pbo732pgr

July 28, 2023

```
[1]: import numpy as np
     import pandas as pd
     import matplotlib.pyplot as plt
     import seaborn as sns
[2]: df=pd.read_csv("/content/4_drug200.csv")
     df
[2]:
          Age Sex
                        BP Cholesterol
                                          Na_to_K
                                                     Drug
     0
           23
                 F
                                    HIGH
                                           25.355
                                                    drugY
                      HIGH
     1
            47
                 М
                       LOW
                                    HIGH
                                           13.093
                                                    drugC
     2
            47
                       LOW
                                    HIGH
                                           10.114
                                                    drugC
     3
           28
                                            7.798
                 F
                    NORMAL
                                    HIGH
                                                    drugX
     4
                 F
                       LOW
                                    HIGH
                                           18.043
                                                    drugY
     195
           56
                 F
                       LOW
                                    HIGH
                                           11.567
                                                    drugC
     196
                       LOW
                                    HIGH
                                           12.006
                                                    drugC
            16
                 М
                                            9.894
     197
            52
                                                    drugX
                    NORMAL
                                    HIGH
     198
            23
                    NORMAL
                                 NORMAL
                                           14.020
                                                    drugX
     199
            40
                 F
                       LOW
                                           11.349
                                                    drugX
                                 NORMAL
     [200 rows x 6 columns]
[3]:
    df.head()
[3]:
        Age Sex
                      BP Cholesterol
                                        Na_to_K
                                                   Drug
         23
               F
                                         25.355
                                                  drugY
     0
                    HIGH
                                 HIGH
         47
     1
               М
                     LOW
                                 HIGH
                                         13.093
                                                  drugC
     2
         47
               М
                     LOW
                                 HIGH
                                         10.114
                                                  drugC
     3
         28
               F
                  NORMAL
                                 HIGH
                                          7.798
                                                  drugX
         61
               F
                     LOW
                                 HIGH
                                         18.043
                                                  drugY
```

1 DATA CLEANING AND DATA PREPROCESSING

```
[4]: df.info()
```

```
RangeIndex: 200 entries, 0 to 199
    Data columns (total 6 columns):
         Column
                       Non-Null Count
                                        Dtype
                       _____
     0
         Age
                       200 non-null
                                        int64
     1
         Sex
                       200 non-null
                                        object
     2
         BP
                       200 non-null
                                        object
     3
         Cholesterol 200 non-null
                                        object
     4
         Na_to_K
                       200 non-null
                                        float64
     5
         Drug
                       200 non-null
                                        object
    dtypes: float64(1), int64(1), object(4)
    memory usage: 9.5+ KB
[5]: df.describe()
[5]:
                    Age
                            Na_to_K
            200.000000
     count
                         200.000000
     mean
             44.315000
                          16.084485
     std
             16.544315
                           7.223956
     min
             15.000000
                           6.269000
     25%
             31.000000
                          10.445500
     50%
             45.000000
                          13.936500
     75%
             58.000000
                          19.380000
     max
             74.000000
                          38.247000
[6]:
     df.columns
[6]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
[7]: df1=df.dropna(axis=1)
     df1
[7]:
                        BP Cholesterol
                                         Na_to_K
          Age Sex
                                                   Drug
     0
           23
                F
                      HIGH
                                  HIGH
                                          25.355
                                                  drugY
     1
           47
                М
                       LOW
                                  HIGH
                                          13.093
                                                  drugC
     2
           47
                М
                       LOW
                                  HIGH
                                          10.114
                                                  drugC
     3
           28
                F
                    NORMAL
                                  HIGH
                                           7.798
                                                  drugX
     4
                F
           61
                       LOW
                                  HIGH
                                          18.043
                                                  drugY
     195
                F
                                          11.567
           56
                       LOW
                                  HIGH
                                                  drugC
     196
                                          12.006
                                                  drugC
           16
                М
                       LOW
                                  HIGH
     197
                                           9.894
           52
                Μ
                    NORMAL
                                  HIGH
                                                  drugX
     198
           23
                Μ
                    NORMAL
                                NORMAL
                                          14.020
                                                  drugX
     199
           40
                F
                       LOW
                                NORMAL
                                          11.349
                                                  drugX
     [200 rows x 6 columns]
```

<class 'pandas.core.frame.DataFrame'>

```
[8]: df1.columns
```

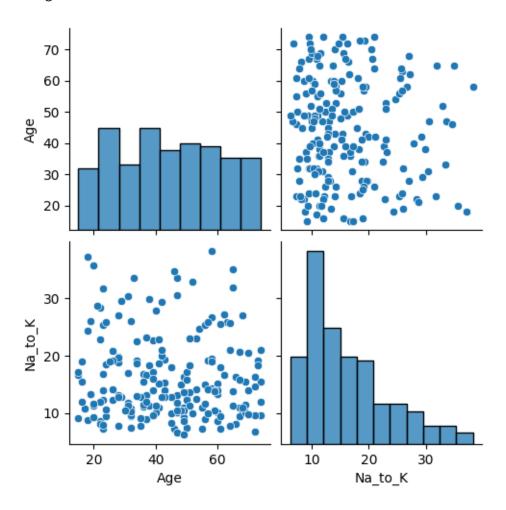
[8]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')

```
[9]: df1=df1[['Age','Na_to_K']]
```

2 EDA AND VISUALIZATION

[10]: sns.pairplot(df1)

[10]: <seaborn.axisgrid.PairGrid at 0x7fb37e492530>



[11]: sns.distplot(df1['Na_to_K'])

<ipython-input-11-4b6a442fe97b>:1: UserWarning:

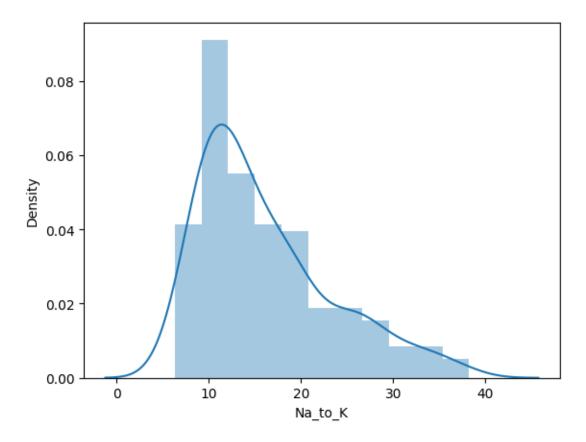
`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751

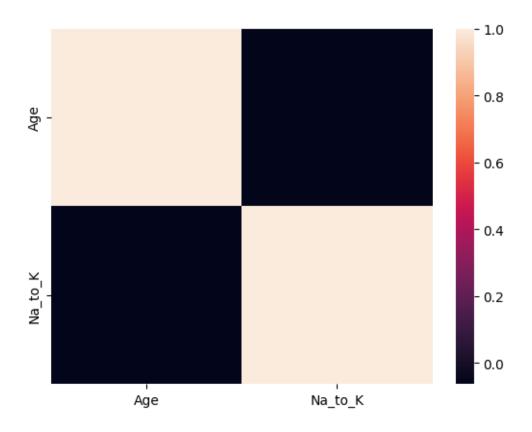
sns.distplot(df1['Na_to_K'])

[11]: <Axes: xlabel='Na_to_K', ylabel='Density'>



[12]: sns.heatmap(df1.corr())

[12]: <Axes: >



3 TO TRAIN THE MODEL AND MODEL BULDING

```
[13]: x=df[['Age','Na_to_K']]
y=df['Na_to_K']

[14]: from sklearn.model_selection import train_test_split
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.2)

[15]: from sklearn.linear_model import LinearRegression
lr=LinearRegression()
lr.fit(x_train,y_train)

[15]: LinearRegression()

[16]: lr.intercept_

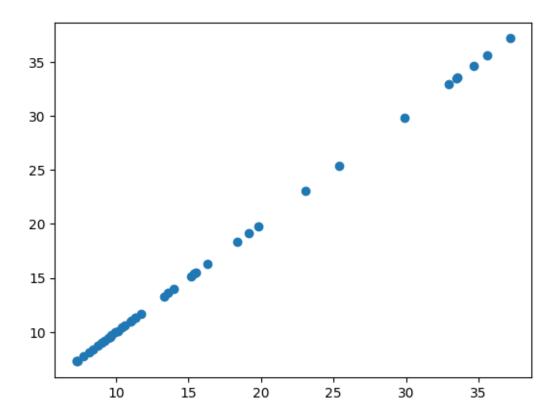
[16]: -3.552713678800501e-15

[17]: coeff=pd.DataFrame(lr.coef_,x.columns,columns=['Co-efficient'])
coeff
```

```
[17]: Co-efficient
Age -8.099643e-19
Na_to_K 1.000000e+00
```

```
[18]: prediction =lr.predict(x_test)
plt.scatter(y_test,prediction)
```

[18]: <matplotlib.collections.PathCollection at 0x7fb37904eb00>



4 ACCURACY