Supervised machine learning algorithms application on dataset with binomial outcome

```
In [45]:
             import warnings
             warnings.filterwarnings("ignore")
             import numpy as np
             import pandas as pd
             import matplotlib.pyplot as plt
             import seaborn as sns
             %matplotlib inline
             from sklearn.linear_model import LogisticRegression
             from sklearn.model selection import train test split
             from sklearn.model selection import KFold
             from sklearn.ensemble import RandomForestClassifier
             from sklearn.naive bayes import GaussianNB
             from sklearn.neighbors import KNeighborsClassifier
             from sklearn.tree import DecisionTreeClassifier
             from sklearn import svm
             from sklearn import metrics
             from sklearn import linear model
```

Loading data called data.csv and call it data, while data m b was created only for plotting purpose

```
In [3]:  data = pd.read_csv("data.csv",header=0)
  data_m_b=pd.read_csv("data.csv",header=0)
```

Getting first info about data


```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 31 columns):
diagnosis
                           569 non-null object
radius mean
                           569 non-null float64
texture mean
                           569 non-null float64
                           569 non-null float64
perimeter_mean
                           569 non-null float64
area mean
                           569 non-null float64
smoothness mean
compactness mean
                           569 non-null float64
concavity mean
                           569 non-null float64
concave points_mean
                           569 non-null float64
symmetry mean
                           569 non-null float64
fractal dimension mean
                           569 non-null float64
                           569 non-null float64
radius se
texture se
                           569 non-null float64
                           569 non-null float64
perimeter_se
area_se
                           569 non-null float64
                           569 non-null float64
smoothness se
                           569 non-null float64
compactness se
concavity se
                           569 non-null float64
concave points_se
                           569 non-null float64
                           569 non-null float64
symmetry_se
fractal dimension se
                           569 non-null float64
radius worst
                           569 non-null float64
texture worst
                           569 non-null float64
                           569 non-null float64
perimeter worst
                           569 non-null float64
area worst
smoothness_worst
                           569 non-null float64
compactness worst
                           569 non-null float64
concavity worst
                           569 non-null float64
concave points worst
                           569 non-null float64
symmetry worst
                           569 non-null float64
fractal dimension worst
                           569 non-null float64
dtypes: float64(30), object(1)
memory usage: 135.6+ KB
```

Name of parameters of the dataset

Snapshot of the data

```
In [7]: ► data.head()
```

Out[7]:

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	cor
0	М	17.99	10.38	122.80	1001.0	0.11840	,
1	М	20.57	17.77	132.90	1326.0	0.08474	
2	М	19.69	21.25	130.00	1203.0	0.10960	
3	М	11.42	20.38	77.58	386.1	0.14250	
4	М	20.29	14.34	135.10	1297.0	0.10030	

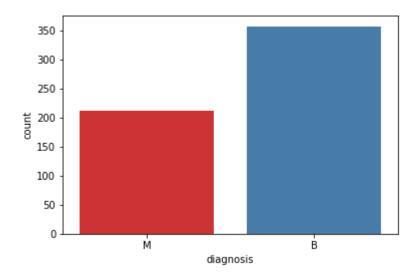
5 rows × 31 columns

→

Countplot of Malignant vs Benign observations

```
In [8]:  sns.countplot(x=data['diagnosis'],label="Count",palette="Set1")
```

Out[8]: <matplotlib.axes._subplots.AxesSubplot at 0x1172bb0>



Defining all predictors as variabl all predictors that will be used later

Chaning M and B at diagnosis column on 1 and 0

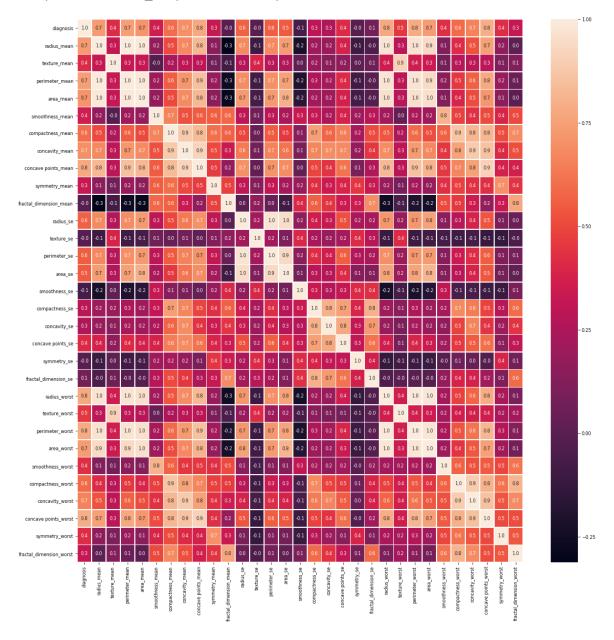
Creating variable for dependant variable

```
In [17]:
                import matplotlib.pyplot as plt
                 import matplotlib.gridspec as gridspec
                features mean=list(data.columns[1:31])
                dfM=data[data['diagnosis'] ==1]
                dfB=data[data['diagnosis'] ==0]
                plt.rcParams.update({'font.size': 8})
                fig, axes = plt.subplots(nrows=15, ncols=2, figsize=(15,30))
                axes = axes.ravel()
                for idx,ax in enumerate(axes):
                      ax.figure
                     binwidth= (max(data[features_mean[idx]]) - min(data[features_mean[idx]]))
                     ax.hist([dfM[features_mean[idx]],dfB[features_mean[idx]]], bins=np.arange
                     ax.legend(loc='upper right')
                     ax.set_title(f"Histogram of {features_mean[idx]} for Bening/Malignant Tun
                     plt.ylabel("Frequency")
                plt.ylabel("Frequency")
                plt.tight_layout()
                plt.show()
                 0.10
                                                                0.04
                                        st for Bening/Malignant Tumors
                                                               0.0015
                 0.015
                                                               0.0010
                 0.010
                                                               0.0005
                  15
                            Histogram of concavity_worst for Bening/Malignant Tumors
                                                                         Histogram of concave points_worst for Bening/Malignant Tumors
                                                          ____ М
____ В
                                                                5.0
                            Histogram of symmetry_worst for Bening/Malignant Tumors
```

Correalation Heatmap for all predictors

```
In [18]:  #correlation map
f,ax = plt.subplots(figsize=(18, 18))
sns.heatmap(data.corr(), annot=True, linewidths=.5, fmt= '.1f',ax=ax)
```

Out[18]: <matplotlib.axes._subplots.AxesSubplot at 0xada7d90>



Defining a function that will do the same process for each algorithm. First calculated accuracy for training data, then divides data into 4 folds and gives 4-fold CV scores.

```
In [23]:
          ▶ def classification model(model,data,prediction input,output):
                 model.fit(data[prediction input],data[output])
                 predictions = model.predict(data[prediction input])
                 accuracy = metrics.accuracy score(predictions,data[output])
                 print("Accuracy : %s" % "{0:.3%}".format(accuracy))
                 kf = KFold(n splits=4,random state=41,shuffle=True)
                 print (kf)
                 error = []
                 for train, test in kf.split(data):
                     train X = (data[prediction input].iloc[train,:])
                     train_y = data[output].iloc[train]
                     model.fit(train X, train y)
                     test X=data[prediction input].iloc[test,:]
                     test y=data[output].iloc[test]
                     error.append(model.score(test_X,test_y))
                     print("Cross-Validation Score : %s" % "{0:.3%}".format(error[-1]))
                 print(f"Average accuracy score is : %s" % "{0:.3%}".format(np.mean(error)
```

6 Most significant predictors chosen by Logitic regression with stepwise selction using SAS are defined as prediction_var

Results of Decision Tree Classifier using all predictors

Results of Decision Tree Classifier using 6 predictors

```
In [27]:
              model = DecisionTreeClassifier()
              classification model(model,data,prediction var,outcome var)
              Accuracy : 100.000%
              KFold(n splits=4, random state=41, shuffle=True)
              Cross-Validation Score: 97.902%
              Cross-Validation Score: 90.845%
              Cross-Validation Score: 90.141%
              Cross-Validation Score: 93.662%
              Average accuracy score is: 93.137%
          Results of Random Forest Classifier using 6 predictors
              model = RandomForestClassifier(n_estimators=100, max_features=2)
In [30]:
              classification model(model, data,prediction var,outcome var)
              Accuracy : 100.000%
              KFold(n splits=4, random state=41, shuffle=True)
              Cross-Validation Score: 99.301%
              Cross-Validation Score: 95.775%
              Cross-Validation Score: 92.254%
              Cross-Validation Score: 96.479%
              Average accuracy score is: 95.952%
          Results of Random Forest Classifier using all predictors
              model = RandomForestClassifier(n estimators=100, max depth=30, max features=2)
In [31]:
              classification model(model, data,all predictors,outcome var)
              Accuracy : 100.000%
              KFold(n_splits=4, random_state=41, shuffle=True)
              Cross-Validation Score: 98.601%
              Cross-Validation Score: 93.662%
              Cross-Validation Score: 89.437%
              Cross-Validation Score: 98.592%
              Average accuracy score is: 95.073%
          Results of SVM with gaussian rbf kernel using 6 predictors
In [459]:
              model = svm.SVC(kernel='rbf')
              classification_model(model,data,prediction_var,outcome_var)
              Accuracy: 99.473%
              KFold(n_splits=4, random_state=41, shuffle=True)
              Cross-Validation Score: 67.133%
              Cross-Validation Score: 61.972%
              Cross-Validation Score: 67.606%
              Cross-Validation Score: 65.493%
              [0.6713286713286714, 0.6197183098591549, 0.676056338028169, 0.6549295774647
              887]
              Average accuracy score is: 65.551%
```

Results of SVM with gaussian rbf kernel using all predictors

```
In [32]:  | model = svm.SVC(kernel='rbf')
    classification_model(model,data,all_predictors,outcome_var)

Accuracy : 100.000%
    KFold(n_splits=4, random_state=41, shuffle=True)
    Cross-Validation Score : 62.937%
    Cross-Validation Score : 59.155%
    Cross-Validation Score : 64.085%
    Cross-Validation Score : 64.789%
    Average accuracy score is : 62.741%
```

Results of SVM with linear kernel using 6 predictors

Results of SVM with linear kernel using all predictors

Results of SVM with sigmoid kernel using 6 predictors

Results of SVM with sigmoid kernel using all predictors

Results of Kneigbors classifier using Euclidean distances using 6 predictors

Results of Kneigbors classifier using Euclidean distances using all predictors

Results of Kneigbors classifier using Manhattan distances using 6 predictors

Results of Kneighbors classifier using Manhattan distances using all predictors

Results of Kneighbors classifier using Minkowski p=3 distances using all predictors

Results of Kneighbors classifier using Minkowski p=3 distances using 6 predictors

Unfortunately my computer coulndt run polynomial kernel of SVM

```
In [41]: | #model = svm.SVC(kernel='poly')
#classification_model(model,X,prediction_var2,outcome_var2)
```

Results of Logistic Regression with ridge penalty using 6 predictors

Results of Logistic Regression with ridge penalty using 6 predictors

```
In [483]: M model=LogisticRegression(penalty='12')
    classification_model(model,data,all_predictors,outcome_var)

Accuracy: 95.958%
    KFold(n_splits=4, random_state=41, shuffle=True)
    Cross-Validation Score: 97.902%
    Cross-Validation Score: 92.254%
    Cross-Validation Score: 92.958%
    Cross-Validation Score: 95.775%
    [0.9790209790209791, 0.9225352112676056, 0.9295774647887324, 0.9577464788732394]
    Average accuracy score is: 94.722%
```

Results of Logistic Regression with lasso penalty using 6 predictors and coefficients of predictors

Results of Logistic Regression with lasso penalty using all predictors and coefficients of predictors

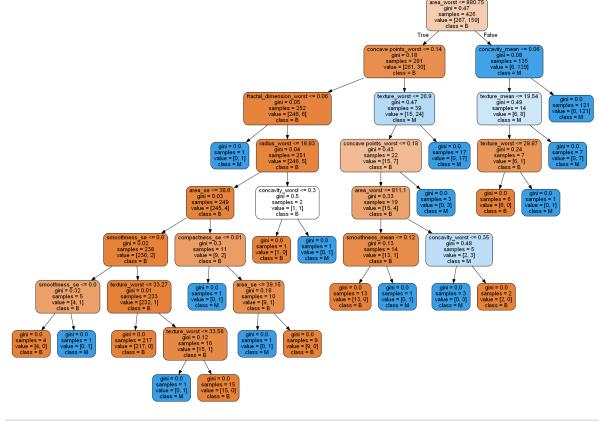
```
In [46]:
             model=LogisticRegression(penalty='l1')
             classification model(model,data,all predictors,outcome var)
             model.coef
             Accuracy : 95.958%
             KFold(n_splits=4, random_state=41, shuffle=True)
             Cross-Validation Score: 98.601%
             Cross-Validation Score: 92.958%
             Cross-Validation Score: 92.254%
             Cross-Validation Score: 95.775%
             Average accuracy score is: 94.897%
   Out[46]: array([[-4.0958732 , -0.06593394, 0.09680937,
                                                            0.01957743,
                      0.
                                               0.
                                                                          0.
                               , 0.
                                                            0.
                      0.
                                , -0.93429995, -0.1148191 ,
                                                            0.11208174,
                                , 0.
                                               0.
                                                                         0.
                                               0.18648275,
                     -0.19824339, 0.30622498,
                                                            0.01283658,
                                                                         0.
                                   2.7244296 ,
                                                                                    11)
```

Decision Tree and Random forest plots

```
In [49]:
             model = DecisionTreeClassifier()
             model.fit(x_train,y_train)
             predictions = model.predict(data_m_b[all_predictors])
             accuracy = metrics.accuracy score(predictions,data m b[outcome var])
             print("Accuracy : %s" % "{0:.3%}".format(accuracy))
             error=[]
             error.append(model.score(x_test,y_test))
             print(error)
             dot_data = tree.export_graphviz(model, out_file=None,
                             feature_names = x.columns,class_names = model.classes_, round
                             precision = 2, filled = True)
             graph = graphviz.Source(dot_data)
             graph = pydotplus.graph_from_dot_data(dot_data)
             # Show graph
             Image(graph.create_png())
             # Create PDF
```

Accuracy: 99.473% [0.9790209790209791]

Out[49]:



Out[50]: <pydotplus.graphviz.Dot at 0xe27c470>


```
In [52]:  # Create PDF
graph.write_pdf("random_forest2.pdf")

# Create PNG
graph.write_png("random_forest2.png")
```

Out[52]: True