## Feature Transformation\*

# **Feature Scaling**

Definition: A technique to standardize the independent features present in the data in a fixed range.

Note: use feature scaling just before model building.

### Normalization

Definition: A technique often applied as a part of data preparation for Machine Learning. The goal of normalization is to change the values of numeric columns in the dataset to use common scale, without distorting differences in the range of values or loosing information.

#### Types:

- 1. MinMax Scaling(most popular, sometimes it is only called normalization)
- 2. Mean Normalization
- 3. Max absolute scaling
- 4. Robust Scaling

MinMax Scaler: x\_new=(x\_old-x\_min)/(x\_max-x\_min), range in [0,1], class-->MinMaxScaler

Geometrical intuition: data is set between range(depends on type of Normalization) and squeezed/expanded depends on data size, in unit square/rectangle(for 2D) and so on.

Mean Normalization : x\_new=(x\_old-x\_mean)/(x\_max-x\_min), range in [-1,1]

Here we are doing mean centering as we use to do in Standardization(so, this technique is used very rerely, instead we use Standardization). There is no separate class for this technique in Scikit-learn library, we have to code it manually.

Max absolute scaling: x\_new=(x\_old)/|x\_max|, class-->MaxAbsScaler

Mostly used when we have sparse data(means, data having many zeros)

Robust Scaling: x\_new=(x\_old-x\_median)/IQR, class-->RobustScaler

3 1 13.24 2.59 2.87 21.0 118 2.80 2.69 0.39 1.82 4.32 1.04 2.93 735 4 1 14.20 1.76 2.45 15.2 112 3.27 3.39 0.34 1.97 6.75 1.05 2.85 1450

This scaling is Robust to outliers(generally it performs well with outliers)

## **Topics Covered:**

- 1. How to Normalize(MinMaxScaler)
- 2. Effect of Normalization on outliers
- 3. Normalization Vs Standardization

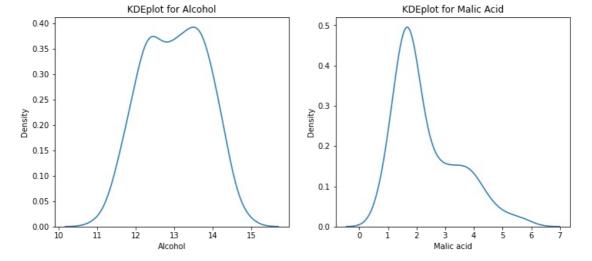
### Importing Dependencies

```
In [1]:
         import pandas as pd
         import numpy as np
         import matplotlib.pyplot as plt
         import seaborn as sns
In [2]:
         #load data
         df1 = pd.read csv('wine data.csv')
         #first 5 rows
         df1.head()
         1 14.23 1.71 2.43 15.6 127 2.8 3.06
                                                .28 2.29 5.64 1.04 3.92 1065
         0 1 13.20 1.78 2.14 11.2 100 2.65 2.76 0.26 1.28 4.38 1.05 3.40
                                                                        1050
         1 1 13.16 2.36 2.67 18.6 101 2.80 3.24 0.30 2.81 5.68 1.03 3.17 1185
         2 1 14.37 1.95 2.50 16.8 113 3.85 3.49 0.24 2.18 7.80 0.86 3.45
```

```
#we will use only first 3 columns in this session
df1 = pd.read_csv('wine_data.csv', header=None, usecols=[0,1,2])
#name features
df1.columns=['Class label', 'Alcohol', 'Malic acid']
#first 5 rows
df1.head()
```

# Out[3]: Class label Alcohol Malic acid 0 1 14.23 1.71 1 1 13.20 1.78 2 1 13.16 2.36 3 1 14.37 1.95 4 1 13.24 2.59

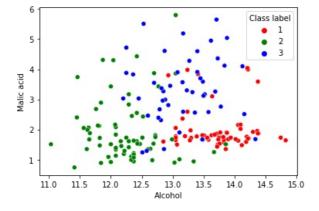
```
In [4]:
#Kernal Density Estimator graphs(kdeplot) for 'Alcohol' and 'Malic acid'
fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(12,5))
sns.kdeplot(df1['Alcohol'], ax=ax1)
ax1.set_title("KDEplot for Alcohol")
sns.kdeplot(df1['Malic acid'], ax=ax2)
ax2.set_title("KDEplot for Malic Acid")
plt.show()
```



```
In [5]:
#Scatter plots for 'Alcohol' and 'Malic acid', colored feature 'Class label'
color_dict={1:'red', 2:'green', 3:'blue'}
sns.scatterplot(df1['Alcohol'], df1['Malic acid'], hue=df1['Class label'], palette=color_dict)
```

C:\Users\RAM\Anaconda\lib\site-packages\seaborn\\_decorators.py:36: FutureWarning: Pass the following variables as
keyword args: x, y. From version 0.12, the only valid positional argument will be `data`, and passing other argum
ents without an explicit keyword will result in an error or misinterpretation.
 warnings.warn(

Out[5]: <AxesSubplot:xlabel='Alcohol', ylabel='Malic acid'>



[0.16935484, 0.20798319],

[0.21774194, 0.

```
In [6]:
         #recommended(train-test split)
         from sklearn.model_selection import train_test_split
         X_train, X_test, y_train, y_test=train_test_split(df1.drop('Class label', axis=1), df1['Class label'], test size=
         #shapes
         X_train.shape, X_test.shape, y_train.shape, y_test.shape
Out[6]: ((124, 2), (54, 2), (124,), (54,))
In [7]:
         #Normalization(MinMaxScaler)
         from sklearn.preprocessing import MinMaxScaler
         scaler=MinMaxScaler()
         # fit the scaler to the train set, it will learn the parameters
         scaler.fit(X train)
         # transform train and test sets
         X_train_scaled = scaler.transform(X_train)
         X test scaled = scaler.transform(X test)
In [9]:
         #by-default standardization converts dataframe into array
         X train scaled
Out[9]: array([[0.72043011, 0.20378151],
                [0.31989247, 0.08403361],
                [0.60215054, 0.71218487],
                [0.57258065, 0.56302521],
                [0.76075269, 0.1302521],
                [0.48924731, 0.5
                                       1,
                [0.75537634, 0.67857143],
                [0.61021505, 0.17436975],
                [0.54301075, 0.62394958],
                [0.39784946, 0.07352941],
                [0.33870968, 0.1092437],
                [0.46774194, 0.53361345],
                [0.5188172 , 0.53781513],
                [0.70967742, 0.07563025],
                [0.57258065, 0.30882353],
                [0.36021505, 0.0105042],
                [0.38709677, 0.13235294],
                [0.20967742, 0.25840336],
                [0.59408602, 0.64915966],
                [0.82526882, 0.26680672],
                [0.15591398, 0.09663866],
                [0.52688172, 0.16386555],
                [0.46774194, 0.31512605],
                [0.65860215, 0.16386555],
                [0.1155914 , 0.5987395 ],
                [0.27956989, 0.26680672],
                [0.21236559, 0.12184874],
                [0.65053763, 0.59033613],
                [0.31451613, 0.44957983],
                [0.54301075, 0.17647059],
                [0.57526882, 0.90336134],
                [0.28225806, 0.09243697],
                [0.30645161, 0.11764706],
                [0.37365591, 0.74369748],
                [0.22580645, 0.71848739],
                [0.3172043 , 0.06302521],
                [0.78225806, 0.17647059],
                [0.82526882, 0.23739496],
                [0.44086022, 0.01890756],
                          , 0.17647059],
                [0.32795699, 0.80462185],
                [0.48924731, 0.14915966],
                [0.72580645, 0.7289916],
                [0.81451613, 0.15546218],
                [0.69892473, 0.19327731],
                [0.34946237, 0.0210084],
                          , 0.18067227],
                [0.75
                [0.28225806, 0.25
                [0.53763441, 0.00210084],
                [0.89784946, 0.22268908],
                [0.57526882, 0.35714286],
                [0.45430108, 0.18067227],
                [0.90322581, 0.20588235],
                [0.62096774, 0.0105042],
```

```
[0.59139785, 0.50630252],
       [0.90053763, 0.56722689],
       [0.83064516, 0.12394958],
       [0.26075269, 0.53361345],
       [0.35215054, 0.32773109],
       [0.65860215, 0.19327731],
       [0.60752688, 0.22689076],
       [0.63978495, 0.6197479],
       [0.72043011, 1.
       [0.36021505, 0.06722689],
       [0.58333333, 0.18697479],
       [0.11290323, 0.31722689],
       [0.69623656, 0.85294118],
       [0.76612903, 0.21008403],
       [0.37365591, 0.3487395],
       [0.47849462, 0.29831933],
       [0.41666667, 0.08403361],
       [0.75268817, 0.14285714],
       [0.5483871, 0.12815126],
       [0.44892473, 0.55882353],
       [0.73655914, 0.21218487],
       [0.48655914, 0.43487395],
       [0.36021505, 0.15546218],
       [0.66666667, 0.19117647],
       [0.76344086, 0.21218487],
       [0.28225806, 0.10504202],
       [0.68548387, 0.16176471],
       [0.55107527, 0.63235294],
       [0.20430108, 0.2605042],
       [0.38172043, 0.44957983],
       [0.71236559, 0.19747899],
       [0.66935484, 0.4789916],
       [0.66397849, 0.46848739],
       [0.49462366, 0.78151261],
       [0.80376344, 0.16596639],
       [0.33870968, 0.4789916],
       [0.28225806, 0.05042017],
       [0.44892473, 0.62605042],
       [0. , 0.1302521],
[0.6155914 , 0.49369748],
       [0.83333333, 0.67436975],
       [0.66129032, 0.16176471],
       [0.21774194, 0.42016807],
       [0.54301075, 0.24369748],
       [0.45430108, 0.19327731],
       [0.4811828 , 0.5210084 ],
       [0.63709677, 0.77941176],
       [0.85752688, 0.6512605],
       [0.72311828, 0.11344538],
       [0.51075269, 0.40336134],
       [0.16397849, 0.24579832],
       [0.33870968, 0.1512605],
       [0.16666667, 0.16386555],
       [0.60483871, 0.15756303],
       [0.51075269, 0.61134454],
       [0.76075269, 0.09663866],
       [0.21236559, 0.17436975],
       [0.36021505, 0.05882353],
       [0.37365591, 0.1512605],
       [0.77150538, 0.16596639],
       [0.84139785, 0.34033613]])
#so we need to convert back array into dataframe
X train scaled = pd.DataFrame(X train scaled, columns=X train.columns)
X_test_scaled = pd.DataFrame(X_test_scaled, columns=X_test.columns)
#first 5 rows of X_train_scaled
X train scaled.head()
  Alcohol Malic acid
```

[0.49193548, 0.09663866], [0.7311828, 0.17647059], [0.40322581, 0.97058824], [0.55913978, 0.21218487], [0.90053763, 0.20588235], [0.86021505, 0.17226891], [0.26075269, 0.1302521],

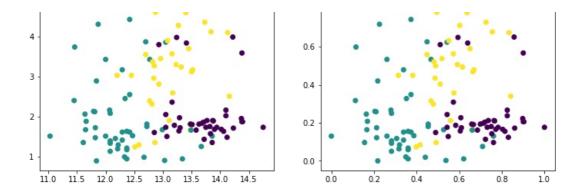
In [10]:

In [11]:

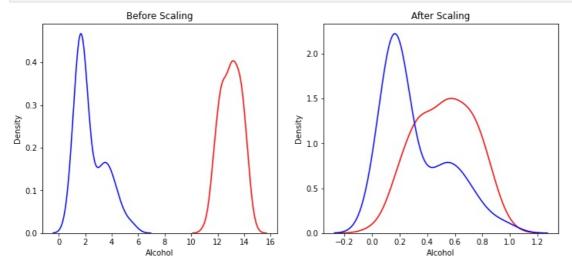
0 0.720430 0.203782

```
0.712185
          2 0.602151
          3 0.572581
                      0.563025
          4 0.760753
                      0.130252
In [12]:
           #first 5 rows of X_test_scaled
           X test scaled.head()
             Alcohol Malic acid
Out[12]:
          0 0.728495
                      0.163866
          1 0.473118
                     0.373950
          2 0.360215
                      0.050420
          3 0.680108
                      0.176471
          4 0.543011
                      1.031513
In [13]:
           #scatistical info before scaling
           X_train.describe()
                   Alcohol
                            Malic acid
          count 124.000000
                          124.000000
                 12.983065
                             2.383710
          mean
                  0.801340
                             1.136696
            std
           min
                 11.030000
                             0.890000
           25%
                 12.362500
                             1.607500
                 13.040000
           50%
                             1.885000
           75%
                 13.640000
                             3.247500
                 14.750000
                             5.650000
           max
In [14]:
           #scatistical info after scaling
           np.round(X_train_scaled.describe(), 1)
Out[14]:
                Alcohol Malic acid
                  124.0
                            124.0
          count
          mean
                    0.5
                              0.3
            std
                    0.2
                              0.2
                    0.0
                              0.0
           min
           25%
                    0.4
                              0.2
           50%
                    0.5
                              0.2
           75%
                    0.7
                              0.5
                    1.0
                              1.0
           max
In [15]:
           #Scatter plots for Alcohol, Malic acid colored by Class label, before and after scaling
           fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(12,5))
           ax1.scatter(X_train['Alcohol'], X_train['Malic acid'], c=y_train)
           ax1.set_title("Before Scaling")
           ax2.scatter(X train scaled['Alcohol'], X train scaled['Malic acid'], c=y train)
           ax2.set_title("After Scaling")
           plt.show()
           #observation: just range of data is changed(taken into [0,1]), distribution looks all same(most of times), but it
                                                                                After Scaling
                           Before Scaling
                                                            1.0
                                                            08
```

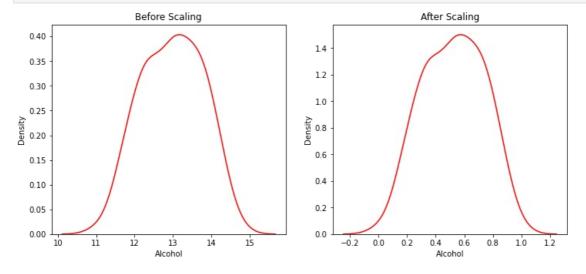
**1** 0.319892 0.084034



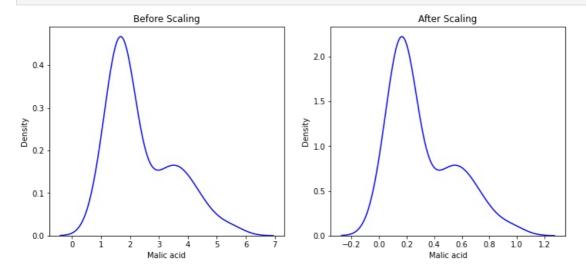
```
In [16]:
#Combined kernel Density Estimator(kde) plots for Alcohol, Malic acid colored by Class label, before and after so
fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(12,5))
sns.kdeplot(X_train['Alcohol'], ax=ax1, color='red')
sns.kdeplot(X_train['Malic acid'], ax=ax1, color='blue')
ax1.set_title("Before Scaling")
sns.kdeplot(X_train_scaled['Alcohol'], ax=ax2, color='red')
sns.kdeplot(X_train_scaled['Malic acid'], ax=ax2, color='blue')
ax2.set_title("After Scaling")
plt.show()
#observation : before scaling both features have different ranges, but after scaling both ae taken to same range
```



```
In [17]: #Kernal Density Estimator graphs(kdeplot) for 'Alcohol' before and after scaling
    fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(12,5))
    sns.kdeplot(X_train['Alcohol'], ax=ax1, color='red')
    ax1.set_title("Before Scaling")
    sns.kdeplot(X_train_scaled['Alcohol'], ax=ax2, color='red')
    ax2.set_title("After Scaling")
    plt.show()
    #observation : before and after scaling shapes looks same(most of times), but its not necessary, it depends on data
```



```
#Kernal Density Estimator graphs(kdeplot) for 'Malic acid' before and after scaling
fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(12,5))
sns.kdeplot(X_train['Malic acid'], ax=ax1, color='blue')
ax1.set_title("Before Scaling")
sns.kdeplot(X_train_scaled['Malic acid'], ax=ax2, color='blue')
ax2.set_title("After Scaling")
plt.show()
#observation : before and after scaling shapes looks same(most of times), but its not necessary, it depends on data
```



#### Effect of Normalization on Outliers

Since, we sqeeze range as a result impact of outlier also squeezes, we need to handle outliers separetely.

### Normalization Vs Standardization

- 1. Depends on type of data
- 2. Most of the problems are solved by using Standardization
- 3. Normalization(MinMaxScaler) is mostly used when we already know min and max values, example: CNN(image processing)
- 4. when have outliers use Robust scaling
- 5. when have sparse data try with MaxAbs scaling
- 6. when have no idea simply use Standardization

These are just tips, but not necessary that data will work best with these only, afterall feature engineering is all about exploration. So, we may try to apply all and check which one works best for your given data. And doing this we will learn more about data and scaling.

Note: Before applying scaling, first ask if scaling is really required? It depends on the algorithm we are going to apply, as some algorithms works good with scaling while others have no effect on them.

**END** of Documentation.

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