

# BREAST CANCER DETECTION

## DATA SOURCE

<https://www.kaggle.com/datasets/uciml/breast-cancer-wisconsin-data>

## Pycaret Module

In [1]: `!pip install pycaret`

```
Collecting pycaret
  Obtaining dependency information for pycaret from https://files.pythonhosted.org/packages/f5/4b/2002980b046ac396618dfc152d384b812a78182b776ca77fe0ae5f80deac/pycaret-3.0.4-py3-none-any.whl.metadata
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Collecting ipywidgets>=7.6.5 (from pycaret)
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  Downloading ipywidgets-8.1.0-py3-none-any.whl.metadata (2.4 kB)
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Requirement already satisfied: pandas<2.0.0,>=1.3.0 in e:\anaconda3\lib\site-packages (from pycaret) (1.4.2)
Requirement already satisfied: jinja2>=1.2 in e:\anaconda3\lib\site-packages (from pycaret) (3.1.2)
Requirement already satisfied: scipy<2.0.0 in e:\anaconda3\lib\site-packages (from pycaret) (1.7.3)
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  Preparing metadata (setup.py): finished with status 'done'
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```

```

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Requirement already satisfied: numba>=0.55.0 in e:\anaconda3\lib\site-packages (from pycaret) (0.55.1)
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Requirement already satisfied: importlib-metadata>=4.12.0 in e:\anaconda3\lib\site-packages (from pycaret) (6.0.0)
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Requirement already satisfied: threadpoolctl>=2.0.0 in e:\anaconda3\lib\site-packages (from imbalanced-learn>=0.8.1->pycaret) (2.2.0)
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>=5.5.0->pycaret) (5.1.1)  
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Requirement already satisfied: python-dateutil>=2.7 in e:\anaconda3\lib\site-packages (from matplotlib>=3.3.0->pycaret) (2.8.2)  
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Requirement already satisfied: jupyter-core in e:\anaconda3\lib\site-packages (from nbformat>=4.2.0->pycaret) (5.3.0)  
Requirement already satisfied: llvmlite<0.39,>=0.38.0rc1 in e:\anaconda3\lib\site-packages (from numba>=0.55.0->pycaret) (0.38.0)  
Requirement already satisfied: setuptools in e:\anaconda3\lib\site-packages (from numba>=0.55.0->pycaret) (68.0.0)  
Requirement already satisfied: pytz>=2020.1 in e:\anaconda3\lib\site-packages (from pandas<2.0.0,>=1.3.0->pycaret) (2022.7)  
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Requirement already satisfied: urllib3 in e:\anaconda3\lib\site-packages (from pmdarima!=1.8.1,<3.0.0,>=1.8.0->pycaret) (1.26.16)  
Requirement already satisfied: charset-normalizer<4,>=2 in e:\anaconda3\lib\site-packages (from requests>=2.27.1->pycaret) (2.0.4)  
Requirement already satisfied: idna<4,>=2.5 in e:\anaconda3\lib\site-packages (from requests>=2.27.1->pycaret) (2.10)  
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Requirement already satisfied: Flask<2.3.0,>=1.0.4 in e:\anaconda3\lib\site-packages (from dash<3.0.0,>=2.11.0->plotly-resampler>=0.8.3.1->pycaret) (1.1.2)  
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Requirement already satisfied: executing in e:\anaconda3\lib\site-packages (from stack-data->ipython>=5.5.0->pycaret) (0.8.3)  
Requirement already satisfied: asttokens in e:\anaconda3\lib\site-packages (from stack-data->ipython>=5.5.0->pycaret) (2.0.5)  
Requirement already satisfied: pure-eval in e:\anaconda3\lib\site-packages (from stack-data->ipython>=5.5.0->pycaret) (0.2.2)  
Requirement already satisfied: itsdangerous>=0.24 in e:\anaconda3\lib\site-packages (from

```
m Flask<2.3.0,>=1.0.4->dash<3.0.0,>=2.11.0->plotly-resampler>=0.8.3.1->pycaret) (2.0.1)
Requirement already satisfied: click>=5.1 in e:\anaconda3\lib\site-packages (from Flask<
2.3.0,>=1.0.4->dash<3.0.0,>=2.11.0->plotly-resampler>=0.8.3.1->pycaret) (8.0.4)
Downloading pycaret-3.0.4-py3-none-any.whl (484 kB)
----- 484.4/484.4 kB 843.6 kB/s eta 0:00:00
Downloading category_encoders-2.6.2-py2.py3-none-any.whl (81 kB)
----- 81.8/81.8 kB 1.2 MB/s eta 0:00:00
Downloading imbalanced_learn-0.11.0-py3-none-any.whl (235 kB)
----- 235.6/235.6 kB 1.2 MB/s eta 0:00:00
Downloading ipywidgets-8.1.0-py3-none-any.whl (139 kB)
----- 139.3/139.3 kB 921.9 kB/s eta 0:00:00
```

## Import Modules

```
In [3]: import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
import warnings
from pycaret.classification import *
%matplotlib inline
warnings.filterwarnings('ignore')
```

```
In [4]: # Load Dataset

df = pd.read_csv(r"E:\PORTFOLIO\1. DATA ANALYTICS\5. BREAST CANCER DETECTION\DATA\data.c
```

```
In [5]: df.head()
```

```
Out[5]:
```

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

5 rows × 33 columns

```
In [7]: # delete unnecessary columns

df = df.drop(columns=['id', 'Unnamed: 32'], axis=1)
```

```
In [8]: # Statistical Information

df.describe()
```

```
Out[8]:
```

	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity
count	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569
mean	14.127292	19.289649	91.969033	654.889104	0.096360	0.104341	0
std	3.524049	4.301036	24.298981	351.914129	0.014064	0.052813	0
min	6.981000	9.710000	43.790000	143.500000	0.052630	0.019380	0

<b>25%</b>	11.700000	16.170000	75.170000	420.300000	0.086370	0.064920	0
<b>50%</b>	13.370000	18.840000	86.240000	551.100000	0.095870	0.092630	0
<b>75%</b>	15.780000	21.800000	104.100000	782.700000	0.105300	0.130400	0
<b>max</b>	28.110000	39.280000	188.500000	2501.000000	0.163400	0.345400	0

8 rows × 30 columns

In [9]: `# datatype info`

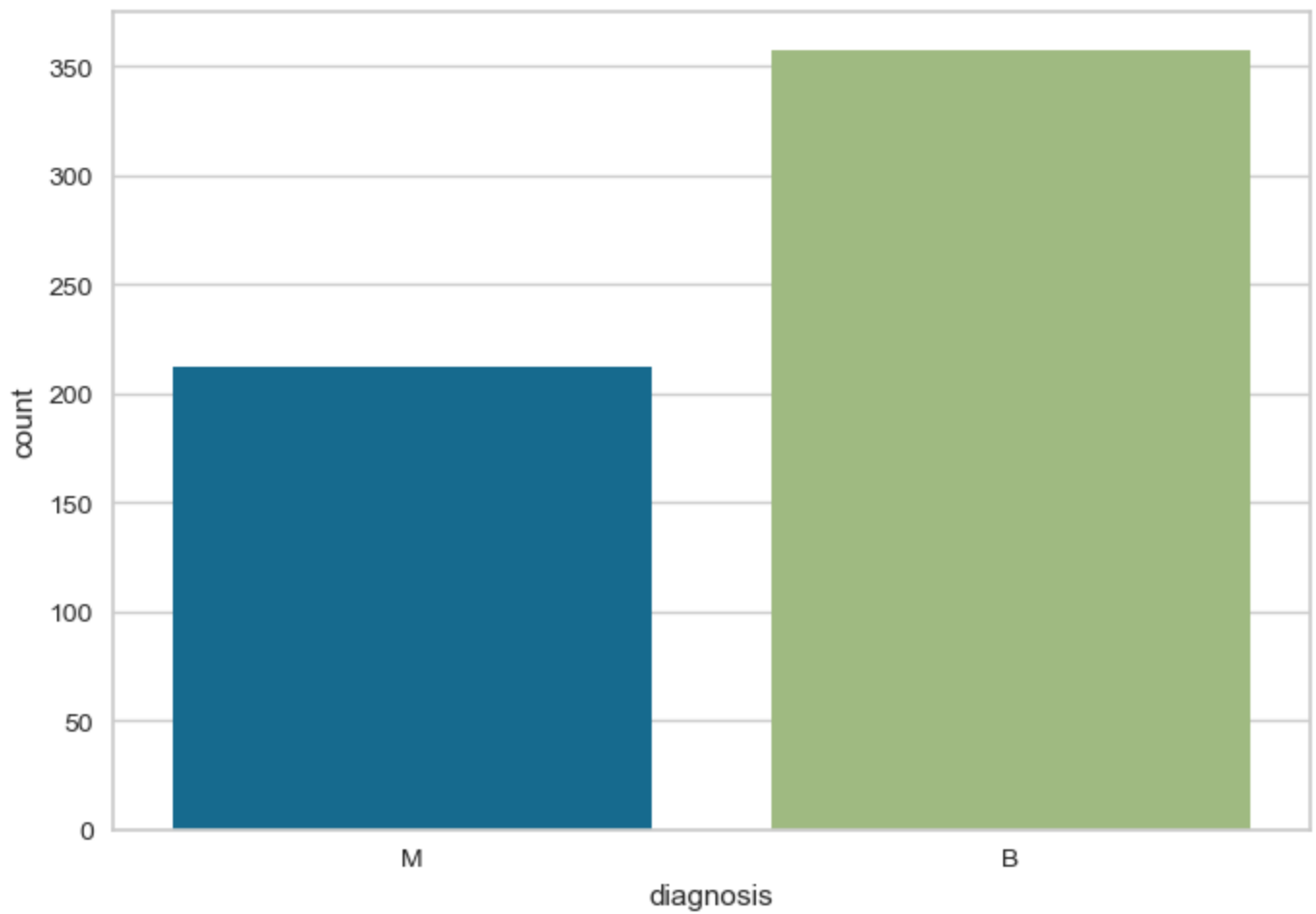
`df.info()`

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 31 columns):
#   Column                                     Non-Null Count  Dtype
---  -
0   diagnosis                                 569 non-null    object
1   radius_mean                             569 non-null    float64
2   texture_mean                             569 non-null    float64
3   perimeter_mean                           569 non-null    float64
4   area_mean                                569 non-null    float64
5   smoothness_mean                           569 non-null    float64
6   compactness_mean                           569 non-null    float64
7   concavity_mean                             569 non-null    float64
8   concave points_mean                       569 non-null    float64
9   symmetry_mean                             569 non-null    float64
10  fractal_dimension_mean                    569 non-null    float64
11  radius_se                                 569 non-null    float64
12  texture_se                                 569 non-null    float64
13  perimeter_se                               569 non-null    float64
14  area_se                                   569 non-null    float64
15  smoothness_se                             569 non-null    float64
16  compactness_se                             569 non-null    float64
17  concavity_se                               569 non-null    float64
18  concave points_se                         569 non-null    float64
19  symmetry_se                               569 non-null    float64
20  fractal_dimension_se                      569 non-null    float64
21  radius_worst                              569 non-null    float64
22  texture_worst                             569 non-null    float64
23  perimeter_worst                           569 non-null    float64
24  area_worst                                569 non-null    float64
25  smoothness_worst                          569 non-null    float64
26  compactness_worst                         569 non-null    float64
27  concavity_worst                           569 non-null    float64
28  concave points_worst                      569 non-null    float64
29  symmetry_worst                            569 non-null    float64
30  fractal_dimension_worst                   569 non-null    float64
dtypes: float64(30), object(1)
memory usage: 137.9+ KB
```

## Exploratory Data

In [10]: `sns.countplot(df['diagnosis'])`

Out[10]: `<AxesSubplot:xlabel='diagnosis', ylabel='count'>`

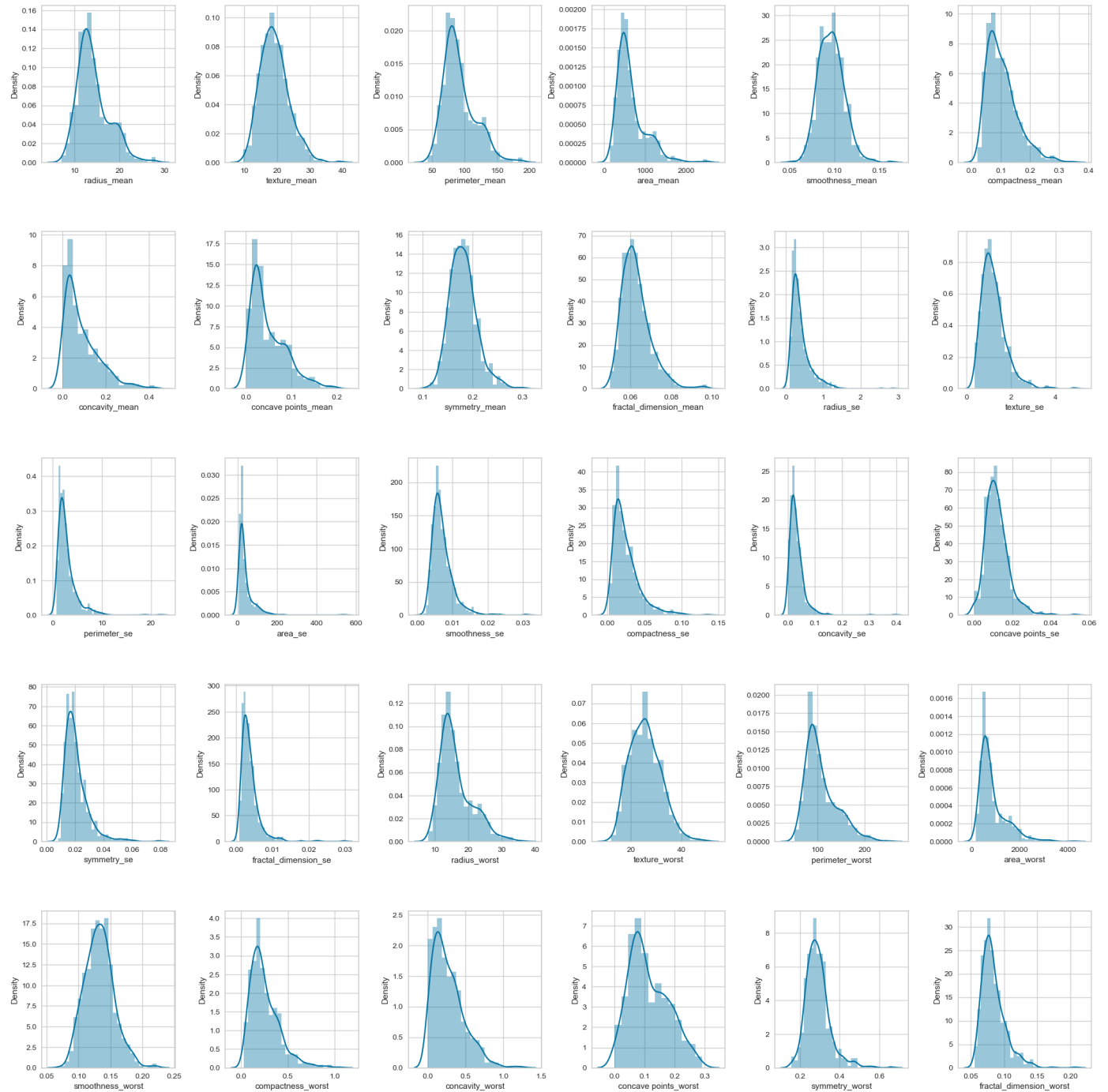


M = Malignant , B = Benign

```
In [11]: df_temp = df.drop(columns=['diagnosis'], axis=1)
```

```
In [13]: # create dist plot
fig, ax = plt.subplots(ncols=6, nrow=5, figsize=(20, 20))
index = 0
ax = ax.flatten()

for col in df_temp.columns:
    sns.distplot(df[col], ax=ax[index])
    index+=1
plt.tight_layout(pad=0.5, w_pad=0.7, h_pad=5.0)
```

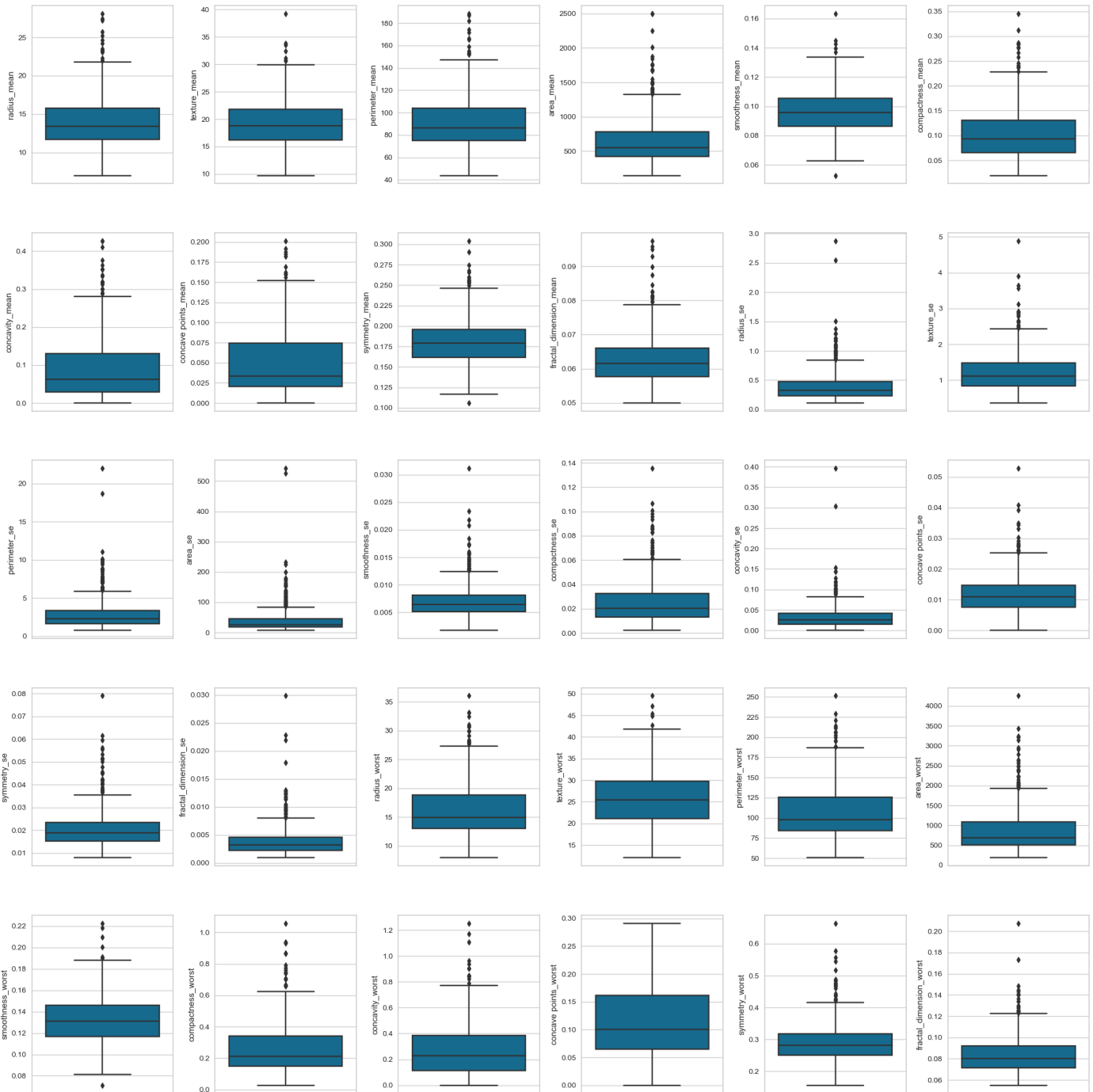


In [14]: `# create box plot`

```
fig, ax = plt.subplots(ncols=6, nrows=5, figsize=(20, 20))
index = 0
ax = ax.flatten()

for col in df_temp.columns:
    sns.boxplot(y=col, data=df, ax=ax[index])
    index+=1
plt.tight_layout(pad=0.5, w_pad=0.7, h_pad=5.0)
```





## Create and Train the Model

```
In [22]: # Setup the Data

clf = setup(df, target='diagnosis')
```

	Description	Value
0	Session id	262
1	Target	diagnosis
2	Target type	Binary
3	Target mapping	B: 0, M: 1
4	Original data shape	(569, 31)
5	Transformed data shape	(569, 31)
6	Transformed train set shape	(398, 31)

7	Transformed test set shape	(171, 31)
8	Numeric features	30
9	Preprocess	True
10	Imputation type	simple
11	Numeric imputation	mean
12	Categorical imputation	mode
13	Fold Generator	StratifiedKFold
14	Fold Number	10
15	CPU Jobs	-1
16	Use GPU	False
17	Log Experiment	False
18	Experiment Name	clf-default-name
19	USI	ca23

```
In [23]: # train and test the models
compare_models()
```

	Model	Accuracy	AUC	Recall	Prec.	F1	Kappa	MCC	TT (Sec)
<b>ada</b>	Ada Boost Classifier	0.9724	0.9933	0.9529	0.9750	0.9629	0.9410	0.9422	0.3550
<b>gbc</b>	Gradient Boosting Classifier	0.9649	0.9952	0.9529	0.9585	0.9540	0.9257	0.9280	0.5100
<b>lightgbm</b>	Light Gradient Boosting Machine	0.9624	0.9938	0.9595	0.9464	0.9512	0.9208	0.9230	0.3960
<b>et</b>	Extra Trees Classifier	0.9623	0.9953	0.9324	0.9670	0.9477	0.9183	0.9206	0.4650
<b>lda</b>	Linear Discriminant Analysis	0.9574	0.9906	0.8929	0.9923	0.9378	0.9059	0.9110	0.1680
<b>rf</b>	Random Forest Classifier	0.9548	0.9933	0.9390	0.9467	0.9401	0.9039	0.9074	0.4810
<b>qda</b>	Quadratic Discriminant Analysis	0.9547	0.9887	0.9390	0.9440	0.9386	0.9029	0.9063	0.1390
<b>ridge</b>	Ridge Classifier	0.9524	0.0000	0.8857	0.9866	0.9305	0.8948	0.9007	0.1330
<b>lr</b>	Logistic Regression	0.9447	0.9908	0.9195	0.9335	0.9237	0.8806	0.8838	1.4840
<b>dt</b>	Decision Tree Classifier	0.9422	0.9373	0.9186	0.9279	0.9203	0.8751	0.8785	0.1440
<b>nb</b>	Naive Bayes	0.9347	0.9873	0.8857	0.9390	0.9081	0.8579	0.8624	0.1400
<b>knn</b>	K Neighbors Classifier	0.9072	0.9518	0.8314	0.9173	0.8668	0.7964	0.8042	0.1640
<b>svm</b>	SVM - Linear Kernel	0.9021	0.0000	0.7833	0.9513	0.8525	0.7811	0.7946	0.1320
<b>dummy</b>	Dummy Classifier	0.6282	0.5000	0.0000	0.0000	0.0000	0.0000	0.0000	0.1760

```
Out[23]: AdaBoostClassifier(algorithm='SAMME.R', base_estimator=None, learning_rate=1.0,
                             n_estimators=50, random_state=262)
```

```
In [28]: # select the best model
model = create_model('ada')
```

Accuracy    AUC    Recall    Prec.    F1    Kappa    MCC

Fold

0	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
1	0.9750	1.0000	0.9333	1.0000	0.9655	0.9459	0.9473
2	0.9500	0.9787	0.8667	1.0000	0.9286	0.8904	0.8958
3	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
4	0.9250	0.9787	0.9333	0.8750	0.9032	0.8421	0.8433
5	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
6	0.9750	1.0000	0.9333	1.0000	0.9655	0.9459	0.9473
7	0.9250	0.9787	0.9333	0.8750	0.9032	0.8421	0.8433
8	0.9744	0.9971	0.9286	1.0000	0.9630	0.9434	0.9449
9	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
Mean	0.9724	0.9933	0.9529	0.9750	0.9629	0.9410	0.9422
Std	0.0284	0.0096	0.0429	0.0500	0.0372	0.0601	0.0593

```
In [29]: # hyperparameter tuning

best_model = tune_model(model)
```

	Accuracy	AUC	Recall	Prec.	F1	Kappa	MCC
Fold							
0	0.9750	0.9973	1.0000	0.9375	0.9677	0.9474	0.9487
1	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
2	0.9500	0.9813	0.8667	1.0000	0.9286	0.8904	0.8958
3	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
4	0.9500	0.9840	0.9333	0.9333	0.9333	0.8933	0.8933
5	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
6	0.9750	0.9973	1.0000	0.9375	0.9677	0.9474	0.9487
7	0.9500	0.9893	0.9333	0.9333	0.9333	0.8933	0.8933
8	0.9487	0.9943	0.9286	0.9286	0.9286	0.8886	0.8886
9	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
Mean	0.9749	0.9944	0.9662	0.9670	0.9659	0.9460	0.9468
Std	0.0225	0.0067	0.0451	0.0331	0.0309	0.0486	0.0480

Fitting 10 folds for each of 10 candidates, totalling 100 fits

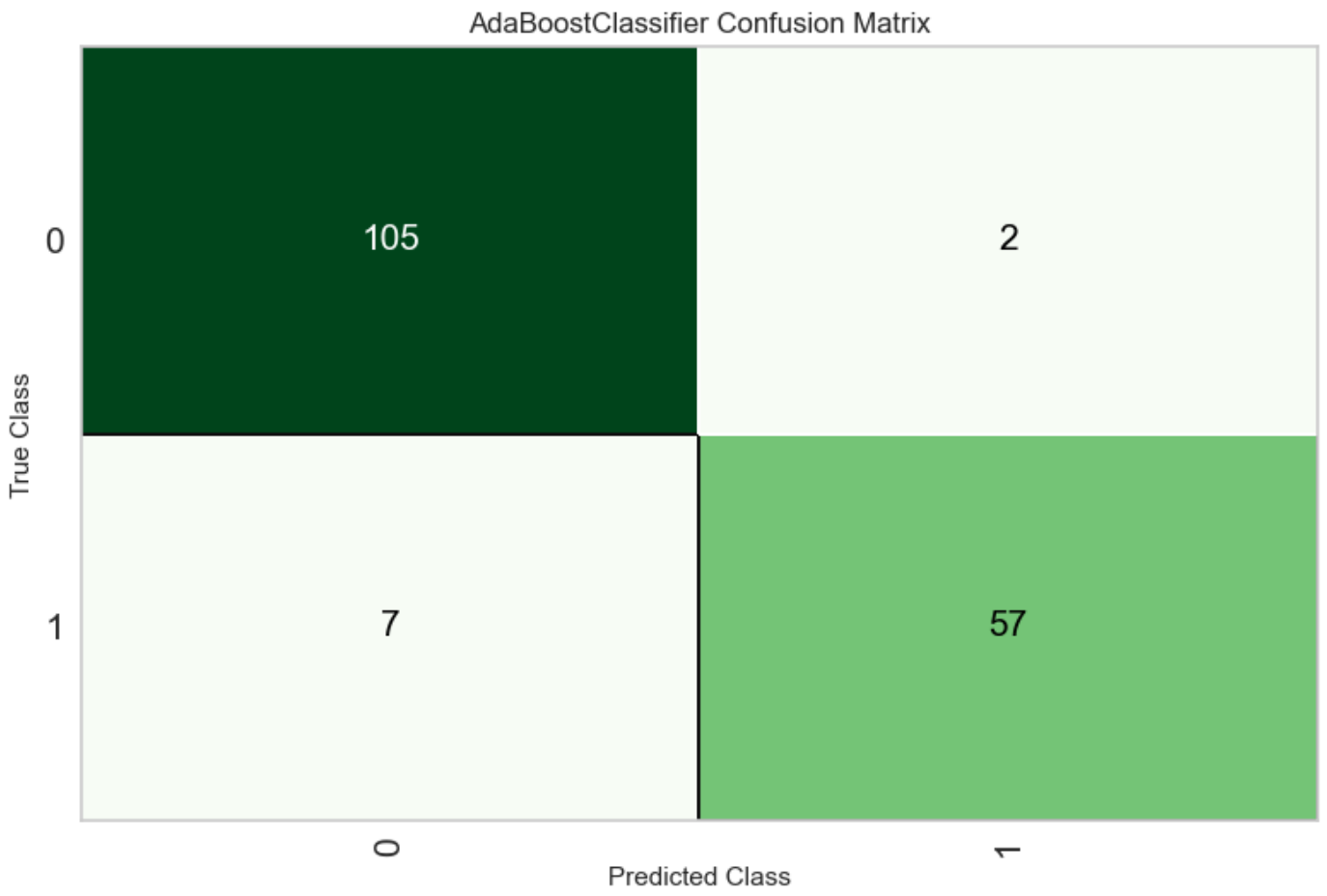
```
In [30]: evaluate_model(best_model)
```



```
interactive(children=(ToggleButtons(description='Plot Type:', icons=('',)), options=('', 'Pi
pipeline Plot', 'pipelin...
```

```
In [31]: # plot the result

plot_model(estimator=best_model, plot='confusion_matrix')
```



In [ ]: