00utqdtnw

July 31, 2023

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

1 DATA CLEANING AND DATA PREPROCESSING

```
[]: 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
[]: df.describe()
[]:
                    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
[]:
     df.columns
[]: Index(['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K', 'Drug'], dtype='object')
[]: df1=df.dropna(axis=1)
     df1
[]:
          Age Sex
                        BP Cholesterol
                                        Na_to_K
                                                   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
                                          18.043
           61
                      LOW
                                  HIGH
                                                  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'>

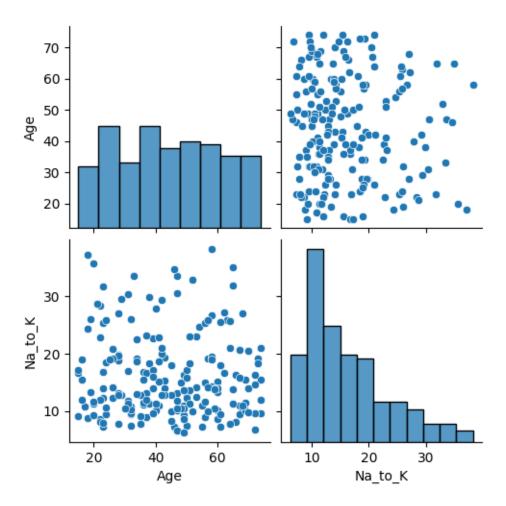
```
[]: df1.columns

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

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

2 EDA AND VISUALIZATION

- []: sns.pairplot(df1)
- []: <seaborn.axisgrid.PairGrid at 0x7e9127466770>



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

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

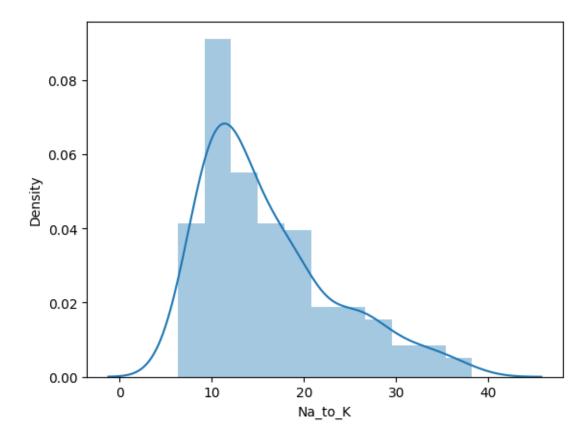
`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

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

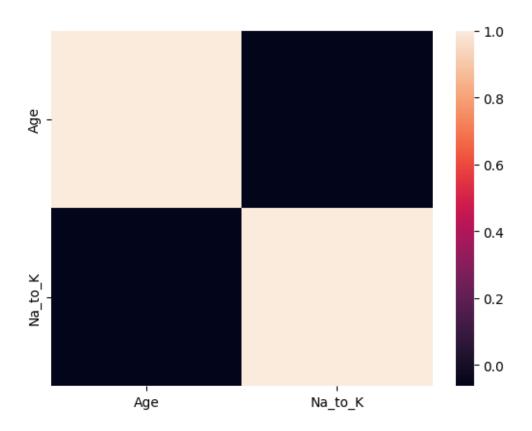
sns.distplot(df1['Na_to_K'])

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



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

[]: <Axes: >



3 TO TRAIN THE MODEL AND MODEL BULDING

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

[ ]: 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)

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

[ ]: LinearRegression()

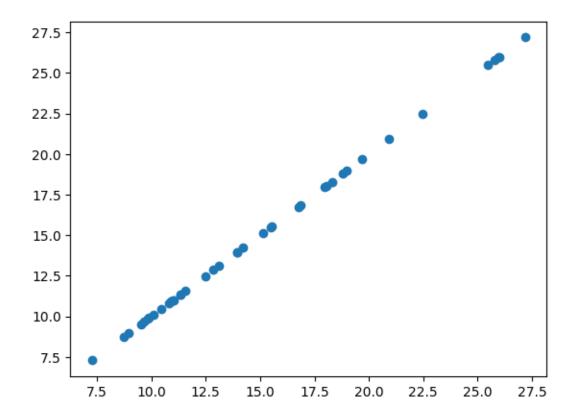
[ ]: lr.intercept_
[ ]: -3.552713678800501e-15

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

```
[]: Co-efficient
Age 1.629640e-18
Na_to_K 1.000000e+00
```

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

[]: <matplotlib.collections.PathCollection at 0x7e912221cc70>



4 ACCURACY

```
rr.fit(x_train,y_train)
[]: Ridge(alpha=10)
[]: rr.score(x_test,y_test)
[]: 0.999998773006138
[]: rr.score(x_train,y_train)
[]: 0.9999988058082202
[]: la=Lasso(alpha=10)
    la.fit(x_train,y_train)
[]: Lasso(alpha=10)
[]: la.score(x_test,y_test)
[]: 0.968148097167346
[]: la.score(x_train,y_train)
[]: 0.9693674618474863
[]: from sklearn.linear_model import ElasticNet
    en=ElasticNet()
    en.fit(x_train,y_train)
[]: ElasticNet()
[]: print(en.coef_)
    print(en.intercept_)
    [-0.
                  0.98264968]
    0.28287061102032496
[]: prediction = en.predict(x_test)
    prediction
[]: array([20.82909278, 18.94437069, 15.52966305, 15.50411416, 11.10577419,
            10.22138948, 25.62147527, 9.96098731, 9.09330765, 22.34925183,
           14.25221847, 19.61650307, 11.42906594, 13.97511126, 10.00520655,
            7.44147353, 17.92241502, 11.06941615, 12.9138496 , 11.64917946,
            14.0075387 , 10.92693195, 10.99178683, 13.18211297, 16.74520071,
            11.42611799, 10.54762917, 8.88105531, 25.80130016, 9.63179967,
            18.01281879, 12.56107837, 18.76552845, 16.84051772, 9.79197157,
```

25.80621341, 18.26044651, 26.99423687, 15.17590917, 25.31587122])

```
[]: en.score(x_test,y_test)
```

[]: 0.9996869834073963

```
[]: from sklearn import metrics
print("Mean Absolute Error: ", metrics.mean_absolute_error(y_test,prediction))
print("Mean Squared Error: ", metrics.mean_squared_error(y_test,prediction))
print("Root Mean Squared Error: ", np.sqrt(metrics.

-mean_squared_error(y_test,prediction)))
```

Mean Absolute Error: 0.08368614917403745
Mean Squared Error: 0.009428228438214944
Root Mean Squared Error: 0.09709906507384582