

Cognorise_Infotech_Project_Task1

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✓ BREAST CANCER CLASSIFICATION

A MACHINE LEARNING PROJECT FOR CLASSIFICATION OF BREAST CANCER INTO MALIGNANT AND BENIGN TUMOR

MALIGNANT : CANCEROUS , FAST GROWING , SPREAD TO OTHER PARTS OF THE BODY.

BENIGN : NON-CANCEROUS , SLOW GROWING , DO NOT SPREAD TO OTHER PARTS OF THE BODY.

```
#Importing the sufficient python libraries
```

```
import numpy as np
```

```
import pandas as pd
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
#Importing the dataset
```

```
df=pd.read_csv('/content/breast_cancer.csv')
```

```
df
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean
0	842302	M	17.99	10.38	122.80	1001.0	0.11840	0.27760
1	842517	M	20.57	17.77	132.90	1326.0	0.08474	0.07864
2	84300903	M	19.69	21.25	130.00	1203.0	0.10960	0.15990
3	84248201	M	11.42	20.28	77.58	286.1	0.14250	0.28220

3	84348301	M	11.42	20.38	77.58	386.1	0.14250	0.28390
4	84358402	M	20.29	14.34	135.10	1297.0	0.10030	0.13280
...
564	926424	M	21.56	22.39	142.00	1479.0	0.11100	0.11590
565	926682	M	20.13	28.25	131.20	1261.0	0.09780	0.10340
566	926954	M	16.60	28.08	108.30	858.1	0.08455	0.10230
567	927241	M	20.60	29.33	140.10	1265.0	0.11780	0.27700
568	92751	B	7.76	24.54	47.92	181.0	0.05263	0.04362

569 rows × 33 columns

```
#To print the first 5 rows of the dataframe
df.head()
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	co
0	842302	M	17.99	10.38	122.80	1001.0	0.11840	0.27760	
1	842517	M	20.57	17.77	132.90	1326.0	0.08474	0.07864	
2	84300903	M	19.69	21.25	130.00	1203.0	0.10960	0.15990	
3	84348301	M	11.42	20.38	77.58	386.1	0.14250	0.28390	
4	84358402	M	20.29	14.34	135.10	1297.0	0.10030	0.13280	

5 rows × 33 columns

```
#To print the last 5 rows of the dataframe
df.tail()
```

```
df.head(5),
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	cc
564	926424	M	21.56	22.39	142.00	1479.0	0.11100	0.11590	
565	926682	M	20.13	28.25	131.20	1261.0	0.09780	0.10340	
566	926954	M	16.60	28.08	108.30	858.1	0.08455	0.10230	
567	927241	M	20.60	29.33	140.10	1265.0	0.11780	0.27700	
568	92751	B	7.76	24.54	47.92	181.0	0.05263	0.04362	

```
5 rows x 33 columns
```

```
#To print the diamension of the dataframe in (rows, columns) format
df.shape
```

```
(569, 33)
```

```
#To print column names of the dataframe
df.columns
```

```
Index(['id', 'diagnosis', 'radius_mean', 'texture_mean', 'perimeter_mean',
       'area_mean', 'smoothness_mean', 'compactness_mean', 'concavity_mean',
       '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', 'Unnamed: 32'],
      dtype='object')
```

```
#To print the datatype of each column
df.dtypes
```

dtypes

id	int64
diagnosis	object
radius_mean	float64
texture_mean	float64
perimeter_mean	float64
area_mean	float64
smoothness_mean	float64
compactness_mean	float64
concavity_mean	float64
concave points_mean	float64
symmetry_mean	float64
fractal_dimension_mean	float64
radius_se	float64
texture_se	float64
perimeter_se	float64
area_se	float64
smoothness_se	float64
compactness_se	float64
concavity_se	float64
concave points_se	float64
symmetry_se	float64
fractal_dimension_se	float64
radius_worst	float64
texture_worst	float64
perimeter_worst	float64
area_worst	float64
smoothness_worst	float64
compactness_worst	float64
concavity_worst	float64
concave points_worst	float64
symmetry_worst	float64
fractal_dimension_worst	float64
Unnamed: 32	float64
dtype:	object

```
#To check whether there is any missing values in the dataframe  
df.isna().sum()
```

```
id          0
diagnosis   0
radius_mean 0
texture_mean 0
perimeter_mean 0
area_mean   0
smoothness_mean 0
compactness_mean 0
concavity_mean 0
concave points_mean 0
symmetry_mean 0
fractal_dimension_mean 0
radius_se   0
texture_se   0
perimeter_se 0
area_se      0
smoothness_se 0
compactness_se 0
concavity_se 0
concave points_se 0
symmetry_se  0
fractal_dimension_se 0
radius_worst 0
texture_worst 0
perimeter_worst 0
area_worst    0
smoothness_worst 0
compactness_worst 0
concavity_worst 0
concave points_worst 0
symmetry_worst 0
fractal_dimension_worst 0
Unnamed: 32    569
dtype: int64
```

```
#To get a concise summary of the dataframe
df.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	fractal_dimension_se	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

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

memory usage: 146.8+ KB

```
#To get statistical measures about the dataset  
df.describe()
```

	id	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	coi
count	5.690000e+02	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	
mean	3.037183e+07	14.127292	19.289649	91.969033	654.889104	0.096360	0.104341	
std	1.250206e+08	3.524049	4.301036	24.298981	351.914129	0.014064	0.052813	
min	8.670000e+03	6.981000	9.710000	43.790000	143.500000	0.052630	0.019380	
25%	8.692180e+05	11.700000	16.170000	75.170000	420.300000	0.086370	0.064920	
50%	9.060240e+05	13.370000	18.840000	86.240000	551.100000	0.095870	0.092630	
75%	8.813129e+06	15.780000	21.800000	104.100000	782.700000	0.105300	0.130400	
max	9.113205e+08	28.110000	39.280000	188.500000	2501.000000	0.163400	0.345400	

8 rows × 32 columns

Count: The number of non-null values for each numerical column.

Mean: The mean (average) value.

Std: The standard deviation, a measure of the amount of variation or dispersion.

Min: The minimum value.

25% (Q1): The first quartile, which represents the 25th percentile.

50% (median): The median or 50th percentile.

75% (Q3): The third quartile, which represents the 75th percentile.

Max: The maximum value.

```
#Since ID is not required for prediction we can drop that column
#Also all the values in the column 'Unnamed: 32' are missing, so we can drop that column to
df.drop(['id','Unnamed: 32'],axis=1,inplace=True)
df
```

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_
0	M	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3
1	M	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0
2	M	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.1
3	M	11.42	20.38	77.58	386.1	0.14250	0.28390	0.2
4	M	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1
...
564	M	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.2
565	M	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.1
566	M	16.60	28.08	108.30	858.1	0.08455	0.10230	0.0
567	M	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.3
568	B	7.76	24.54	47.92	181.0	0.05263	0.04362	0.0

569 rows × 31 columns

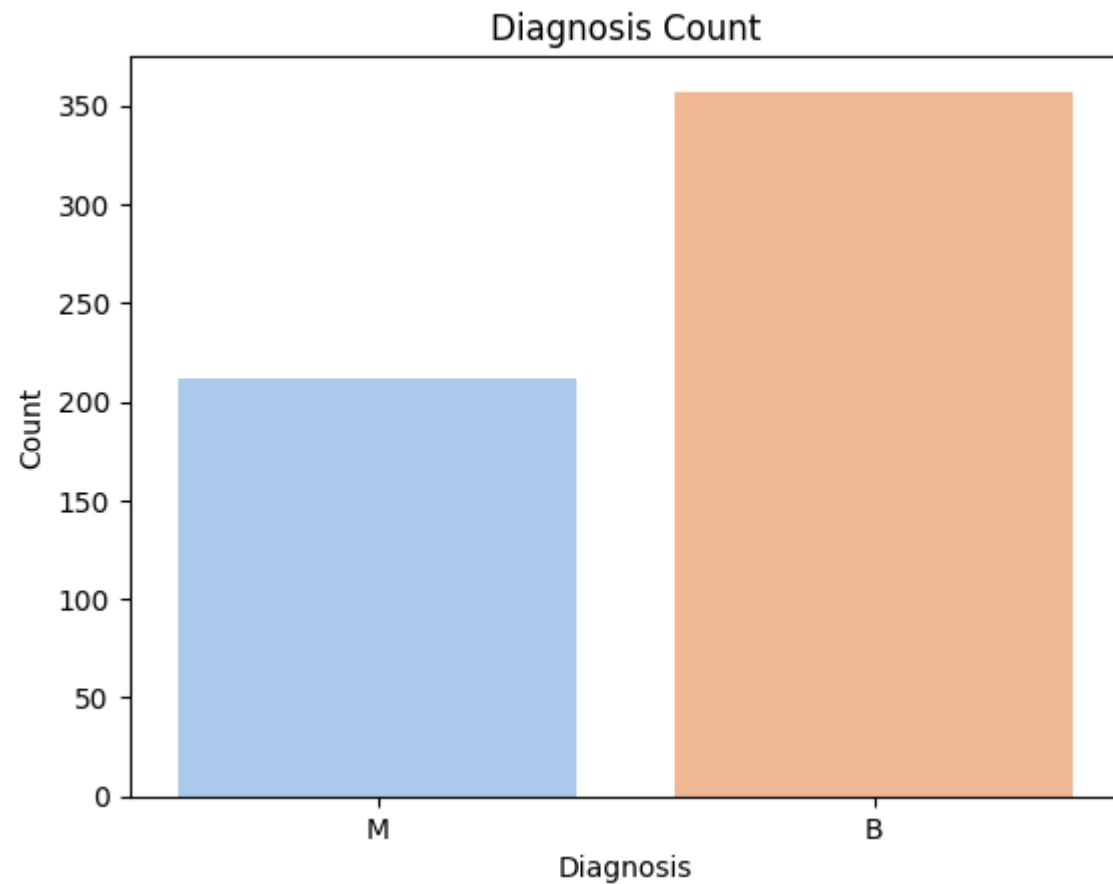
```
#Checking the distribution of the target variable
df['diagnosis'].value_counts()
```

```
B    357
M    212
```



```
Name: diagnosis, dtype: int64
```

```
sns.countplot(x='diagnosis',data=df,hue='diagnosis',palette='pastel')  
plt.title('Diagnosis Count')  
plt.xlabel('Diagnosis')  
plt.ylabel('Count')  
Text(0, 0.5, 'Count')
```



```
df['diagnosis']=df['diagnosis'].map({'M':1,'B':0})  
df
```

#1 represents Malignant (riskier)

#0 represents Benign

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_
0	1	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3
1	1	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0
2	1	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.1
3	1	11.42	20.38	77.58	386.1	0.14250	0.28390	0.2
4	1	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1
...
564	1	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.2
565	1	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.1
566	1	16.60	28.08	108.30	858.1	0.08455	0.10230	0.0
567	1	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.3
568	0	7.76	24.54	47.92	181.0	0.05263	0.04362	0.0

569 rows × 31 columns

#To check how much riskier is Malignant than Benign

df.groupby('diagnosis').mean()

	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean
diagnosis							
0	12.146524	17.914762	78.075406	462.790196	0.092478	0.080085	0.046058

1	17.462830	21.604906	115.365377	978.376415	0.102898	0.145188	0.160775
---	-----------	-----------	------------	------------	----------	----------	----------

2 rows × 30 columns

#Since this dataset has no object input features, no need to perform encoding, also there a

✓ OUTLIER DETECTION AND REMOVAL

OUTLIER : A datapoint which is significantly far from other data points

IQR : Inter Quartile Range

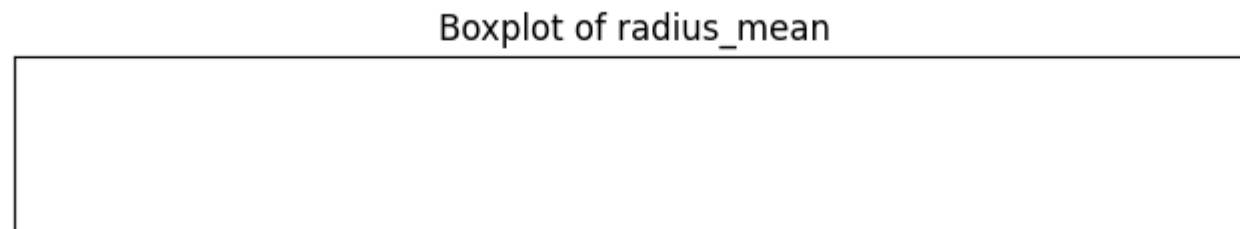
IQR = Q3-Q1

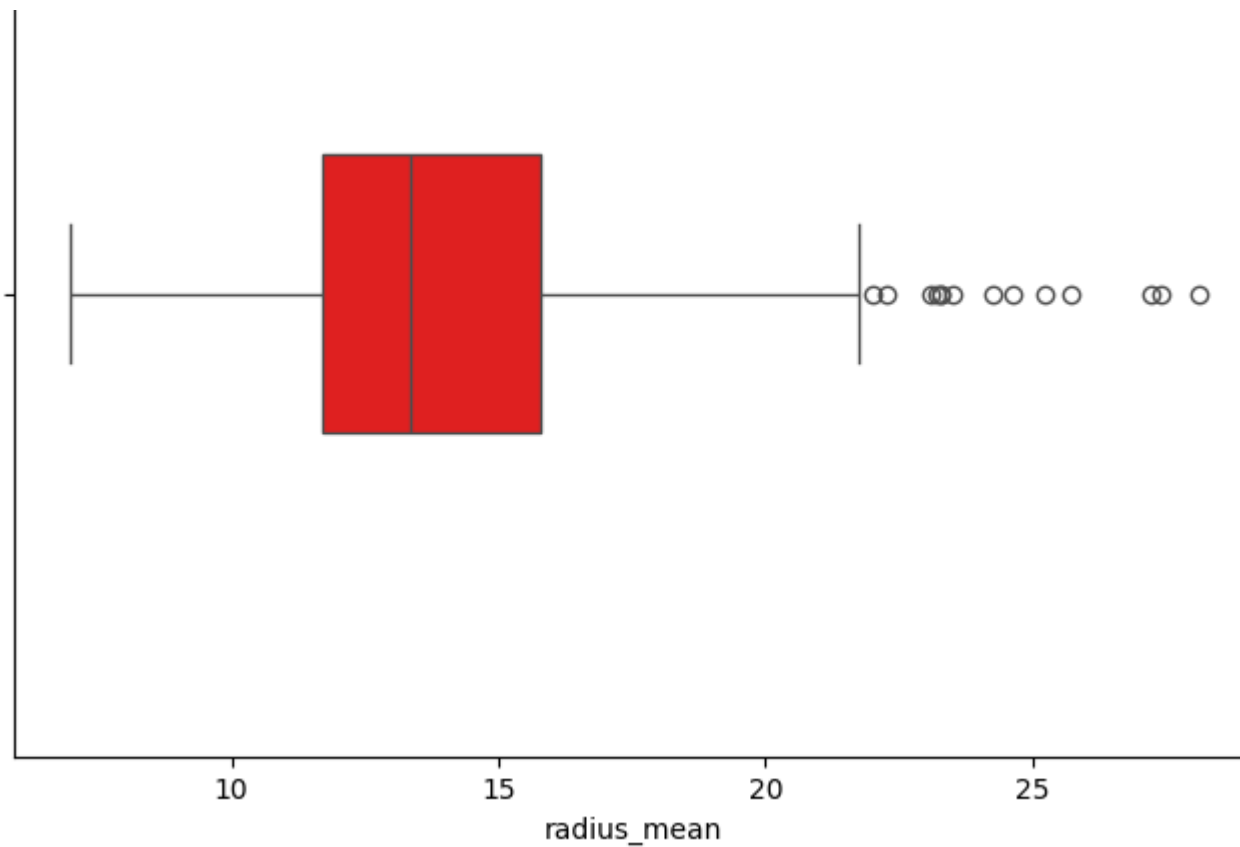
Top = Q3+1.5*IQR

Bottom = Q1-1.5*IQR

Boxplot

```
plt.figure(figsize=(8,6))
sns.boxplot(x='radius_mean',data=df,orient='h',width=0.3,color='red')
plt.title('Boxplot of radius_mean')
plt.show()
```





```
Q1=df.quantile(0.25)
Q3=df.quantile(0.75)
IQR=Q3-Q1
threshold=1.5
df=df[~((df<(Q1-threshold*IQR)) | (df>(Q3+threshold*IQR))).any(axis=1)]
df
```

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_
6	1	18.25	19.98	119.60	1040.0	0.09463	0.10900	0.1
7	1	12.71	20.83	82.30	577.0	0.11800	0.16450	0.0

7	1	13.71	20.83	90.20	577.9	0.11890	0.16430	0.0
10	1	16.02	23.24	102.70	797.8	0.08206	0.06669	0.0
11	1	15.78	17.89	103.60	781.0	0.09710	0.12920	0.0
13	1	15.85	23.95	103.70	782.7	0.08401	0.10020	0.0
...
554	0	12.88	28.92	82.50	514.3	0.08123	0.05824	0.0
555	0	10.29	27.61	65.67	321.4	0.09030	0.07658	0.0
558	0	14.59	22.68	96.39	657.1	0.08473	0.13300	0.1
560	0	14.05	27.15	91.38	600.4	0.09929	0.11260	0.0
566	1	16.60	28.08	108.30	858.1	0.08455	0.10230	0.0

398 rows × 31 columns

✓ FEATURE SELECTION

```
#To print the correlation matrix
#Checking correlations of different features with the target variable.
df.corr()
```

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactn
diagnosis	1.000000	0.680705	0.388596	0.695835	0.698005	0.271570	
radius_mean	0.680705	1.000000	0.280796	0.998113	0.992047	0.055311	
texture_mean	0.388596	0.280796	1.000000	0.285880	0.288688	-0.046879	
perimeter_mean	0.695835	0.998113	0.285880	1.000000	0.999335	0.002106	

perimeter_mean	0.695835	0.998113	0.285880	1.000000	0.990235	0.093196
area_mean	0.698005	0.992047	0.288688	0.990235	1.000000	0.058280
smoothness_mean	0.271570	0.055311	-0.046879	0.093196	0.058280	1.000000
compactness_mean	0.514346	0.442360	0.181078	0.492915	0.428703	0.675183
concavity_mean	0.710567	0.658031	0.295111	0.696240	0.664847	0.463982
concave points_mean	0.752190	0.755607	0.239147	0.785954	0.758145	0.522678
symmetry_mean	0.243588	0.075653	0.051037	0.102371	0.077918	0.494062
fractal_dimension_mean	-0.086631	-0.436397	-0.110214	-0.393736	-0.423006	0.614693
radius_se	0.571842	0.534801	0.290195	0.542837	0.561732	0.224720
texture_se	0.007755	-0.149935	0.474188	-0.144895	-0.132103	0.056968
perimeter_se	0.565909	0.530347	0.294101	0.546622	0.554631	0.213312
area_se	0.703715	0.758127	0.325637	0.763671	0.786897	0.181729
smoothness_se	-0.036030	-0.301325	0.089100	-0.283129	-0.260649	0.330804
compactness_se	0.229104	0.165214	0.212798	0.207785	0.161122	0.240507
concavity_se	0.325974	0.265486	0.242464	0.302232	0.268137	0.230219
concave points_se	0.387972	0.363046	0.198599	0.394612	0.354173	0.378191
symmetry_se	-0.194890	-0.327179	0.011039	-0.318263	-0.308046	0.077346
fractal_dimension_se	0.062799	-0.100183	0.095912	-0.063267	-0.093091	0.283411
radius_worst	0.761081	0.971286	0.311152	0.972912	0.969353	0.122928
texture_worst	0.437909	0.273847	0.914721	0.280552	0.280953	0.020158
perimeter_worst	0.768194	0.964851	0.316562	0.971558	0.962721	0.146367
area_worst	0.773626	0.957772	0.317755	0.959710	0.971053	0.125280
smoothness_worst	0.396847	0.087209	0.069886	0.121237	0.102255	0.803945

```

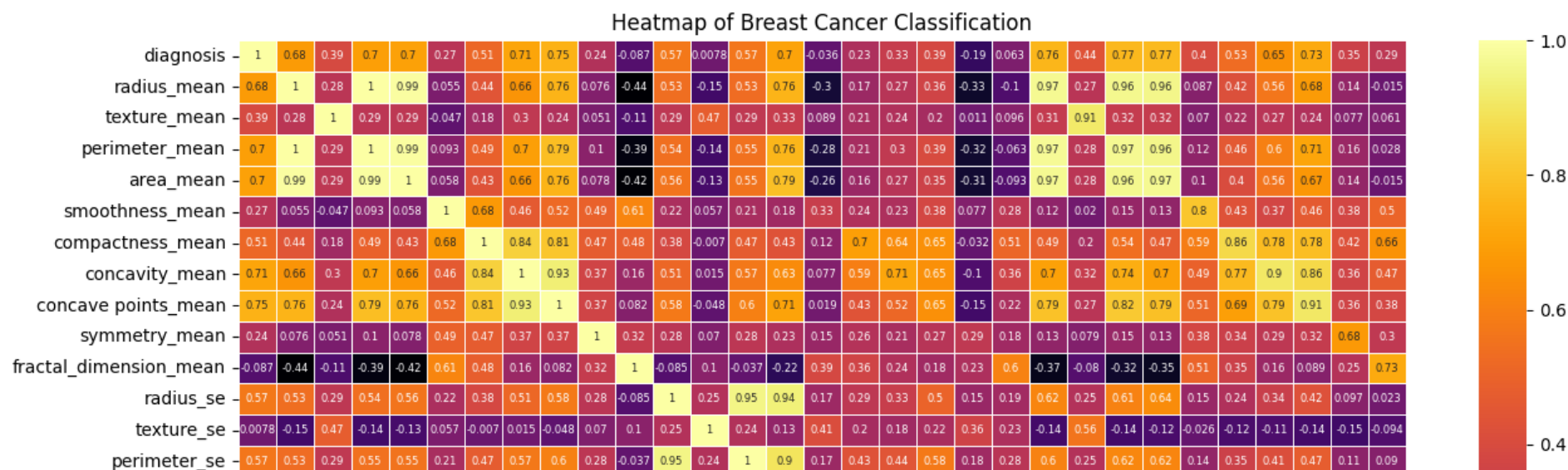
-
compactness_worst    0.533996    0.416464    0.216026    0.460901    0.400703    0.432375
concavity_worst      0.653300    0.563240    0.274070    0.599672    0.559618    0.372220
concave points_worst 0.726639    0.682235    0.238186    0.712906    0.671695    0.463910
symmetry_worst       0.347406    0.137655    0.076977    0.160178    0.136700    0.375969
fractal_dimension_worst 0.290821   -0.014583    0.060915    0.027877   -0.014754    0.504310
31 rows x 31 columns

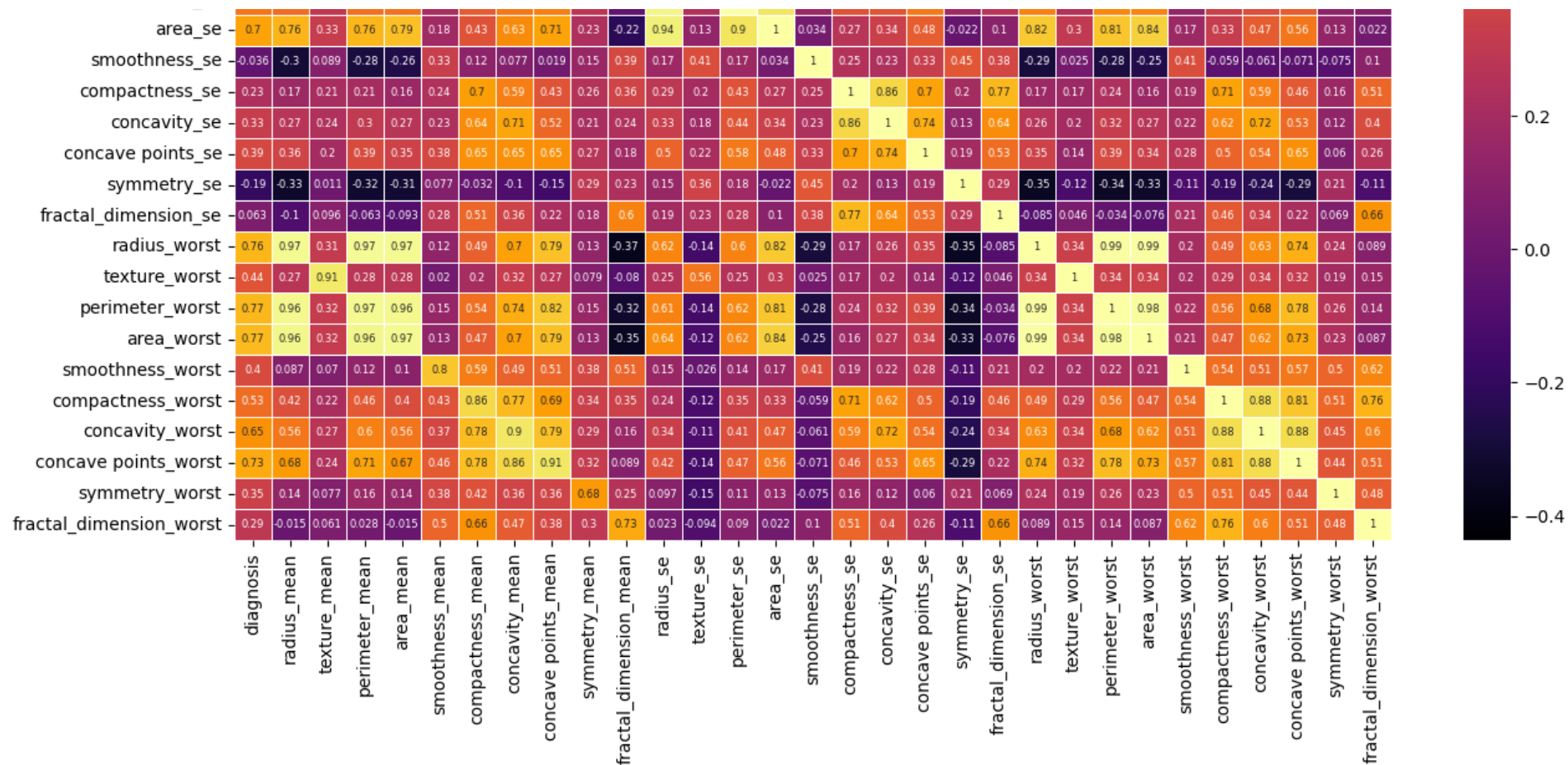
```

```

#To visualize the correlation matrix graphically
#Heatmap
plt.figure(figsize=(15,10))
corr=df.corr()
sns.heatmap(corr,cmap='inferno',linewidths=.5,annot=True,annot_kws={"fontsize":6})
plt.title("Heatmap of Breast Cancer Classification")
plt.show()

```





✓ Applying **ANNOVA TEST** for Feature Selection

```
df_copy=df.copy()
x_copy=df_copy.drop(['diagnosis'],axis=1)
y_copy=df['diagnosis']
from sklearn.feature_selection import f_classif
score2=f_classif(x_copy,y_copy)
score2
(array([3.41923458e+02, 7.04350576e+01, 3.71719551e+02, 3.76248046e+02,
        3.15303773e+01, 1.42446996e+02, 4.03848026e+02, 5.15999640e+02,
        2.49788576e+01, 2.99440304e+00, 1.92412626e+02, 2.38191066e-02,
        1.86569790e+02, 3.88492502e+02, 5.14731747e-01, 2.19369677e+01,
        4.70812572e+01, 7.01688565e+01, 1.56347914e+01, 1.56789246e+00,
        5.45163293e+02, 9.39559734e+01, 5.70142066e+02, 5.90294605e+02,
        7.40225711e+01, 1.57963547e+02, 2.94858852e+02, 4.42989671e+02,
        5.43535467e+01, 3.65869335e+01]),
array([1.76202025e-55, 8.51880326e-16, 6.80281885e-59, 2.11648603e-59,
        3.70131299e-08, 2.92362441e-28, 1.98753990e-62, 9.78048268e-74,
        8.71811279e-07, 8.43317446e-02, 6.16508492e-36, 8.77424701e-01,
        4.49299726e-35, 9.31925157e-61, 4.73520649e-01, 3.87829076e-06,
        2.63532505e-11, 9.55311414e-16, 9.10031278e-05, 2.11252375e-01,
        1.89923667e-76, 4.46169928e-20, 1.05225985e-78, 1.75342370e-80,
        1.83097645e-16, 1.01615082e-30, 8.52732130e-50, 1.51484069e-66,
        9.87189587e-13, 3.38223856e-09]))

f_value2=pd.Series(score2[0],index=x_copy.columns)
f_value2.sort_values(ascending=False)

area_worst          590.294605
perimeter_worst     570.142066
radius_worst        545.163293
concave points mean  515.000610
```

```
concave points_mean      315.999840
concave points_worst     442.989671
concavity_mean           403.848026
area_se                  388.492502
area_mean                376.248046
perimeter_mean          371.719551
radius_mean              341.923458
concavity_worst          294.858852
radius_se                192.412626
perimeter_se             186.569790
compactness_worst        157.963547
compactness_mean         142.446996
texture_worst             93.955973
smoothness_worst         74.022571
texture_mean             70.435058
concave points_se        70.168856
symmetry_worst           54.353547
concavity_se             47.081257
fractal_dimension_worst  36.586934
smoothness_mean          31.530377
symmetry_mean            24.978858
compactness_se           21.936968
symmetry_se              15.634791
fractal_dimension_mean   2.994403
fractal_dimension_se     1.567892
smoothness_se            0.514732
texture_se               0.023819
dtype: float64
```

```
p_value2=pd.Series(score2[1],index=x_copy.columns)
p_value2.sort_values(ascending=False)
```

```
texture_se      8.774247e-01
smoothness_se  4.735206e-01
fractal_dimension_se  2.112524e-01
fractal_dimension_mean  8.433174e-02
symmetry_se     9.100313e-05
compactness_se  3.878291e-06
symmetry_mean   8.718113e-07
smoothness mean 3.701313e-08
```

```

fractal_dimension_worst    3.382239e-09
concavity_se               2.635325e-11
symmetry_worst             9.871896e-13
concave_points_se         9.553114e-16
texture_mean              8.518803e-16
smoothness_worst          1.830976e-16
texture_worst              4.461699e-20
compactness_mean          2.923624e-28
compactness_worst         1.016151e-30
perimeter_se              4.492997e-35
radius_se                 6.165085e-36
concavity_worst           8.527321e-50
radius_mean               1.762020e-55
perimeter_mean            6.802819e-59
area_mean                 2.116486e-59
area_se                   9.319252e-61
concavity_mean            1.987540e-62
concave_points_worst      1.514841e-66
concave_points_mean       9.780483e-74
radius_worst              1.899237e-76
perimeter_worst           1.052260e-78
area_worst                1.753424e-80
dtype: float64

```

#Anova test : we can drop the columns 'fractal_dimension_se','smoothness_se','fractal_dimension_mean','texture_se',axis=1
df.drop(['fractal_dimension_se','smoothness_se','fractal_dimension_mean','texture_se'],axis=1,inplace=True)
df

```

<ipython-input-24-c4208603c380>:2: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame

```

See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
df.drop(['fractal_dimension_se','smoothness_se','fractal_dimension_mean','texture_se'],axis=1,inplace=True)

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean
6	1	18.25	19.98	119.60	1040.0	0.09463	0.10900	0.1
7	1	13.71	20.83	90.20	577.9	0.11890	0.16450	0.0

10	1	16.02	23.24	102.70	797.8	0.08206	0.06669	0.0
11	1	15.78	17.89	103.60	781.0	0.09710	0.12920	0.0
13	1	15.85	23.95	103.70	782.7	0.08401	0.10020	0.0
...
554	0	12.88	28.92	82.50	514.3	0.08123	0.05824	0.0
555	0	10.29	27.61	65.67	321.4	0.09030	0.07658	0.0
558	0	14.59	22.68	96.39	657.1	0.08473	0.13300	0.1
560	0	14.05	27.15	91.38	600.4	0.09929	0.11260	0.0
566	1	16.60	28.08	108.30	858.1	0.08455	0.10230	0.0

398 rows × 27 columns

✓ SEPARATING THE INPUT AND OUTPUT FEATURES AS X AND Y

```
x=df.drop(['diagnosis'],axis=1)
y=df['diagnosis']
```

✓ BALANCING THE DATASET

```
from imblearn.over_sampling import SMOTE
smote=SMOTE()
x_resampled,y_resampled=smote.fit_resample(x,y)
```

```
from collections import Counter
print("Before Smote :",Counter(y))
print("After Smote :",Counter(y_resampled))

Before Smote : Counter({0: 300, 1: 98})
After Smote : Counter({1: 300, 0: 300})
```

✓ SPLITTING INTO TRAINING AND TESTING DATA

```
from sklearn.model_selection import train_test_split
x_train,x_test,y_train,y_test=train_test_split(x_resampled,y_resampled,test_size=0.30,random_state=42)
y_train
```

108	1
272	0
599	1
479	1
436	1
	..
71	0
106	0
270	0
435	1
102	0

Name: diagnosis, Length: 420, dtype: int64

✓ NORMALIZATION USING STANDARDSCALER

```
from sklearn.preprocessing import StandardScaler
scaler=StandardScaler()
scaler.fit(x_train)
```

```
nx_train=scaler.transform(x_train)
nx_test=scaler.transform(x_test)
```

✓ MODEL CREATION USING K-NEAREST NEIGHBORS CLASSIFIER

GRIDSEARCHCV

```
from sklearn.neighbors import KNeighborsClassifier
knn1=KNeighborsClassifier()
param={'n_neighbors':[3,5,7,9], 'weights':['uniform','distance']}
```

```
from sklearn.model_selection import GridSearchCV
clf=GridSearchCV(knn1,param,cv=10,scoring='accuracy')
clf.fit(nx_train,y_train)
print(clf.best_params_)
{'n_neighbors': 5, 'weights': 'distance'}
```

#Model creation

```
from sklearn.neighbors import KNeighborsClassifier
knn=KNeighborsClassifier(n_neighbors=3,weights='distance')
knn.fit(nx_train,y_train)
y_prediction=knn.predict(nx_test)
y_prediction
array([0, 1, 1, 0, 1, 0, 1, 1, 1, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1, 0, 1, 1,
       1, 1, 0, 1, 0, 0, 1, 0, 1, 0, 0, 0, 0, 1, 1, 1, 1, 0, 1, 1, 1, 0,
       0, 0, 1, 1, 0, 1, 0, 1, 0, 1, 1, 0, 0, 1, 0, 0, 1, 1, 0, 1, 1, 1,
       1, 1, 1, 0, 0, 1, 1, 0, 1, 1, 1, 1, 1, 1, 0, 1, 1, 0, 0, 0, 0, 0,
       0, 0, 0, 1, 0, 1, 0, 1, 0, 1, 1, 1, 0, 0, 1, 1, 0, 1, 0, 0, 0, 1,
```

```

0, 0, 1, 1, 0, 1, 0, 1, 1, 1, 0, 0, 0, 1, 0, 0, 1, 0, 1, 1, 0, 1,
0, 1, 1, 0, 1, 0, 0, 0, 1, 0, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 0,
1, 0, 0, 1, 1, 0, 1, 0, 1, 1, 0, 1, 0, 0, 1, 1, 1, 0, 1, 1, 1,
1, 1, 0, 0])

```

✓ PERFORMANCE EVALUATION

```

from sklearn.metrics import accuracy_score, confusion_matrix, classification_report, ConfusionMatrix
#confusion matrix
matr=confusion_matrix(y_test,y_prediction)
print(matr)

[[75  3]
 [ 6 96]]

```

```

#accuracy score
score=accuracy_score(y_test,y_prediction)
score

0.95

```

```

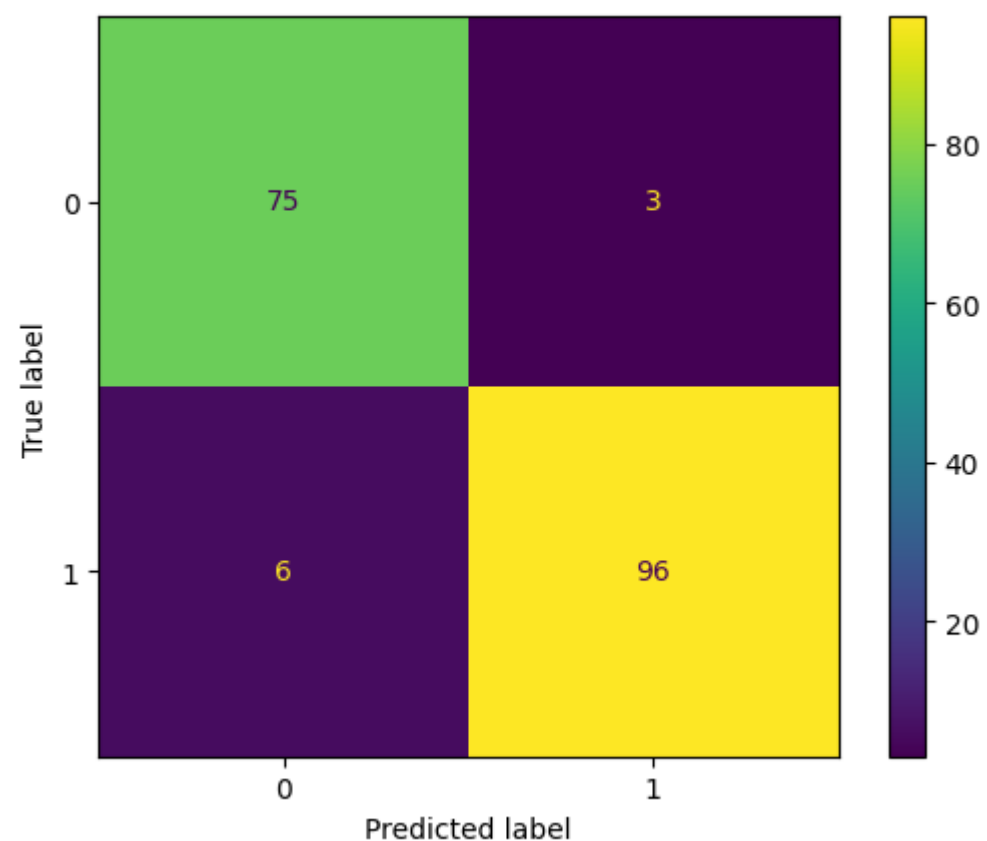
#classification report
report=classification_report(y_test,y_prediction)
print(report)

```

	precision	recall	f1-score	support
0	0.93	0.96	0.94	78
1	0.97	0.94	0.96	102
accuracy			0.95	180
macro avg	0.95	0.95	0.95	180
weighted avg	0.95	0.95	0.95	180

```
#confusion matrix display
labels=[0,1]
cmd=ConfusionMatrixDisplay(matr,display_labels=labels)
cmd.plot()
#0 : Benign
#1 : Malignant(riskier)

<sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x7b3ed84c5d20>
```



✓ BUILDING A PREDICTIVE SYSTEM


```
-----  
input_data=[[13,21.82,87.5,519.8,0.1273,0.1932,0.1859,0.09353,0.235,0.3063,2.406,24.32,0.03  
prediction=knn.predict(input_data)  
if (prediction[0]==1):  
    print("The Breast Cancer is Malignant")  
else:  
    print("The Breast Cancer is Benign")  
    The Breast Cancer is Malignant
```

✓ MODEL CREATION USING DECISION TREE CLASSIFIER, RANDOMFOREST CLASSIFIER, LOGISTIC REGRESSION AND THEN PERFORMANCE EVALUATION

```
from sklearn.tree import DecisionTreeClassifier  
from sklearn.ensemble import RandomForestClassifier  
from sklearn.linear_model import LogisticRegression  
  
dtc=DecisionTreeClassifier(criterion='entropy')  
rfc=RandomForestClassifier(n_estimators=100,criterion='entropy')  
lr=LogisticRegression(solver='saga',max_iter=100,class_weight='balanced')  
  
lst=[dtc,rfc,lr]  
for i in lst:  
    print(i)  
    i.fit(x_train,y_train)  
    y_prediction=i.predict(x_test)  
    print(confusion_matrix(y_test,y_prediction))
```

```
print(accuracy_score(y_test,y_prediction))
print(classification_report(y_test,y_prediction))
```

```
DecisionTreeClassifier(criterion='entropy')
```

```
[[76  2]
```

```
 [ 7 95]]
```

```
0.95
```

	precision	recall	f1-score	support
0	0.92	0.97	0.94	78
1	0.98	0.93	0.95	102
accuracy			0.95	180
macro avg	0.95	0.95	0.95	180
weighted avg	0.95	0.95	0.95	180

```
RandomForestClassifier(criterion='entropy')
```

```
[[74  4]
```

```
 [ 4 98]]
```

```
0.9555555555555556
```

	precision	recall	f1-score	support
0	0.95	0.95	0.95	78
1	0.96	0.96	0.96	102
accuracy			0.96	180
macro avg	0.95	0.95	0.95	180
weighted avg	0.96	0.96	0.96	180

```
LogisticRegression(class_weight='balanced', solver='saga')
```

```
[[71  7]
```

```
 [11 91]]
```

```
0.9
```

	precision	recall	f1-score	support
0	0.87	0.91	0.89	78
1	0.93	0.89	0.91	102
accuracy			0.90	180
macro avg	0.90	0.90	0.90	180
weighted avg	0.90	0.90	0.90	180

```
weighted avg      0.90      0.90      0.90      100
```

```
/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_sag.py:350: ConvergenceWarning: The max_iter was re  
warnings.warn(
```