3
                                                                                                Hide
# Train and evaluate the kNN regression on the imputed original data
knn_original <- knn.reg(train = train_imputed, test = test_imputed, y = y_train)</pre>
# Calculate the mean squared error (MSE) and root mean squared error (RMSE)
mse_original <- mean((knn_original$pred - y_test)^2)</pre>
경고: longer object length is not a multiple of shorter object length
                                                                                                Hide
rmse_original <- sqrt(mse_original)</pre>
# Calculate the difference in performance metrics
rmse_diff_pca <- rmse_original - rmse_pca</pre>
rmse diff lda <- rmse original - rmse lda
cat("RMSE for kNN regression on original imputed data:", rmse_original, "\n")
RMSE for kNN regression on original imputed data: 503804.7
                                                                                               Hide
cat("RMSE difference (PCA):", rmse diff pca, "\n")
RMSE difference (PCA): 194590.3
```

```
cat("RMSE difference (LDA):", rmse_diff_lda, "\n")
```

RMSE difference (LDA): 297500.4

When you save the notebook, an HTML file containing the code and output will be saved alongside it (click the *Preview* button or press *Ctrl+Shift+K* to preview the HTML file).

The preview shows you a rendered HTML copy of the contents of the editor. Consequently, unlike *Knit*, *Preview* does not run any R code chunks. Instead, the output of the chunk when it was last run in the editor is displayed.