# **Initial Import for Standard Libraries**

```
In [0]: # importing libraries.!!

import pandas as pd
import numpy as np
import os
import seaborn as sns
from matplotlib import pyplot as plt

import warnings
warnings.filterwarnings('ignore')
In [0]: #Load the Data

df = pd.read_csv('./heart.csv')
```

In [183]: #Evaluating Non Zero Columns

df.isnull().sum()

Out[183]: 0 age sex 0 0 ср trestbps 0 chol 0 fbs 0 restecg 0 thalach 0 exang 0 oldpeak slope 0 ca 0

thal

target 0
dtype: int64

0

## There are No Columns that have Non Zero Value

In [184]: df.head()

Out[184]:

	age	sex	ср	trestbps	chol	fbs	restecg	thalach	exang
0	63	1	3	145	233	1	0	150	0
1	37	1	2	130	250	0	1	187	0
2	41	0	1	130	204	0	0	172	0
3	56	1	1	120	236	0	1	178	0
4	57	0	0	120	354	0	1	163	1

```
In [185]: f,ax=plt.subplots(1,2,figsize=(8,4))

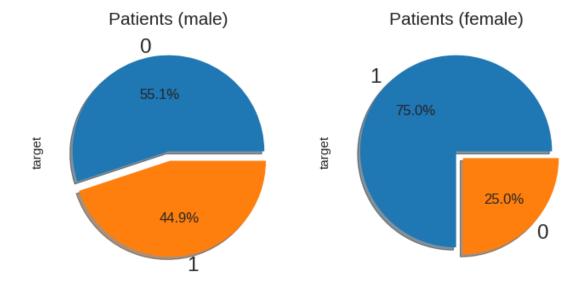
sns.set_context("paper", font_scale = 2, rc = {"font.size": 12,
    "axes.titlesize": 15,"axes.labelsize": 12})

df.loc[df['sex']==1, 'target'].value_counts().plot.pie(explode=[
    0,0.10],autopct='%1.1f%%',ax=ax[0],shadow=True)

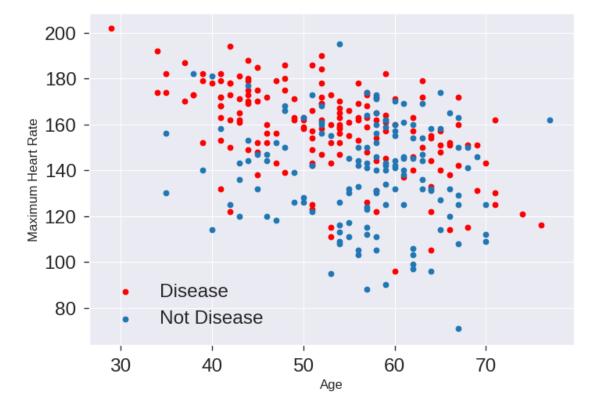
df.loc[df['sex']==0, 'target'].value_counts().plot.pie(explode=[
    0,0.10],autopct='%1.1f%%',ax=ax[1],shadow=True)

ax[0].set_title('Patients (male)')
ax[1].set_title('Patients (female)')

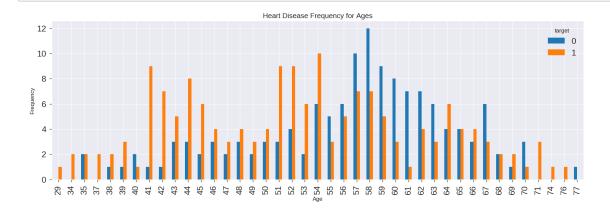
plt.show()
```



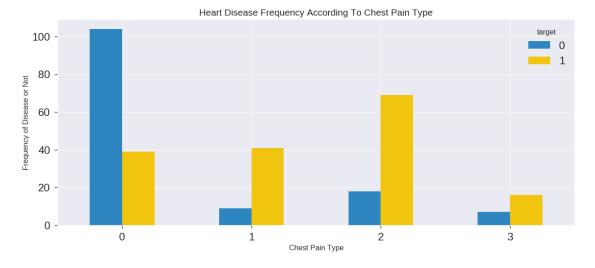
```
In [186]: plt.scatter(x=df.age[df.target==1], y=df.thalach[(df.target==1)], c="red")
    plt.scatter(x=df.age[df.target==0], y=df.thalach[(df.target==0)])
    plt.legend(["Disease", "Not Disease"])
    plt.xlabel("Age")
    plt.ylabel("Maximum Heart Rate")
    plt.show()
```



```
In [187]: pd.crosstab(df.age,df.target).plot(kind="bar",figsize=(20,6))
    plt.title('Heart Disease Frequency for Ages')
    plt.xlabel('Age')
    plt.ylabel('Frequency')
    plt.savefig('heartDiseaseAndAges.png')
    plt.show()
```



```
In [188]: pd.crosstab(df.cp,df.target).plot(kind="bar",figsize=(15,6),colo
    r=['#2E86C1','#F1C40F' ])
    plt.title('Heart Disease Frequency According To Chest Pain Type'
    )
    plt.xlabel('Chest Pain Type')
    plt.xticks(rotation = 0)
    plt.ylabel('Frequency of Disease or Not')
    plt.show()
```



### Use Pandas Profiling to Improve EDA for data.

In [189]:

pip install pandas-profiling

Requirement already satisfied: pandas-profiling in /usr/loca l/lib/python3.6/dist-packages (1.4.1)

Requirement already satisfied: pandas>=0.19 in /usr/local/lib/python3.6/dist-packages (from pandas-profiling) (0.25.3)

Requirement already satisfied: jinja2>=2.8 in /usr/local/lib/python3.6/dist-packages (from pandas-profiling) (2.10.3)

Requirement already satisfied: matplotlib>=1.4 in /usr/local/lib/python3.6/dist-packages (from pandas-profiling) (3.1.2)

Requirement already satisfied: six>=1.9 in /usr/local/lib/pyt hon3.6/dist-packages (from pandas-profiling) (1.12.0)

Requirement already satisfied: python-dateutil>=2.6.1 in /us r/local/lib/python3.6/dist-packages (from pandas>=0.19->panda s-profiling) (2.6.1)

Requirement already satisfied: numpy>=1.13.3 in /usr/local/lib/python3.6/dist-packages (from pandas>=0.19->pandas-profiling) (1.17.4)

Requirement already satisfied: pytz>=2017.2 in /usr/local/li b/python3.6/dist-packages (from pandas>=0.19->pandas-profilin g) (2018.9)

Requirement already satisfied: MarkupSafe>=0.23 in /usr/loca l/lib/python3.6/dist-packages (from jinja2>=2.8->pandas-profi ling) (1.1.1)

Requirement already satisfied: pyparsing!=2.0.4,!=2.1.2,!=2. 1.6,>=2.0.1 in /usr/local/lib/python3.6/dist-packages (from m atplotlib>=1.4->pandas-profiling) (2.4.5)

Requirement already satisfied: kiwisolver>=1.0.1 in /usr/loca l/lib/python3.6/dist-packages (from matplotlib>=1.4->pandas-p rofiling) (1.1.0)

Requirement already satisfied: cycler>=0.10 in /usr/local/li b/python3.6/dist-packages (from matplotlib>=1.4->pandas-profiling) (0.10.0)

Requirement already satisfied: setuptools in /usr/local/lib/p ython3.6/dist-packages (from kiwisolver>=1.0.1->matplotlib>= 1.4->pandas-profiling) (42.0.2)

In [0]: import pandas\_profiling

In [191]:

pandas profiling.ProfileReport(df)

Out[191]:

# **Overview**

Dataset info

Number of variables 14

Number of observations 303

Total Missing (%) 0.0%

Total size in memory 33.3 KiB

Average record size in memory 112.4 B

Numeric 10

Categorical 0

Boolean 4

**Date** 0

Text (Unique) 0

**Rejected** 0

**Unsupported** 0

Warnings

- <u>cp</u> has 143 / 47.2% zeros **Zeros**
- restecg has 147 / 48.5% zeros Zeros
- oldpeak has 99 / 32.7% zeros Zeros
- slope has 21 / 6.9% zeros Zeros
- <u>ca</u> has 175 / 57.8% zeros **Z**eros
- Dataset has 1 duplicate rows Warning

# **Variables**

#### age Numeric

**Distinct count** 41

**Unique (%)** 13.5%

Missing (%) 0.0%

Missing (n)

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 54.366

Minimum 29

Maximum 77

**Zeros (%)** 0.0%



## Toggle details

#### sex

Boolean

**Distinct count** 2

**Unique (%)** 0.7%

Missing (%) 0.0%

Missing (n) 0

**Mean** 0.68317

1 207

0 96

### Toggle details

cp Numeric

Distinct count 4

**Unique (%)** 1.3%

Missing (%) 0.0%

Missing (n) 0

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 0.967

**Minimum** 0

Maximum 3

**Zeros (%)** 47.2%



Toggle details

### trestbps

Numeric

**Distinct count** 49

**Unique (%)** 16.2%

**Missing (%)** 0.0%

Missing (n) 0

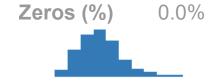
**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 131.62

Minimum 94

Maximum 200



## Toggle details

chol Numeric

**Distinct count** 152

**Unique (%)** 50.2%

Missing (%) 0.0%

Missing (n)

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 246.26

Minimum 126

Maximum 564

**Zeros (%)** 0.0%



Toggle details

fbs

Boolean

**Distinct count** 2

**Unique (%)** 0.7%

**Missing (%)** 0.0%

Missing (n) 0

**Mean** 0.14851

0 258

1 45

## Toggle details

restecg

Numeric

**Distinct count** 3

**Unique (%)** 1.0%

Missing (%) 0.0%

Missing (n)

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 0.52805

**Minimum** 0

Maximum 2

**Zeros (%)** 48.5%



Toggle details

thalach

Numeric

**Distinct count** 91

**Unique (%)** 30.0%

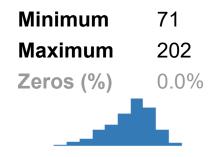
Missing (%) 0.0%

Missing (n) 0

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 149.65



## Toggle details

exang

Boolean

**Distinct count** 2

**Unique (%)** 0.7%

Missing (%) 0.0%

Missing (n) 0

**Mean** 0.32673

0 204

1 99

Toggle details

oldpeak

Numeric

**Distinct count** 40

**Unique (%)** 13.2%

**Missing (%)** 0.0%

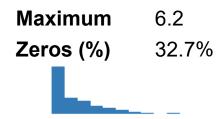
Missing (n) 0

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 1.0396

Minimum 0



Toggle details

slope

Numeric

**Distinct count** 3

**Unique (%)** 1.0%

Missing (%) 0.0%

Missing (n) 0

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 1.3993

Minimum 0

Maximum 2

**Zeros (%)** 6.9%

Toggle details

ca

Numeric

**Distinct count** 5

**Unique (%)** 1.7%

**Missing (%)** 0.0%

Missing (n) 0

**Infinite (%)** 0.0%

 Infinite (n)
 0

 Mean
 0.72937

 Minimum
 0

 Maximum
 4

 Zeros (%)
 57.8%

Toggle details

thal

Numeric

Distinct count 4

**Unique (%)** 1.3%

Missing (%) 0.0%

Missing (n) 0

**Infinite (%)** 0.0%

Infinite (n) 0

**Mean** 2.3135

**Minimum** 0

Maximum 3

**Zeros (%)** 0.7%

. . l ı

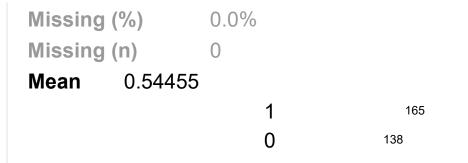
Toggle details

target

Boolean

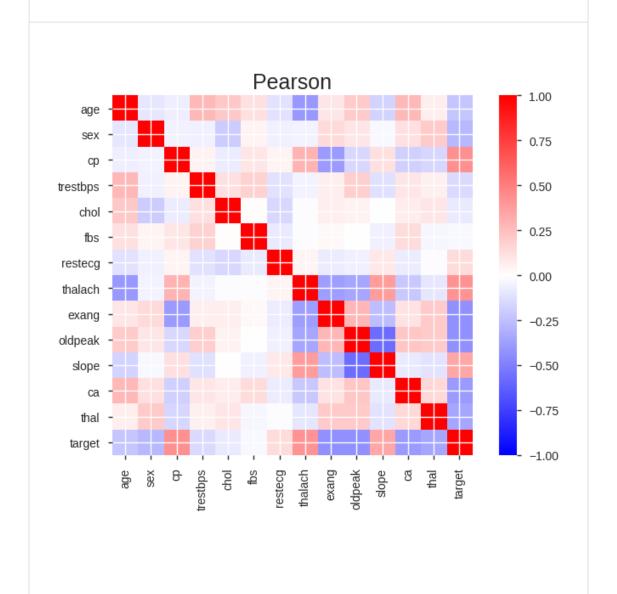
**Distinct count** 2

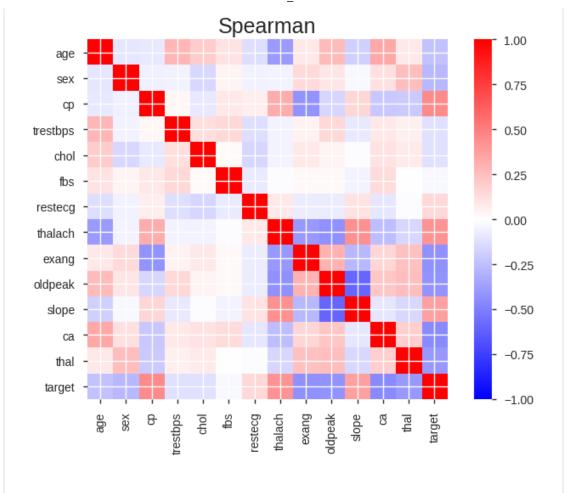
**Unique (%)** 0.7%



Toggle details

# **Correlations**





# **Sample**

	age	sex	ср	trestbps	chol	fbs	reste
0	63	1	3	145	233	1	0
1	37	1	2	130	250	0	1
2	41	0	1	130	204	0	0
3	56	1	1	120	236	0	1
4	57	0	0	120	354	0	1

## Plot the Split of Diseases across Male and Female

75.0 % of Women Suffer from Heart Diseases 45.0 % of Men Suffer from Heart Diseases

## **Train Test Split**

```
In [0]: X = df.iloc[:, :-1].values
y = df.iloc[:, -1].values
```

```
In [0]: from sklearn.model_selection import train_test_split
    X_train, X_test, y_train, y_test = train_test_split(X, y, test_s
    ize = 0.2, random_state = 0)
```

```
Model Evaluation

In [0]: # Importing Models that needs to be evaluated

from sklearn import model_selection
    from sklearn.naive_bayes import MultinomialNB
    from sklearn.tree import DecisionTreeClassifier
    from sklearn.svm import SVC
    from sklearn.calibration import CalibratedClassifierCV
    from sklearn.ensemble import AdaBoostClassifier
    from sklearn.linear_model import LogisticRegression
    from sklearn.ensemble import RandomForestClassifier
In [0]: # prepare models

model_list = []
```

```
In [0]: # Variable to Score Results
  results = []
  names = []
  scoring = 'accuracy'
  seed = 5
```

### **Cross Validation Score**

```
In [200]: # Evaluation of Each Model One by One (Cross Validation Score)

for name, model in model_list:
    kfold = model_selection.KFold(n_splits=10, random_state=seed)

    cv_results = model_selection.cross_val_score(model, X_train, y_train, cv=kfold, scoring=scoring)
    results.append(cv_results)
    names.append(name)
    msg = "%s: %f (%f)" % (name, cv_results.mean(), cv_results.s
td())
    print(msg)
```

```
Multinomial NB: 0.740000 (0.086165)

Decision Tree: 0.764833 (0.073088)

SVM: 0.831000 (0.066396)

ADA Boost With Decision Tree: 0.802333 (0.060755)

Logistic Regression: 0.831167 (0.070392)

Random Forest: 0.814667 (0.067814)
```

### Preparing a DataFrame to Plot the Scores

```
In [0]: def insert(df, row):
    insert_loc = df.index.max()
    if np.isnan(insert_loc):
        df.loc[0] = row
    else:
        df.loc[insert_loc + 1] = row
```

Out[202]:

Model\_Name CV\_Score

In [204]: cv\_score

Out[204]:

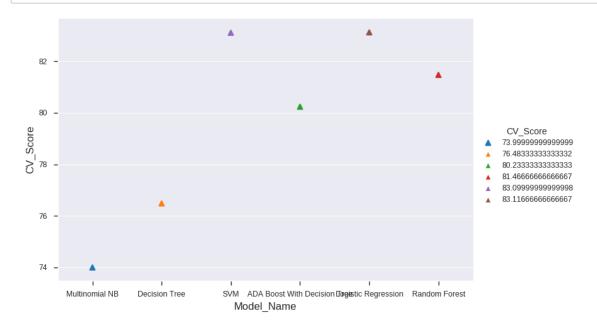
	Model_Name	CV_Score
0	Multinomial NB	74.000000
1	Decision Tree	76.483333
2	SVM	83.100000
3	ADA Boost With Decision Tree	80.233333
4	Logistic Regression	83.116667
5	Random Forest	81.466667

In [205]:

```
sns.set_context("paper", font_scale = 1, rc = {"font.size": 12,

"axes.titlesize": 15,"axes.labelsize": 12})

plot = sns.catplot(x="Model_Name", y="CV_Score", hue="CV_Score",
kind="point", data=cv_score,height=5,aspect=1.5,markers="^")
```



# Libaries to explain the Models

```
In [214]:
```

pip install eli5

Requirement already satisfied: eli5 in /usr/local/lib/python 3.6/dist-packages (0.10.1)

Requirement already satisfied: numpy>=1.9.0 in /usr/local/li b/python3.6/dist-packages (from eli5) (1.17.4)

Requirement already satisfied: graphviz in /usr/local/lib/pyt hon3.6/dist-packages (from eli5) (0.10.1)

Requirement already satisfied: scikit-learn>=0.18 in /usr/loc al/lib/python3.6/dist-packages (from eli5) (0.21.3)

Requirement already satisfied: tabulate>=0.7.7 in /usr/local/lib/python3.6/dist-packages (from eli5) (0.8.6)

Requirement already satisfied: scipy in /usr/local/lib/python 3.6/dist-packages (from eli5) (1.3.3)

Requirement already satisfied: attrs>16.0.0 in /usr/local/lib/python3.6/dist-packages (from eli5) (19.3.0)

Requirement already satisfied: six in /usr/local/lib/python3. 6/dist-packages (from eli5) (1.12.0)

Requirement already satisfied: jinja2 in /usr/local/lib/pytho n3.6/dist-packages (from eli5) (2.10.3)

Requirement already satisfied: joblib>=0.11 in /usr/local/lib/python3.6/dist-packages (from scikit-learn>=0.18->eli5) (0.14.0)

Requirement already satisfied: MarkupSafe>=0.23 in /usr/loca l/lib/python3.6/dist-packages (from jinja2->eli5) (1.1.1)

In [0]:

#Libraries for Explaning ML Models

import eli5 #for purmutation importance
from eli5.sklearn import PermutationImportance

In [0]:

df1 = df[df.columns.difference(['target'])]

In [0]:

perm\_list =[]

Out[220]:

```
In [0]:
         # Evaluation of Each Model One by One (Cross Validation Score)
         for name, model in model list:
             model.fit(X_train, y_train)
             perm = PermutationImportance(model, random state=1).fit(X te
         st, y_test)
             perm_list.append(perm)
```

In [0]: for index,value in enumerate(perm\_list): eli5.show\_weights(perm\_list[index], feature\_names = df1.colu mns.tolist())

In [220]: # Feature Importance using MultinomialNB eli5.show weights(perm list[0], feature names = df1.columns.toli st())

**Feature** 

Weight  $0.0230 \pm 0.0445$ restecg  $0.0131 \pm 0.0131$ ca  $0.0098 \pm 0.0533$ chol  $0.0033 \pm 0.0382$ slope  $0 \pm 0.0000$ trestbps  $0 \pm 0.0000$ thal  $0 \pm 0.0000$ fbs  $-0.0033 \pm 0.0321$ sex

> $-0.0098 \pm 0.0161$ oldpeak  $-0.0131 \pm 0.0131$ exang  $-0.0131 \pm 0.0245$ ср  $-0.0262 \pm 0.0334$ age  $-0.0393 \pm 0.0533$ thalach

In [221]: # Feature Importance using DecisionTreeClassifier eli5.show\_weights(perm\_list[1], feature\_names = df1.columns.toli st())

Out[221]:	Weight	Feature
	$0.0787 \pm 0.0564$	chol
	$0.0754 \pm 0.0445$	restecg
	0.0590 ± 0.0161	trestbps
	$0.0459 \pm 0.0435$	age
	0.0164 ± 0.0415	exang
	0.0131 ± 0.0131	thal
	0.0131 ± 0.0131	oldpeak
	0.0131 ± 0.0435	thalach
	0.0066 ± 0.0161	slope
	$0.0033 \pm 0.0245$	ср
	$0 \pm 0.0000$	sex
	$0 \pm 0.0000$	fbs
	-0.0131 ± 0.0482	ca

st())

Out[222]: Weight Feature

 $0.0164 \pm 0.0293$  trestbps

 $0.0164 \pm 0.0464$  ca

 $0.0131 \pm 0.0759$  slope

 $0.0131 \pm 0.0482$  chol

0.0033 ± 0.0525 thalach

 $0 \pm 0.0000$  thal

 $0 \pm 0.0000$  exang

 $0 \pm 0.0000$  age

 $-0.0033 \pm 0.0131$  cp

-0.0066 ± 0.0161 oldpeak

 $-0.0098 \pm 0.0161$  fbs

 $-0.0164 \pm 0.0207$  sex

 $-0.0197 \pm 0.0131$  restecg

In [223]: # Feature Importance using AdaBoostClassifier eli5.show\_weights(perm\_list[3], feature\_names = df1.columns.toli st())

Out[223]:	Weight	Feature
	$0.0787 \pm 0.0382$	chol
	$0.0754 \pm 0.0675$	trestbps
	$0.0656 \pm 0.0549$	age
	$0.0492 \pm 0.0359$	slope
	$0.0492 \pm 0.0207$	ca
	$0.0328 \pm 0.0994$	thalach
	$0.0164 \pm 0.0415$	restecg
	$0.0131 \pm 0.0435$	exang
	$0.0131 \pm 0.0245$	fbs
	$0.0066 \pm 0.0262$	thal
	0.0066 ± 0.0161	sex
	$0.0033 \pm 0.0321$	oldpeak
	$-0.0066 \pm 0.0572$	ср

In [224]: # Feature Importance using LogisticRegression eli5.show\_weights(perm\_list[4], feature\_names = df1.columns.toli st())

Out[224]:	Weight	Feature
	$0.0918 \pm 0.0334$	chol
	$0.0492 \pm 0.0464$	ca
	$0.0328 \pm 0.0207$	trestbps
	$0.0262 \pm 0.0334$	restecg
	$0.0197 \pm 0.0525$	slope
	$0.0164 \pm 0.0549$	thalach
	$0.0164 \pm 0.0207$	thal
	0.0131 ± 0.0131	exang
	0.0066 ± 0.0161	ср
	$0 \pm 0.0000$	fbs
	$0 \pm 0.0000$	age
	-0.0033 ± 0.0245	oldpeak

 $-0.0066 \pm 0.0262$ 

sex

In [225]: # Feature Importance using RandomForestClassifier
 eli5.show\_weights(perm\_list[5], feature\_names = df1.columns.toli
 st())

Weight Out[225]: **Feature**  $0.1082 \pm 0.0894$ trestbps thalach  $0.0262 \pm 0.0572$  $0.0066 \pm 0.0262$ sex  $0.0066 \pm 0.0262$ exang  $0.0066 \pm 0.0262$ ca  $0.0033 \pm 0.0382$ restecg  $-0.0066 \pm 0.0334$ chol oldpeak  $-0.0098 \pm 0.0161$ fbs  $-0.0098 \pm 0.0161$  $-0.0098 \pm 0.0262$ age thal  $-0.0131 \pm 0.0382$ slope  $-0.0131 \pm 0.0321$  $-0.0492 \pm 0.0359$ ср

# **Accuracy & RoC Curve**

```
In [0]: from sklearn.ensemble import RandomForestClassifier from sklearn.metrics import confusion_matrix from sklearn.metrics import accuracy_score from sklearn.metrics import roc_curve from sklearn.metrics import f1_score from sklearn.metrics import auc

from sklearn.metrics import precision_recall_curve from sklearn.metrics import average_precision_score
```

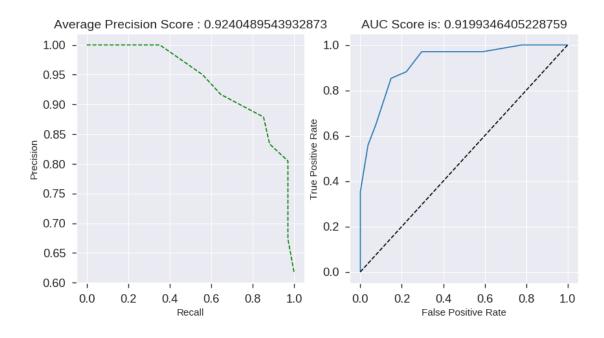
```
In [0]:
        def plotting(true,pred):
             fig,ax=plt.subplots(1,2,figsize=(10,5))
             precision,recall,threshold = precision recall curve(true,pre
        d[:,1])
             ax[0].plot(recall, precision, 'g--')
             ax[0].set_xlabel('Recall')
             ax[0].set_ylabel('Precision')
             ax[0].set_title("Average Precision Score : {}".format(average)
         e_precision_score(true,pred[:,1])))
             fpr,tpr,threshold = roc curve(true,pred[:,1])
             ax[1].plot(fpr,tpr)
             ax[1].set_title("AUC Score is: {}".format(auc(fpr,tpr)))
             ax[1].plot([0,1],[0,1],'k--')
             ax[1].set_xlabel('False Positive Rate')
             ax[1].set ylabel('True Positive Rate')
```

```
In [0]: sns.set_context("paper", font_scale = 1.5, rc = {"font.size": 11
,"axes.titlesize": 14,"axes.labelsize": 11})
```

### **Random Forest**

In [230]: plotting(y\_test,RandomForestClassifier.predict\_proba(X\_test))
 plt.figure()

Out[230]: <Figure size 800x550 with 0 Axes>



<Figure size 800x550 with 0 Axes>

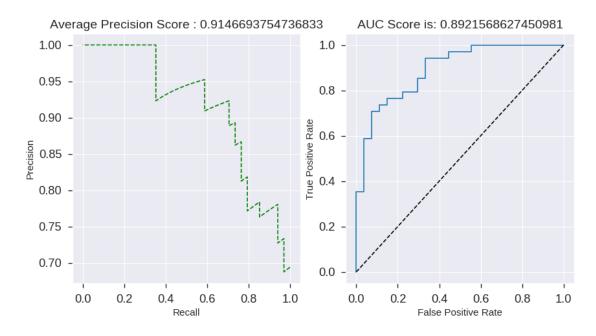
### **MultinomialNB**

In [231]: MultinomialNB = MultinomialNB(alpha=0.1)
 MultinomialNB.fit(X\_train,y\_train)

Out[231]: MultinomialNB(alpha=0.1, class\_prior=None, fit\_prior=True)

In [232]: plotting(y\_test,MultinomialNB.predict\_proba(X\_test))
 plt.figure()

Out[232]: <Figure size 800x550 with 0 Axes>



<Figure size 800x550 with 0 Axes>

### **DecisionTreeClassifier**

In [233]: DecisionTreeClassifier = DecisionTreeClassifier()
 DecisionTreeClassifier.fit(X\_train,y\_train)

Out[233]: DecisionTreeClassifier(class\_weight=None, criterion='gini', m ax\_depth=None,

max\_features=None, max\_leaf\_nodes=None, max\_lea

max\_features=None, max\_leaf\_nodes=Non

e,

min\_impurity\_decrease=0.0, min\_impurit

y\_split=None,

min samples leaf=1, min samples split=

2,

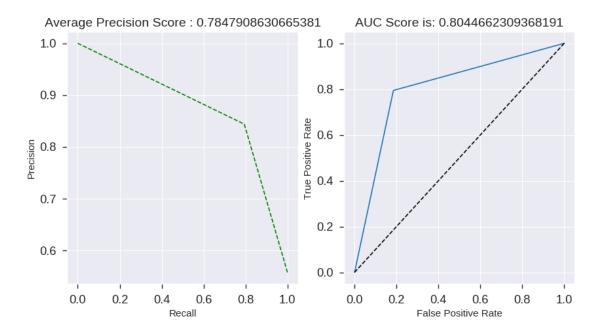
min\_weight\_fraction\_leaf=0.0, presort=

False,

random\_state=None, splitter='best')

In [234]: plotting(y\_test,DecisionTreeClassifier.predict\_proba(X\_test))
 plt.figure()

Out[234]: <Figure size 800x550 with 0 Axes>



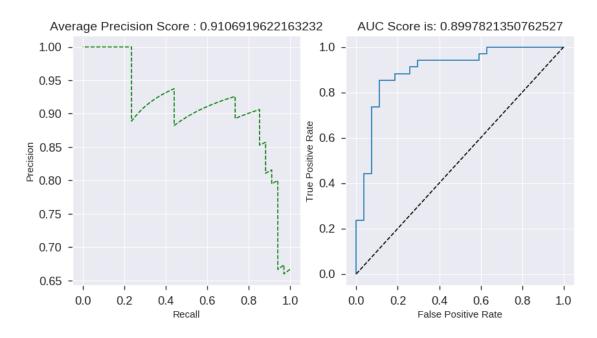
<Figure size 800x550 with 0 Axes>

### **Linear SVC**

In [235]: LinearSVC = SVC(kernel='linear',probability=True)
 LinearSVC.fit(X\_train,y\_train)

In [236]: plotting(y\_test,LinearSVC.predict\_proba(X\_test))
 plt.figure()

Out[236]: <Figure size 800x550 with 0 Axes>



<Figure size 800x550 with 0 Axes>

### AdaBoostClassifier

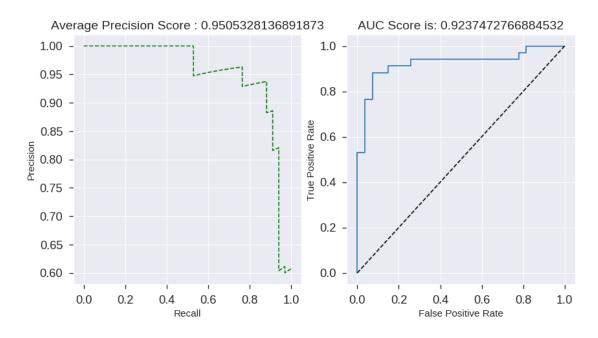
In [237]: AdaBoostClassifier = AdaBoostClassifier()
 AdaBoostClassifier.fit(X\_train,y\_train)

Out[237]: AdaBoostClassifier(algorithm='SAMME.R', base\_estimator=None, learning\_rate=1.0,

n\_estimators=50, random\_state=None)

In [238]: plotting(y\_test,AdaBoostClassifier.predict\_proba(X\_test))
 plt.figure()

Out[238]: <Figure size 800x550 with 0 Axes>



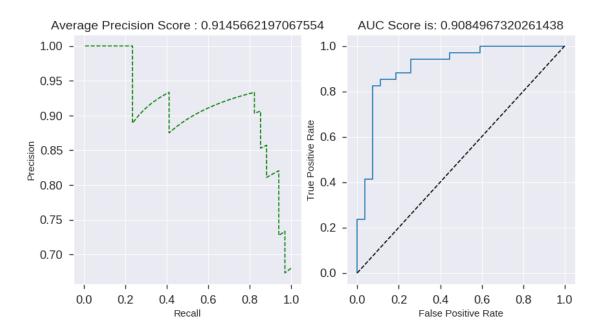
<Figure size 800x550 with 0 Axes>

# LogisticRegression

In [239]: LogisticRegression = LogisticRegression(solver='liblinear')
 LogisticRegression.fit(X\_train,y\_train)

In [240]: plotting(y\_test,LogisticRegression.predict\_proba(X\_test))
 plt.figure()

Out[240]: <Figure size 800x550 with 0 Axes>



<Figure size 800x550 with 0 Axes>