```
In [1]:
         import numpy as np
         import pandas as pd
         import warnings
         warnings.filterwarnings("ignore")
         data = pd.read_csv("breat_cancer_data.csv")
In [2]:
         len(data.index), len(data.columns)
In [3]:
         (569, 33)
Out[3]:
         data.shape
         (569, 33)
Out[4]:
In [5]:
         data.head
```

0+ [ [ ] .	<bou< td=""><td>nd method NDFra</td><td>me.head of</td><td>f</td><td>id diagnosis</td><td>radius mean</td><td>texture_mean</td><td>per</td></bou<>	nd method NDFra	me.head of	f	id diagnosis	radius mean	texture_mean	per
Out[5]:	imet	er_mean area_m	nean \			_	_	•
				17.00	10 20	122 00	1001 0	
	0	842302	М	17.99	10.38	122.80		
	1	842517	Μ	20.57	17.77	132.90	1326.0	
	2	84300903	Μ	19.69	21.25	130.00	1203.0	
	3	84348301	M	11.42	20.38	77.58	386.1	
	4							
	4	84358402	М	20.29	14.34	135.10	1297.0	
	• •	• • •	• • •	• • •	• • •	• • •	• • •	
	564	926424	Μ	21.56	22.39	142.00	1479.0	
	565	926682	Μ	20.13	28.25	131.20	1261.0	
	566	926954	М	16.60	28.08	108.30		
	567	927241	М	20.60	29.33	140.10		
	568	92751	В	7.76	24.54	47.92	181.0	
		smoothness_mea	n compact	tness mean d	concavity mean	concave no	ints mean \	
	0			0.27760	0.30010			
	0	0.1184					0.14710	
	1	0.0847	'4	0.07864	0.08690		0.07017	
	2	0.1096	10	0.15990	0.19740		0.12790	
	3	0.1425		0.28390	0.24140		0.10520	
	4						0.10320	
		0.1003		0.13280	0.19800		Ø.10430	
	• •			• • •			• • •	
	564	0.1110	10	0.11590	0.24390		0.13890	
	565	0.0978	80	0.10340	0.14400		0.09791	
	566	0.0845		0.10230	0.09251		0.05302	
	567	0.1178		0.27700	0.35140		0.15200	
	568	0.0526	53	0.04362	0.00000		0.00000	
		texture w	orst peri	imeter worst	area_worst	smoothness w	orst \	
	0	_	7.33	184.60	2019.0		6220	
	1		3.41	158.80	1956.0		2380	
	2	2	5.53	152.50	1709.0	0.1	4440	
	3	2	6.50	98.87	567.7	0.2	0980	
	4	1	.6.67	152.20	1575.0		3740	
						0.1		
	• •	• • •	• • •	• • • •	• • • •		• • •	
	564	2	6.40	166.10	2027.0	0.1	4100	
	565	3	8.25	155.00	1731.0	0.1	1660	
	566	3	4.12	126.70	1124.0		1390	
	567		9.42	184.60	1821.0		6500	
	568	3	0.37	59.16	268.6	0.0	8996	
		compactness_wo	rst conca	avity_worst	concave point	s_worst sym	metry_worst \	
	0	0.66		0.7119	•	0.2654	0.4601	
						0.1860	0.2750	
	1	0.18		0.2416				
	2	0.42		0.4504		0.2430	0.3613	
	3	0.86	630	0.6869		0.2575	0.6638	
	4	0.20	500	0.4000		0.1625	0.2364	
	564	0.21	120	0.4107		0.2216	0.2060	
	565	0.19	1220	0.3215		0.1628	0.2572	
	566	0.30	1940	0.3403		0.1418	0.2218	
	567	0.86	810	0.9387		0.2650	0.4087	
	568	0.06		0.0000		0.0000	0.2871	
	200	0.00		0.0000		0.0000	0.20/1	
		fractal_dimens	_					
	0		0.11890	NaN	V			
	1		0.08902	NaN				
	2		0.08758	Nan				
	3		0.17300	Nan				
	4		0.07678	NaN	V			

564	0.07115	NaN	
565	0.06637	NaN	
566	0.07820	NaN	
567	0.12400	NaN	
568	0.07039	NaN	

[569 rows x 33 columns]>

In [7]: data.tail()

Out[7]:

	Id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean
204	wenaea	r.a	e1.50	act act a con too	1-64-000	1479.0	W-11100
~~~	rennee	Peri	#14: T (5)	with all to with the	101-00	1 40 1 10	No. HOLDER STATE
	*******	F=0	4.85.8588	20.00	1 028-20	25.75.25.7	101-101-20-20-25-25
	*******	~	20.00	28.50.75.75	4-880-410	1 (0.05.05.10)	88.4.4.9.888
~~~	Mark to 1		Y-Y-0	10 ch - 15 ch	40 Y 100 M	101.0	111 - 111 TO 40 TO 110 TO

5 rows × 33 columns

4

# **Data Analysis**

In [8]: data.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 33 columns):

#	Column	Non-Null Count	Dtype
0	id	569 non-null	int64
1	diagnosis	569 non-null	object
2	radius_mean	569 non-null	float64
3	texture_mean	569 non-null	float64
4	perimeter_mean	569 non-null	float64
5	area_mean	569 non-null	float64
6	smoothness_mean	569 non-null	float64
7	compactness_mean	569 non-null	float64
8	concavity_mean	569 non-null	float64
9	concave points_mean	569 non-null	float64
10	symmetry_mean	569 non-null	float64
11	fractal_dimension_mean	569 non-null	float64
12	radius_se	569 non-null	float64
13	texture_se	569 non-null	float64
14	perimeter_se	569 non-null	float64
15	area_se	569 non-null	float64
16	smoothness_se	569 non-null	float64
17	compactness_se	569 non-null	float64
18	concavity_se	569 non-null	float64
19	concave points_se	569 non-null	float64
20	symmetry_se	569 non-null	float64
21	<pre>fractal_dimension_se</pre>	569 non-null	float64
22	radius_worst	569 non-null	float64
23	texture_worst	569 non-null	float64
24	perimeter_worst	569 non-null	float64
25	area_worst	569 non-null	float64
26	smoothness_worst	569 non-null	float64
27	compactness_worst	569 non-null	float64
28	concavity_worst	569 non-null	float64
29	concave points_worst	569 non-null	float64
30	symmetry_worst	569 non-null	float64
31	fractal_dimension_worst	569 non-null	float64
32	Unnamed: 32	0 non-null	float64
dtvp	es: float64(31), int64(1)	. object(1)	

dtypes: float64(31), int64(1), object(1)

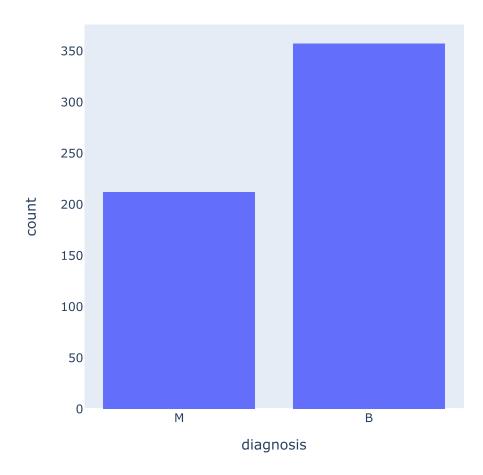
memory usage: 146.8+ KB

In [9]: data.isnull().sum()

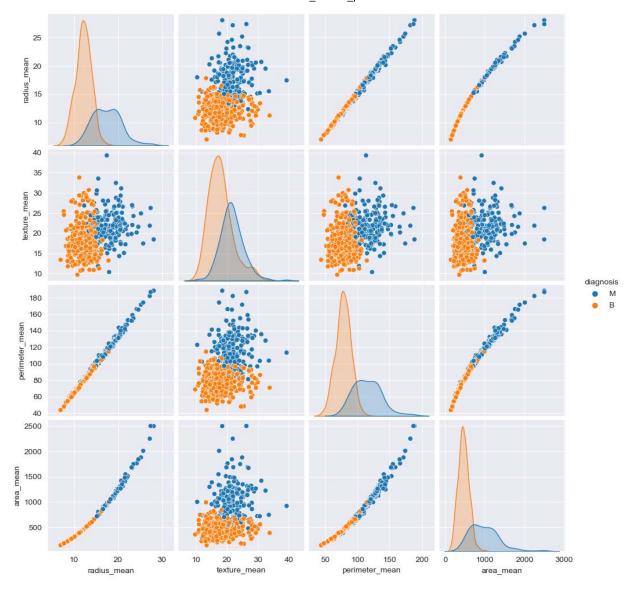
```
0
         id
Out[9]:
         diagnosis
                                        0
         radius_mean
                                        0
         texture_mean
                                        0
         perimeter mean
                                        0
         area_mean
                                        0
         smoothness mean
         compactness_mean
                                        0
         concavity_mean
                                        0
         concave points mean
                                        0
         symmetry_mean
                                        0
         fractal_dimension_mean
                                        0
         radius_se
                                        0
         texture_se
         perimeter se
                                        0
                                        0
         area se
         smoothness_se
         compactness_se
                                        0
         concavity_se
                                        0
         concave points_se
         symmetry_se
                                        0
         fractal_dimension_se
         radius worst
         texture worst
                                        0
         perimeter_worst
                                        0
                                        0
         area_worst
         smoothness\_worst
                                        0
         compactness_worst
                                        0
         concavity worst
                                        0
         concave points_worst
         symmetry_worst
                                        0
         fractal_dimension_worst
                                        0
         Unnamed: 32
                                      569
         dtype: int64
In [10]:
         data = data.dropna(axis='columns')
          data.describe(include="0")
Out[10]:
                 diagnosis
                      569
           count
                        2
          unique
                        В
             top
                      357
            freq
In [11]:
         data.diagnosis.value_counts()
               357
Out[11]:
               212
         Name: diagnosis, dtype: int64
         diagnosis unique = data.diagnosis.unique()
In [12]:
          diagnosis unique
         array(['M', 'B'], dtype=object)
Out[12]:
```

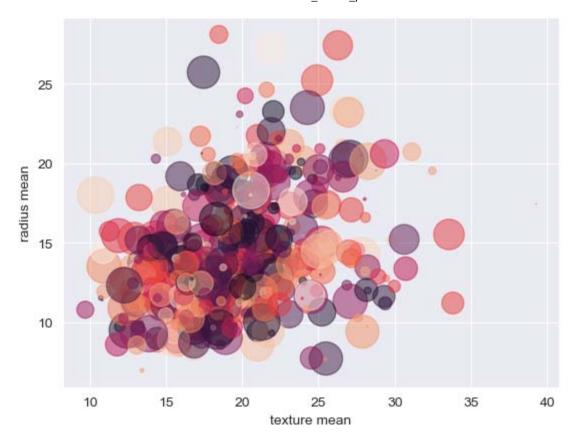
#### **Data Visualization**

```
In [13]: !pip install plotly
         Requirement already satisfied: plotly in c:\users\admin\anaconda3\lib\site-packages
          (5.9.0)
         Requirement already satisfied: tenacity>=6.2.0 in c:\users\admin\anaconda3\lib\site-p
         ackages (from plotly) (8.0.1)
In [16]: import matplotlib.pyplot as plt
          import seaborn as sns
          import plotly.express as px
          import plotly.graph_objects as go
          %matplotlib inline
          sns.set_style('darkgrid')
In [19]: plt.figure(figsize=(15, 5))
          plt.subplot(1, 2, 1)
          plt.hist( data.diagnosis)
          plt.title("Counts of Diagnosis")
          plt.xlabel("Diagnosis")
          plt.subplot(1, 2, 2)
          sns.countplot('diagnosis', data=data)
          TypeError
                                                     Traceback (most recent call last)
         Cell In[19], line 10
                6 plt.xlabel("Diagnosis")
                8 plt.subplot(1, 2, 2)
          ---> 10 sns.countplot('diagnosis', data=data)
         TypeError: countplot() got multiple values for argument 'data'
                          Counts of Diagnosis
         350
         300
         250
         200
          100
          50
                             Diagnosis
         # plt.figure(figsize=(7,12))
          px.histogram(data, x='diagnosis')
          # plt.show()
```



```
In [21]: cols = ["diagnosis", "radius_mean", "texture_mean", "perimeter_mean", "area_mean"]
    sns.pairplot(data[cols], hue="diagnosis")
    plt.show()
```





```
from sklearn.preprocessing import LabelEncoder
In [23]:
         labelencoder_Y = LabelEncoder()
In [24]:
         data.diagnosis = labelencoder_Y.fit_transform(data.diagnosis)
         print(data.diagnosis.value_counts())
In [25]:
         print("\n", data.diagnosis.value_counts().sum())
         0
              357
              212
         Name: diagnosis, dtype: int64
          569
         cols = ['diagnosis', 'radius_mean', 'texture_mean', 'perimeter_mean',
In [26]:
                 'area_mean', 'smoothness_mean', 'compactness_mean', 'concavity_mean',
                 'concave points_mean', 'symmetry_mean', 'fractal_dimension_mean']
         print(len(cols))
         data[cols].corr()
         11
```

Out[26]:

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothn
diagnosis	1.0000000	0.230029	10.00 1 50 1 80 5	W-74-8-8-8	D. ZUBSBA	
radius mean		4				
***************************************	11.415165	Harares	7-00000000	M-0 4 M 0 0 0	H- 041 THOM	
per <sup>imer</sup> er_mean		10.000 V M S S		4.00000000	ELL STATE IS AS DESCRIPTION	
A-Ed-III Edit	nvuestes	Hamaraar	H-3-6 1 H63-D	HI-MARINDERY		
smoothness_mean	0.358560	0.170581	-0.023389	0.207278	0.177028	
compactness_mean	0.596534	0.506124	0.236702	0.556936	0.498502	
concavity_mean	0.696360	0.676764	0.302418	0.716136	0.685983	
concave points_mean	0.776614	0.822529	0.293464	0.850977	0.823269	
symmetry_mean	0.330499	0.147741	0.071401	0.183027	0.151293	
fractal_dimension_mean	-0.012838	-0.311631	-0.076437	-0.261477	-0.283110	
mlt figure/figging/	(12 0))					

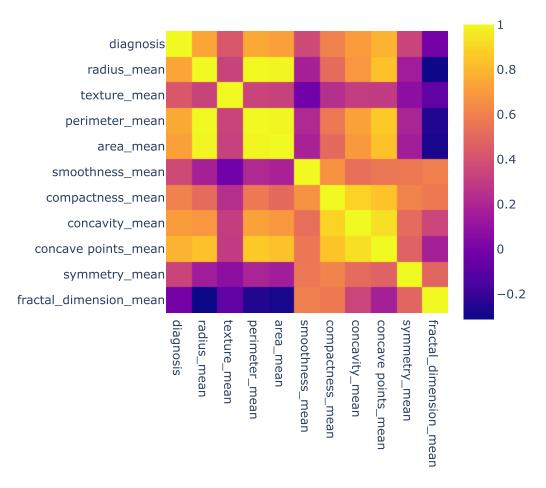
```
In [27]: plt.figure(figsize=(12, 9))
    plt.title("Correlation Graph")

cmap = sns.diverging_palette( 1000, 120, as_cmap=True)
    sns.heatmap(data[cols].corr(), annot=True, fmt='.1%', linewidths=.05, cmap=cmap);
```

					Corr	elation G	raph					_		1.0
diagnosis	100.0%	73.0%	41.5%	74.3%	70.9%	35.9%	59.7%	69.6%	77.7%	33.0%	-1.3%			1.0
radius_mean	73.0%	100.0%	32.4%	99.8%	98.7%	17.1%	50.6%	67.7%	82.3%	14.8%	-31.2%		-	0.8
texture_mean	41.5%	32.4%	100.0%	33.0%	32.1%	-2.3%	23.7%	30.2%	29.3%	7.1%	-7.6%			
perimeter_mean	74.3%	99.8%	33.0%	100.0%	98.7%	20.7%	55.7%	71.6%	85.1%	18.3%	-26.1%		-	0.6
area_mean	70.9%	98.7%	32.1%	98.7%	100.0%	17.7%	49.9%	68.6%	82.3%	15.1%	-28.3%			
smoothness_mean	35.9%	17.1%	-2.3%	20.7%	17.7%	100.0%	65.9%	52.2%	55.4%	55.8%	58.5%		-	0.4
compactness_mean	59.7%	50.6%	23.7%	55.7%	49.9%	65.9%	100.0%	88.3%	83.1%	60.3%	56.5%		-	0.2
concavity_mean	69.6%	67.7%	30.2%	71.6%	68.6%	52.2%	88.3%	100.0%	92.1%	50.1%	33.7%			
concave points_mean	77.7%	82.3%	29.3%	85.1%	82.3%	55.4%	83.1%	92.1%	100.0%	46.2%	16.7%		-	0.0
symmetry_mean	33.0%	14.8%	7.1%	18.3%	15.1%	55.8%	60.3%	50.1%	46.2%	100.0%	48.0%			
fractal_dimension_mean	-1.3%	-31.2%	-7.6%	-26.1%	-28.3%	58.5%	56.5%	33.7%	16.7%	48.0%	100.0%		-	-0.2
	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	symmetry_mean	ctal_dimension_mean			

```
In [28]: plt.figure(figsize=(15, 10))

fig = px.imshow(data[cols].corr());
fig.show()
```



<Figure size 1500x1000 with 0 Axes>

## **Model Implementation**

```
from sklearn.model_selection import train_test_split
In [29]:
         from sklearn.preprocessing import StandardScaler
         from sklearn.linear model import LogisticRegression
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.ensemble import RandomForestClassifier
         from sklearn.naive bayes import GaussianNB
         from sklearn.neighbors import KNeighborsClassifier
         from sklearn.metrics import accuracy_score, confusion_matrix, f1_score
         from sklearn.metrics import classification_report
         from sklearn.model selection import KFold
         from sklearn.model selection import cross validate, cross val score
         from sklearn.svm import SVC
         from sklearn import metrics
In [30]:
         data.columns
```

```
Index(['id', 'diagnosis', 'radius_mean', 'texture_mean', 'perimeter_mean',
Out[30]:
                 'concave points_mean', 'symmetry_mean', 'fractal_dimension_mean',
                 'radius_se', 'texture_se', 'perimeter_se', 'area_se', 'smoothness_se', 'compactness_se', 'concavity_se', 'concave points_se', 'symmetry_se',
                 'fractal_dimension_se', 'radius_worst', 'texture_worst',
                 'perimeter_worst', 'area_worst', 'smoothness_worst',
                 'compactness_worst', 'concavity_worst', 'concave points_worst',
                 'symmetry_worst', 'fractal_dimension_worst'],
                dtype='object')
         prediction_feature = [ "radius_mean", 'perimeter_mean', 'area_mean', 'symmetry_mean']
In [31]:
          targeted_feature = 'diagnosis'
          # len(prediction feature)
Out[31]:
In [32]: X = data[prediction_feature]
Out[32]:
                                                                                        concave
              radius_mean perimeter_mean area_mean symmetry_mean compactness_mean
                                                                                    points_mean
                    17.00
                                             1001.0
                                                                                         0.14710
                                   140.10
                                             12500.00
         569 rows × 6 columns
In [33]: y = data.diagnosis
         У
```

```
1
Out[33]:
         1
                1
         2
                1
         3
                1
         4
                1
         564
                1
         565
                1
         566
                1
         567
                1
         568
                0
         Name: diagnosis, Length: 569, dtype: int32
In [34]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.33, random_state
In [35]: sc = StandardScaler()
         X_train = sc.fit_transform(X_train)
         X_test = sc.fit_transform(X_test)
In [36]: def model_building(model, X_train, X_test, y_train, y_test):
             Model Fitting, Prediction And Other stuff
             return ('score', 'accuracy_score', 'predictions' )
             model.fit(X_train, y_train)
             score = model.score(X_train, y_train)
             predictions = model.predict(X_test)
             accuracy = accuracy_score(predictions, y_test)
             return (score, accuracy, predictions)
         models_list = {
In [37]:
              "LogisticRegression" : LogisticRegression(),
              "RandomForestClassifier" : RandomForestClassifier(n_estimators=10, criterion='ent
              "DecisionTreeClassifier" : DecisionTreeClassifier(criterion='entropy', random_sta
              "SVC" : SVC(),
         }
In [38]: print(list(models_list.keys()))
         print(list(models_list.values()))
         ['LogisticRegression', 'RandomForestClassifier', 'DecisionTreeClassifier', 'SVC']
         [LogisticRegression(), RandomForestClassifier(criterion='entropy', n_estimators=10, r
         andom_state=5), DecisionTreeClassifier(criterion='entropy', random_state=0), SVC()]
In [39]: def cm_metrix_graph(cm):
             sns.heatmap(cm,annot=True,fmt="d")
              plt.show()
In [40]: df prediction = []
         confusion matrixs = []
         df_prediction_cols = [ 'model_name', 'score', 'accuracy_score' , "accuracy_percentage'
         for name, model in zip(list(models_list.keys()), list(models_list.values())):
```

```
(score, accuracy, predictions) = model_building(model, X_train, X_test, y_train, y
print("\n\nClassification Report of '"+ str(name), "'\n")
print(classification_report(y_test, predictions))

df_prediction.append([name, score, accuracy, "{0:.2%}".format(accuracy)])

# For Showing Metrics
confusion_matrixs.append(confusion_matrix(y_test, predictions))

df_pred = pd.DataFrame(df_prediction, columns=df_prediction_cols)
```

Classification Report of 'LogisticRegression'

	precision	recall	f1-score	support
0 1	0.90 0.92	0.96 0.84	0.93 0.88	115 73
accuracy macro avg weighted avg	0.91 0.91	0.90 0.91	0.91 0.90 0.91	188 188 188

Classification Report of 'RandomForestClassifier'

	precision	recall	f1-score	support
0 1	0.92 0.93	0.96 0.88	0.94 0.90	115 73
accuracy macro avg weighted avg	0.93 0.93	0.92 0.93	0.93 0.92 0.93	188 188 188

Classification Report of 'DecisionTreeClassifier '

	precision	recall	f1-score	support
0 1	0.90 0.92	0.96 0.84	0.93 0.88	115 73
accuracy macro avg weighted avg	0.91 0.91	0.90 0.91	0.91 0.90 0.91	188 188 188

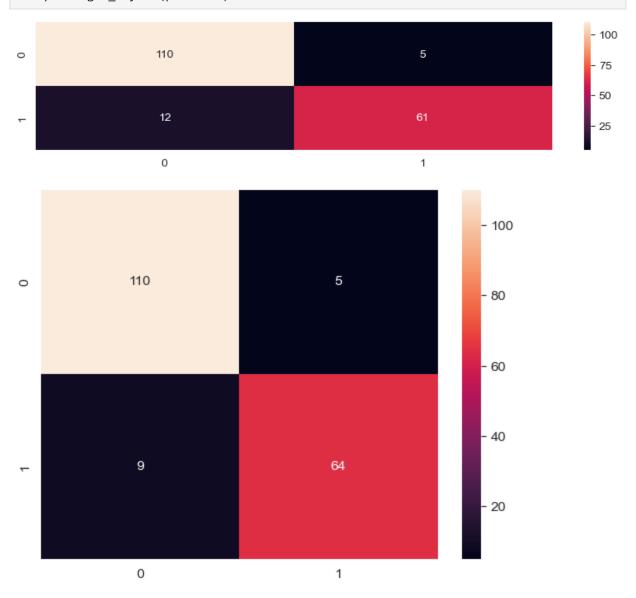
Classification Report of 'SVC '

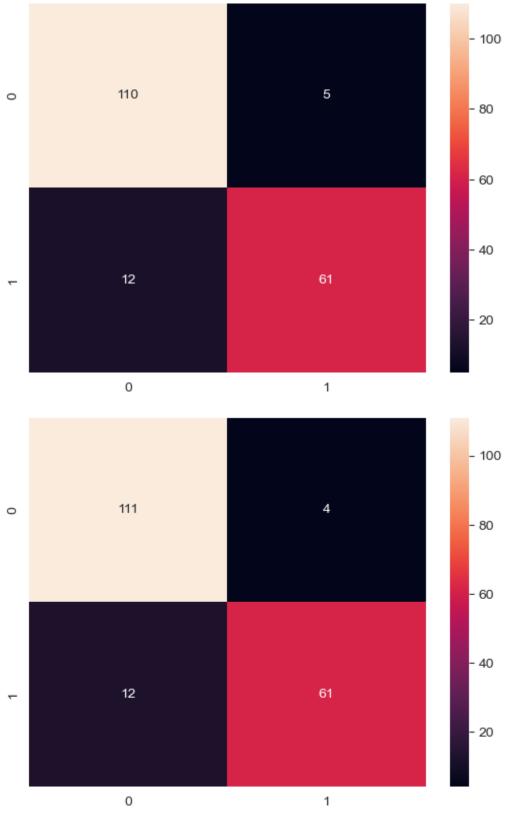
```
precision recall f1-score support
                  0.90
                           0.97
                                     0.93
          0
                                               115
          1
                 0.94
                           0.84
                                     0.88
                                                73
   accuracy
                                     0.91
                                               188
                 0.92
                           0.90
                                     0.91
                                               188
  macro avg
weighted avg
                 0.92
                           0.91
                                     0.91
                                               188
```

```
In [41]: plt.figure(figsize=(10, 2))
# plt.title("Confusion Metric Graph")

for index, cm in enumerate(confusion_matrixs):
# plt.xlabel("Negative Positive")
# plt.ylabel("True Positive")
# Show The Metrics Graph
```

cm\_metrix\_graph(cm) # Call the Confusion Metrics Graph
plt.tight\_layout(pad=True)





<Figure size 640x480 with 0 Axes>

In [42]: df\_pred

Out[42]:		model_name	score	accuracy_score	accuracy_percentage
	0	LogisticRegression	0.916010	0.909574	90.96%
	1	Random Forest Classifier	0.992126	0.925532	92.55%
	2	DecisionTreeClassifier	1.000000	0.909574	90.96%
	3	SVC	0.923885	0.914894	91.49%
In [43]:	df	f_pred.sort_values('	score'.	ascending <b>=Fal</b>	se)
[] .		(	500.0	ascending rule	
Out[43]:		model_name			accuracy_percentage
		model_name			
		model_name	score 1.000000	accuracy_score	accuracy_percentage
	2	model_name  DecisionTreeClassifier  RandomForestClassifier	score 1.000000	accuracy_score 0.909574	accuracy_percentage 90.96%
	2	model_name  DecisionTreeClassifier  RandomForestClassifier	1.000000 0.992126 0.923885	0.909574 0.925532	accuracy_percentage 90.96% 92.55%

#### K-fold

```
len(data)
In [44]:
         569
Out[44]:
In [45]: cv_score = cross_validate(LogisticRegression(), X, y, cv=3,
                                   scoring=('r2', 'neg_mean_squared_error'),
                                   return_train_score=True)
          pd.DataFrame(cv_score).describe().T
Out[45]:
           est_neg_mean_squared_error
                                       3.0 -0.108902 0.043669 -0.157895 -0.126316 -0.094737 -0.084405
          train_neg_mean_squared_error
                                       3.0 -0.106321 0.012102 -0.113456 -0.113307 -0.113158 -0.102753
In [46]: def cross_val_scorring(model):
                (score, accuracy, predictions) = model_building(model, X_train, X_test, y_train,
              model.fit(data[prediction_feature], data[targeted_feature])
              # score = model.score(X_train, y_train)
              predictions = model.predict(data[prediction_feature])
              accuracy = accuracy_score(predictions, data[targeted_feature])
```

```
print("\nFull-Data Accuracy:", round(accuracy, 2))
print("Cross Validation Score of'"+ str(name), "'\n")
# Initialize K folds.
kFold = KFold(n splits=5) # define 5 diffrent data folds
err = []
for train_index, test_index in kFold.split(data):
    # print("TRAIN:", train index, "TEST:", test index)
    # Data Spliting via fold indexes
   X_train = data[prediction_feature].iloc[train_index, :] # train_index = rows or
   y_train = data[targeted_feature].iloc[train_index] # all targeted features tree
   X_test = data[prediction_feature].iloc[test_index, :] # testing all rows and d
   y test = data[targeted feature].iloc[test index] # all targeted tests
   # Again Model Fitting
   model.fit(X_train, y_train)
   err.append(model.score(X_train, y_train))
   print("Score:", round(np.mean(err), 2) )
```

```
Full-Data Accuracy: 0.9
Cross Validation Score of LogisticRegression '
Score: 0.91
Score: 0.91
Score: 0.9
Score: 0.9
Score: 0.9
Full-Data Accuracy: 1.0
Cross Validation Score of RandomForestClassifier '
Score: 0.99
Score: 0.99
Score: 0.99
Score: 1.0
Score: 1.0
Full-Data Accuracy: 1.0
Cross Validation Score of DecisionTreeClassifier '
Score: 1.0
Score: 1.0
Score: 1.0
Score: 1.0
Score: 1.0
Full-Data Accuracy: 0.89
Cross Validation Score of 'SVC '
Seece: 8.89
acece: e.ee
```

```
print("\n Best Score is ")
print(gsc.best_score_)

print("\n Best Estinator is ")
print(gsc.best_estimator_)

print("\n Best Parametes are")
print(gsc.best_params_)

Best Score is
0.9185560053981108

Best Estinator is
DecisionTreeClassifier(max_features='log2', min_samples_leaf=9)

Best Parametes are
{'max_features': 'log2', 'min_samples_leaf': 9, 'min_samples_split': 2}
```

# (2) KNeighborsClassifier

```
In [50]: # Pick the model
         model = KNeighborsClassifier()
          # Tunning Params
          param_grid = {
              'n_neighbors': list(range(1, 30)),
              'leaf_size': list(range(1,30)),
              'weights': [ 'distance', 'uniform' ]
          }
          # Implement GridSearchCV
          gsc = GridSearchCV(model, param_grid, cv=10)
          # Model Fitting
          gsc.fit(X_train, y_train)
          print("\n Best Score is ")
          print(gsc.best_score_)
          print("\n Best Estinator is ")
          print(gsc.best_estimator_)
          print("\n Best Parametes are")
          print(gsc.best params )
          Best Score is
         0.9159244264507423
          Best Estinator is
         KNeighborsClassifier(leaf_size=1, n_neighbors=10)
          Best Parametes are
         {'leaf_size': 1, 'n_neighbors': 10, 'weights': 'uniform'}
```

### (3) SVC

model = SVC()

In [51]: # Pick the model

```
# Tunning Params
          param grid = [
                        {'C': [1, 10, 100, 1000],
                         'kernel': ['linear']
                        {'C': [1, 10, 100, 1000],
                         'gamma': [0.001, 0.0001],
                         'kernel': ['rbf']
                        }
          ]
          # Implement GridSearchCV
          gsc = GridSearchCV(model, param_grid, cv=10) # 10 Cross Validation
          # Model Fitting
          gsc.fit(X_train, y_train)
          print("\n Best Score is ")
          print(gsc.best_score_)
          print("\n Best Estinator is ")
          print(gsc.best_estimator_)
          print("\n Best Parametes are")
         print(gsc.best_params_)
          Best Score is
         0.9184885290148447
          Best Estinator is
         SVC(C=10, gamma=0.001)
          Best Parametes are
         {'C': 10, 'gamma': 0.001, 'kernel': 'rbf'}
In [55]: # Pick the model
         model = RandomForestClassifier()
          # Tunning Params
          random grid = {'bootstrap': [True, False],
           'max depth': [40, 50, None], # 10, 20, 30, 60, 70, 100,
           'max_features': ['auto', 'sqrt'],
           'min_samples_leaf': [1, 2], # , 4
           'min_samples_split': [2, 5], # , 10
           'n_estimators': [200, 400]} # , 600, 800, 1000, 1200, 1400, 1600, 1800, 2000
```

```
# Implement GridSearchCV
gsc = GridSearchCV(model, random_grid, cv=10) # 10 Cross Validation
# Model Fitting
gsc.fit(X_train, y_train)
print("\n Best Score is ")
print(gsc.best_score_)
print("\n Best Estinator is ")
print(gsc.best_estimator_)
print("\n Best Parametes are")
print(gsc.best_params_)
Best Score is
0.9105937921727396
Best Estinator is
RandomForestClassifier(min_samples_split=5, n_estimators=400)
Best Parametes are
{'bootstrap': True, 'max_depth': None, 'max_features': 'sqrt', 'min_samples_leaf': 1,
'min_samples_split': 5, 'n_estimators': 400}
```