# **Logistic Regression Classifier**

## In [1]:

```
import numpy as np
import pandas as pd
import statsmodels.api as sm
import matplotlib.pyplot as plt
import seaborn as sns
```

# **Dataset**

#### In [2]:

```
dta = sm.datasets.fair.load_pandas().data
dta.head(5)
```

## Out[2]:

	rate_marriage	age	yrs_married	children	religious	educ	occupation	occupation_husb	
0	3.0	32.0	9.0	3.0	3.0	17.0	2.0	5.0	0
1	3.0	27.0	13.0	3.0	1.0	14.0	3.0	4.0	3
2	4.0	22.0	2.5	0.0	1.0	16.0	3.0	5.0	1.
3	4.0	37.0	16.5	4.0	3.0	16.0	5.0	5.0	0.
4	5.0	27.0	9.0	1.0	1.0	14.0	3.0	4.0	4.
4									•

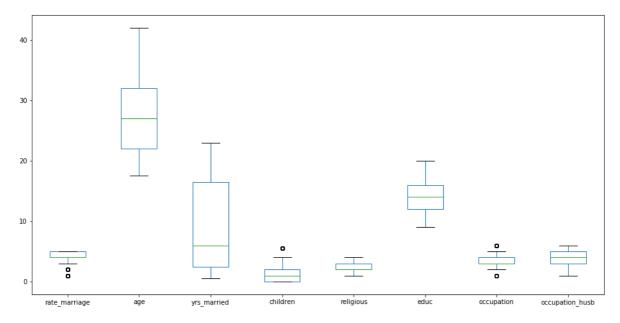
Checking for outliers using box plot

# In [3]:

```
1 dta.drop(['affairs'],axis=1).plot(kind='box',figsize=(16,8))
```

# Out[3]:

<matplotlib.axes.\_subplots.AxesSubplot at 0x187078d18b0>



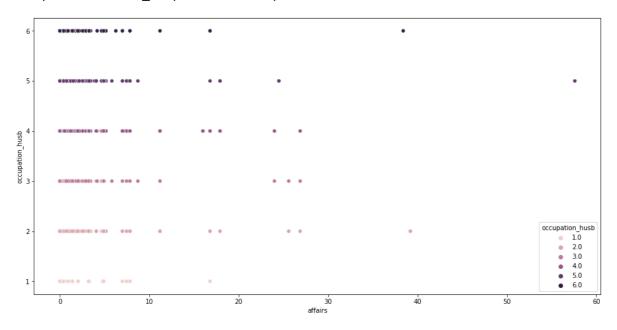
# Relation between Affair and Occupation of Husband

#### In [4]:

```
fig = plt.figure(figsize=(16,8))
sns.scatterplot(x=dta['affairs'],y=dta['occupation_husb'],hue=dta['occupation_husb'])
```

#### Out[4]:

<matplotlib.axes.\_subplots.AxesSubplot at 0x18765fea4c0>



We can see that the chances of Divorce differes with the occupation hunbands is in.

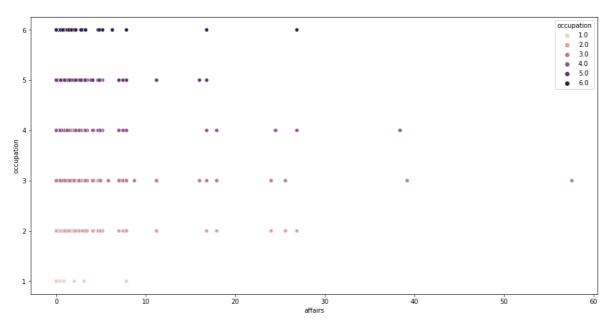
# Relation between Affair and Occupation of Women.

#### In [5]:

```
fig = plt.figure(figsize=(16,8))
sns.scatterplot(x=dta['affairs'],y=dta['occupation'],hue=dta['occupation'])
```

## Out[5]:

<matplotlib.axes.\_subplots.AxesSubplot at 0x18707a698e0>



Women having White Collar Job show high chances of Affair.

# **Information about Dataset**

The dataset contains 6366 observations of 9 variables:

```
rate_marriage: woman's rating of her marriage

(1 = very poor, 5 = very good)
```

age: woman's age

yrs\_married: number of years married

children: number of children

religious: woman's rating of how religious she is

(1 = not religious, 4 =strongly religious)

educ: level of education

(9 = grade school, 12 = high school, 14 = some college, 16 = college graduate, 17 = some graduate school, 20 = advanced degree)

occupation: woman's occupation

(1 = student, 2 = farming/semi- skilled/unskilled, 3 = "white collar", 4 = teacher/nurse/writer/technician/skilled, 5 = managerial/business, 6 = professional with advanced degree)

occupation\_husb: husband's occupation (same coding as above)

(1 = student, 2 = farming/semi- skilled/unskilled, 3 = "white collar", 4 = teacher/nurse/writer/technician/skilled, 5 = managerial/business, 6 = professional with advanced degree)

affairs: time spent in extra-marital affairs

Making a new Column that will contain Affairs in Binary Form

0 - No Affairs

1 - Having Affairs

#### In [6]:

```
1 dta['Affairs'] = (dta.affairs> 0).astype(int)
 dta.head(3)
```

### Out[6]:

	rate_marriage	age	yrs_married	children	religious	educ	occupation	occupation_husb	
0	3.0	32.0	9.0	3.0	3.0	17.0	2.0	5.0	0
1	3.0	27.0	13.0	3.0	1.0	14.0	3.0	4.0	3.
2	4.0	22.0	2.5	0.0	1.0	16.0	3.0	5.0	1.
4									•

## **Checking For Null Values**

#### In [7]:

```
1 print(dta.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 6366 entries, 0 to 6365
Data columns (total 10 columns):
```

rate\_marriage 6366 non-null float64 6366 non-null float64 age yrs\_married 6366 non-null float64 6366 non-null float64 children religious 6366 non-null float64 educ 6366 non-null float64 occupation 6366 non-null float64 occupation\_husb 6366 non-null float64 6366 non-null float64 affairs Affairs 6366 non-null int32

dtypes: float64(9), int32(1) memory usage: 472.6 KB

None

No Null values are there in any column. Data type is Float, with affairs as int

# **Creating Design Matrix**

We are creating two design matrices. The first is the martix of endogenous variables(dependent variables ~ 'X'). The second design matrix is of exogenous variables(Independent Variables ~ 'y'). For this we are uisng dmatrices function from patsy

#### In [8]:

```
from patsy import dmatrices
y,X = dmatrices('Affairs ~ rate_marriage+age+yrs_married+children+religious+educ+C(occu
X.head(2)
```

#### Out[8]:

	Intercept	C(occupation) [T.2.0]	C(occupation) [T.3.0]	C(occupation) [T.4.0]	C(occupation) [T.5.0]	C(occupation) [T.6.0]	C(occur
0	1.0	1.0	0.0	0.0	0.0	0.0	
1	1.0	0.0	1.0	0.0	0.0	0.0	
4							•

dmatrices returns a dataframe constituting of categories for categorical variables in our case occupation(female occupation) and occupation husb(Husbands occupations).

It add a contant variable "intercept" to the dataframe.

#### In [9]:

```
1 y.head(2)
```

#### Out[9]:

	Affairs
0	1.0
1	1.0

#### Converting y into a 1 D flattern array.

#### In [10]:

```
1 y = np.ravel(y)
2 y
```

#### Out[10]:

```
array([1., 1., 1., ..., 0., 0., 0.])
```

## Renaming Columns of categorical variables in X

for simplicity

#### In [11]:

```
1  X.rename(inplace=True,columns={'C(occupation)[T.2.0]':'occ_2.0','C(occupation)[T.3.0]':
2  X.head(5)
```

#### Out[11]:

										4
	Intercept	occ_2.0	occ_3.0	occ_4.0	occ_5.0	occ_6.0	occ_husb_2.0	occ_husb_3.0	occ	
0	1.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0		
1	1.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0		
2	1.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0		
3	1.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0		
4	1.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0		_
4									•	

#### In [12]:

```
print("Info of X:\n")
X.info()
```

#### Info of X:

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 6366 entries, 0 to 6365
Data columns (total 17 columns):
Intercept
                 6366 non-null float64
occ_2.0
                 6366 non-null float64
occ_3.0
                 6366 non-null float64
occ_4.0
                 6366 non-null float64
occ_5.0
                 6366 non-null float64
                 6366 non-null float64
occ_6.0
                 6366 non-null float64
occ_husb_2.0
                 6366 non-null float64
occ_husb_3.0
                 6366 non-null float64
occ_husb_4.0
occ_husb_5.0
                 6366 non-null float64
occ husb 6.0
                 6366 non-null float64
                 6366 non-null float64
rate_marriage
                 6366 non-null float64
age
                 6366 non-null float64
yrs_married
children
                 6366 non-null float64
religious
                 6366 non-null float64
                 6366 non-null float64
educ
dtypes: float64(17)
memory usage: 895.2 KB
```

Everything looks good. Now let's check whether anyvalues is null values

#### In [13]:

```
1 X.isna().sum()
```

#### Out[13]:

```
Intercept
                 0
occ_2.0
                 0
occ_3.0
                 0
                 0
occ_4.0
occ_5.0
                 0
occ_6.0
                 0
occ_husb_2.0
                 0
occ_husb_3.0
                 0
occ_husb_4.0
                 0
occ_husb_5.0
                 0
occ_husb_6.0
                 0
rate_marriage
                 0
age
yrs_married
                 0
                 0
children
religious
                 0
                 0
educ
dtype: int64
```

No Null values are there so Moving forward with our next step.

# **Checking Correlation**

#### Correlation

Relation of a variable with other variables.

# In [14]:

1 X.corr()

# Out[14]:

	Intercept	occ_2.0	occ_3.0	occ_4.0	occ_5.0	occ_6.0	occ_husb_2.0	01
Intercept	NaN							
occ_2.0	NaN	1.000000	-0.348075	-0.251243	-0.143237	-0.052128	0.183782	
occ_3.0	NaN	-0.348075	1.000000	-0.560645	-0.319631	-0.116322	-0.000638	
occ_4.0	NaN	-0.251243	-0.560645	1.000000	-0.230712	-0.083962	-0.083123	
occ_5.0	NaN	-0.143237	-0.319631	-0.230712	1.000000	-0.047868	-0.053426	
occ_6.0	NaN	-0.052128	-0.116322	-0.083962	-0.047868	1.000000	-0.046140	
occ_husb_2.0	NaN	0.183782	-0.000638	-0.083123	-0.053426	-0.046140	1.000000	
occ_husb_3.0	NaN	-0.020904	0.090043	-0.043159	-0.044053	-0.029028	-0.146849	
occ_husb_4.0	NaN	-0.009786	0.011248	0.037341	-0.039932	-0.043541	-0.347951	
occ_husb_5.0	NaN	-0.093292	0.003021	-0.001946	0.114903	-0.030926	-0.316693	
occ_husb_6.0	NaN	-0.059107	-0.101673	0.085766	0.006016	0.218824	-0.153248	
rate_marriage	NaN	-0.019697	-0.053082	0.068882	-0.002109	0.008878	-0.038992	
age	NaN	-0.034223	-0.066371	0.040982	0.079533	0.030676	-0.057368	
yrs_married	NaN	0.004668	-0.021261	-0.026816	0.076820	-0.004912	-0.033451	
children	NaN	0.081182	-0.063298	-0.003235	0.033274	-0.026830	0.001190	
religious	NaN	-0.013129	-0.034986	0.043996	0.004260	0.011784	0.009990	
educ	NaN	-0.217719	-0.335615	0.477505	-0.022121	0.226920	-0.160756	
4								•

# VIF Score

VIF Score > 5 Indicates sever correlation.

#### In [15]:

```
# VIF Score
from statsmodels.stats.outliers_influence import variance_inflation_factor as vif
from sklearn.preprocessing import StandardScaler as StdS
scaler = StdS()
X_data = scaler.fit_transform(X=X)
vif_score = [vif(X_data,i) for i in range(X_data.shape[1])]
for i in range(len(X.columns)):
    print(X.columns[i],":",vif_score[i])
datset1 = X
Intercept : nan
```

occ\_2.0 : 19.340780081786384 occ\_3.0 : 39.33561800583917 occ\_4.0 : 32.93190959613787 occ\_5.0 : 17.05716507563386 occ 6.0 : 3.697958521454455 occ\_husb\_2.0 : 5.5662915598555065 occ\_husb\_3.0 : 2.9910696353751987 occ\_husb\_4.0 : 6.930281429175677 occ\_husb\_5.0 : 6.577077423174822 occ\_husb\_6.0 : 3.1852660411536395 rate marriage : 1.0387458559035507 age: 5.477889737628554 yrs\_married : 7.16961101396522 children : 2.5856914303595504 religious : 1.0375560897654312 educ: 1.6357895734289674

C:\Users\adity\anaconda3\lib\site-packages\statsmodels\regression\linear\_mod
el.py:1717: RuntimeWarning: invalid value encountered in double\_scalars
 return 1 - self.ssr/self.uncentered\_tss

#### In [16]:

```
for i in range(len(X.columns)):
    print(X.columns[i],":",vif_score[i])
```

Intercept : nan occ 2.0 : 19.340780081786384 occ\_3.0 : 39.33561800583917 occ 4.0 : 32.93190959613787 occ\_5.0 : 17.05716507563386 occ\_6.0 : 3.697958521454455 occ husb 2.0 : 5.5662915598555065 occ husb 3.0 : 2.9910696353751987 occ husb 4.0 : 6.930281429175677 occ\_husb\_5.0 : 6.577077423174822 occ\_husb\_6.0 : 3.1852660411536395 rate\_marriage : 1.0387458559035507 age: 5.477889737628554 yrs\_married : 7.16961101396522 children: 2.5856914303595504 religious : 1.0375560897654312 educ: 1.6357895734289674

Now from this VIF Score it is clear that there are features having very high correlation values. occ\_2.0,occ\_3.0,occ\_4.0 and occ\_5.0. In occ\_husb - occ\_husb\_2.0, occ\_husb\_4.0, occ\_husb\_5.0 are also having high correlation.

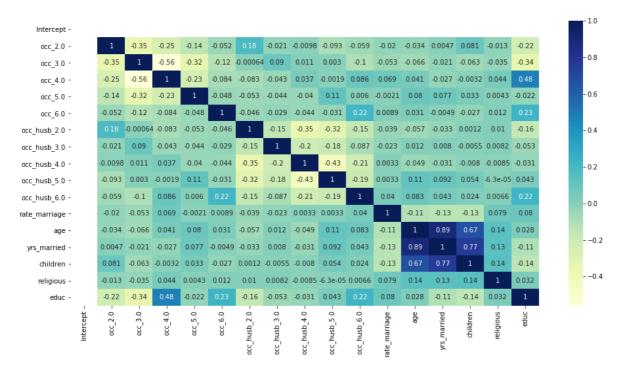
#### Ploting heat map with correlation value with X.

#### In [17]:

```
fig = plt.figure(figsize=(16,8))
sns.heatmap(data=X.corr(),cmap="YlGnBu", annot=True)
```

#### Out[17]:

<matplotlib.axes.\_subplots.AxesSubplot at 0x1870a279280>



As we can observe that in occ 2.0,occ 3.0,occ 4.0 and occ 5.0.

occ\_2.0 with {occ\_2.0,occ\_3.0,occ\_4.0 and occ\_5.0} is {1.000000 -0.348075 -0.251243 -0.143237} respectively.

occ\_3.0 with {occ\_2.0,occ\_3.0,occ\_4.0 and occ\_5.0} is {-0.348075 1.000000 -0.560645 -0.319631} respectively.

occ\_4.0 with {occ\_2.0,occ\_3.0,occ\_4.0 and occ\_5.0} is {-0.251243 -0.560645 1.000000 -0.230712} respectively.

occ\_5.0 with {occ\_2.0,occ\_3.0,occ\_4.0 and occ\_5.0} is {-0.143237 -0.319631 -0.230712 1.000000} respectively.

The feature occ\_3.0 is having high correlation in respect to all other features. So removing occ\_3.0 and lets check the effect on VIF Score.

#### In [18]:

```
1  X_data = scaler.fit_transform(X=X.drop('occ_3.0',axis=1))
2  vif_score = [vif(X_data,i) for i in range(X_data.shape[1])]
3  for i in range(len(X.drop('occ_3.0',axis=1).columns)):
4    print(X.drop('occ_3.0',axis=1).columns[i],":",vif_score[i])
```

Intercept : nan

occ\_4.0 : 1.58134483216193 occ\_5.0 : 1.1518453049129829 occ\_6.0 : 1.166552590194721 occ\_husb\_2.0 : 5.530740700962974 occ\_husb\_3.0 : 2.9761967717120825 occ\_husb\_4.0 : 6.8868032112916735 occ\_husb\_5.0 : 6.533261649523526 occ\_husb\_6.0 : 3.1752920128006568 rate\_marriage : 1.038534992602799

occ\_2.0 : 1.177577731855246

age: 5.474644652197843

yrs\_married : 7.169399081318488
children : 2.5846493806759088
religious : 1.037500793190026
educ : 1.6292132746346843

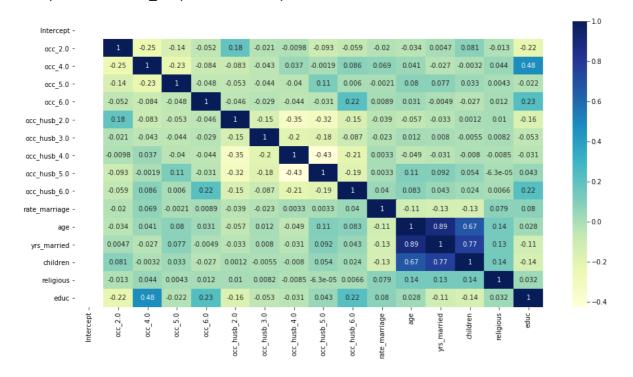
C:\Users\adity\anaconda3\lib\site-packages\statsmodels\regression\linear\_mod
el.py:1717: RuntimeWarning: invalid value encountered in double\_scalars
 return 1 - self.ssr/self.uncentered\_tss

#### In [19]:

```
fig = plt.figure(figsize=(16,8))
sns.heatmap(data=X.drop('occ_3.0',axis=1).corr(),cmap="YlGnBu", annot=True)
```

#### Out[19]:

<matplotlib.axes.\_subplots.AxesSubplot at 0x1870b800070>



#### In [20]:

1 X.head(2)

#### Out[20]:

	Intercept	occ_2.0	occ_3.0	occ_4.0	occ_5.0	occ_6.0	occ_husb_2.0	occ_husb_3.0	occ_hu
0	1.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	_
1	1.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	
4									•

#### In [21]:

```
from sklearn.model_selection import train_test_split as tts
    x_train,x_test,y_train,y_test = tts(X,y,test_size=0.20,random_state=225)
```

# **Using Stochastic Gradient Descent**

#### In [22]:

```
from sklearn.linear_model import SGDClassifier
model1 = SGDClassifier()
model1.fit(x_train,y_train)
```

## Out[22]:

```
SGDClassifier(alpha=0.0001, average=False, class_weight=None, early_stopping=False, epsilon=0.1, eta0=0.0, fit_intercept=True,

l1_ratio=0.15, learning_rate='optimal', loss='hinge', max_iter=1000, n_iter_no_change=5, n_jobs=None, penalty='l2', power_t=0.5, random_state=None, shuffle=True, tol=0.001, validation_fraction=0.1, verbose=0, warm_start=False)
```

#### **Testing Score**

#### In [23]:

```
1 model1.score(x_train,y_train)
```

#### Out[23]:

0.43676355066771405

#### **Traning Score**

```
In [24]:
```

```
1 model1.score(x_test,y_test)
```

#### Out[24]:

0.4340659340659341

# **Using LogisticRegression Classifier**

```
In [25]:
```

```
from sklearn.linear_model import LogisticRegression
model2 = LogisticRegression(solver='liblinear')
model2.fit(x_train,y_train)
```

#### Out[25]:

```
LogisticRegression(C=1.0, class_weight=None, dual=False, fit_intercept=True, intercept_scaling=1, l1_ratio=None, max_iter=100, multi_class='auto', n_jobs=None, penalty='l2', random_state=None, solver='liblinear', tol=0.0001, verbos e=0, warm_start=False)
```

#### **Traning Score**

```
In [26]:
```

```
model2.score(x_train,y_train)*100
```

#### Out[26]:

73.19324430479183

#### **Testing Score**

```
In [27]:
```

```
1 model2.score(x_test,y_test)*100
```

#### Out[27]:

69.78021978021978

#### **Evaluation of Classification Model**

- 1. Accuracy
- 2. Recall
- 3. Precision
- 4. F1 Score
- 5. Specifity
- 6. AUC (Area Under Curve)
- 7. RUC(Receiver Operator Characteristic)

#### **Cross Validation**

#### In [28]:

```
# Performing cross validation for model1 and Model 2

from sklearn.model_selection import cross_val_predict
model1_y_predict = cross_val_predict(model1,x_train,y_train,cv=3)
model2_y_predict = cross_val_predict(model2,x_train,y_train,cv=3)
```

## **Confusion Matrix**

	Actual	Values
Predicted	True Positive	False Positive
Values	False Negative	True Negative

#### For Model 1

#### In [29]:

```
from sklearn.metrics import confusion_matrix,recall_score, precision_score, f1_score, a
conf_matrix_model1 = confusion_matrix(y_train,model1_y_predict)
conf_matrix_model1
```

#### Out[29]:

```
array([[3021, 427], [1088, 556]], dtype=int64)
```

#### For Model 2

#### In [30]:

```
conf_matrix_model2 = confusion_matrix(y_train,model2_y_predict)
conf_matrix_model2
```

#### Out[30]:

```
array([[3099, 349], [1030, 614]], dtype=int64)
```

```
In [31]:
```

```
1
    # Accuracy Recall Precision F1 Score Specifity calculator.
 2
 3
    def calulate(c_matrix):
 4
         tp = c_matrix[0][0]
 5
        fp = c_matrix[0][1]
 6
        fn = c_matrix[1][0]
 7
        tn = c_matrix[1][1]
 8
         acc = (tp+tn)/(tp+fp+fn+tn)
 9
        rell = tp/(tp+fn)
10
        pre = tp/(tp+fp)
        f1 = 2*(pre*rell)/(pre+rell)
11
12
         spe = tn/(tn+fp)
13
        return acc,rell,pre,f1,spe
    .....
14
15
16
    def get_values(model):
         recall = recall_score(y_train,model)
17
18
         precision = precision_score(y_train,model)
19
        f1 = f1_score(y_train,model)
         accuracy = accuracy_score(y_train,model)
20
21
         return print(f'recall: {recall}\nprecision: {precision}\nf1: {f1}\naccuracy: {accur
22
"""m1_accuracy,m1_recall,m1_precision,m1_f1_score,m1_specificity = calulate(conf_matrix_model1)
print(f'Confusion Matrix:\n{conf matrix model1}\nm1 accuracy: {m1 accuracy}\nm1 recall:
{m1 recall}\nm1 precision: {m1 precision}\nm1 f1 score: {m1 f1 score}\nm1 specificity: {m1 specificity}')"""
"""m2 accuracy,m2 recall,m2 precision,m2 f1 score,m2 specificity = calulate(conf matrix model2)
print(f'Confusion Matrix:{conf matrix model2}\nm2 accuracy: {m2 accuracy}\nm2 recall:
{m2 recall}\nm2 precision: {m2 precision}\nm2 f1 score: {m2 f1 score}\nm2 specificity: {m2 specificity}')"""
In [32]:
 1 print("model 1")
    get_values(model1_y_predict)
    print("\nModel 2")
    get_values(model2_y_predict)
model 1
recall: 0.3381995133819951
precision: 0.5656154628687691
f1: 0.4232965359725923
accuracy: 0.7024744697564808
Model 2
recall: 0.3734793187347932
precision: 0.6375908618899273
f1: 0.47103950901419256
accuracy: 0.7291830322073841
```

Now each of these values signifies. Let's find out

Total class values accurately classified by the model. Model 1 Accuracy is about 53 % Model 2 Accuracy is about 72 %

Model 2 is more accurate but in the case of classification only model accuracy is not the cirteria of getting an actual accurate model.

Moving forward let's see recall and presision of the model.

#### Recall

```
In [33]:
```

```
print("model 1")
get_values(model1_y_predict)
print("Model 2")
get_values(model2_y_predict)
```

model 1 recall: 0.3381995133819951 precision: 0.5656154628687691 f1: 0.4232965359725923

accuracy: 0.7024744697564808

Model 2

recall: 0.3734793187347932 precision: 0.6375908618899273 f1: 0.47103950901419256 accuracy: 0.7291830322073841

Recall is defined as the ability of ML Model to correctly predict positive out of all positive results. Taking the recall into consideration we can choose Model 1.

#### **Precision**

It measure of amongst all the positive predictions how many of them were were actaully positive.

Precision of Model 1 is 37% and of Model 2 is 63%. So if we want that our model must accurately predict the outcome we should consider high precision. taking this into consideration we should go with Model 2.

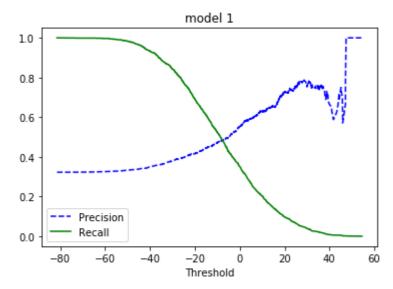
#### **Ploting Precision Recall Curve**

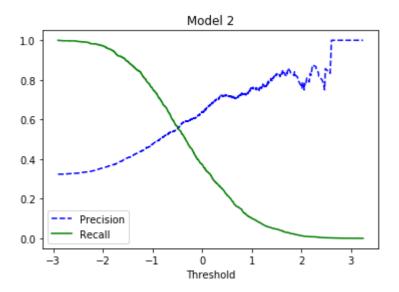
#### In [34]:

```
def plot_precision_recall_vs_threshold(precisions, recalls, thresholds, model):
   plt.title(model)
   plt.plot(thresholds, precisions[:-1], "b--", label="Precision")
   plt.plot(thresholds, recalls[:-1], "g-", label="Recall")
   plt.xlabel('Threshold')
   plt.legend()
```

#### In [35]:

```
from sklearn.metrics import precision_recall_curve as prc
   y_score_m1 = cross_val_predict(model1,x_train,y_train,cv=3,method='decision_function')
   y_score_m2 = cross_val_predict(model2,x_train,y_train,cv=3,method='decision_function')
   pre_m1, rec_m1 , threshol_m1 = prc(y_train,y_score_m1)
 5
   pre_m2, rec_m2 , threshol_m2 = prc(y_train,y_score_m2)
 6
 7
   plot_precision_recall_vs_threshold(pre_m1, rec_m1, threshol_m1, "model 1")
 8
9
   plt.show()
10
11
12
   plot_precision_recall_vs_threshold(pre_m2, rec_m2, threshol_m2, "Model 2")
13
   plt.show()
14
```





We can oberve with change in threshold we can change the precision and recall of both of our model. But for the time being let's go with the default thresold score.

#### F1 Score

It's like a trade off factor . classifier having high precision and recall value will get high f1 score. Considering this we should go with Model 1.

#### **ROC**

ROC curve is grapph of True Positive rate Vs false Positive rate True Positive Rate is Recall/Sensitivity while False Positive rate is out of total false prediction how many were actually false.

#### **AUC**

Aea under the curve, According to it the model is best which encloses maximum area.

#### In [36]:

```
from sklearn.metrics import roc_auc_score

roc_auc_m1 = roc_auc_score(y_train,y_score_m1)
roc_auc_m2 = roc_auc_score(y_train,y_score_m2)

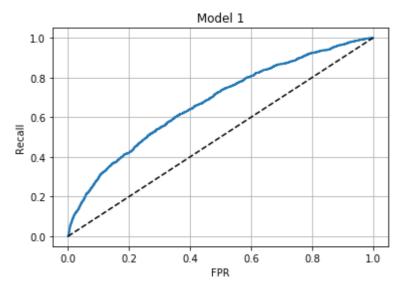
print(f'roc_auc_m1: {roc_auc_m1}\nroc_auc_m2: {roc_auc_m2}')
```

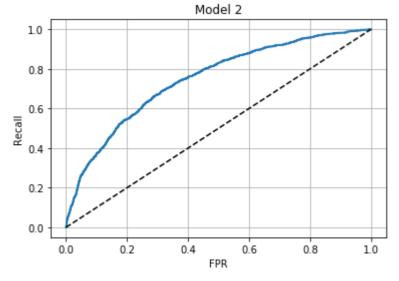
roc\_auc\_m1: 0.6739852539784691 roc\_auc\_m2: 0.7477612290491755

#### Ploting roc\_auc

# In [37]:

```
from sklearn.metrics import roc curve
   fpr_m1, tpr_m1, thresholds_m1 = roc_curve(y_train, y_score_m1)
 2
   fpr_m2, tpr_m2, thresholds_m2 = roc_curve(y_train, y_score_m2)
 4
 5
   def plot_roc_curve(fpr, tpr,model,label=None):
 6
 7
        plt.title(model)
        plt.plot(fpr, tpr, linewidth=2, label=label)
 8
 9
        plt.plot([0, 1], [0, 1], 'k--') # dashed diagonal
        plt.ylabel('Recall')
10
        plt.xlabel('FPR')
11
12
        plt.grid()
13
        plt.show()
14
   plot_roc_curve(fpr_m1, tpr_m1, model="Model 1")
15
   plot_roc_curve(fpr_m2, tpr_m2, model="Model 2")
16
17
18
```





According to this plot the best model is the model which is far away from the dotted line. So in this case both our model is good we can use any of the two based on the precison or recall what we want more.

# Taking this into consideration Using Model 2 for Application Purpose because we want high precision at $37\ \%$ recall

```
In [38]:
 1
    import pickle
   file_name = 'Model.picle'
 3 pickle.dump(model2,open(file_name,'wb'))
In [40]:
    model = pickle.load(open(file_name,'rb'))
   #model.predict([a])
Testing Score
In [41]:
 1 model1.score(x_test,y_test)
Out[41]:
0.4340659340659341
In [42]:
 1 model2.score(x_test,y_test)
Out[42]:
0.6978021978021978
EOF
In [ ]:
 1
```