Assignment 5

Details

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3. Batch : M11 4. Class : TE11

Problem Statement

Perform the following operations using Python on the Air quality and Heart Diseases data sets

- 1. Data cleaning
- 2. Data integration
- 3. Data transformation
- 4. Error correcting
- 5. Data model building

Implementation details

Dataset URL: https://archive.ics.uci.edu/ml/datasets/Heart+Disease)

2. Python version: 3.7.4

3. Imports:

- A. pandas
- B. numpy
- C. matplotlib.pyolot
- D. seaborn
- E. sklearn.linear_model.LogisticRegression

Dataset details

- 1. This database contains 76 attributes, but all published experiments refer to using a subset of 14 of them. In particular, the Cleveland database is the only one that has been used by ML researchers to this date.
- 2. The "goal" field refers to the presence of heart disease in the patient.
- 3. It is integer valued from 0 (no presence) to 4. Experiments with the Cleveland database have concentrated on simply attempting to distinguish presence (values 1,2,3,4) from absence (value 0).
- 4. The names and social security numbers of the patients were recently removed from the database, replaced with dummy values.

Importing required libraries

In [1]:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
```

Loading the dataset

```
In [2]:
```

In [3]:

oldpeak

dtype: int64

slope

thal num

ca

0

0

0

```
dataset = pd.read_csv("./preprocessed_data.csv", index_col=0)
```

Displaying metadata for dataset (Statistical)

```
dataset.shape
Out[3]:
(682, 14)
In [4]:
dataset.isnull().sum()
Out[4]:
               0
age
               0
sex
chest_pain
trestbps
               0
cholestrol
               0
               0
fbs
restecg
               0
               0
thalach
               0
exang
```

In [5]:

dataset.head(15)

Out[5]:

| | age | sex | chest_pain | trestbps | cholestrol | fbs | restecg | thalach | exang | oldpeak | |
|----|----------|-----|------------|----------|------------|-----|---------|----------|-------|----------|--|
| 0 | 0.714286 | 1 | 1 | 0.541667 | 0.386401 | 1.0 | 2.0 | 0.633803 | 0.0 | 0.890909 | |
| 1 | 0.795918 | 1 | 4 | 0.666667 | 0.474295 | 0.0 | 2.0 | 0.338028 | 1.0 | 0.745455 | |
| 2 | 0.795918 | 1 | 4 | 0.333333 | 0.379768 | 0.0 | 2.0 | 0.485915 | 1.0 | 0.945455 | |
| 4 | 0.265306 | 0 | 2 | 0.416667 | 0.338308 | 0.0 | 2.0 | 0.788732 | 0.0 | 0.727273 | |
| 5 | 0.571429 | 1 | 2 | 0.333333 | 0.391376 | 0.0 | 0.0 | 0.830986 | 0.0 | 0.618182 | |
| 7 | 0.591837 | 0 | 4 | 0.333333 | 0.587065 | 0.0 | 0.0 | 0.725352 | 1.0 | 0.581818 | |
| 8 | 0.714286 | 1 | 4 | 0.416667 | 0.421227 | 0.0 | 2.0 | 0.612676 | 0.0 | 0.727273 | |
| 10 | 0.591837 | 1 | 4 | 0.500000 | 0.318408 | 0.0 | 0.0 | 0.619718 | 0.0 | 0.545455 | |
| 11 | 0.571429 | 0 | 2 | 0.500000 | 0.487562 | 0.0 | 2.0 | 0.654930 | 0.0 | 0.709091 | |
| 12 | 0.571429 | 1 | 3 | 0.416667 | 0.424544 | 1.0 | 2.0 | 0.577465 | 1.0 | 0.581818 | |
| 13 | 0.326531 | 1 | 2 | 0.333333 | 0.436153 | 0.0 | 0.0 | 0.795775 | 0.0 | 0.472727 | |
| 14 | 0.489796 | 1 | 3 | 0.766667 | 0.330017 | 1.0 | 0.0 | 0.718310 | 0.0 | 0.563636 | |
| 15 | 0.591837 | 1 | 3 | 0.583333 | 0.278607 | 0.0 | 0.0 | 0.802817 | 0.0 | 0.763636 | |
| 16 | 0.408163 | 1 | 2 | 0.250000 | 0.379768 | 0.0 | 0.0 | 0.760563 | 0.0 | 0.654545 | |
| 17 | 0.530612 | 1 | 4 | 0.500000 | 0.396352 | 0.0 | 0.0 | 0.704225 | 0.0 | 0.690909 | |

Observations:

- 1. There are 682 data points with 14 columns (including target column)
- 2. Null values are removed / replaced and dataset is scaled for numerical variables

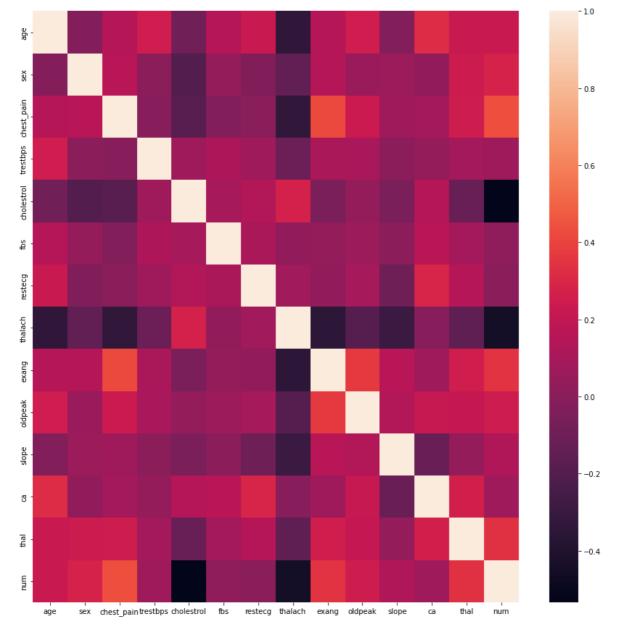
A) Feature selection

In [6]:

```
# Displaying heatmap for correlation matrix
fig = plt.figure(figsize=(15, 15))

# Adds subplot on position 1
ax = fig.add_subplot(111)

sns.heatmap(dataset.corr())
plt.show()
```



In [7]:

dataset.corr()

Out[7]:

| | age | sex | chest_pain | trestbps | cholestrol | fbs | restecg | thal |
|------------|-----------|-----------|------------|-----------|------------|-----------|-----------|--------|
| age | 1.000000 | -0.023624 | 0.145740 | 0.247017 | -0.089696 | 0.154740 | 0.223164 | -0.340 |
| sex | -0.023624 | 1.000000 | 0.163370 | 0.003323 | -0.199176 | 0.037774 | -0.036257 | -0.148 |
| chest_pain | 0.145740 | 0.163370 | 1.000000 | -0.009134 | -0.182046 | -0.026207 | -0.002091 | -0.334 |
| trestbps | 0.247017 | 0.003323 | -0.009134 | 1.000000 | 0.066543 | 0.120023 | 0.074495 | -0.111 |
| cholestrol | -0.089696 | -0.199176 | -0.182046 | 0.066543 | 1.000000 | 0.097576 | 0.143226 | 0.272 |
| fbs | 0.154740 | 0.037774 | -0.026207 | 0.120023 | 0.097576 | 1.000000 | 0.113193 | 0.027 |
| restecg | 0.223164 | -0.036257 | -0.002091 | 0.074495 | 0.143226 | 0.113193 | 1.000000 | 0.082 |
| thalach | -0.340013 | -0.148256 | -0.334507 | -0.111419 | 0.272218 | 0.027782 | 0.082551 | 1.000 |
| exang | 0.146912 | 0.147096 | 0.415408 | 0.118941 | -0.054860 | 0.036641 | 0.024909 | -0.351 |
| oldpeak | 0.246984 | 0.056113 | 0.229230 | 0.106618 | 0.039519 | 0.062684 | 0.098591 | -0.186 |
| slope | -0.024984 | 0.063177 | 0.074927 | 0.000372 | -0.058404 | 0.001531 | -0.104784 | -0.295 |
| ca | 0.321673 | 0.025315 | 0.093902 | 0.037259 | 0.148186 | 0.173009 | 0.289584 | -0.006 |
| thal | 0.222136 | 0.233270 | 0.239361 | 0.084489 | -0.125926 | 0.084525 | 0.147006 | -0.152 |
| num | 0.225453 | 0.279127 | 0.434501 | 0.067330 | -0.533687 | 0.019112 | -0.000813 | -0.450 |
| 4 | | | | | | | | • |

Note:

- 1. The above heatmap and the correlation table suggests that all of the data features are significantly correlated with the target variables
- 2. The following features are negatively correlated with the target variable:
 - A. cholestrol
 - B. thalach

Further action:

- 1. No feature drop is necessary due to significant correlation with target variable
- 2. The target variable is considered to be "num" (last column of the dataset)

B) Building the data model

1) Understanding the target variable

In [8]:

```
# checking the unique values (categories in the dataset)
dataset.num.unique()
Out[8]:
```

Note:

1. The values beyond 0 are indicative of the fact that there is presence of heart disease

Further action:

array([0, 1, 3, 2, 4])

1. Binarize the target variable for the classes as presence or absence of heart disease

2) Binarizing the target variable

```
In [9]:
dataset["num"] = dataset["num"].replace([2, 3, 4], 1)

In [10]:
dataset.num.unique()

Out[10]:
array([0, 1])
```

3) Checking distribution of target variable

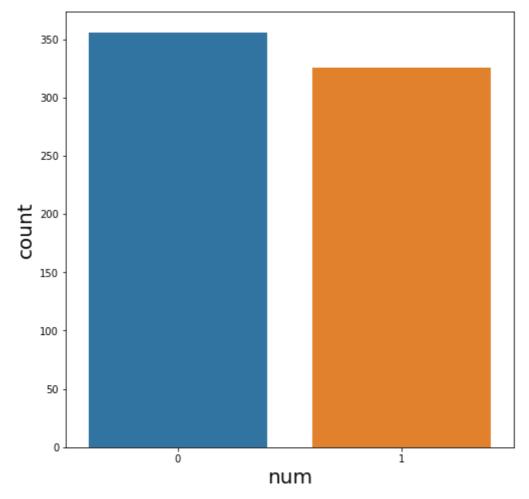
In [11]:

```
# Plotting the count plot for target variable
fig = plt.figure(figsize=(8, 8))

# Adds subplot on position 1
ax = fig.add_subplot(111)

plt.xlabel("num", fontsize=20)
plt.ylabel("count", fontsize=20)

sns.countplot(x=dataset.num)
plt.show()
```



In [12]:

```
# creating subsets for target variables for fair distribution in training and testi
dataset target 0 = dataset[dataset.num == 0]
dataset target 1 = dataset[dataset.num == 1]
print("Shape of target 1 data : ", dataset_target_1.shape)
print("Shape of target 0 data : ", dataset_target_0.shape)
Shape of target 1 data :
                               (326, 14)
Shape of target 0 data: (356, 14)
In [13]:
# Shuffling the data subsets
dataset_target_1 = dataset_target_1.sample(frac=1)
dataset target 0 = dataset target 0.sample(frac=1)
# Confirming shapes for no value loss
print("Shape of target 1 data : ", dataset_target_1.shape)
print("Shape of target 0 data : ", dataset_target_0.shape)
Shape of target 1 data:
                               (326, 14)
Shape of target 0 data :
                               (356, 14)
```

4) Creating training and testing data with 80:20 ratio

In [14]:

```
# Calculating 80 percent mark
target_0_mark = int(dataset_target_0.shape[0]*0.8)
target 1 mark = int(dataset target 1.shape[0]*0.8)
# Generating train data
train data = pd.concat(
    objs=[
        dataset target 0.iloc[:target 0 mark, :],
        dataset_target_1.iloc[:target_1_mark, :]
    ],
    axis=0
)
# Generating test data
test data = pd.concat(
    objs=[
        dataset target 0.iloc[target 0 mark:, :],
        dataset target 1.iloc[target 1 mark:, :]
    axis=0
)
# Shuffling the training and testing data
train data = train data.sample(frac=1)
test data = test data.sample(frac=1)
# Checking data shapes
print("Training data shape : ", train_data.shape)
print("Testing data shape : ", test data.shape)
```

Training data shape : (544, 14) Testing data shape : (138, 14)

5) Splitting training and testing inputs and targets

In [15]:

```
# Splitting training data
train_inputs = train_data.iloc[:, :-1]
train_targets = train_data.iloc[:, -1]

# Splitting testing data
test_inputs = test_data.iloc[:, :-1]
test_targets = test_data.iloc[:, -1]

# Checking shape of data
print("Train inputs shape : ", train_inputs.shape)
print("Train targets shape : ", train_targets.shape)
print("Test inputs shape : ", test_inputs.shape)
print("Test targets shape : ", test_targets.shape)
```

Train inputs shape : (544, 13)
Train targets shape : (544,)
Test inputs shape : (138, 13)
Test targets shape : (138,)

6) Building the data model

```
In [16]:
```

```
# Importing model
from sklearn.linear_model import LogisticRegression
```

In [17]:

```
logReg_model = LogisticRegression()
```

In [18]:

```
# Training the model
logReg_model.fit(train_inputs, train_targets)
print("Model trained")
```

Model trained

7) Checking accuracy of model on testing data

```
In [19]:
```

```
logReg_model.score(test_inputs, test_targets)
```

Out[19]:

0.8985507246376812

In [20]:

```
logReg_model.score(train_inputs, train_targets)
```

Out[20]:

0.8253676470588235

Conclusion

- 1. The logistic regression model was fit on the given dataset
- 2. The model gave 89.85% accuracy on testing data and 82.53% accuracy on testing data

End of Notebook