

# BREAST CANCER DETECTION MODEL

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## OVERVIEW

One of the main causes of cancer-related mortality for women globally is breast cancer. Treatment results and survival rates can be greatly enhanced by early identification. This project's objective is to develop a machine learning model that, using pertinent characteristics extracted from clinical and diagnostic data, reliably classifies tumors as benign (non-cancerous) or malignant (cancerous).

## OBJECTIVE

The objective of this assessment is to evaluate the understanding and ability to apply supervised learning techniques to a real-world dataset.

## DATASET DESCRIPTION:

BREAST CANCER: SKLEARN LIBRARY

## IMPORTING LIBRARIES

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

from sklearn.preprocessing import LabelEncoder, OneHotEncoder
from sklearn.feature_selection import VarianceThreshold
from sklearn.preprocessing import StandardScaler, MinMaxScaler

from sklearn.model_selection import train_test_split

from sklearn.linear_model import LogisticRegression
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from sklearn.neighbors import KNeighborsClassifier

from sklearn.metrics import accuracy_score, classification_report,
confusion_matrix

from sklearn.model_selection import GridSearchCV

import joblib
```

```
import warnings
warnings.filterwarnings("ignore")
```

## IMPORTING DATASET

```
from sklearn.datasets import load_breast_cancer
data = load_breast_cancer()
data
```

{'data': array([[1.799e+01, 1.038e+01, 1.228e+02, ..., 2.654e-01,  
4.601e-01,  
1.189e-01],  
[2.057e+01, 1.777e+01, 1.329e+02, ..., 1.860e-01, 2.750e-01,  
8.902e-02],  
[1.969e+01, 2.125e+01, 1.300e+02, ..., 2.430e-01, 3.613e-01,  
8.758e-02],  
...,  
[1.660e+01, 2.808e+01, 1.083e+02, ..., 1.418e-01, 2.218e-01,  
7.820e-02],  
[2.060e+01, 2.933e+01, 1.401e+02, ..., 2.650e-01, 4.087e-01,  
1.240e-01],  
[7.760e+00, 2.454e+01, 4.792e+01, ..., 0.000e+00, 2.871e-01,  
7.039e-02]]),  
'target': array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,  
0, 0, 1, 1, 1,  
0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0,  
0,  
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0,  
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0,  
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0,  
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0,  
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0,  
0, 0, 1, 1, 1, 1, 1, 1, 0, 1, 0, 1, 1, 0, 1, 1, 0, 1, 0, 0, 1,  
1,  
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1,

```

1,
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0,
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0,
    0, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 1, 0, 0, 0, 1, 0,
0,
    1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 0, 1, 1, 1, 0, 1, 1, 0, 0, 1,
1,
    1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 0, 1, 1,
0,
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1,
    1, 0, 1, 1, 0, 1, 0, 1, 1, 0, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 0,
0,
    1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1,
1,
    1, 1, 1, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 0, 0, 1, 0, 1, 0, 1,
1,
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1,
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1,
    1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 1)),
'frame': None,
'target_names': array(['malignant', 'benign'], dtype='<U9'),
'DESCR': '.. _breast_cancer_dataset:\n\nBreast cancer wisconsin
(diagnostic) dataset\n-----\n\n
**Data Set Characteristics:**\n\n:
Number of Instances: 569\n\n:
Number of Attributes: 30 numeric, predictive attributes and the class\n
n:
Attribute Information:\n
    - radius (mean of distances from center
to points on the perimeter)\n
    - texture (standard deviation of
gray-scale values)\n
    - perimeter\n
    - area\n
    - smoothness
(local variation in radius lengths)\n
    - compactness (perimeter^2 /
area - 1.0)\n
    - concavity (severity of concave portions of the
contour)\n
    - concave points (number of concave portions of the
contour)\n
    - symmetry\n
    - fractal dimension ("coastline
approximation" - 1)\n\n
The mean, standard error, and "worst" or
largest (mean of the three\n
worst/largest values) of these
features were computed for each image,\n
resulting in 30 features.
For instance, field 0 is Mean Radius, field\n
10 is Radius SE,
field 20 is Worst Radius.\n\n
    - class:\n
    - WDBC-
Malignant\n
    - WDBC-Benign\n\n:
Summary Statistics:\n\n
n===== \n
Min    Max\n===== \n
nradius (mean):          6.981  28.11\ntexture (mean):
9.71  39.28\nnperimeter (mean):          43.79  188.5\nnarea
(mean):          143.5  2501.0\nnsmoothness (mean):
0.053  0.163\nncompactness (mean):          0.019  0.345\

```

```

nconcavity (mean):          0.0    0.427\nconcave points
(mean):          0.0    0.201\nsymmetry (mean):
0.106  0.304\nfractal dimension (mean):          0.05    0.097\n
nradius (standard error):          0.112  2.873\ntexture (standard
error):          0.36    4.885\nperimeter (standard error):
0.757  21.98\narea (standard error):          6.802  542.2\n
nsmoothness (standard error):          0.002  0.031\ncompactness
(standard error):          0.002  0.135\nconcavity (standard error):
0.0    0.396\nconcave points (standard error):          0.0    0.053\n
nsymmetry (standard error):          0.008  0.079\nfractal dimension
(standard error):  0.001  0.03\nradius (worst):
7.93   36.04\ntexture (worst):          12.02  49.54\n
nperimeter (worst):          50.41  251.2\narea (worst):
185.2  4254.0\nsmoothness (worst):          0.071  0.223\n
ncompactness (worst):          0.027  1.058\nconcavity
(worst):          0.0    1.252\nconcave points (worst):
0.0    0.291\nsymmetry (worst):          0.156  0.664\n
nfractal dimension (worst):          0.055  0.208\n
n===== \n\nMissing
Attribute Values: None\n\nClass Distribution: 212 - Malignant, 357 -
Benign\n\nCreator: Dr. William H. Wolberg, W. Nick Street, Olvi L.
Mangasarian\n\nDonor: Nick Street\n\nDate: November, 1995\n\nThis is
a copy of UCI ML Breast Cancer Wisconsin (Diagnostic) datasets.\n
nhttps://goo.gl/U2Uwz2\n\nFeatures are computed from a digitized image
of a fine needle\naspirate (FNA) of a breast mass. They describe\n
ncharacteristics of the cell nuclei present in the image.\n\n
nSeparating plane described above was obtained using\nMultisurface
Method-Tree (MSM-T) [K. P. Bennett, "Decision Tree\nConstruction Via
Linear Programming." Proceedings of the 4th\nMidwest Artificial
Intelligence and Cognitive Science Society,\npp. 97-101, 1992], a
classification method which uses linear\nprogramming to construct a
decision tree. Relevant features\nwere selected using an exhaustive
search in the space of 1-4\nfeatures and 1-3 separating planes.\n\nThe
actual linear program used to obtain the separating plane\nin the 3-
dimensional space is that described in:\n[K. P. Bennett and O. L.
Mangasarian: "Robust Linear\nProgramming Discrimination of Two
Linearly Inseparable Sets",\nOptimization Methods and Software 1,
1992, 23-34].\n\nThis database is also available through the UW CS ftp
server:\n\nftp ftp.cs.wisc.edu\ncd math-prog/cpo-dataset/machine-
learn/WDBC/\n\n|details-start|\n**References**\n|details-split|\n\n-
W.N. Street, W.H. Wolberg and O.L. Mangasarian. Nuclear feature
extraction\n for breast tumor diagnosis. IS&T/SPIE 1993 International
Symposium on\n Electronic Imaging: Science and Technology, volume
1905, pages 861-870,\n San Jose, CA, 1993.\n- O.L. Mangasarian, W.N.
Street and W.H. Wolberg. Breast cancer diagnosis and\n prognosis via
linear programming. Operations Research, 43(4), pages 570-577,\n
July-August 1995.\n- W.H. Wolberg, W.N. Street, and O.L. Mangasarian.
Machine learning techniques\n to diagnose breast cancer from fine-
needle aspirates. Cancer Letters 77 (1994)\n 163-171.\n\n|details-

```

```

end|\n',
'feature_names': array(['mean radius', 'mean texture', 'mean
perimeter', 'mean area',
                        'mean smoothness', 'mean compactness', 'mean concavity',
                        'mean concave points', 'mean symmetry', 'mean fractal
dimension',
                        'radius error', 'texture error', 'perimeter error', 'area
error',
                        'smoothness error', 'compactness error', 'concavity error',
                        'concave points error', 'symmetry error',
                        'fractal dimension error', 'worst radius', 'worst texture',
                        'worst perimeter', 'worst area', 'worst smoothness',
                        'worst compactness', 'worst concavity', 'worst concave
points',
                        'worst symmetry', 'worst fractal dimension'], dtype='<U23'),
'filename': 'breast_cancer.csv',
'data_module': 'sklearn.datasets.data'}

```

*# converting to dataframe*

```
df = pd.DataFrame(data.data, columns=data.feature_names)
```

```
df['target'] = data.target
```

```
df.head()
```

	mean radius	mean texture	mean perimeter	mean area	mean smoothness \
0	17.99	10.38	122.80	1001.0	0.11840
1	20.57	17.77	132.90	1326.0	0.08474
2	19.69	21.25	130.00	1203.0	0.10960
3	11.42	20.38	77.58	386.1	0.14250
4	20.29	14.34	135.10	1297.0	0.10030

	mean compactness	mean concavity	mean concave points	mean symmetry \
0	0.27760	0.3001	0.14710	0.2419
1	0.07864	0.0869	0.07017	0.1812
2	0.15990	0.1974	0.12790	0.2069
3	0.28390	0.2414	0.10520	0.2597
4	0.13280	0.1980	0.10430	0.1809

	mean fractal dimension	...	worst texture	worst perimeter	worst
area \					
0	0.07871	...	17.33	184.60	
2019.0					
1	0.05667	...	23.41	158.80	
1956.0					
2	0.05999	...	25.53	152.50	
1709.0					
3	0.09744	...	26.50	98.87	
567.7					
4	0.05883	...	16.67	152.20	
1575.0					

	worst smoothness	worst compactness	worst concavity	worst concave
points \				
0	0.1622	0.6656	0.7119	
0.2654				
1	0.1238	0.1866	0.2416	
0.1860				
2	0.1444	0.4245	0.4504	
0.2430				
3	0.2098	0.8663	0.6869	
0.2575				
4	0.1374	0.2050	0.4000	
0.1625				

	worst symmetry	worst fractal dimension	target
0	0.4601	0.11890	0
1	0.2750	0.08902	0
2	0.3613	0.08758	0
3	0.6638	0.17300	0
4	0.2364	0.07678	0

[5 rows x 31 columns]

df.info()

<class 'pandas.core.frame.DataFrame'>

RangeIndex: 569 entries, 0 to 568

Data columns (total 31 columns):

#	Column	Non-Null Count	Dtype
---	-----	-----	-----
0	mean radius	569 non-null	float64
1	mean texture	569 non-null	float64
2	mean perimeter	569 non-null	float64
3	mean area	569 non-null	float64
4	mean smoothness	569 non-null	float64
5	mean compactness	569 non-null	float64
6	mean concavity	569 non-null	float64
7	mean concave points	569 non-null	float64

8	mean symmetry	569	non-null	float64
9	mean fractal dimension	569	non-null	float64
10	radius error	569	non-null	float64
11	texture error	569	non-null	float64
12	perimeter error	569	non-null	float64
13	area error	569	non-null	float64
14	smoothness error	569	non-null	float64
15	compactness error	569	non-null	float64
16	concavity error	569	non-null	float64
17	concave points error	569	non-null	float64
18	symmetry error	569	non-null	float64
19	fractal dimension error	569	non-null	float64
20	worst radius	569	non-null	float64
21	worst texture	569	non-null	float64
22	worst perimeter	569	non-null	float64
23	worst area	569	non-null	float64
24	worst smoothness	569	non-null	float64
25	worst compactness	569	non-null	float64
26	worst concavity	569	non-null	float64
27	worst concave points	569	non-null	float64
28	worst symmetry	569	non-null	float64
29	worst fractal dimension	569	non-null	float64
30	target	569	non-null	int32

dtypes: float64(30), int32(1)

memory usage: 135.7 KB

df.describe()

	mean radius	mean texture	mean perimeter	mean area \
count	569.000000	569.000000	569.000000	569.000000
mean	14.127292	19.289649	91.969033	654.889104
std	3.524049	4.301036	24.298981	351.914129
min	6.981000	9.710000	43.790000	143.500000
25%	11.700000	16.170000	75.170000	420.300000
50%	13.370000	18.840000	86.240000	551.100000
75%	15.780000	21.800000	104.100000	782.700000
max	28.110000	39.280000	188.500000	2501.000000

	mean smoothness	mean compactness	mean concavity	mean concave points \
count	569.000000	569.000000	569.000000	569.000000
mean	0.096360	0.104341	0.088799	0.048919
std	0.014064	0.052813	0.079720	0.038803
min	0.052630	0.019380	0.000000	0.000000
25%	0.086370	0.064920	0.029560	0.020310

50%	0.095870	0.092630	0.061540
0.033500			
75%	0.105300	0.130400	0.130700
0.074000			
max	0.163400	0.345400	0.426800
0.201200			

	mean symmetry	mean fractal dimension	...	worst texture \
count	569.000000	569.000000	...	569.000000
mean	0.181162	0.062798	...	25.677223
std	0.027414	0.007060	...	6.146258
min	0.106000	0.049960	...	12.020000
25%	0.161900	0.057700	...	21.080000
50%	0.179200	0.061540	...	25.410000
75%	0.195700	0.066120	...	29.720000
max	0.304000	0.097440	...	49.540000

	worst perimeter	worst area	worst smoothness	worst compactness \
count	569.000000	569.000000	569.000000	569.000000
mean	107.261213	880.583128	0.132369	0.254265
std	33.602542	569.356993	0.022832	0.157336
min	50.410000	185.200000	0.071170	0.027290
25%	84.110000	515.300000	0.116600	0.147200
50%	97.660000	686.500000	0.131300	0.211900
75%	125.400000	1084.000000	0.146000	0.339100
max	251.200000	4254.000000	0.222600	1.058000

	worst concavity	worst concave points	worst symmetry \
count	569.000000	569.000000	569.000000
mean	0.272188	0.114606	0.290076
std	0.208624	0.065732	0.061867
min	0.000000	0.000000	0.156500
25%	0.114500	0.064930	0.250400
50%	0.226700	0.099930	0.282200
75%	0.382900	0.161400	0.317900
max	1.252000	0.291000	0.663800

	worst fractal dimension	target
count	569.000000	569.000000
mean	0.083946	0.627417
std	0.018061	0.483918



min	0.055040	0.000000
25%	0.071460	0.000000
50%	0.080040	1.000000
75%	0.092080	1.000000
max	0.207500	1.000000

[8 rows x 31 columns]

df.shape

(569, 31)

## DATA CLEANING AND PRE PROCESSING

*#checking for duplicated*

df.duplicated()

```
0      False
1      False
2      False
3      False
4      False
```

...

```
564     False
565     False
566     False
567     False
568     False
```

Length: 569, dtype: bool

df.duplicated().sum()

0

## Checking for missing values

df.isnull()

	mean radius	mean texture	mean perimeter	mean area	mean
smoothness \					
0	False	False	False	False	
False					
1	False	False	False	False	
False					
2	False	False	False	False	
False					
3	False	False	False	False	
False					
4	False	False	False	False	
False					
..	...	...	...	...	...

```

...
564      False      False      False      False
False
565      False      False      False      False
False
566      False      False      False      False
False
567      False      False      False      False
False
568      False      False      False      False
False

      mean compactness  mean concavity  mean concave points  mean
symmetry \
0      False      False      False
False
1      False      False      False
False
2      False      False      False
False
3      False      False      False
False
4      False      False      False
False
..      ...      ...      ...
...
564      False      False      False
False
565      False      False      False
False
566      False      False      False
False
567      False      False      False
False
568      False      False      False
False

      mean fractal dimension  ...  worst texture  worst perimeter
worst area \
0      False  ...      False      False
False
1      False  ...      False      False
False
2      False  ...      False      False
False
3      False  ...      False      False
False
4      False  ...      False      False
False

```

```

..      ...      ...      ...
...
564      False      ...      False      False
False
565      False      ...      False      False
False
566      False      ...      False      False
False
567      False      ...      False      False
False
568      False      ...      False      False
False

```

```

      worst smoothness      worst compactness      worst concavity \
0      False      False      False
1      False      False      False
2      False      False      False
3      False      False      False
4      False      False      False
..      ...      ...      ...
564      False      False      False
565      False      False      False
566      False      False      False
567      False      False      False
568      False      False      False

```

```

      worst concave points      worst symmetry      worst fractal dimension
target
0      False      False      False
False
1      False      False      False
False
2      False      False      False
False
3      False      False      False
False
4      False      False      False
False
..      ...      ...      ...
...
564      False      False      False
False
565      False      False      False
False
566      False      False      False
False
567      False      False      False
False
568      False      False      False

```

False

[569 rows x 31 columns]

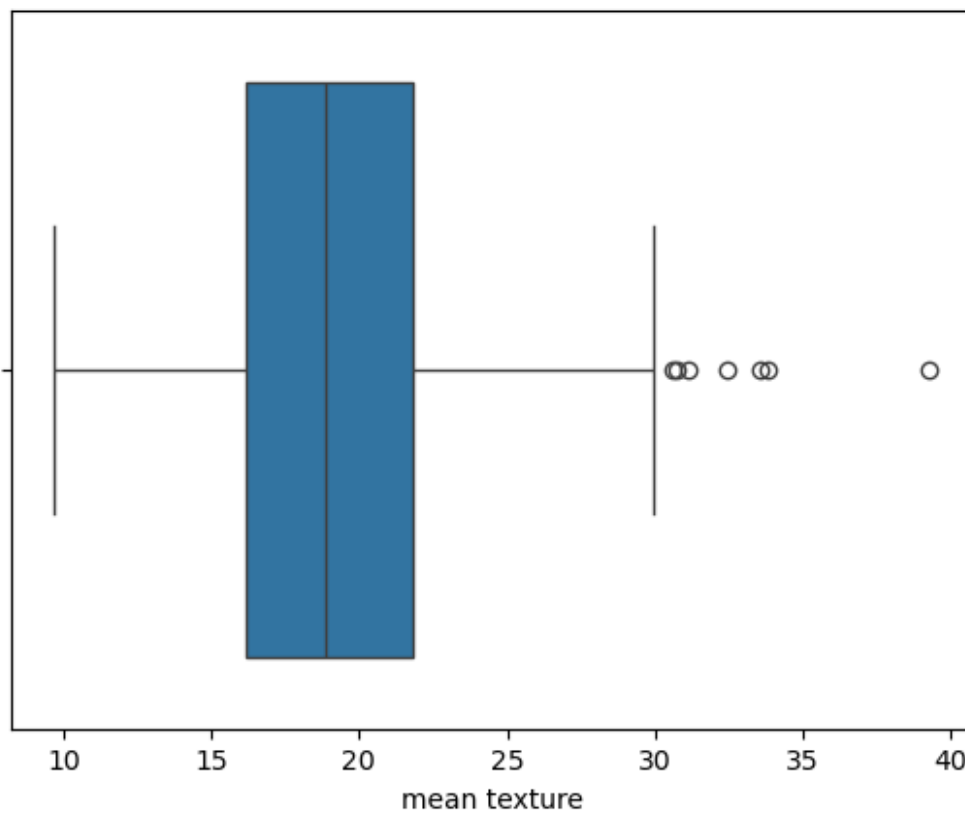
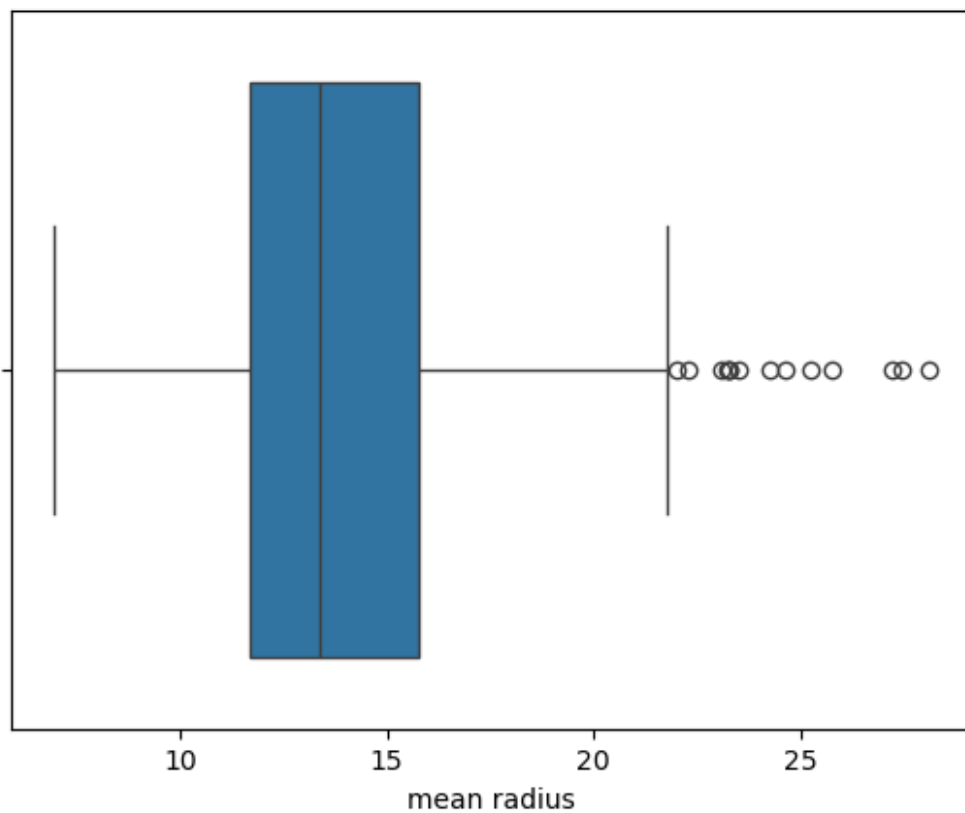
```
df.isnull().sum()
```

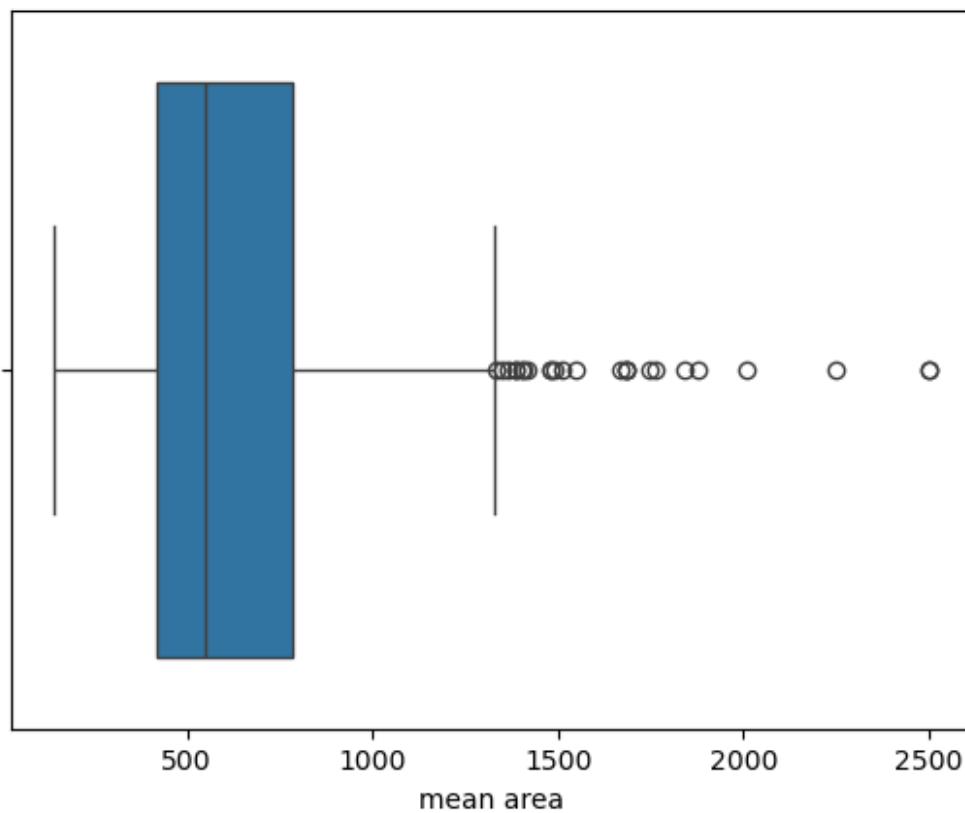
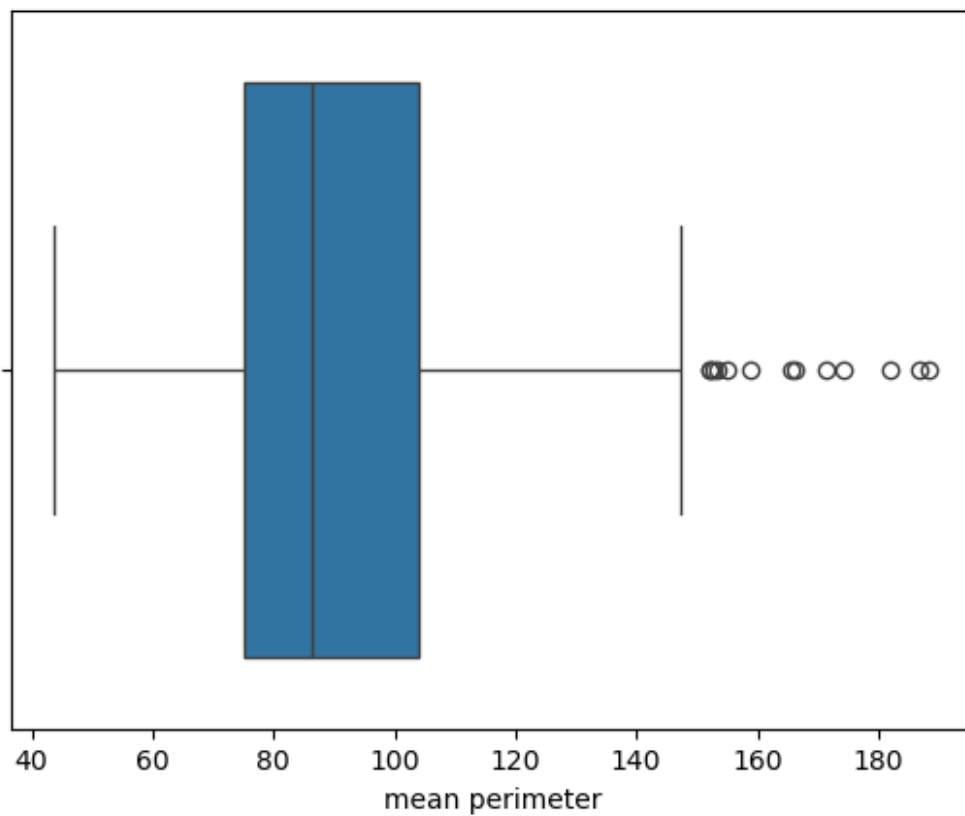
mean radius	0
mean texture	0
mean perimeter	0
mean area	0
mean smoothness	0
mean compactness	0
mean concavity	0
mean concave points	0
mean symmetry	0
mean fractal dimension	0
radius error	0
texture error	0
perimeter error	0
area error	0
smoothness error	0
compactness error	0
concavity error	0
concave points error	0
symmetry error	0
fractal dimension error	0
worst radius	0
worst texture	0
worst perimeter	0
worst area	0
worst smoothness	0
worst compactness	0
worst concavity	0
worst concave points	0
worst symmetry	0
worst fractal dimension	0
target	0

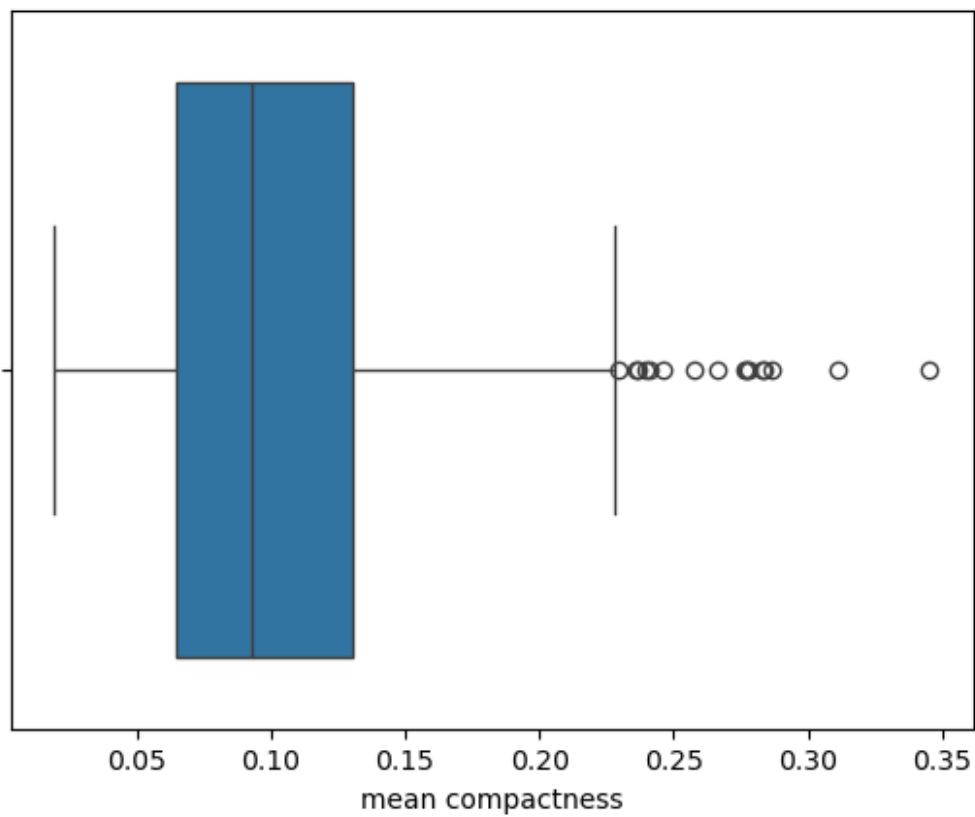
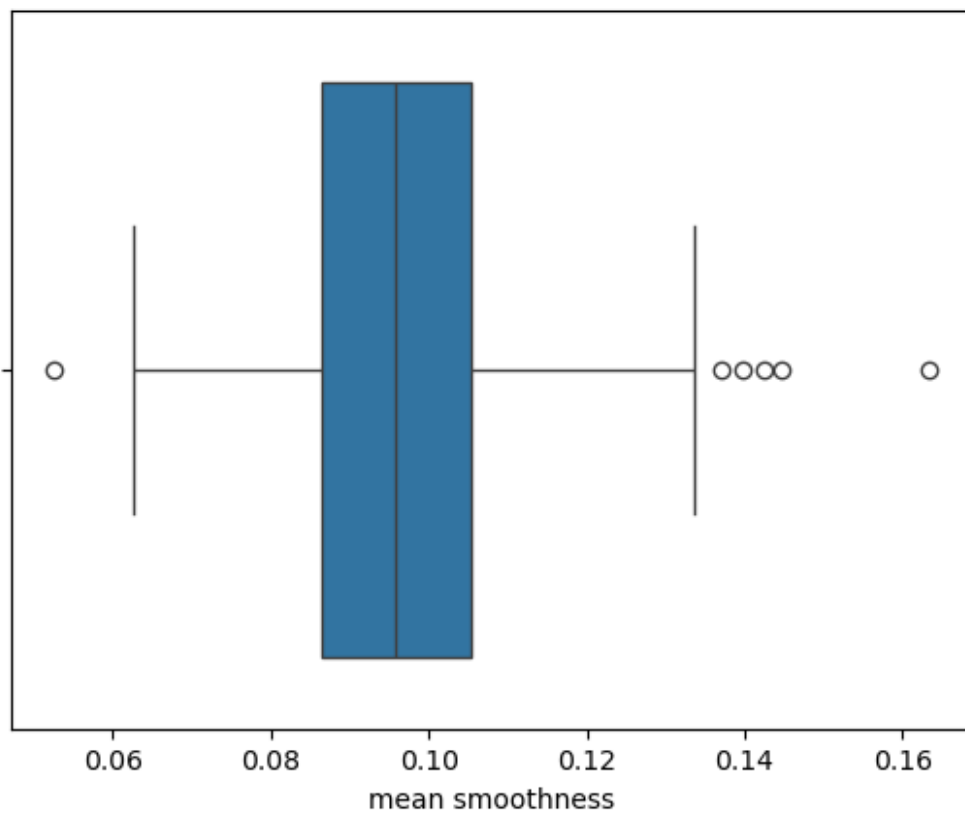
dtype: int64

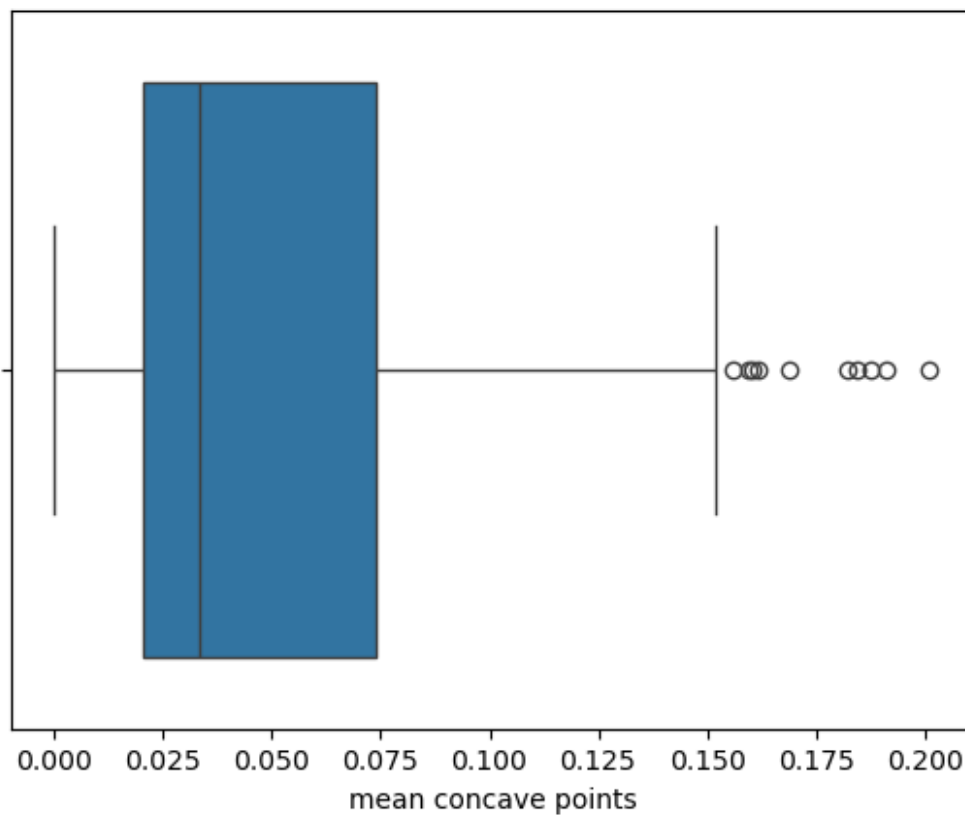
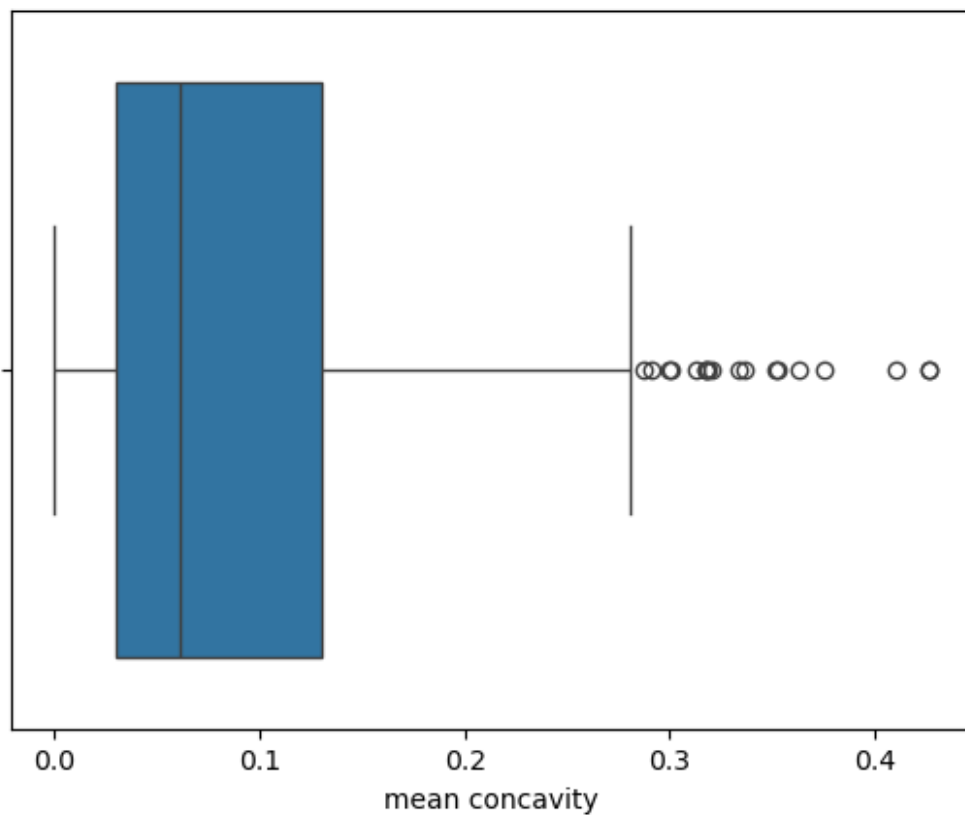
## checking for Outliers

```
# Visualising outliers in each feature using boxplot method
for i in df.columns:
    sns.boxplot(data=df,x=i)
    plt.show()
```

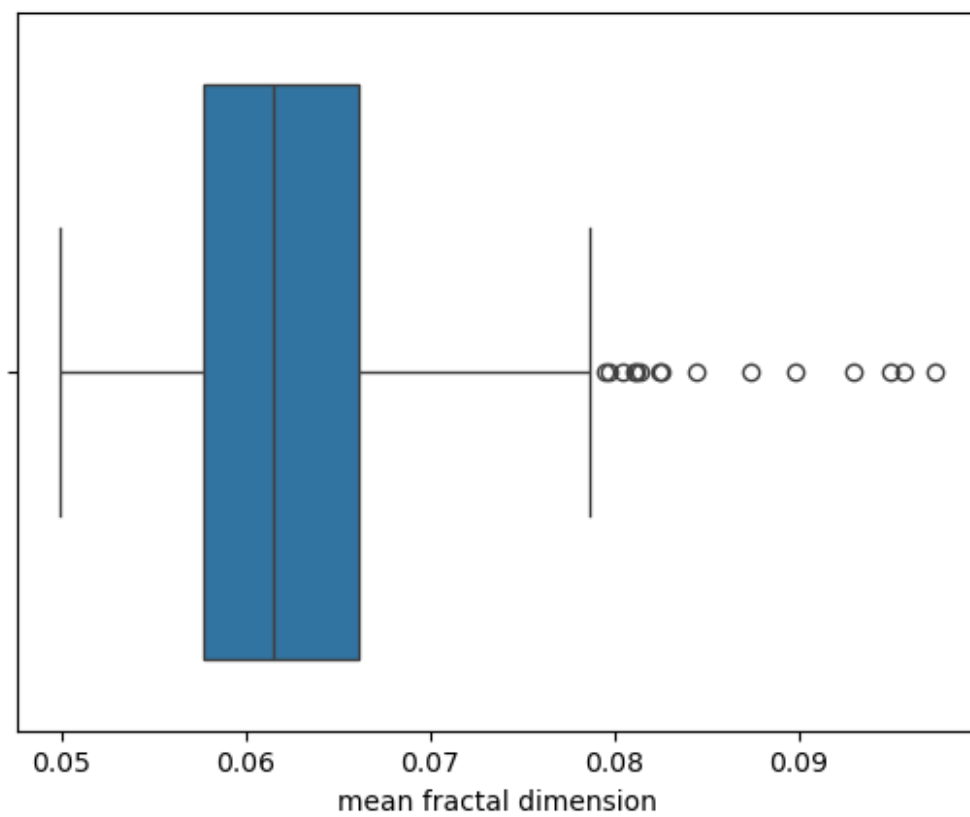
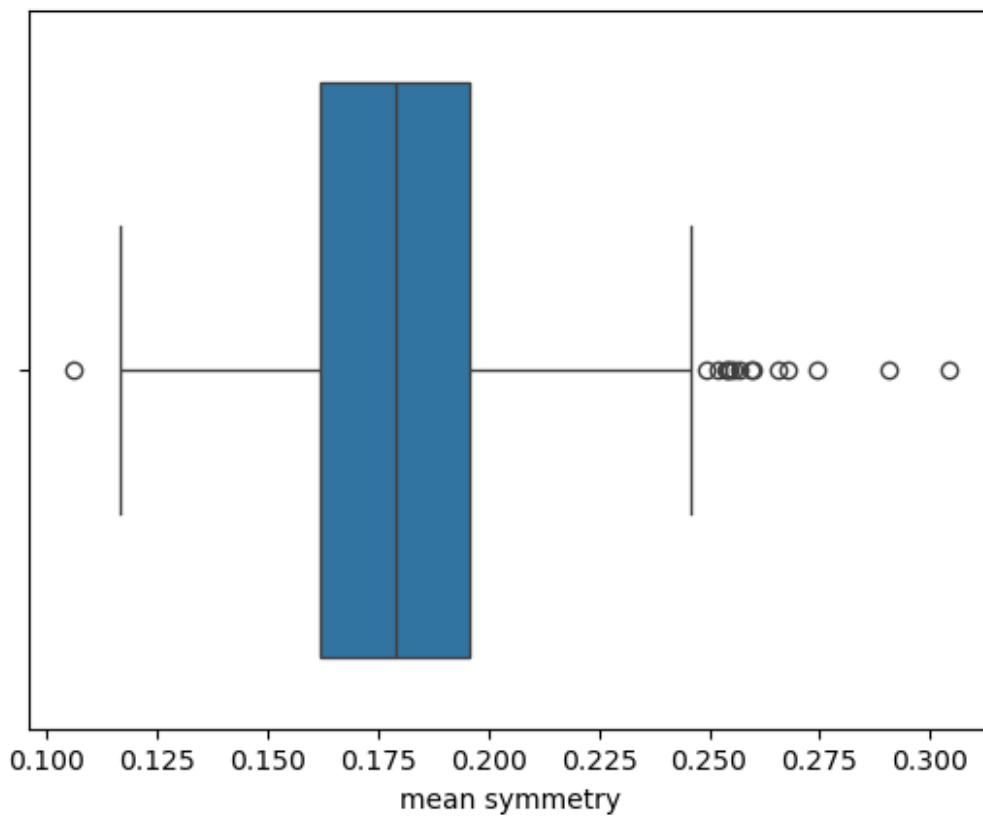


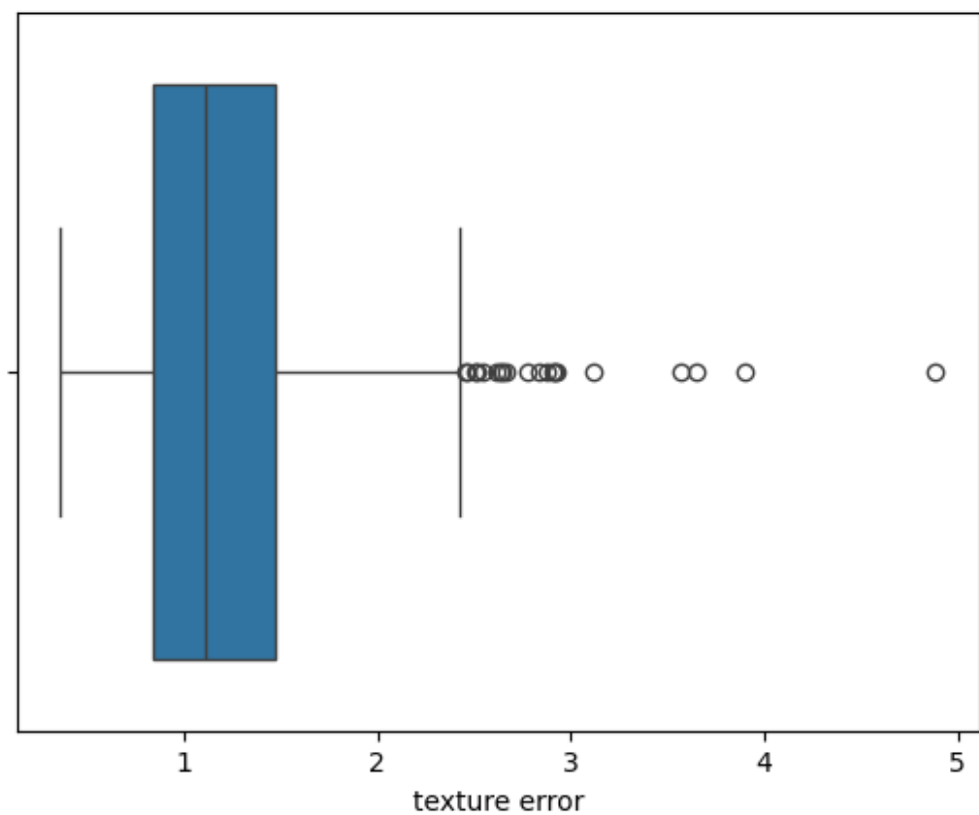
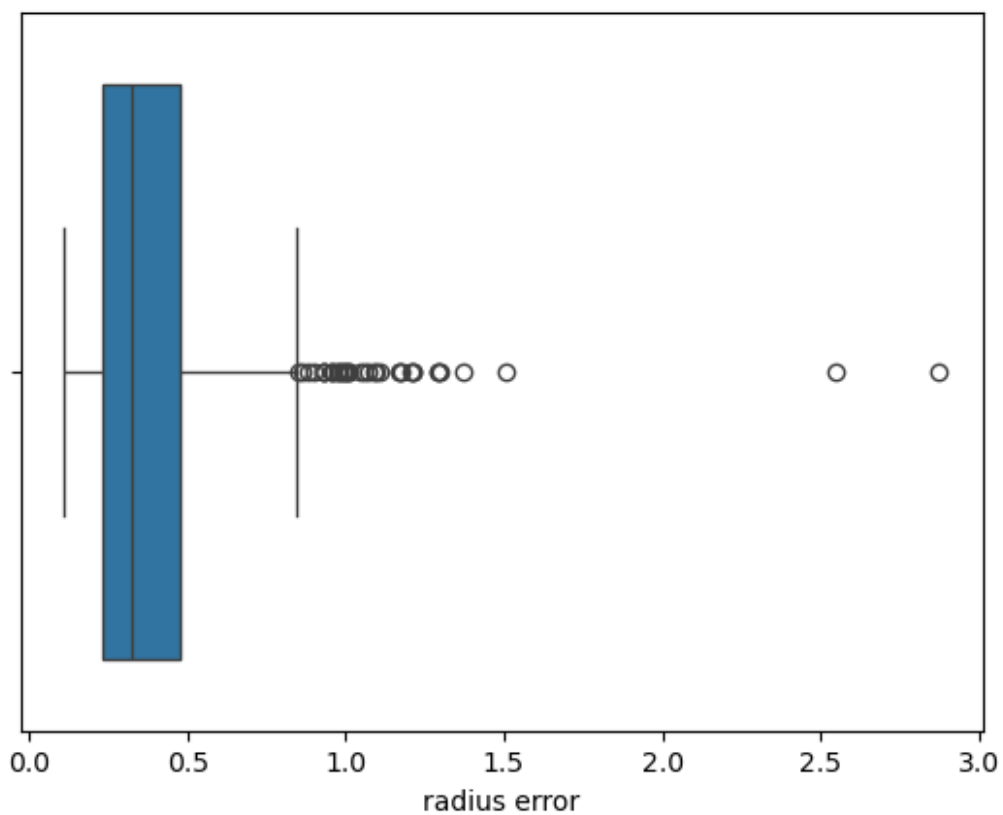


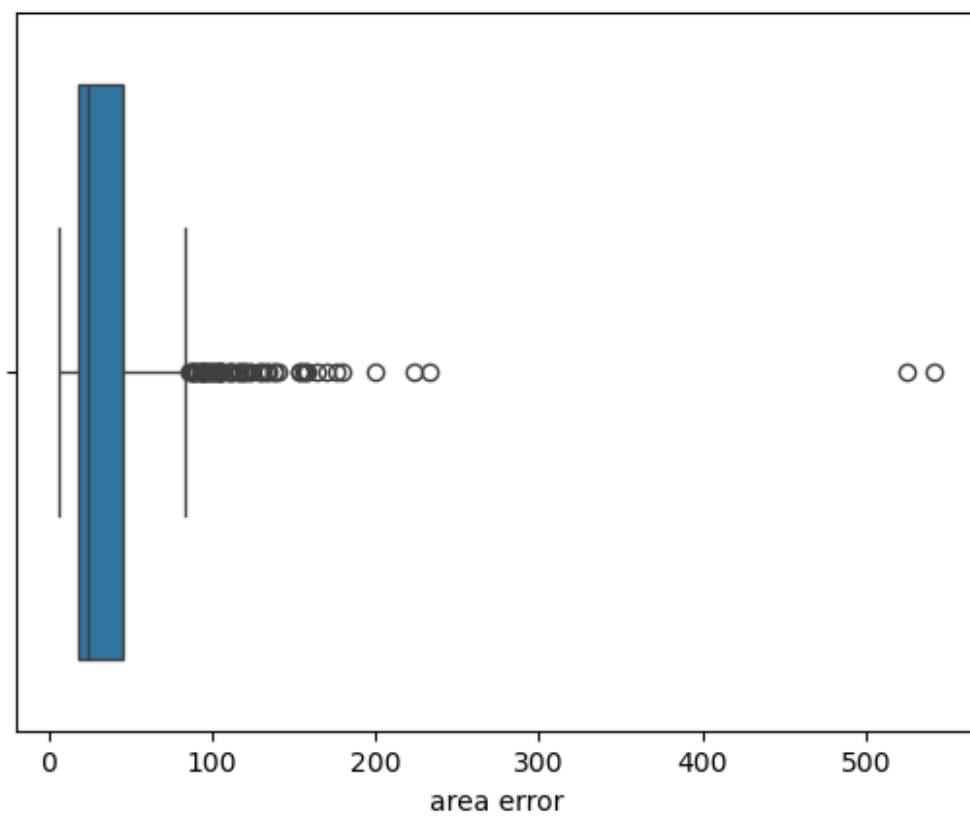
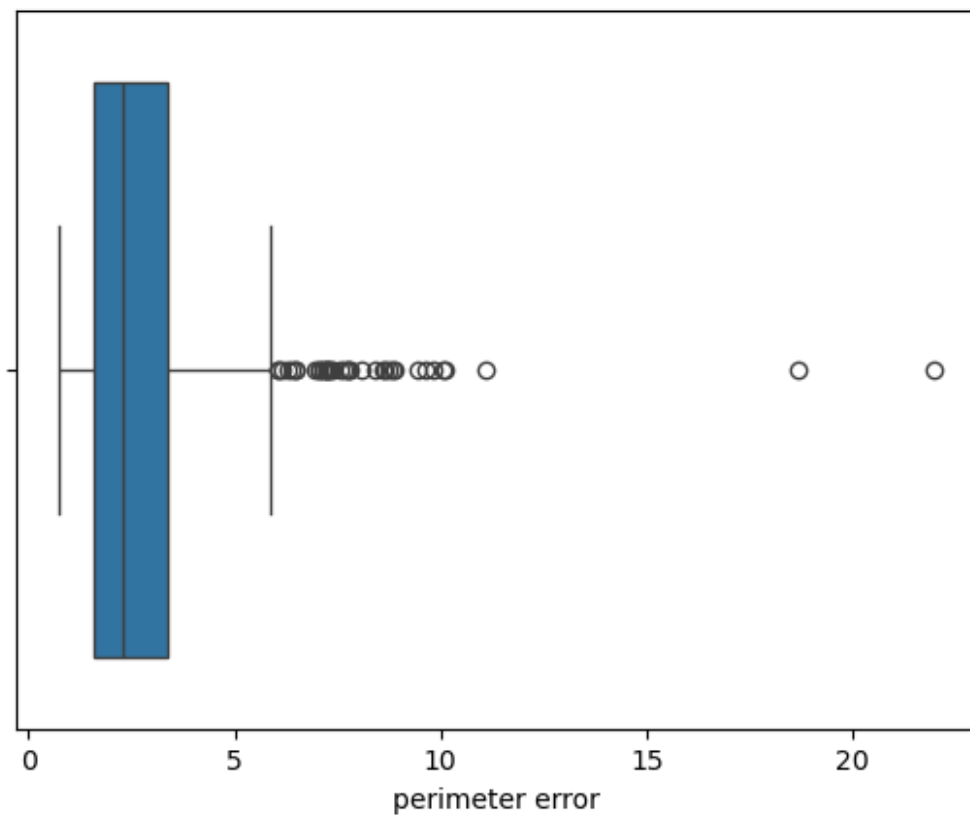


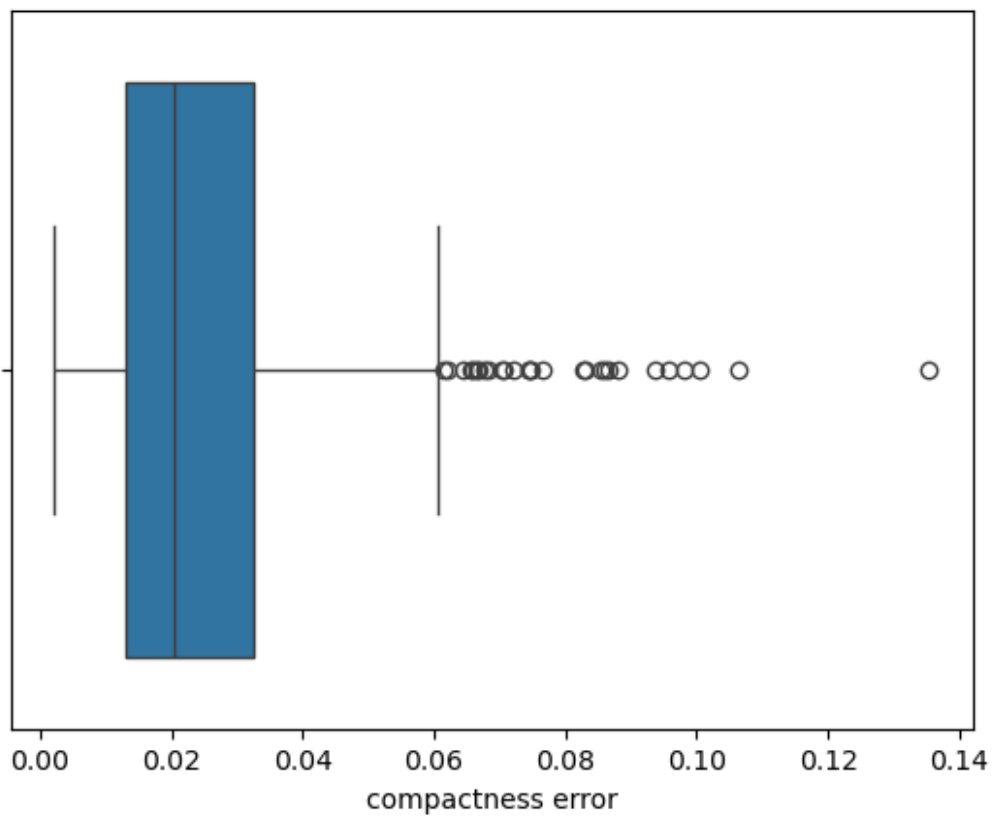
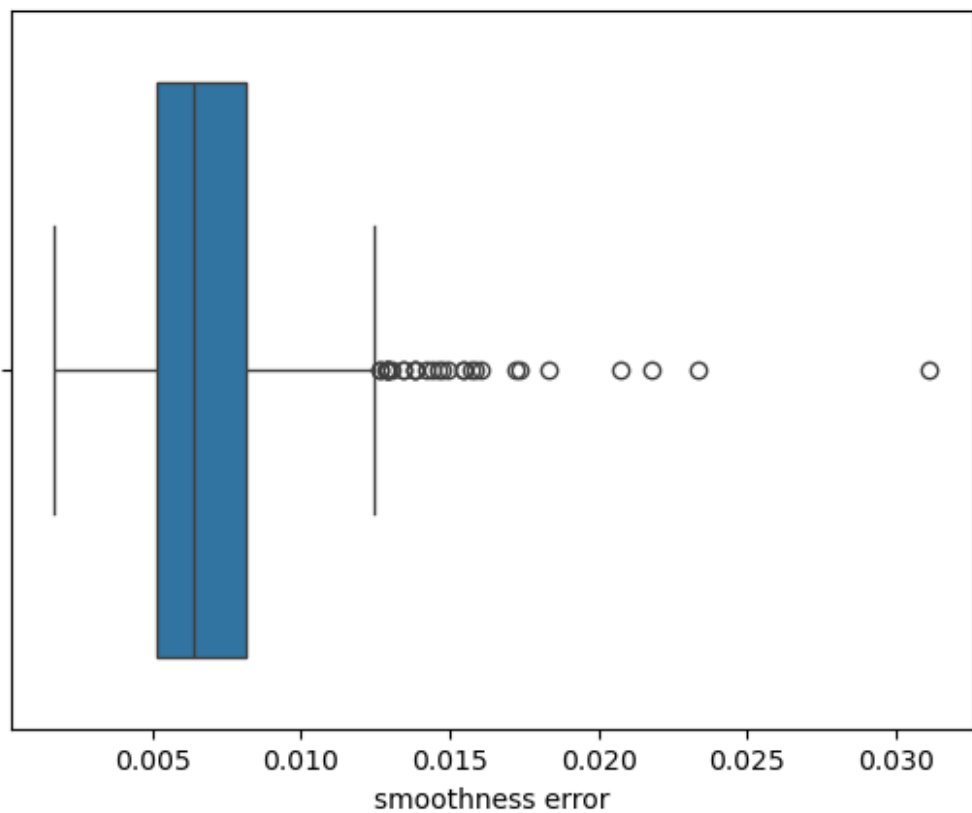


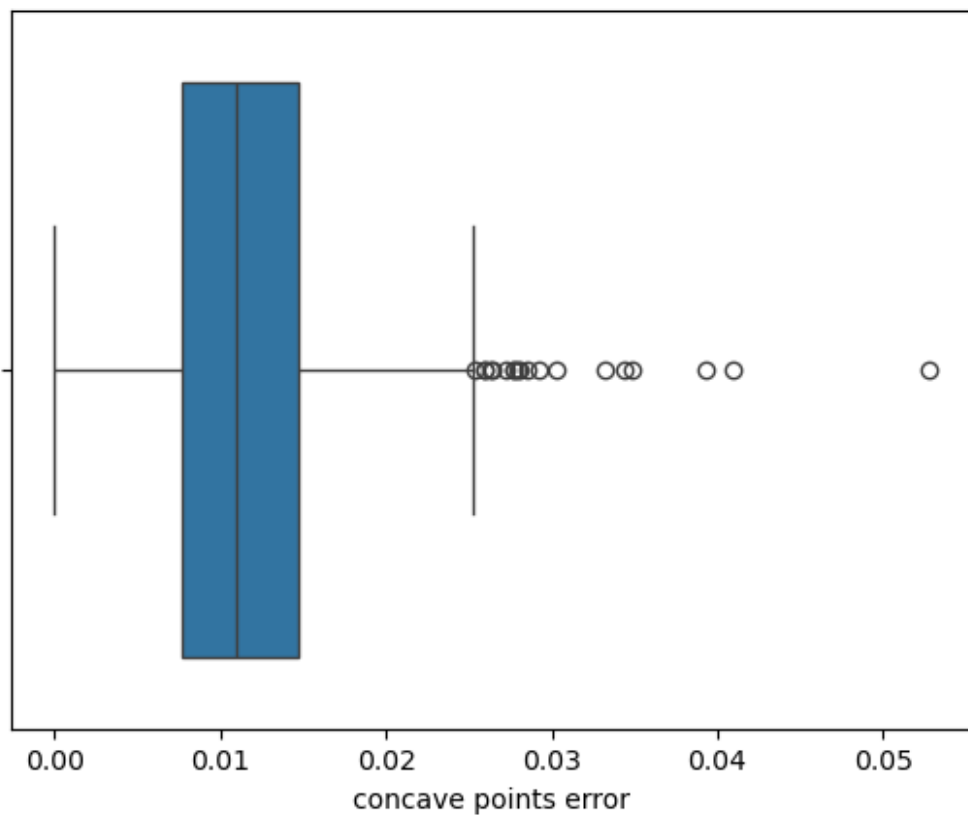
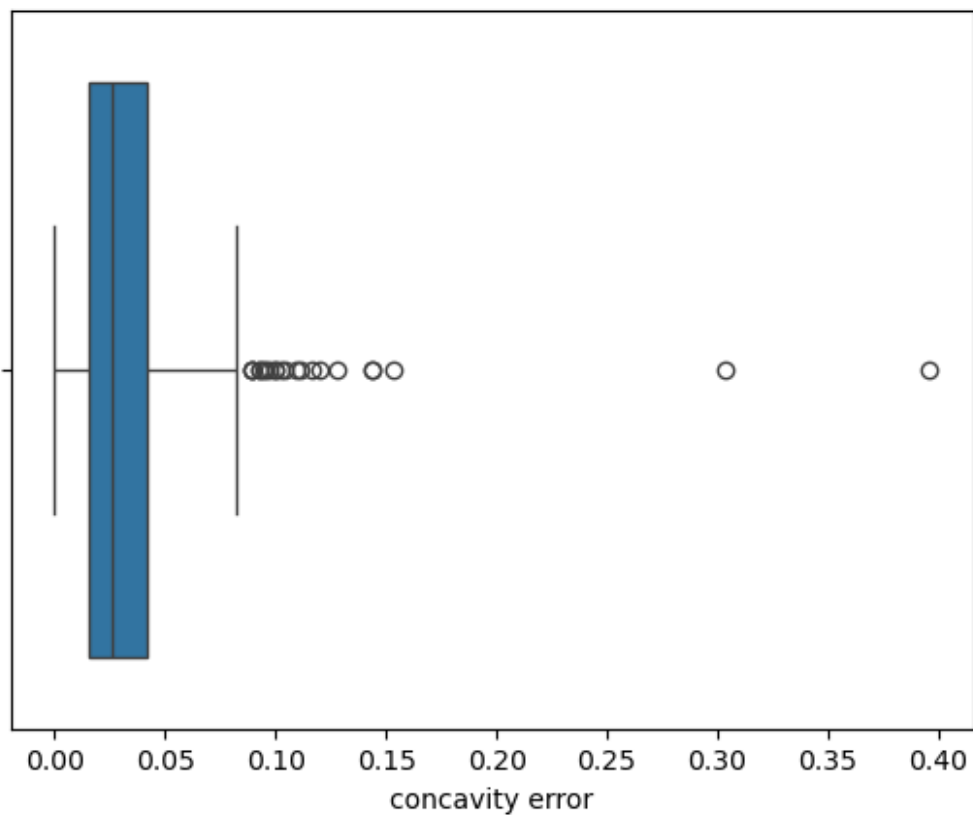


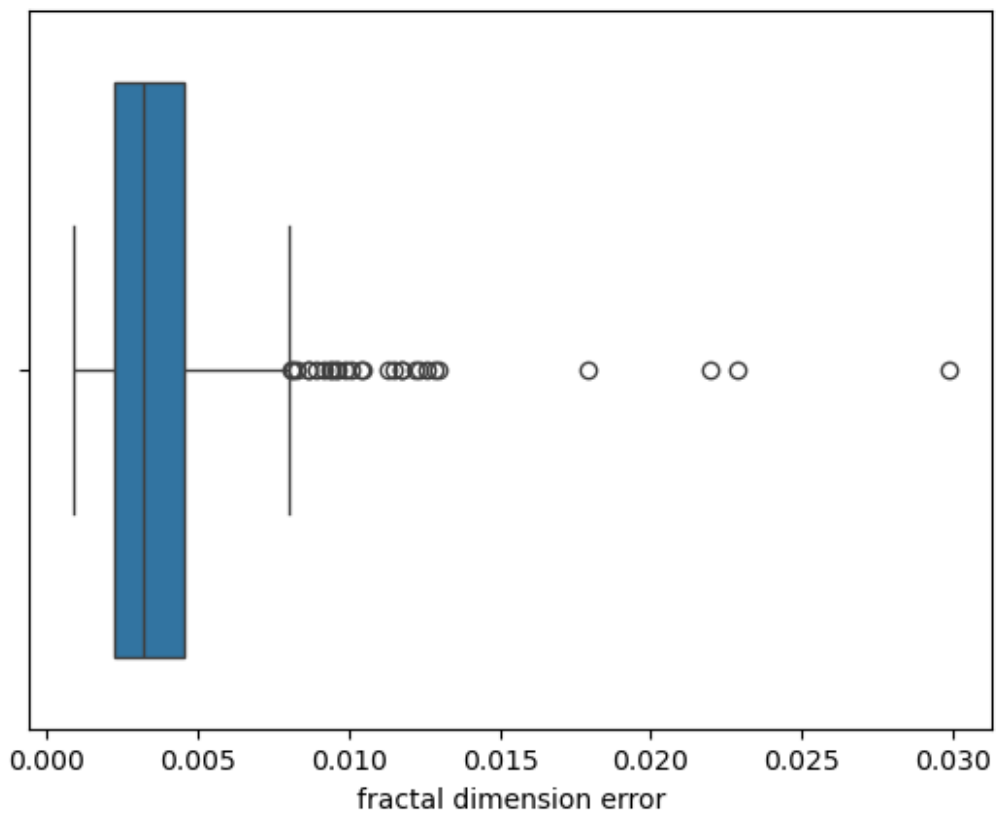
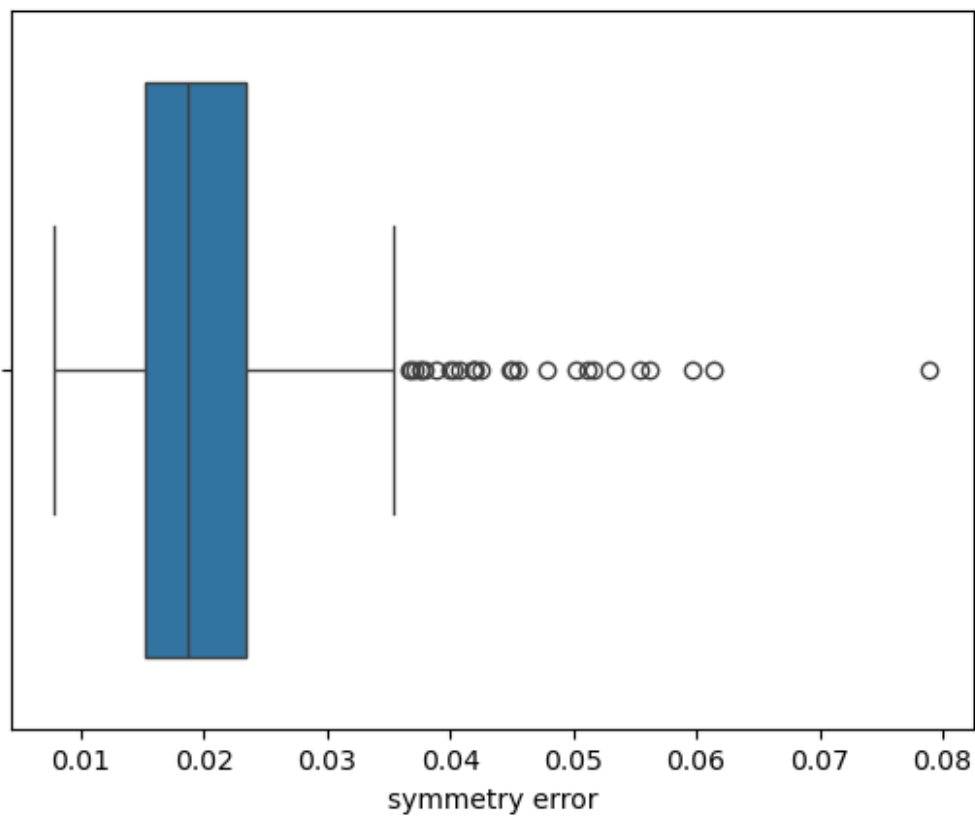


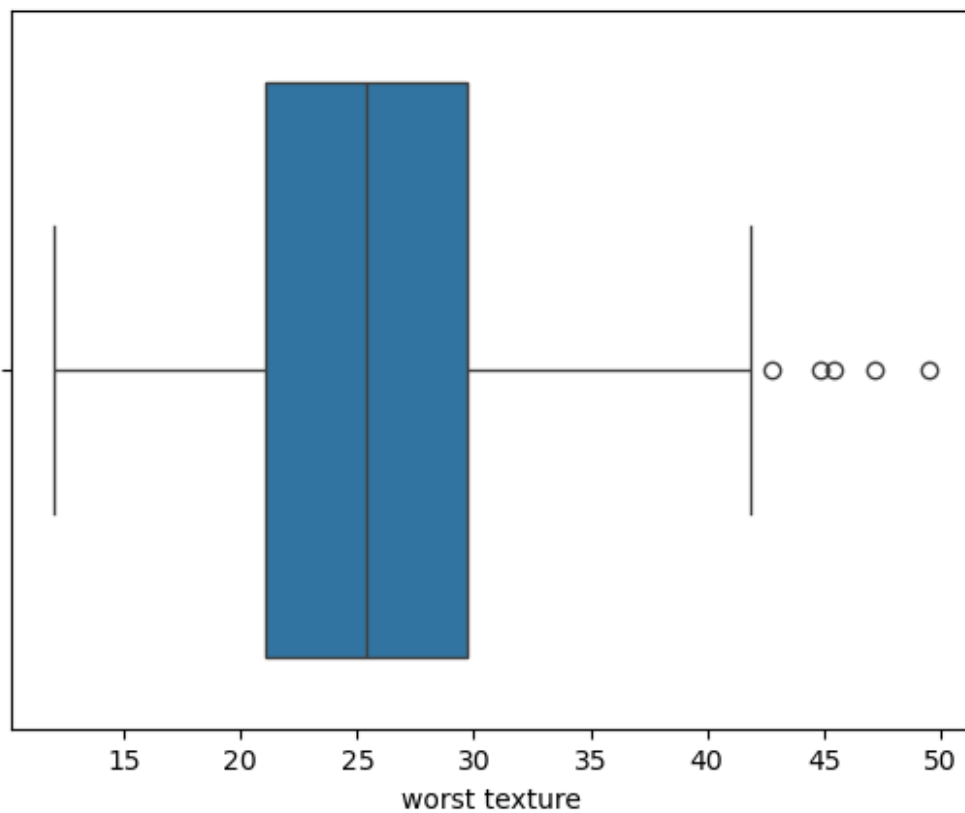
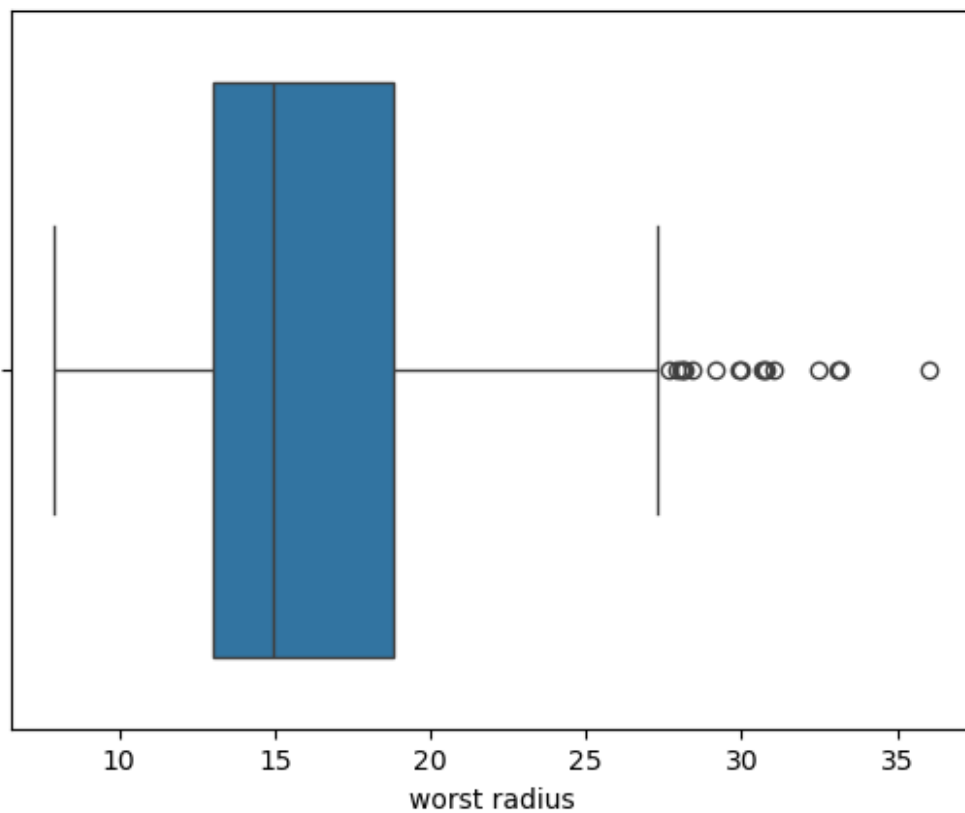


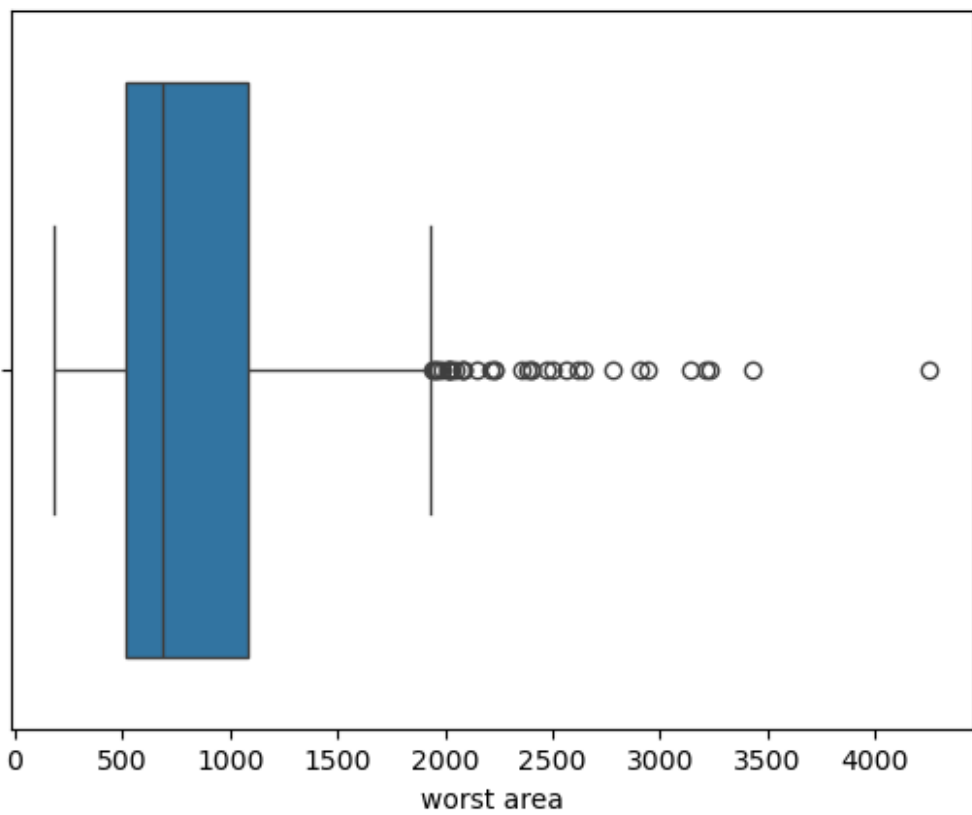
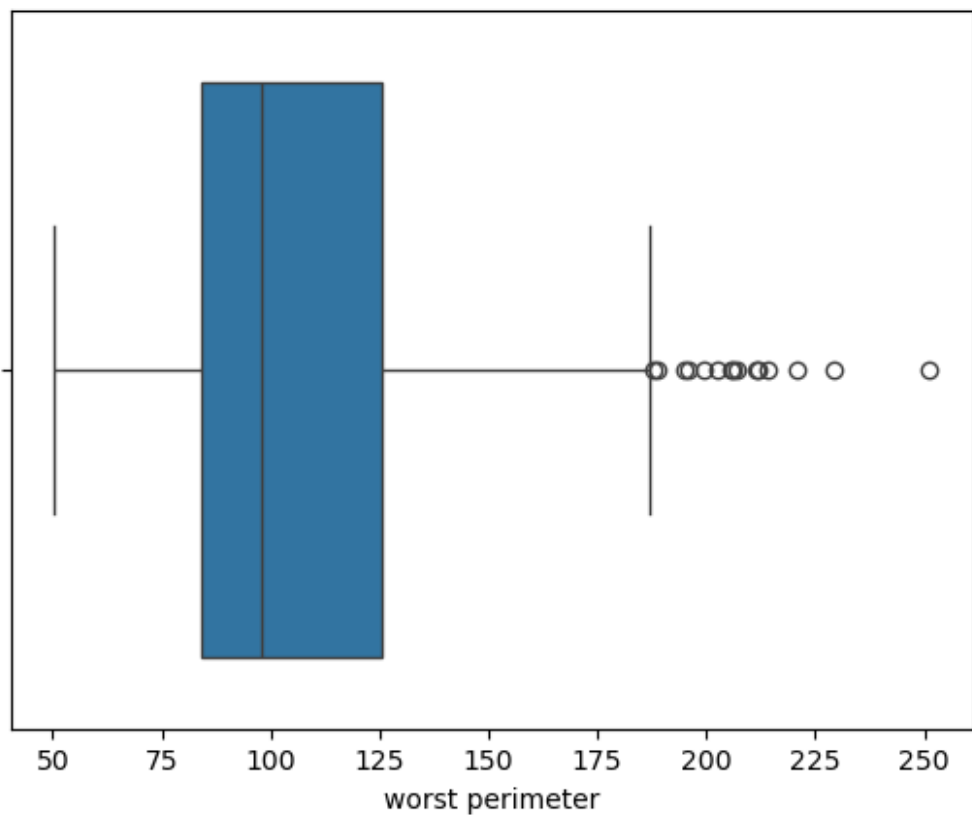




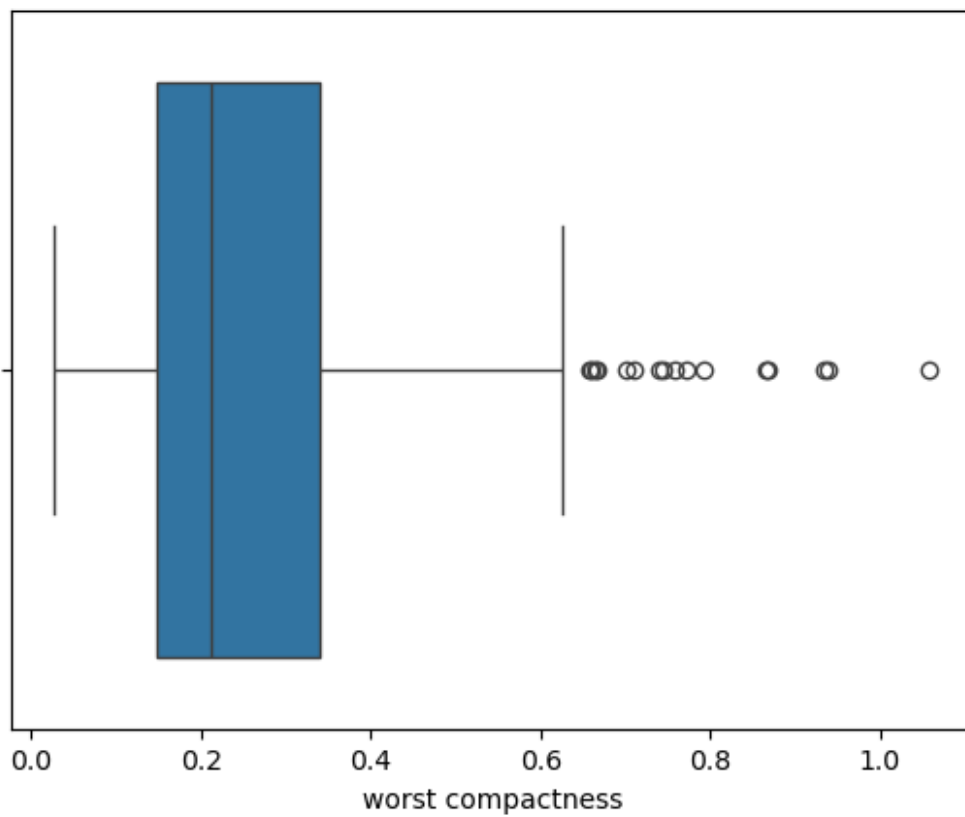
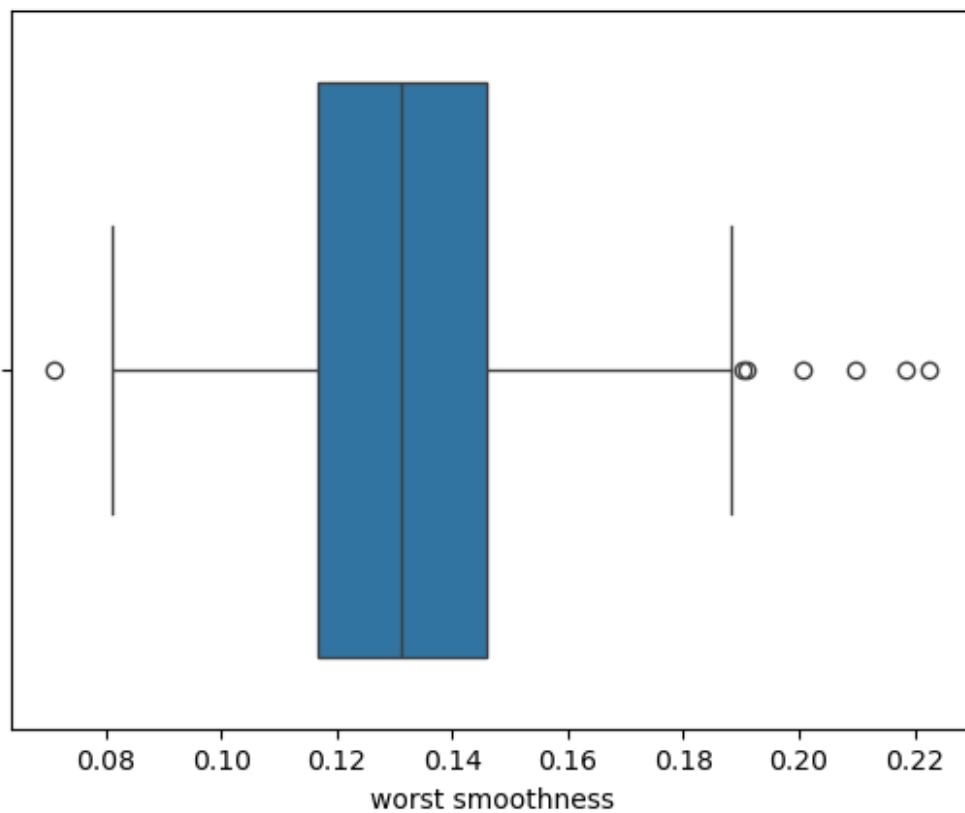


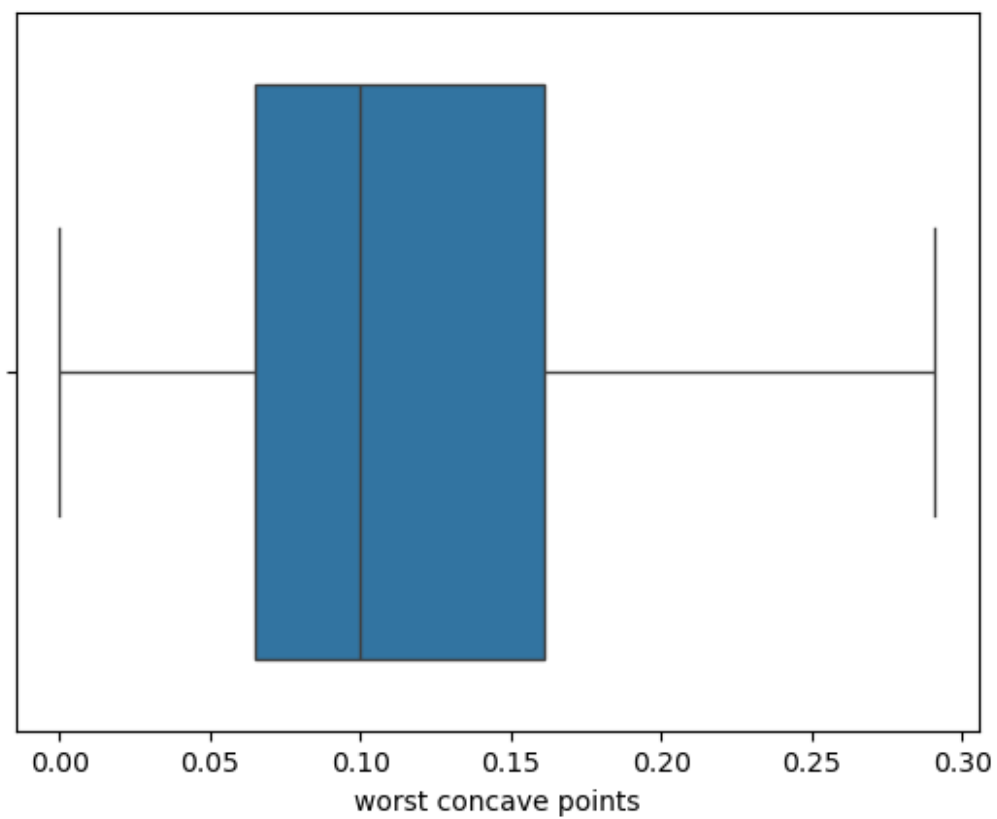
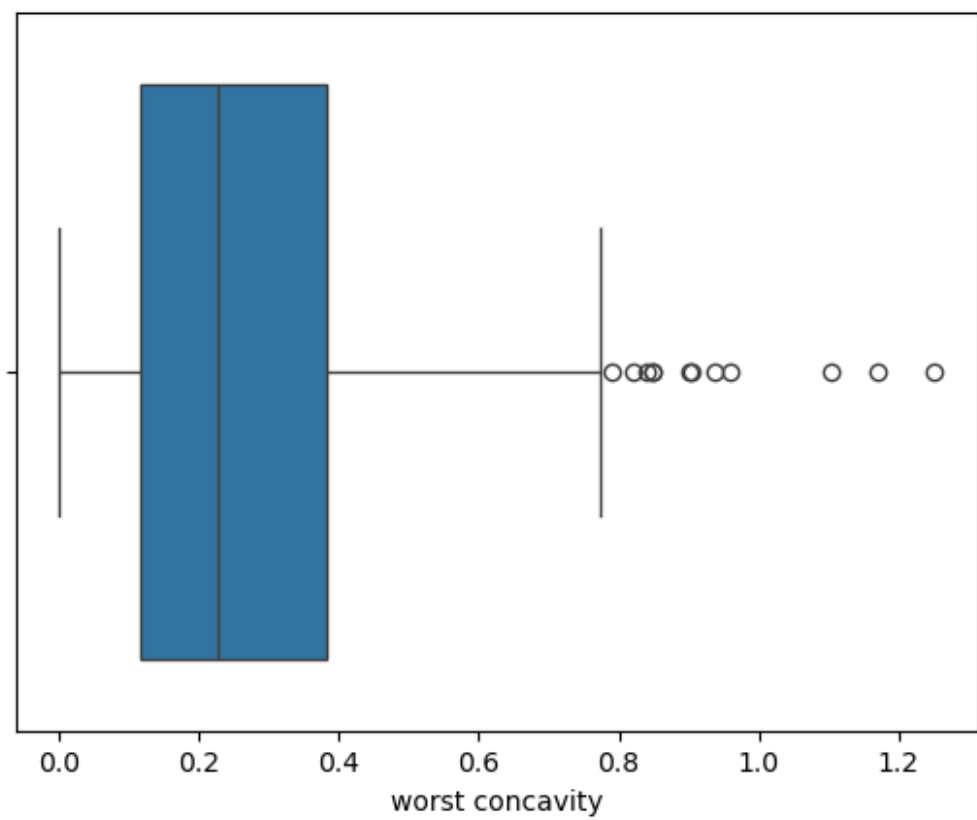


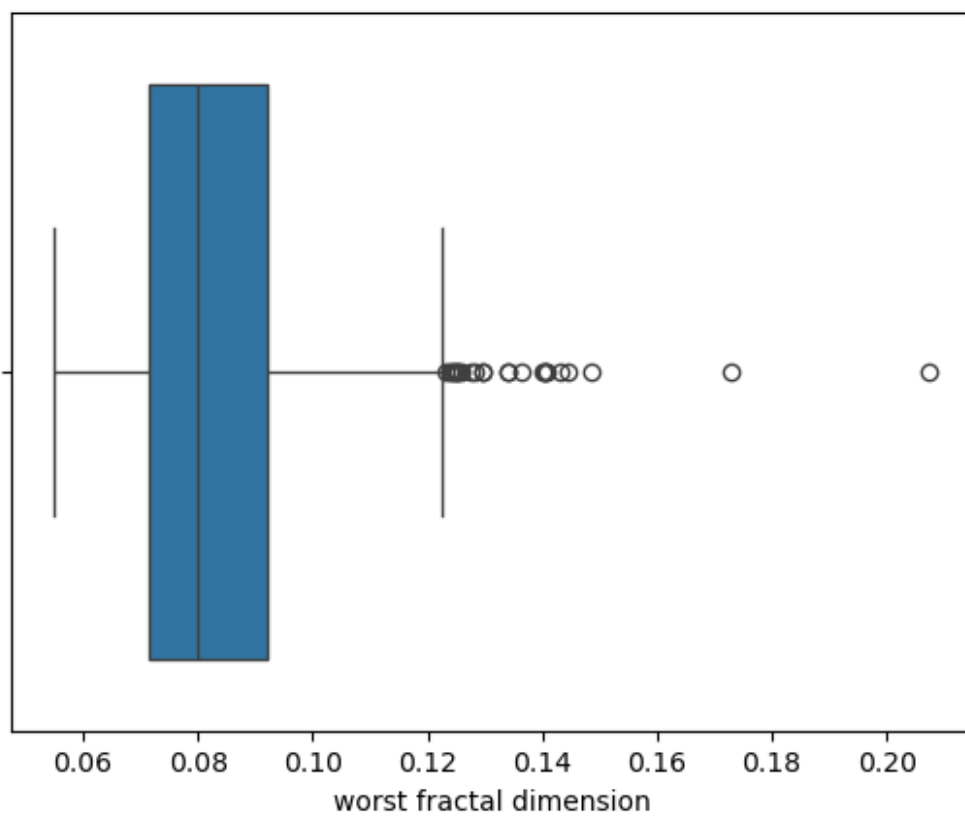
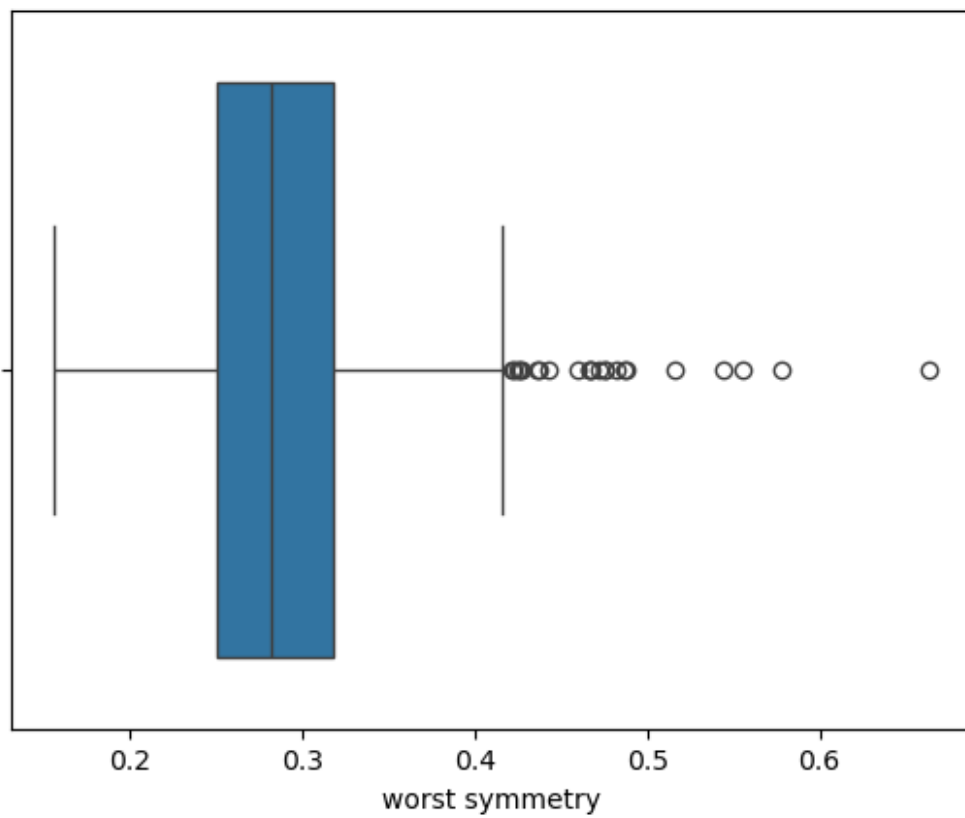


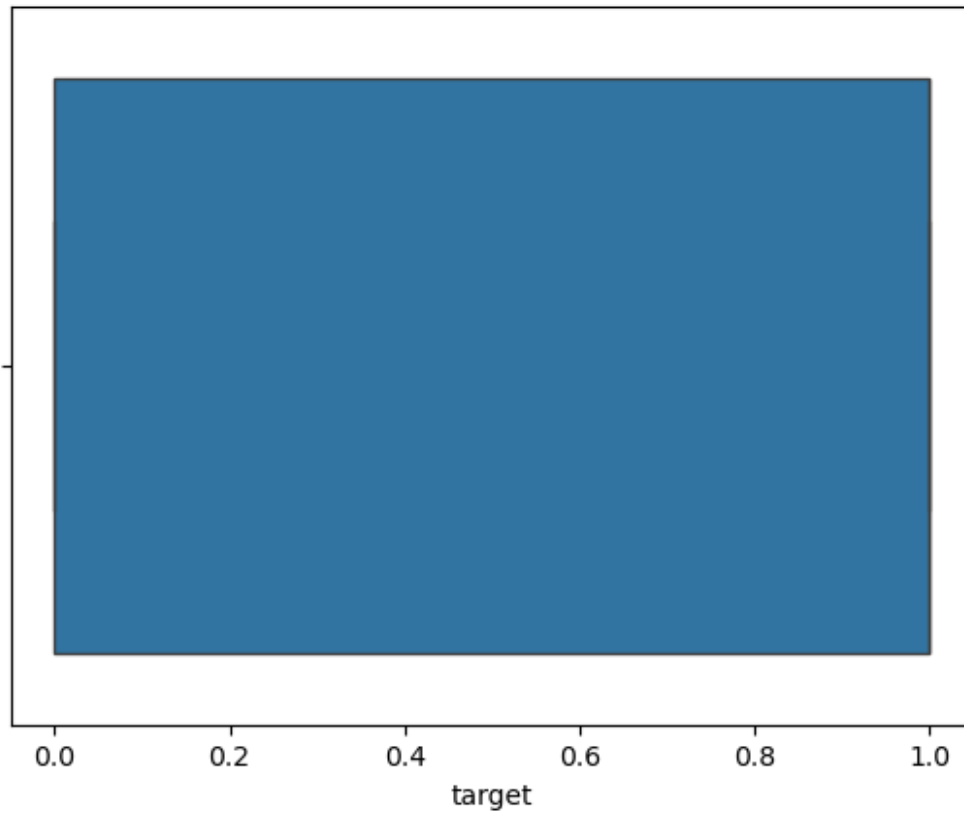








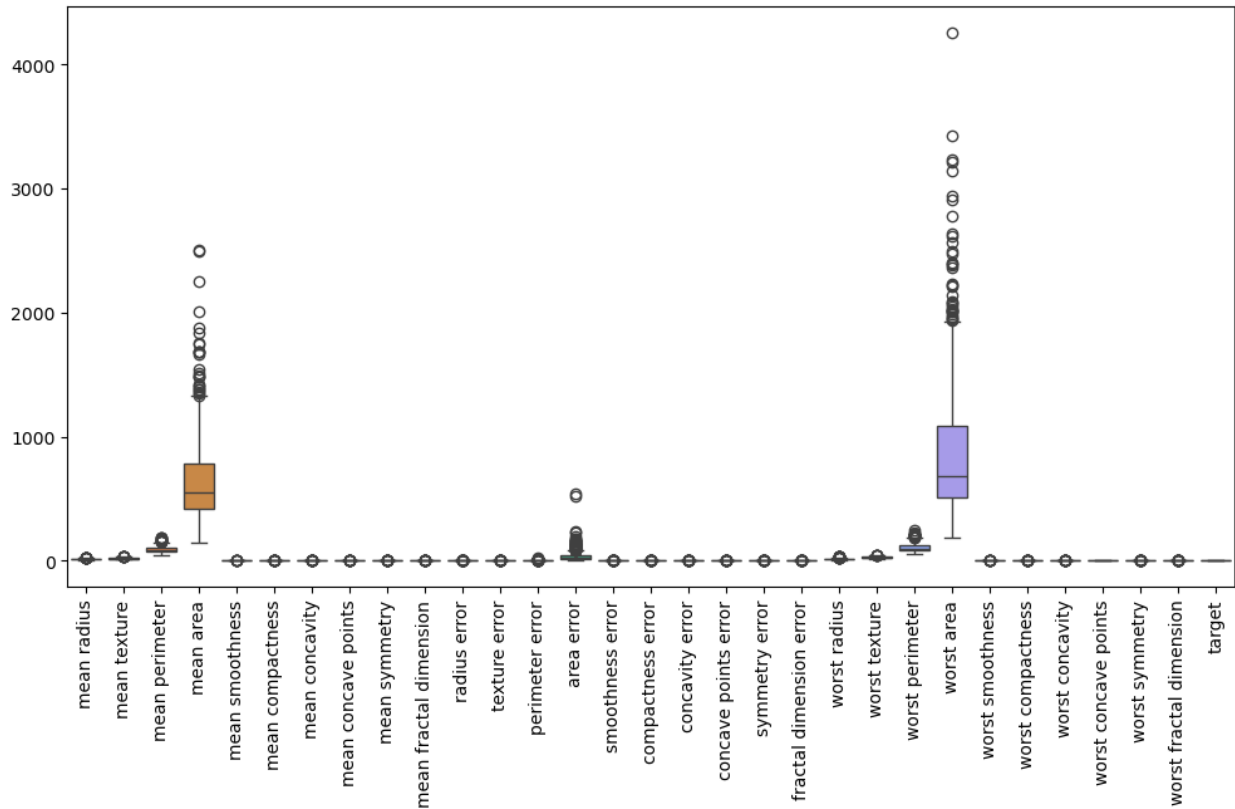




```
# comparing the outliers
value = df.columns
plt.figure(figsize=(12,6))
sns.boxplot(data = df[value])
plt.xticks(rotation=90)
```

```
([0,
 1,
 2,
 3,
 4,
 5,
 6,
 7,
 8,
 9,
10,
11,
12,
13,
14,
15,
16,
17,
```

```
18,  
19,  
20,  
21,  
22,  
23,  
24,  
25,  
26,  
27,  
28,  
29,  
30],  
[Text(0, 0, 'mean radius'),  
Text(1, 0, 'mean texture'),  
Text(2, 0, 'mean perimeter'),  
Text(3, 0, 'mean area'),  
Text(4, 0, 'mean smoothness'),  
Text(5, 0, 'mean compactness'),  
Text(6, 0, 'mean concavity'),  
Text(7, 0, 'mean concave points'),  
Text(8, 0, 'mean symmetry'),  
Text(9, 0, 'mean fractal dimension'),  
Text(10, 0, 'radius error'),  
Text(11, 0, 'texture error'),  
Text(12, 0, 'perimeter error'),  
Text(13, 0, 'area error'),  
Text(14, 0, 'smoothness error'),  
Text(15, 0, 'compactness error'),  
Text(16, 0, 'concavity error'),  
Text(17, 0, 'concave points error'),  
Text(18, 0, 'symmetry error'),  
Text(19, 0, 'fractal dimension error'),  
Text(20, 0, 'worst radius'),  
Text(21, 0, 'worst texture'),  
Text(22, 0, 'worst perimeter'),  
Text(23, 0, 'worst area'),  
Text(24, 0, 'worst smoothness'),  
Text(25, 0, 'worst compactness'),  
Text(26, 0, 'worst concavity'),  
Text(27, 0, 'worst concave points'),  
Text(28, 0, 'worst symmetry'),  
Text(29, 0, 'worst fractal dimension'),  
Text(30, 0, 'target')])
```



```
## Using IQR method to find outliers and capping it
columns_to_process = [col for col in df.columns if col != 'target']
#target doesn't have outliers
```

```
# Fix outliers using the IQR method
```

```
for col in columns_to_process:
    Q1 = df[col].quantile(0.25) # First quartile
    Q3 = df[col].quantile(0.75) # Third quartile
    IQR = Q3 - Q1                # Interquartile range
```

```
lower_limit = Q1 - 1.5 * IQR
upper_limit = Q3 + 1.5 * IQR
```

```
# Clip values outside the bounds
```

```
df[col] = df[col].clip(lower=lower_limit, upper=upper_limit)
```

```
df.head()
```

	mean radius	mean texture	mean perimeter	mean area	mean smoothness
0	17.99	10.38	122.80	1001.0	0.118400
1	20.57	17.77	132.90	1326.0	0.084740
2	19.69	21.25	130.00	1203.0	

0.109600

3	11.42	20.38	77.58	386.1
---	-------	-------	-------	-------

0.133695

4	20.29	14.34	135.10	1297.0
---	-------	-------	--------	--------

0.100300

	mean compactness	mean concavity	mean concave points	mean symmetry \
--	------------------	----------------	---------------------	-----------------

0	0.22862	0.28241	0.14710	
---	---------	---------	---------	--

0.2419

1	0.07864	0.08690	0.07017	
---	---------	---------	---------	--

0.1812

2	0.15990	0.19740	0.12790	
---	---------	---------	---------	--

0.2069

3	0.22862	0.24140	0.10520	
---	---------	---------	---------	--

0.2464

4	0.13280	0.19800	0.10430	
---	---------	---------	---------	--

0.1809

	mean fractal dimension	...	worst texture	worst perimeter	worst area \
--	------------------------	-----	---------------	-----------------	--------------

0	0.07871	...	17.33	184.60	
---	---------	-----	-------	--------	--

1937.05

1	0.05667	...	23.41	158.80	
---	---------	-----	-------	--------	--

1937.05

2	0.05999	...	25.53	152.50	
---	---------	-----	-------	--------	--

1709.00

3	0.07875	...	26.50	98.87	
---	---------	-----	-------	-------	--

567.70

4	0.05883	...	16.67	152.20	
---	---------	-----	-------	--------	--

1575.00

	worst smoothness	worst compactness	worst concavity	worst concave points \
--	------------------	-------------------	-----------------	------------------------

0	0.1622	0.62695	0.7119	
---	--------	---------	--------	--

0.2654

1	0.1238	0.18660	0.2416	
---	--------	---------	--------	--

0.1860

2	0.1444	0.42450	0.4504	
---	--------	---------	--------	--

0.2430

3	0.1901	0.62695	0.6869	
---	--------	---------	--------	--

0.2575

4	0.1374	0.20500	0.4000	
---	--------	---------	--------	--

0.1625

	worst symmetry	worst fractal dimension	target
--	----------------	-------------------------	--------

0	0.41915	0.11890	0
---	---------	---------	---

1	0.27500	0.08902	0
---	---------	---------	---

2	0.36130	0.08758	0
---	---------	---------	---

3	0.41915	0.12301	0
---	---------	---------	---

4	0.23640	0.07678	0
---	---------	---------	---

[5 rows x 31 columns]

*# visualising outliers after fixing*

value = df.columns

plt.figure(figsize=(12,6))

sns.boxplot(data = df[value])

plt.xticks(rotation=90)

([0,  
1,  
2,  
3,  
4,  
5,  
6,  
7,  
8,  
9,  
10,  
11,  
12,  
13,  
14,  
15,  
16,  
17,  
18,  
19,  
20,  
21,  
22,  
23,  
24,  
25,  
26,  
27,  
28,  
29,  
30],

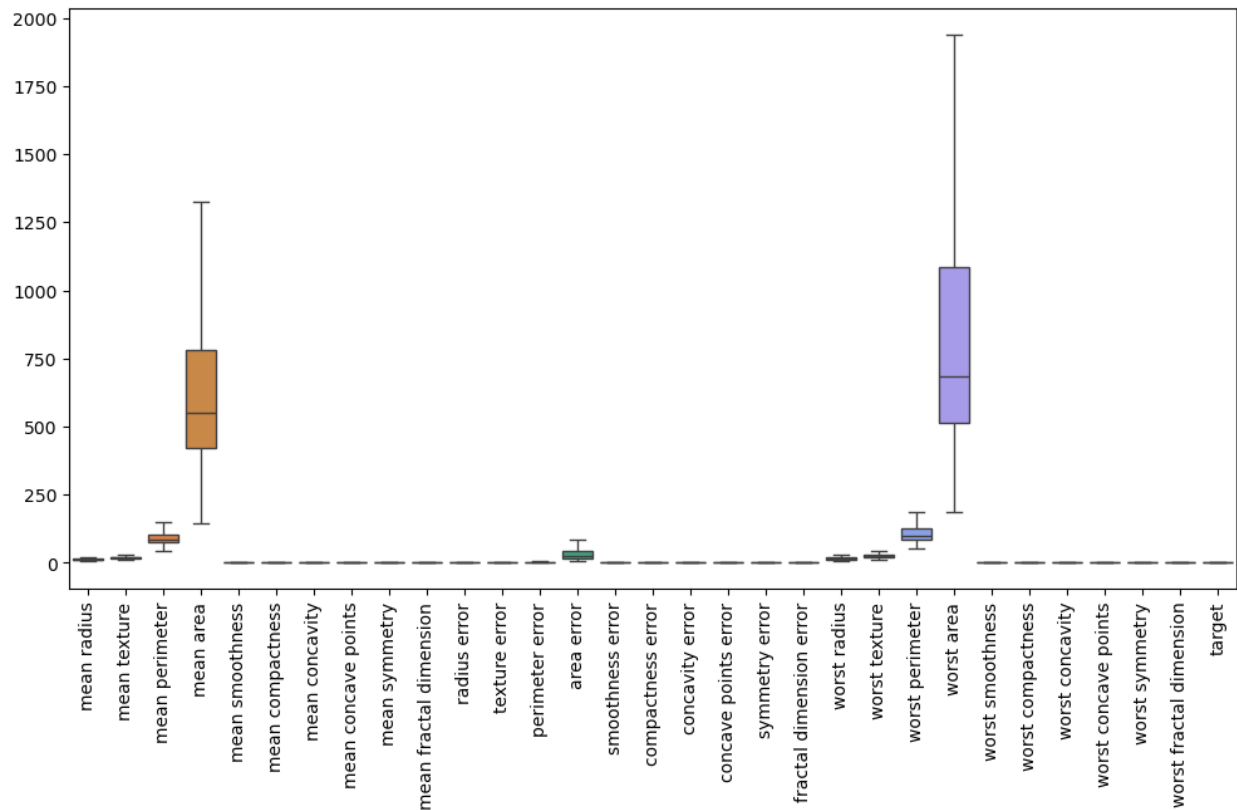
[Text(0, 0, 'mean radius'),  
Text(1, 0, 'mean texture'),  
Text(2, 0, 'mean perimeter'),  
Text(3, 0, 'mean area'),  
Text(4, 0, 'mean smoothness'),  
Text(5, 0, 'mean compactness'),  
Text(6, 0, 'mean concavity'),  
Text(7, 0, 'mean concave points'),  
Text(8, 0, 'mean symmetry'),



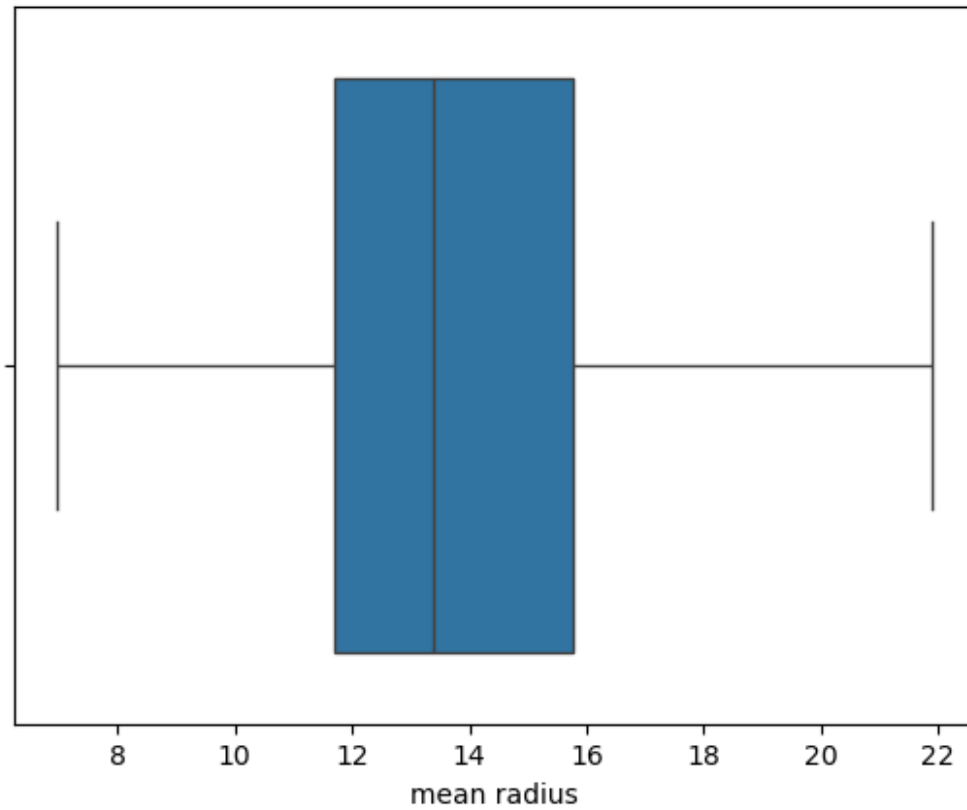
```

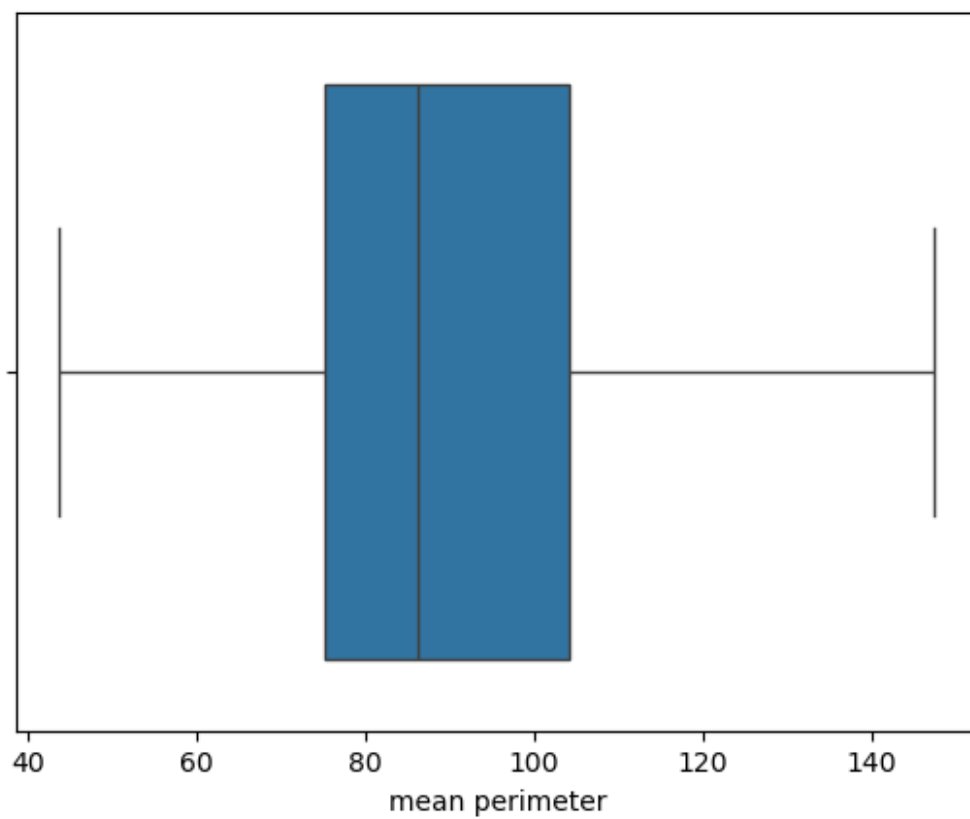
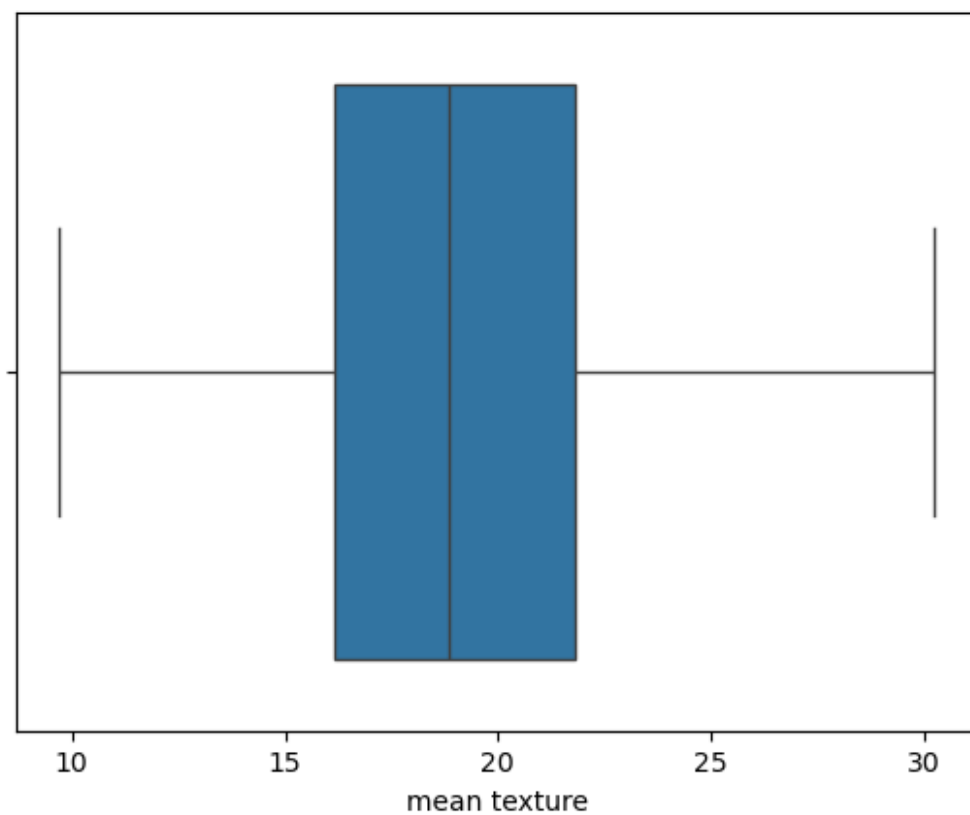
Text(9, 0, 'mean fractal dimension'),
Text(10, 0, 'radius error'),
Text(11, 0, 'texture error'),
Text(12, 0, 'perimeter error'),
Text(13, 0, 'area error'),
Text(14, 0, 'smoothness error'),
Text(15, 0, 'compactness error'),
Text(16, 0, 'concavity error'),
Text(17, 0, 'concave points error'),
Text(18, 0, 'symmetry error'),
Text(19, 0, 'fractal dimension error'),
Text(20, 0, 'worst radius'),
Text(21, 0, 'worst texture'),
Text(22, 0, 'worst perimeter'),
Text(23, 0, 'worst area'),
Text(24, 0, 'worst smoothness'),
Text(25, 0, 'worst compactness'),
Text(26, 0, 'worst concavity'),
Text(27, 0, 'worst concave points'),
Text(28, 0, 'worst symmetry'),
Text(29, 0, 'worst fractal dimension'),
Text(30, 0, 'target']]

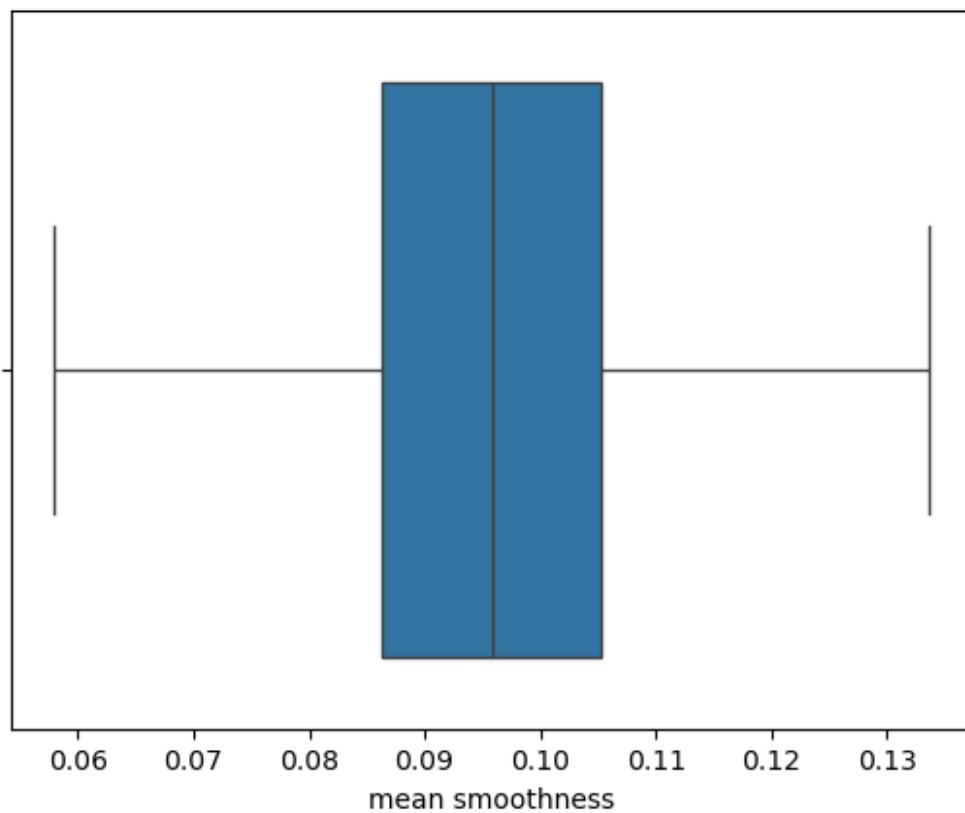
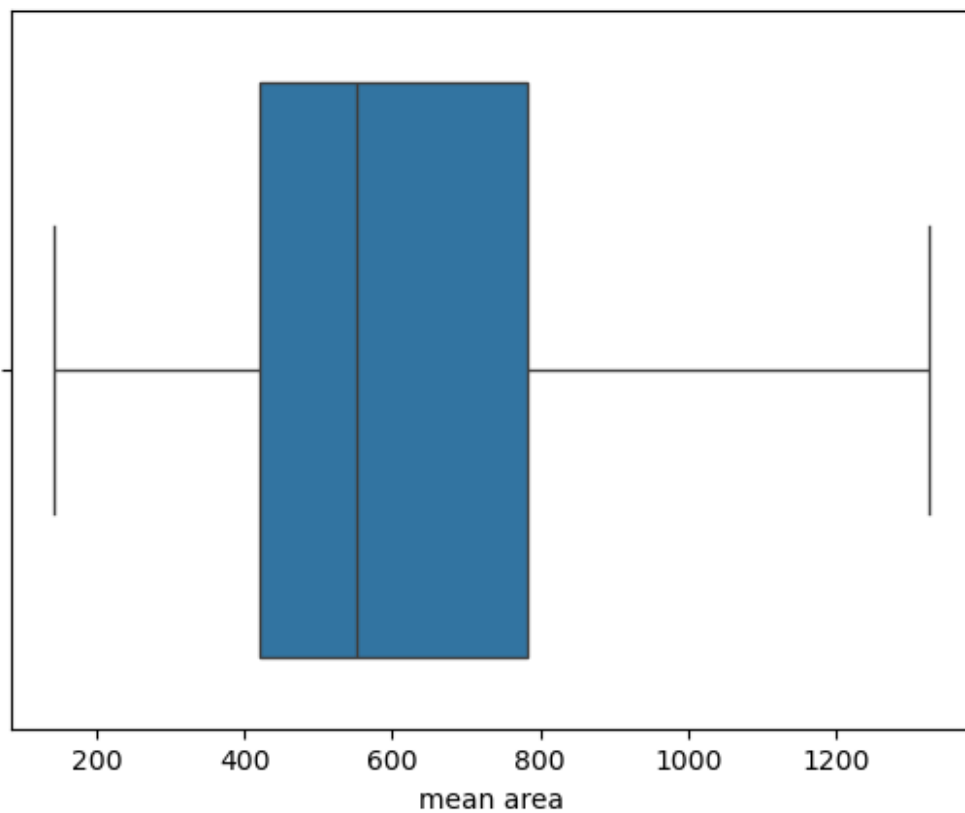
```

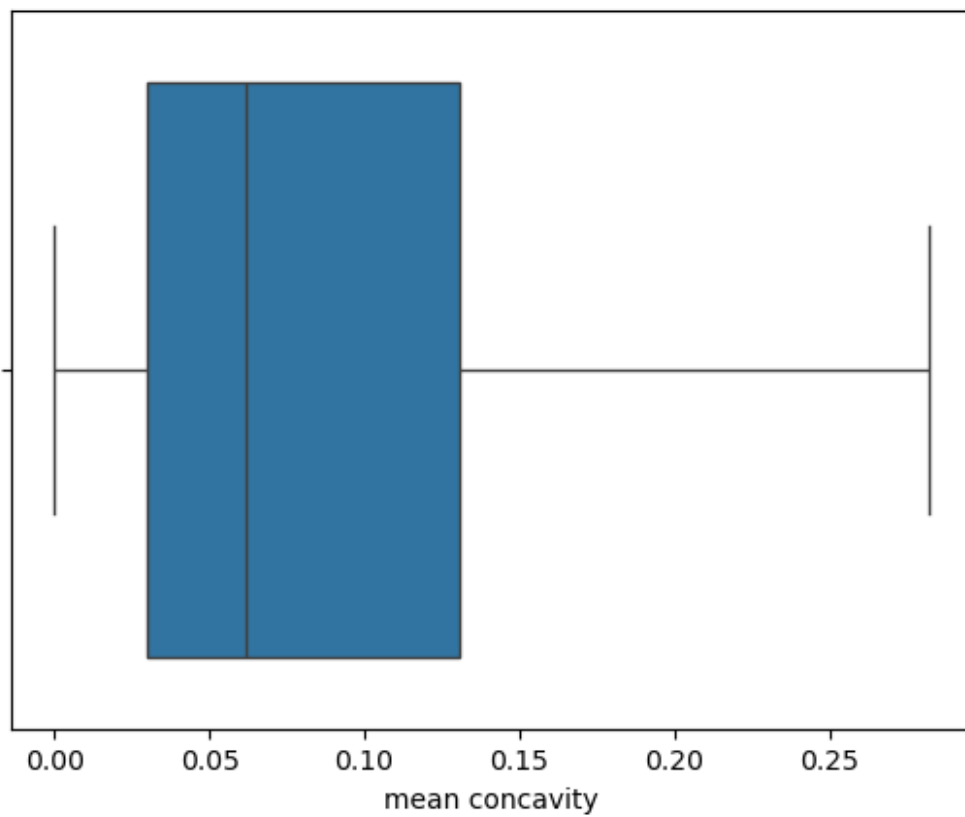
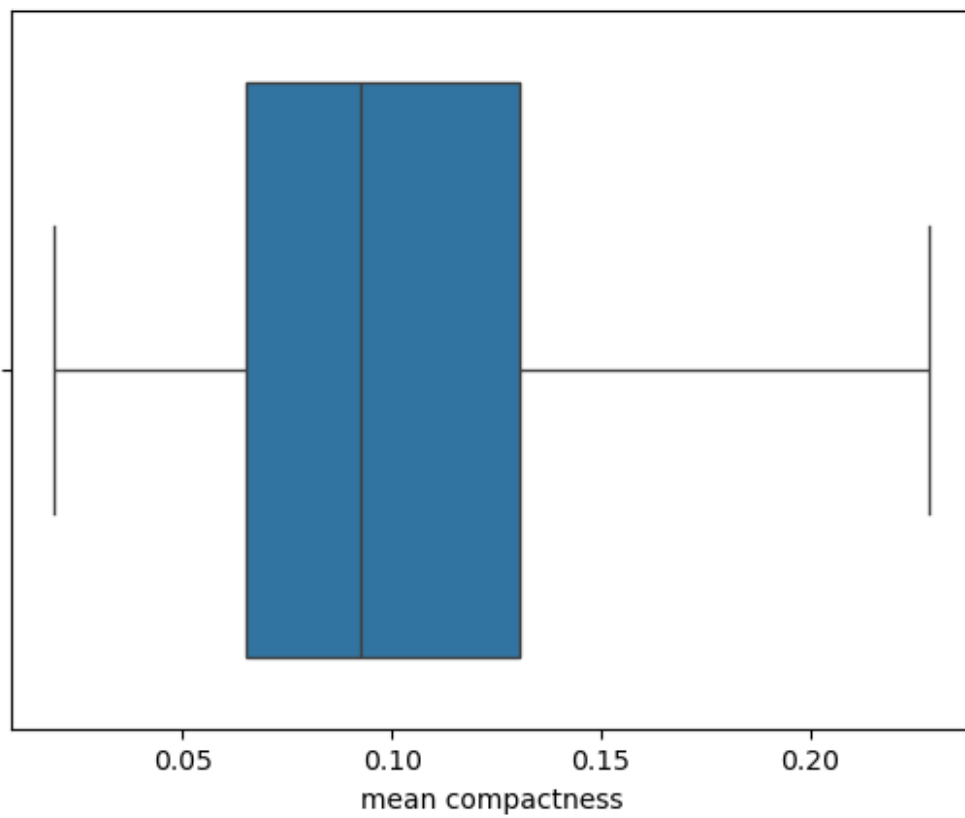


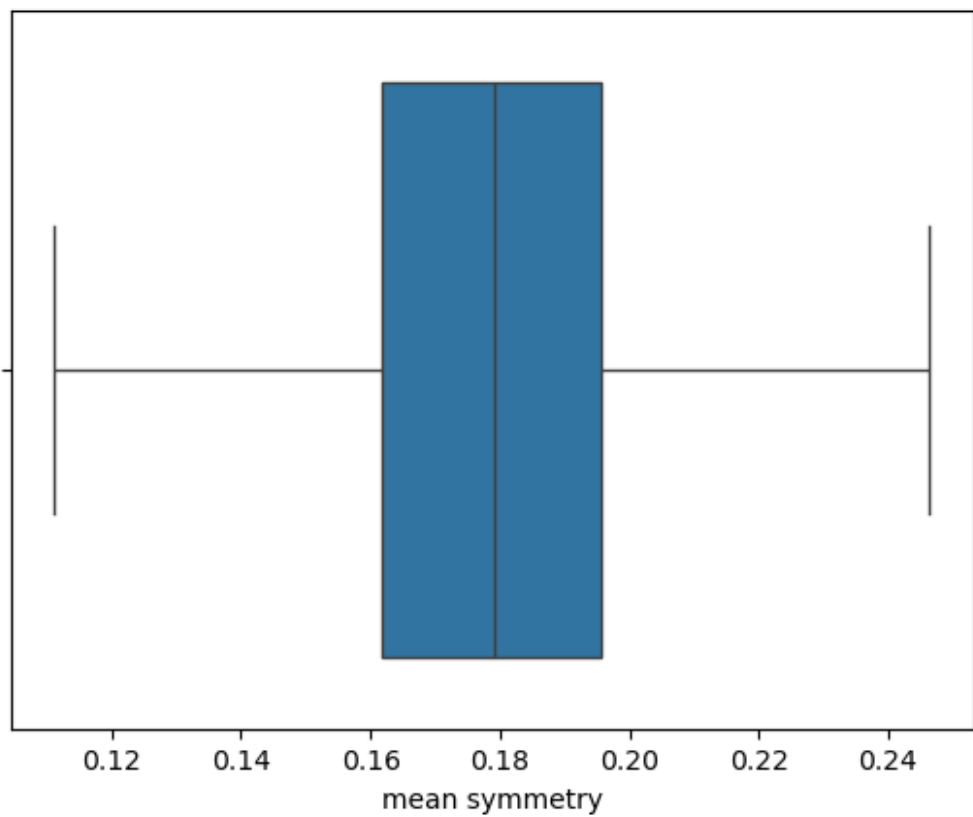
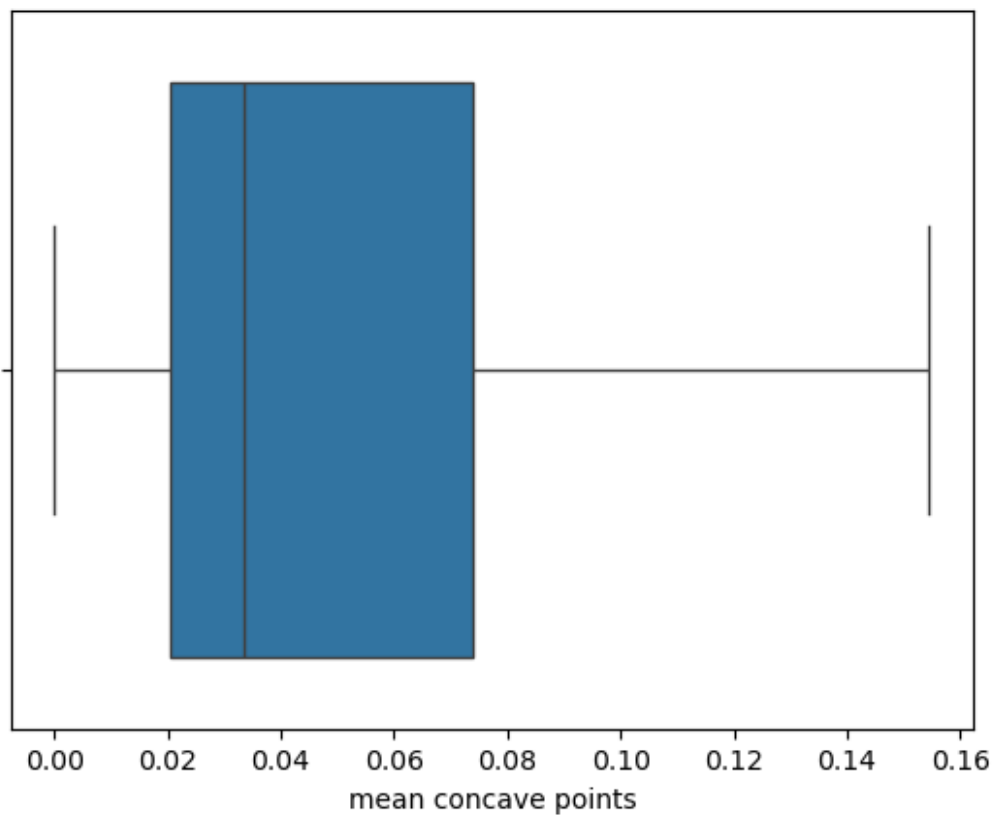
```
# Box plot for each column after fixing outliers
for i in df.columns:
    sns.boxplot(data=df, x=i)
    plt.show()
```

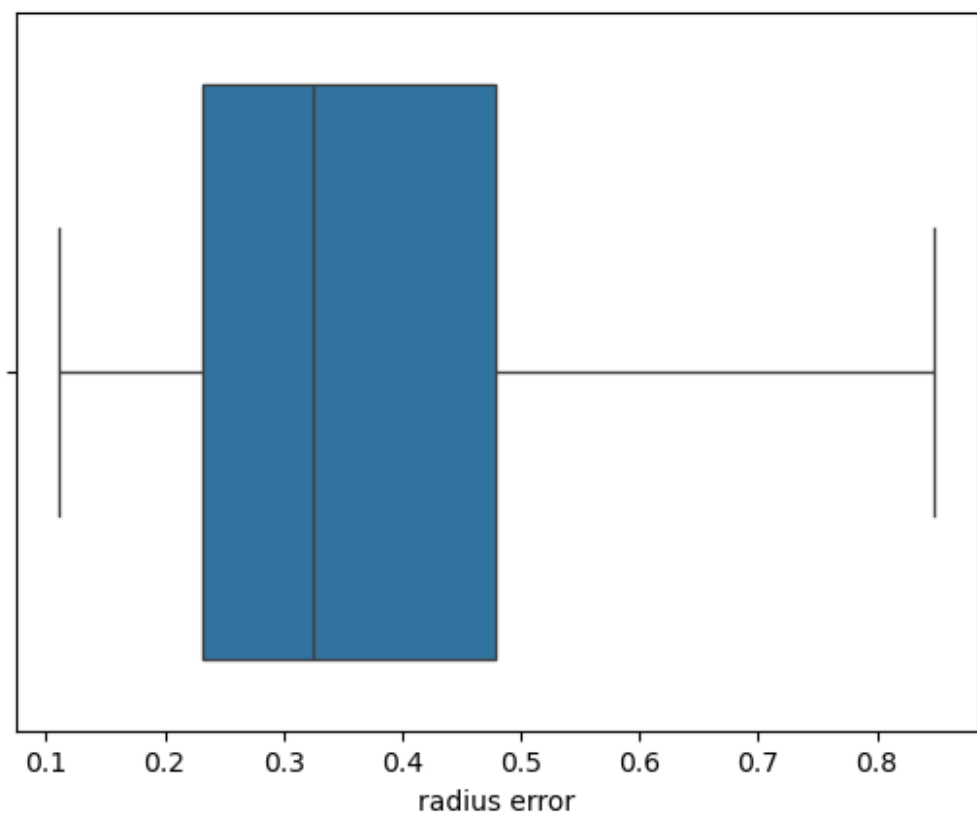
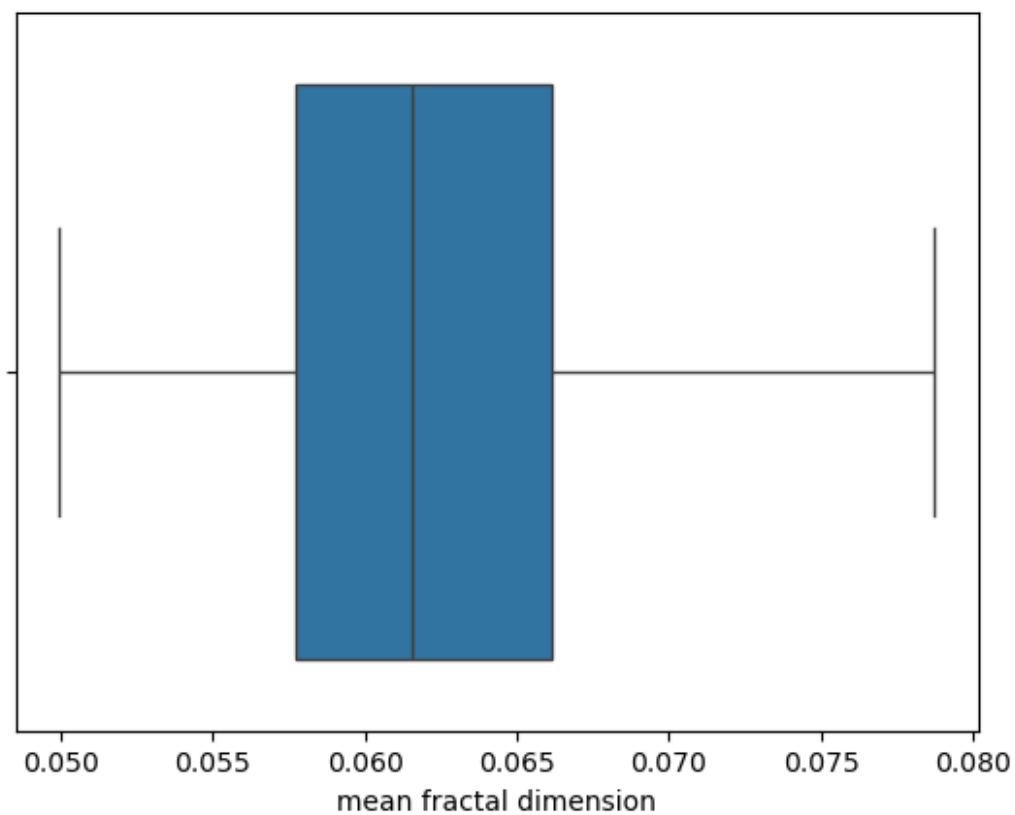


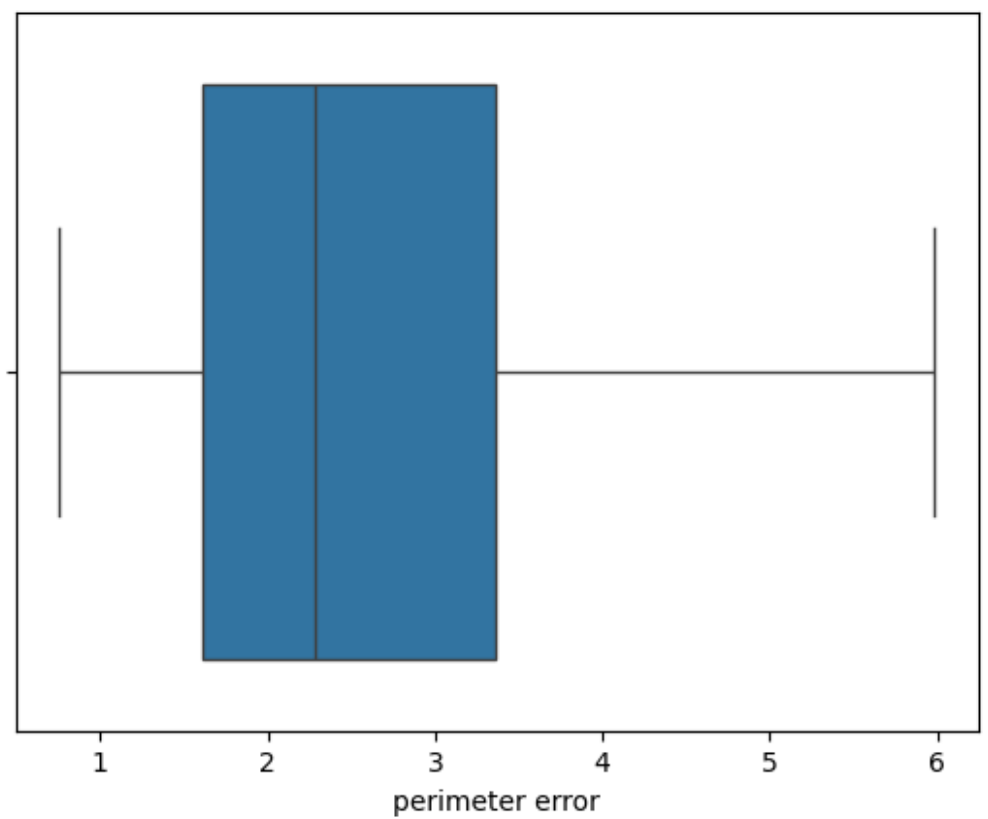
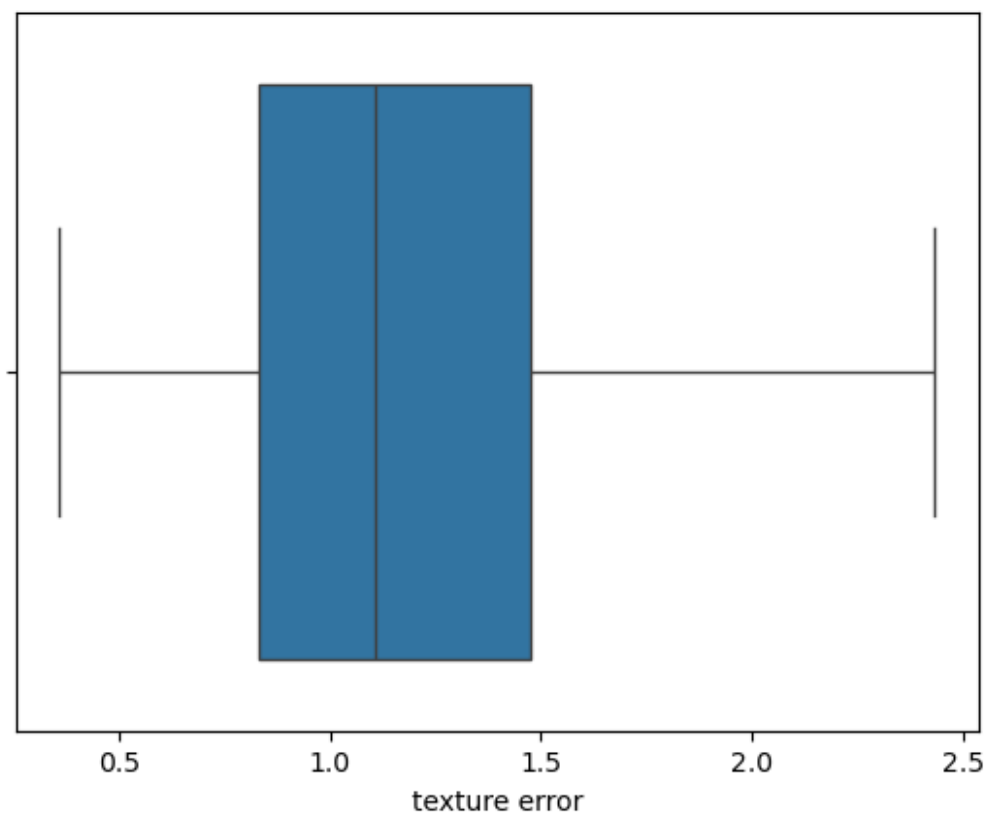




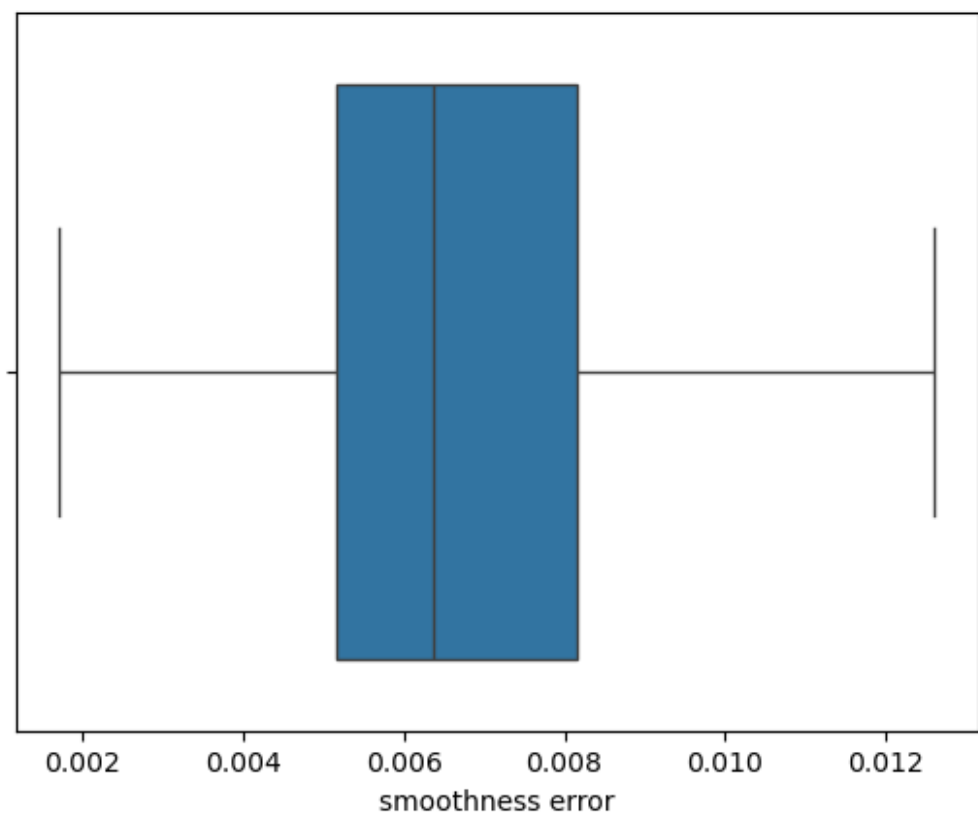
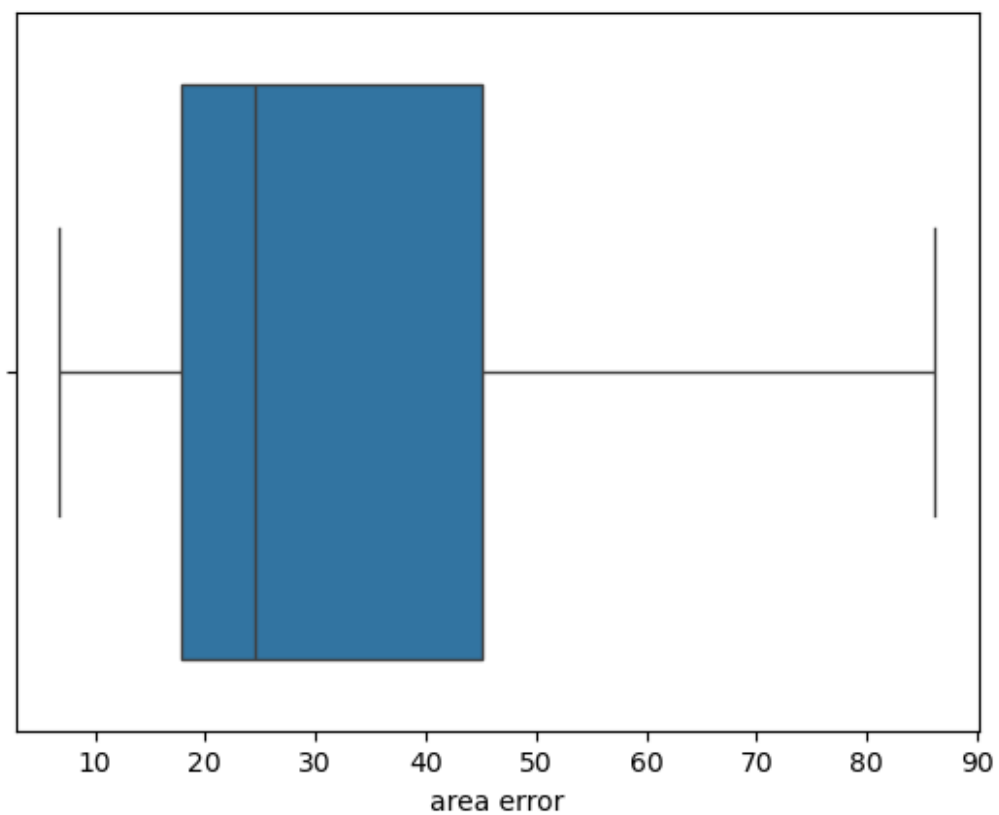


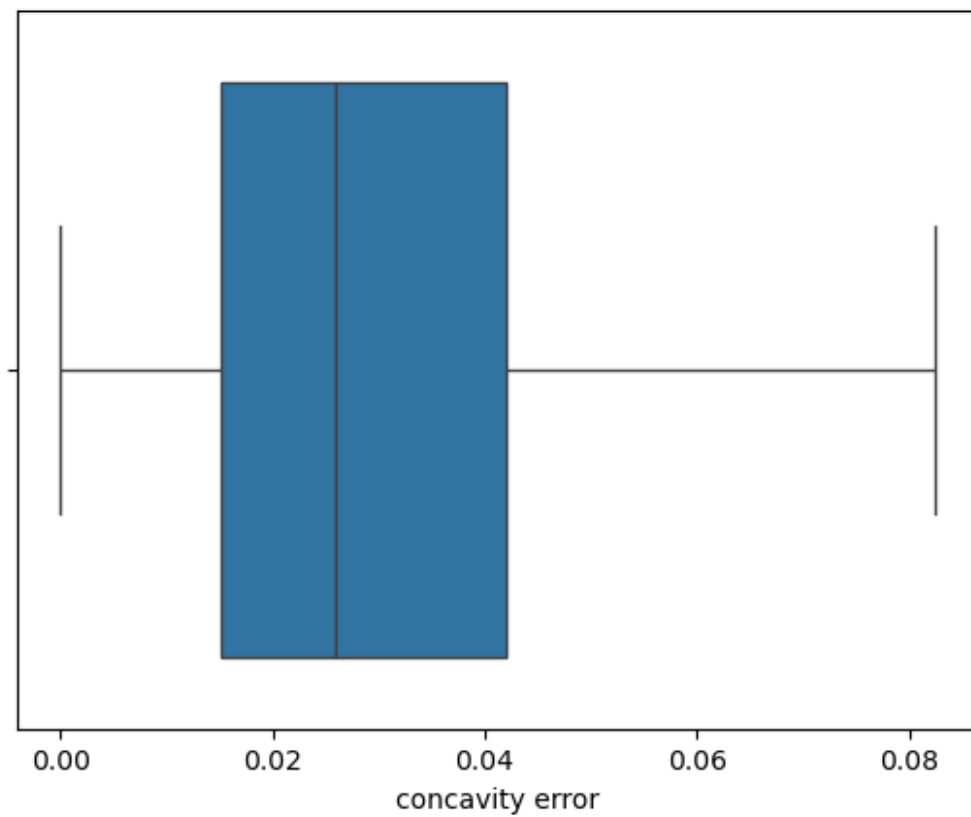
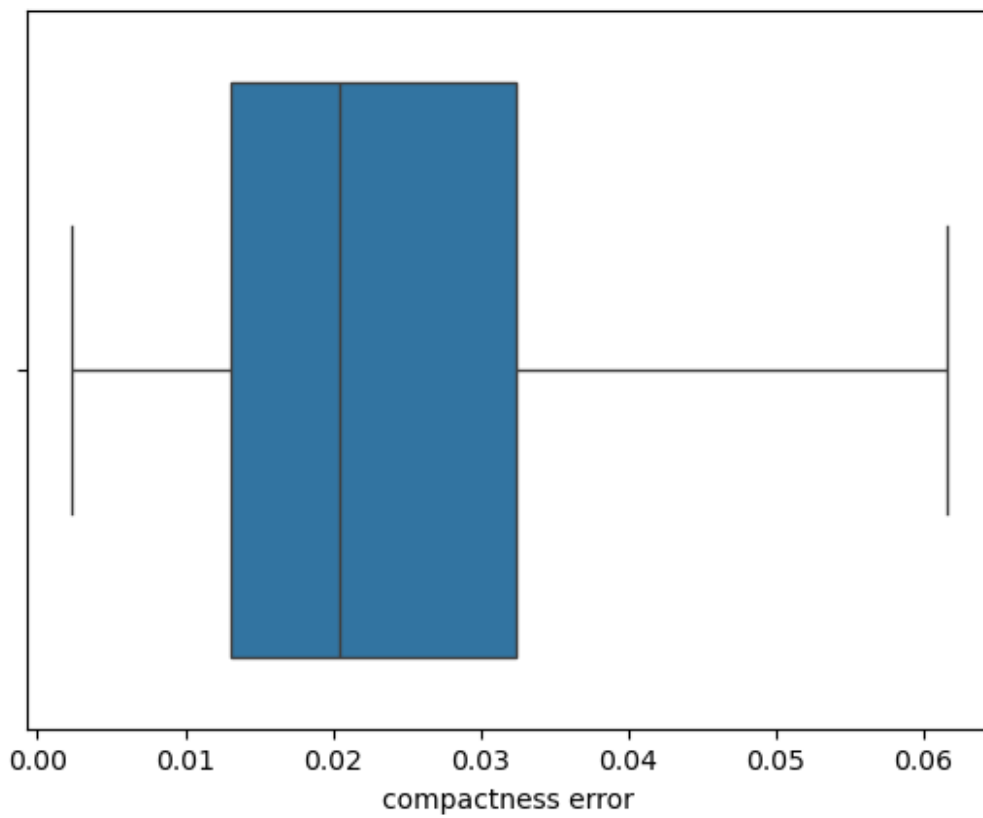


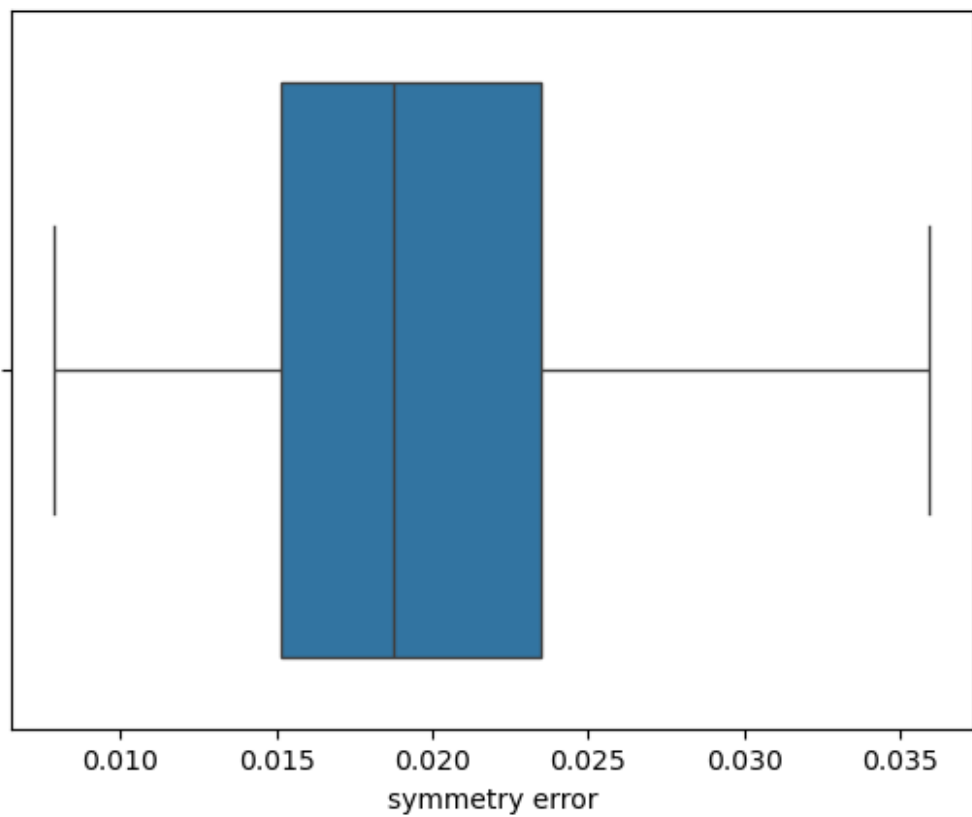
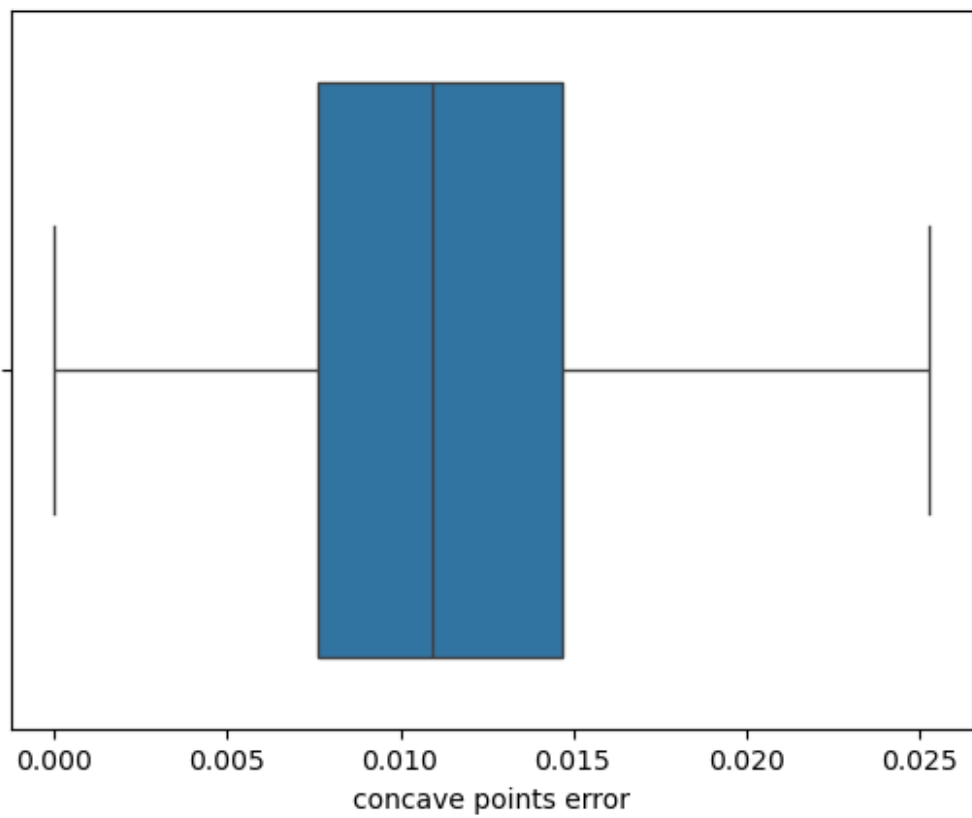


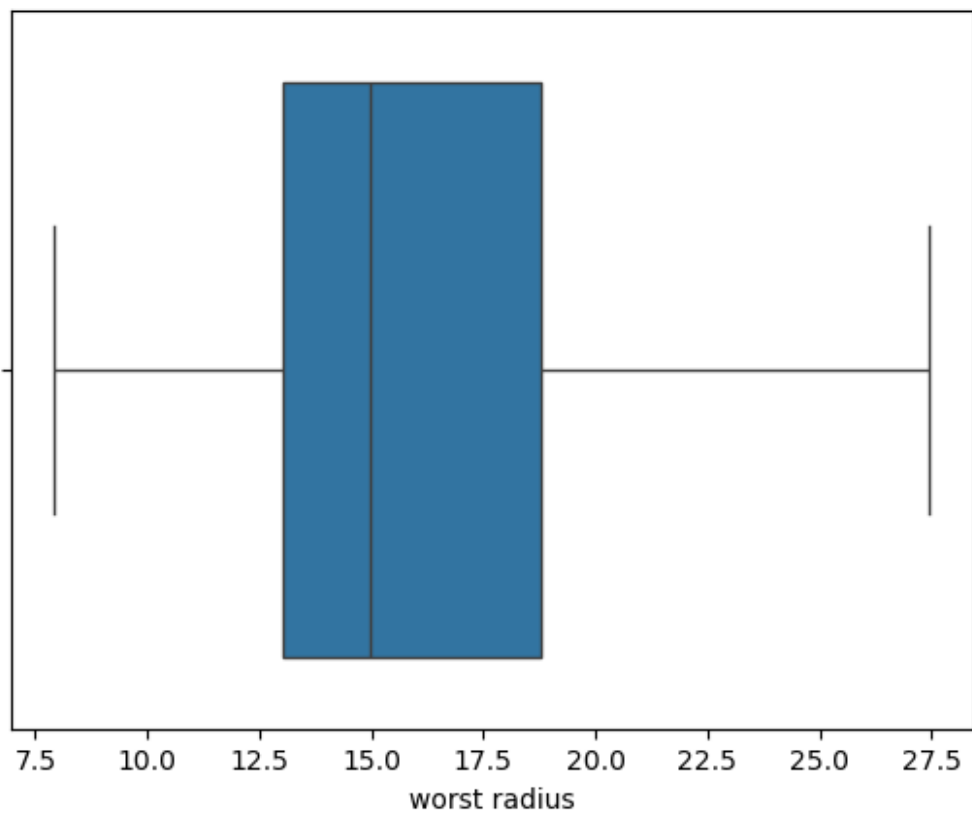
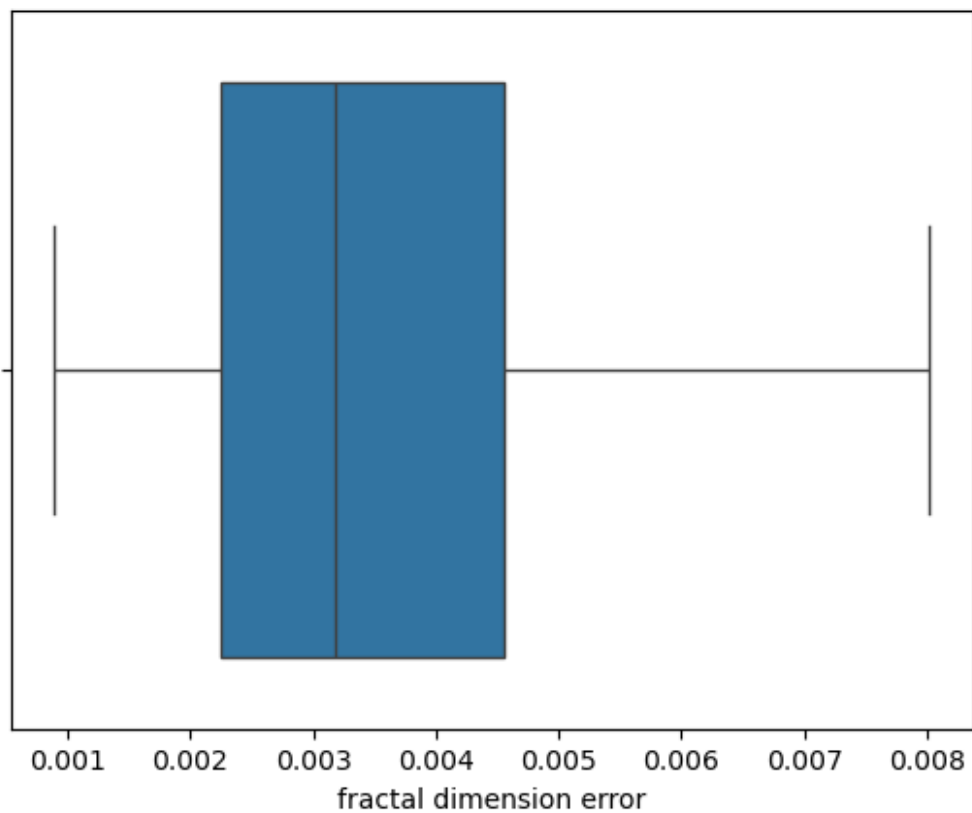


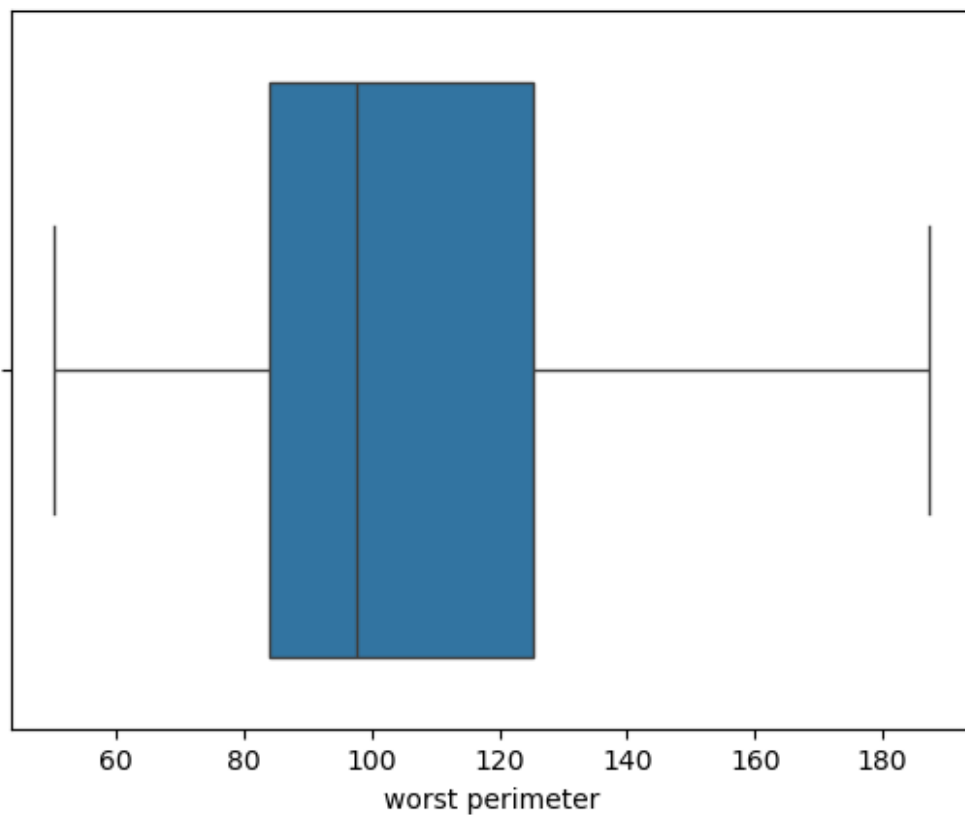
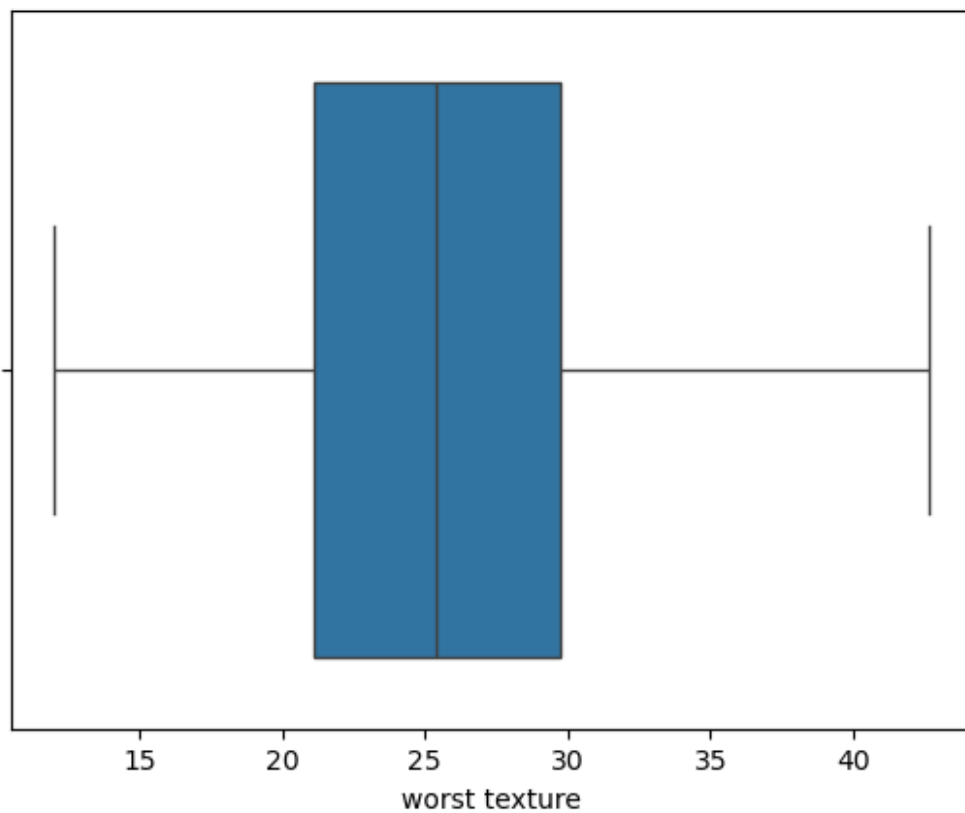


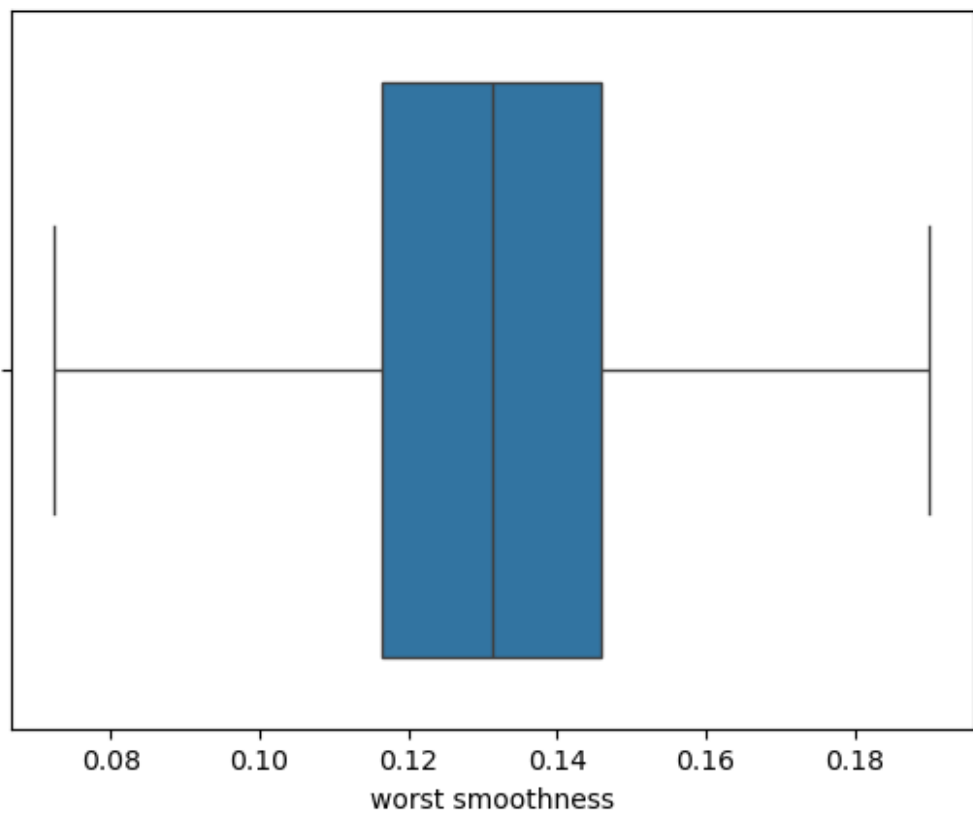
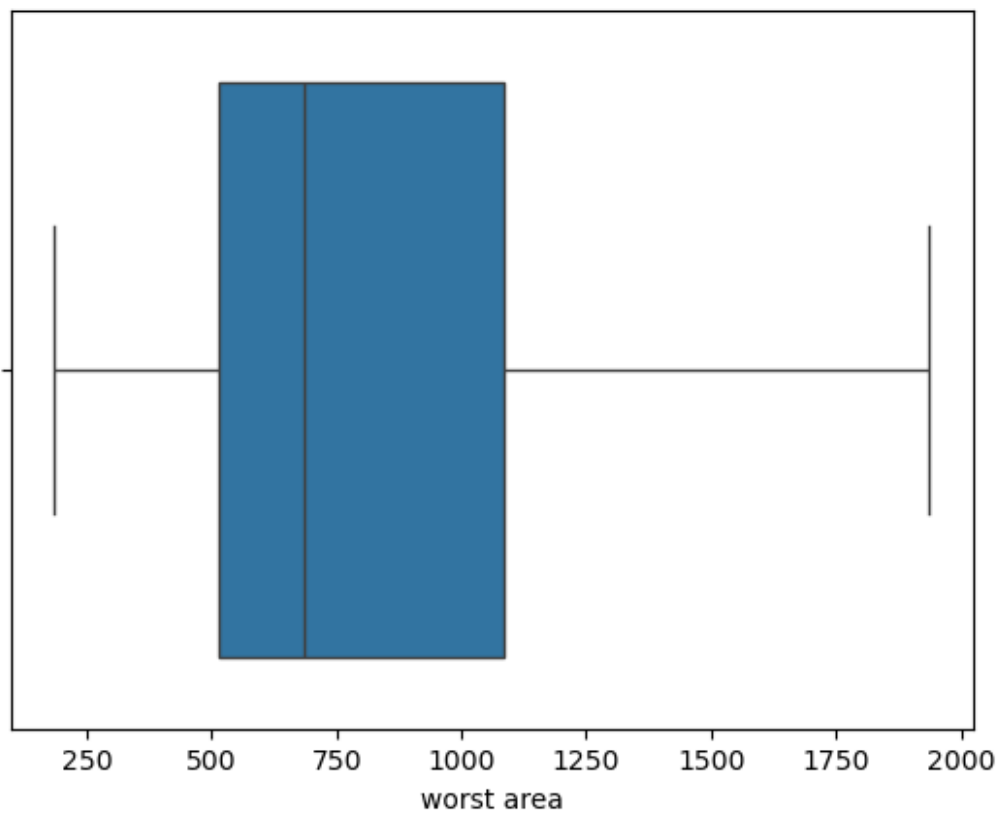


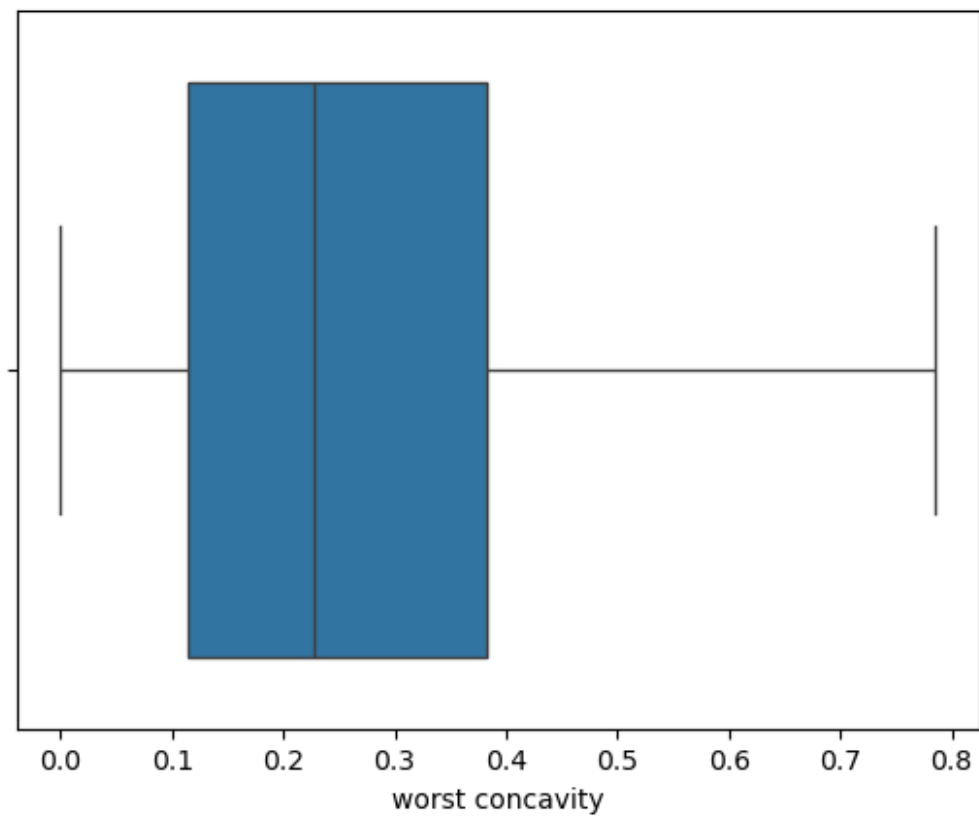
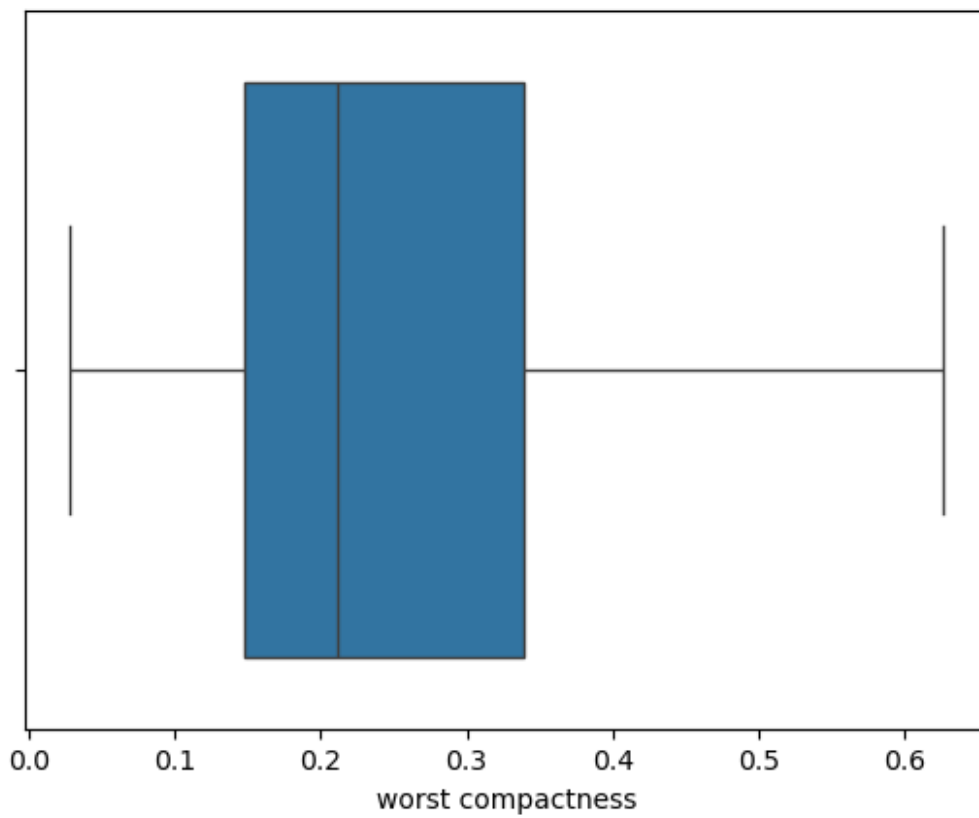


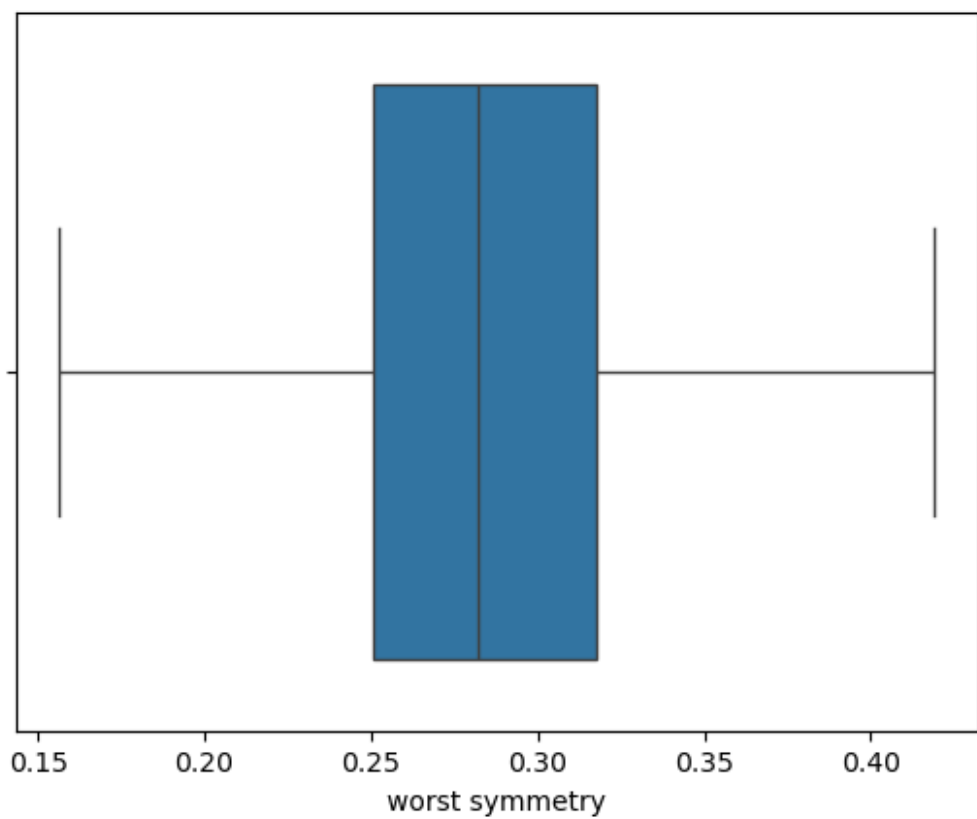
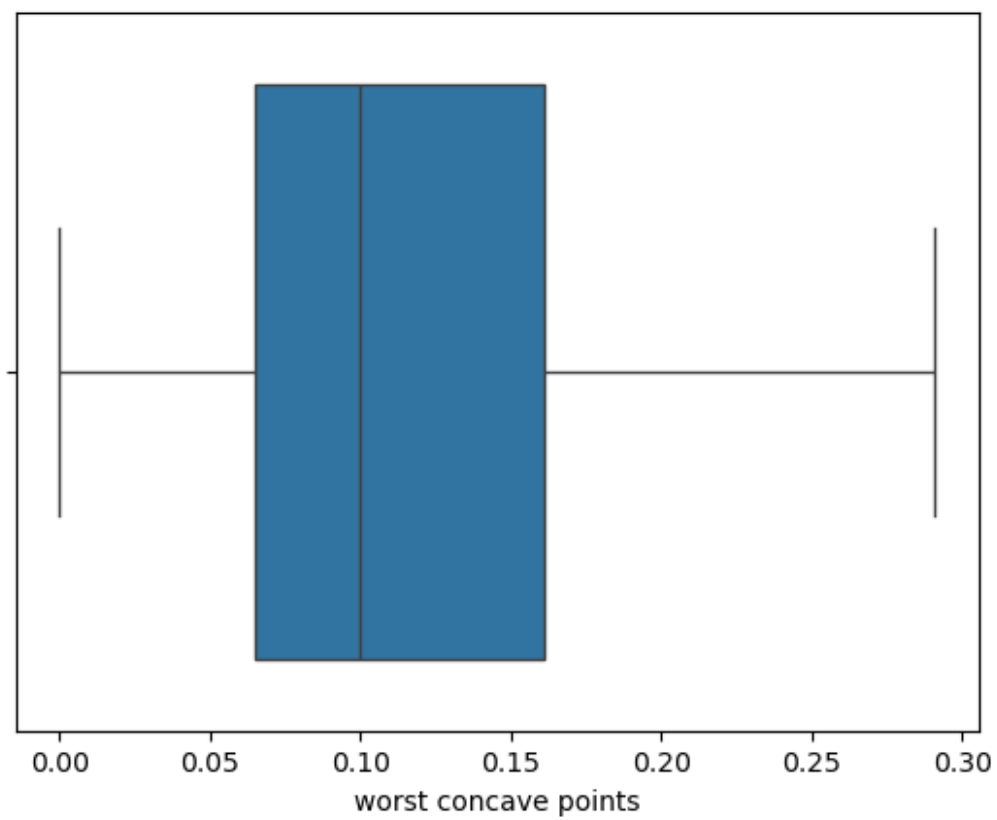




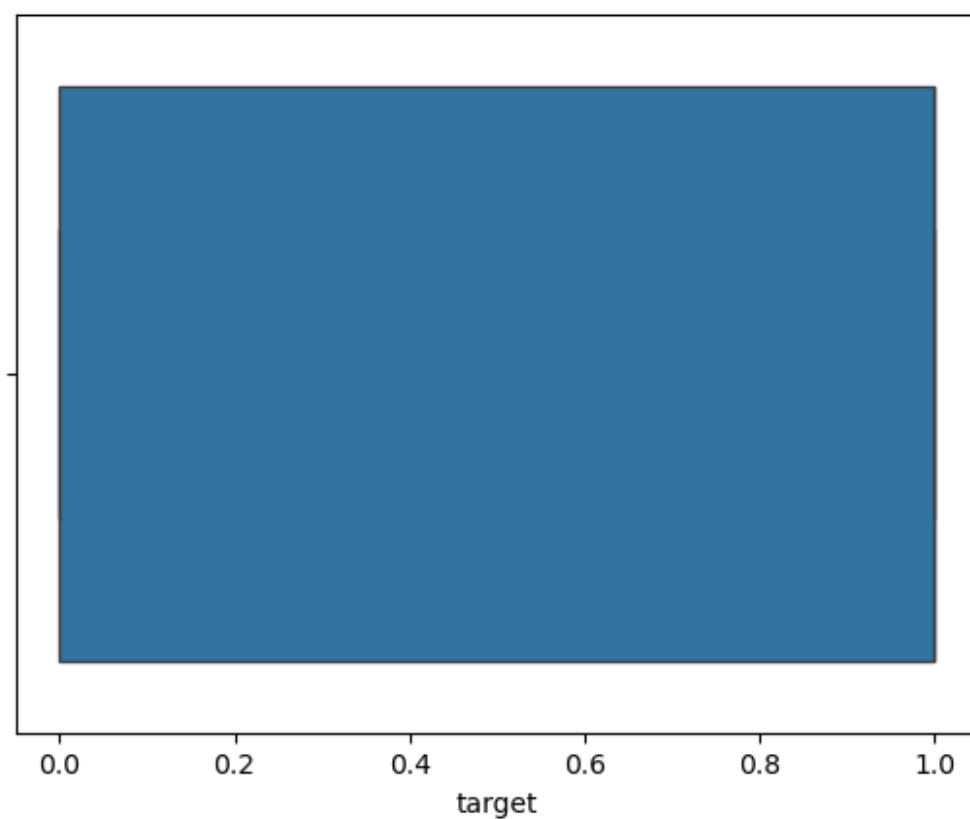
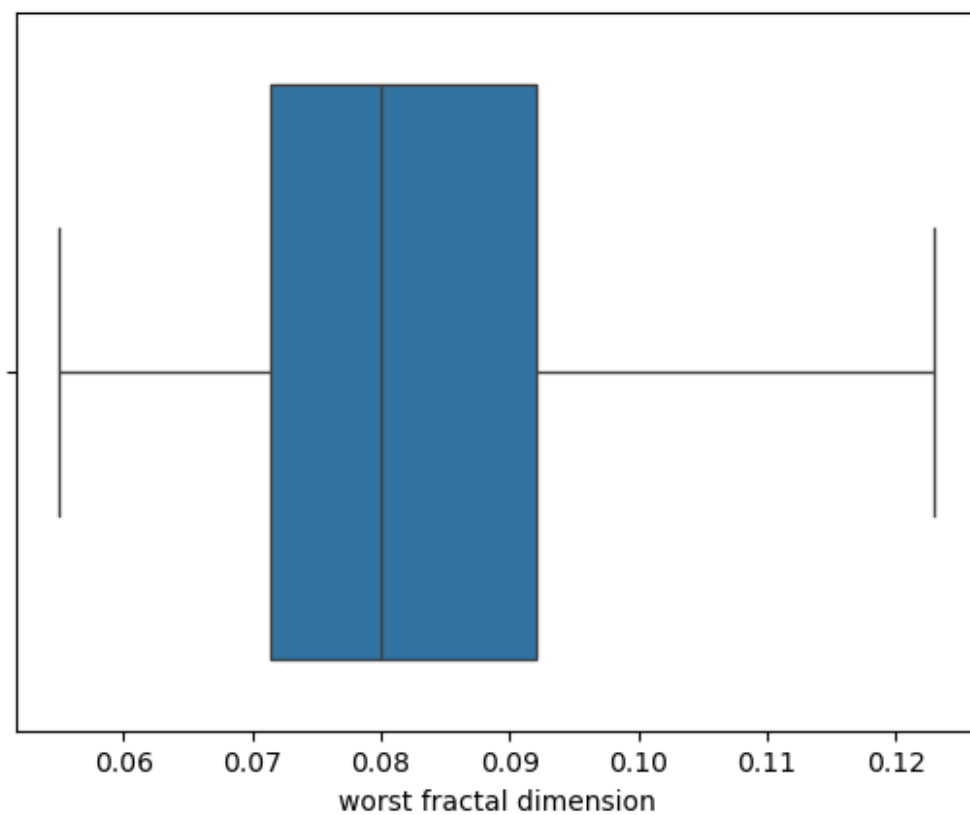












```
# checking skewness
```

```
df.skew()
```

```
mean radius      0.655953
mean texture     0.449700
mean perimeter   0.701081
mean area        0.922884
mean smoothness  0.257712
mean compactness 0.826755
mean concavity   1.023859
mean concave points 1.004049
mean symmetry    0.403621
mean fractal dimension 0.682430
radius error     1.025031
texture error    0.740987
perimeter error  1.034389
area error       1.130940
smoothness error 0.780923
compactness error 0.990285
concavity error  0.916740
concave points error 0.539571
symmetry error   0.869297
fractal dimension error 0.979344
worst radius     0.849779
worst texture    0.386858
worst perimeter  0.874870
worst area       1.048970
worst smoothness 0.247199
worst compactness 0.915295
worst concavity  0.809174
worst concave points 0.492616
worst symmetry   0.521772
worst fractal dimension 0.831581
target          -0.528461
dtype: float64
```

## EDA

```
df1 = df.copy()
```

```
df1.head()
```

	mean radius	mean texture	mean perimeter	mean area	mean smoothness \
0	17.99	10.38	122.80	1001.0	0.118400
1	20.57	17.77	132.90	1326.0	0.084740
2	19.69	21.25	130.00	1203.0	0.109600

3	11.42	20.38	77.58	386.1
0.133695				
4	20.29	14.34	135.10	1297.0
0.100300				

	mean compactness	mean concavity	mean concave points	mean symmetry \
0	0.22862	0.28241	0.14710	0.2419
1	0.07864	0.08690	0.07017	0.1812
2	0.15990	0.19740	0.12790	0.2069
3	0.22862	0.24140	0.10520	0.2464
4	0.13280	0.19800	0.10430	0.1809

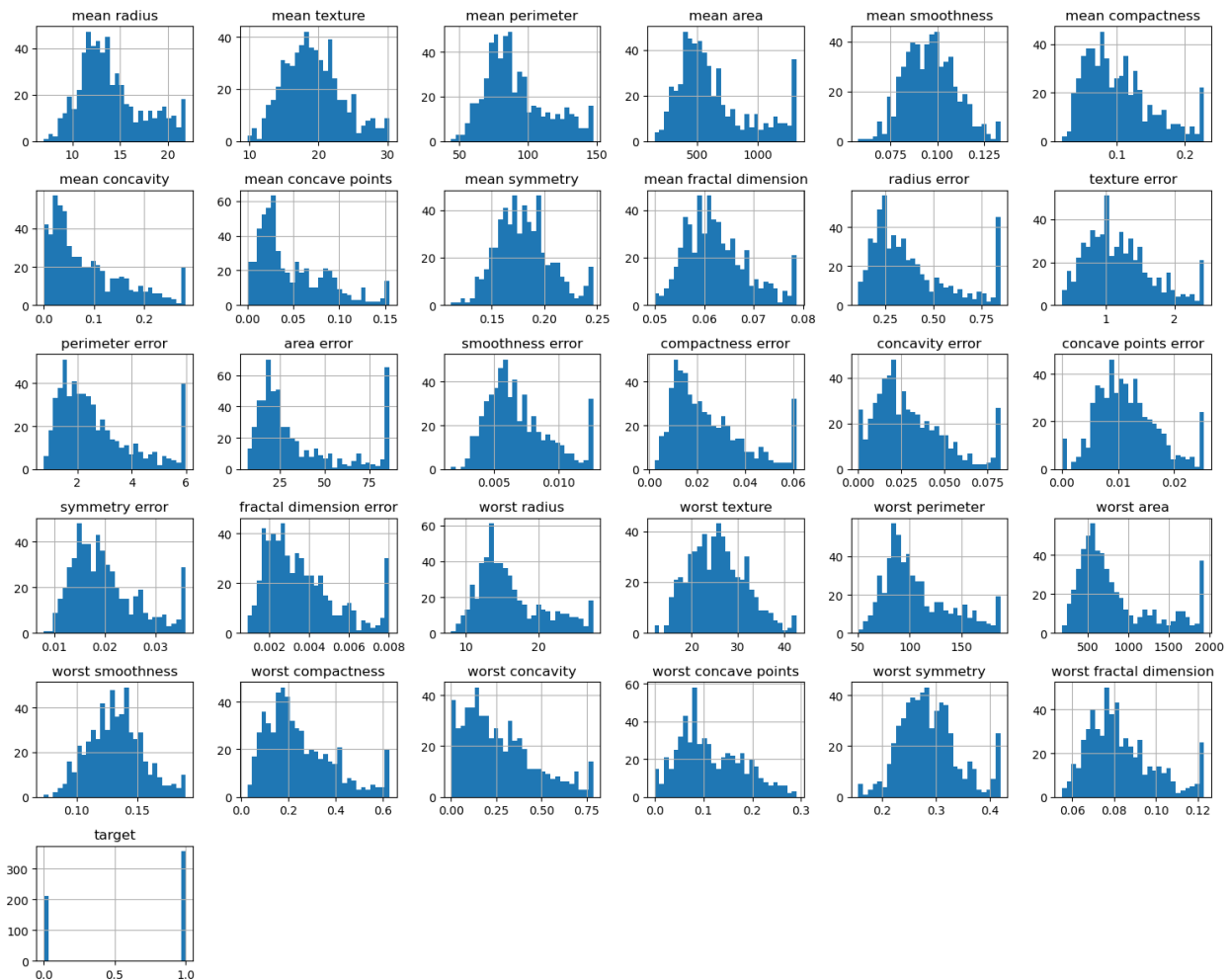
	mean fractal dimension	...	worst texture	worst perimeter	worst area \
0	0.07871	...	17.33	184.60	1937.05
1	0.05667	...	23.41	158.80	1937.05
2	0.05999	...	25.53	152.50	1709.00
3	0.07875	...	26.50	98.87	567.70
4	0.05883	...	16.67	152.20	1575.00

	worst smoothness	worst compactness	worst concavity	worst concave points \
0	0.1622	0.62695	0.7119	0.2654
1	0.1238	0.18660	0.2416	0.1860
2	0.1444	0.42450	0.4504	0.2430
3	0.1901	0.62695	0.6869	0.2575
4	0.1374	0.20500	0.4000	0.1625

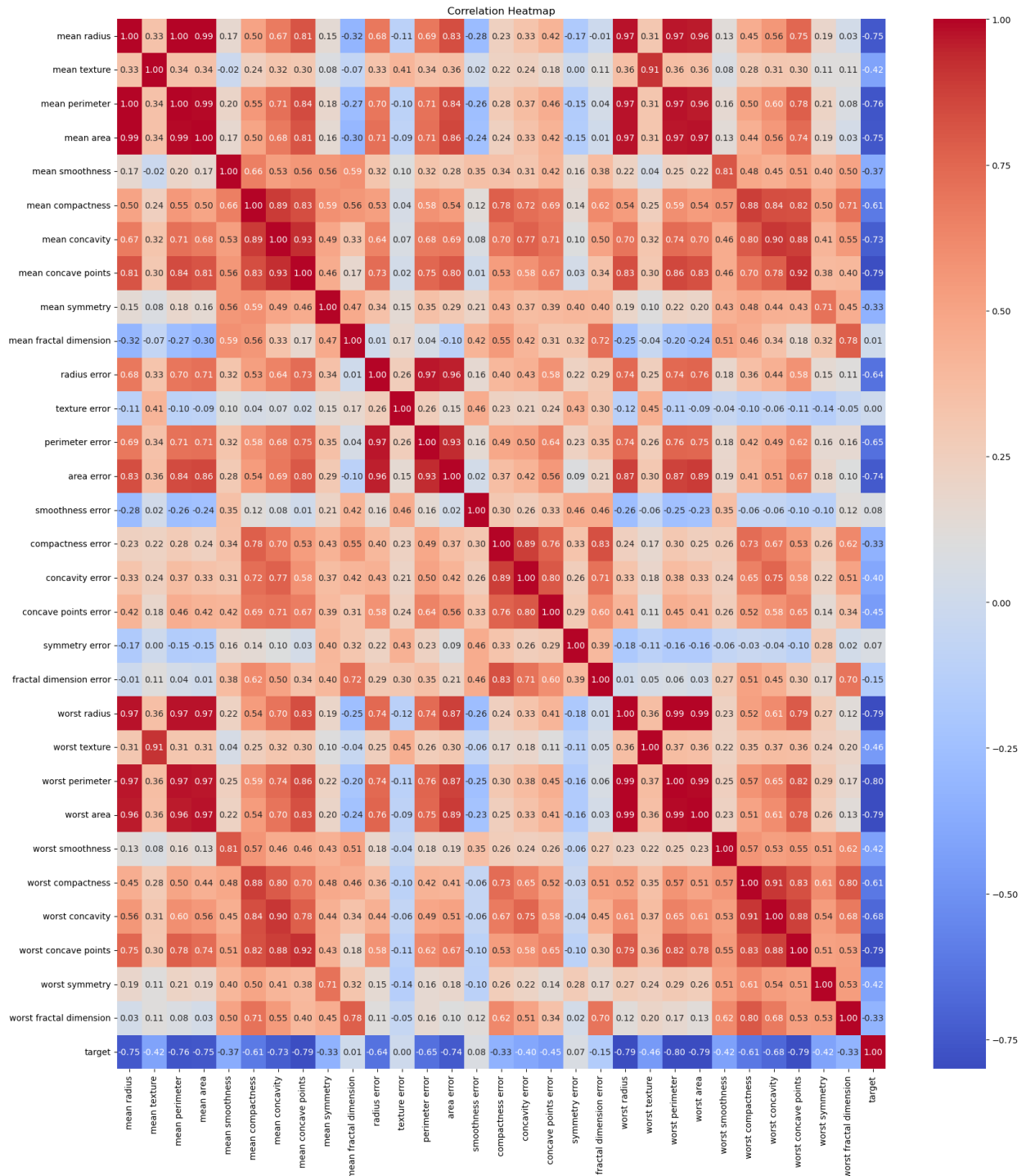
	worst symmetry	worst fractal dimension	target
0	0.41915	0.11890	0
1	0.27500	0.08902	0
2	0.36130	0.08758	0
3	0.41915	0.12301	0
4	0.23640	0.07678	0

```
[5 rows x 31 columns]
```

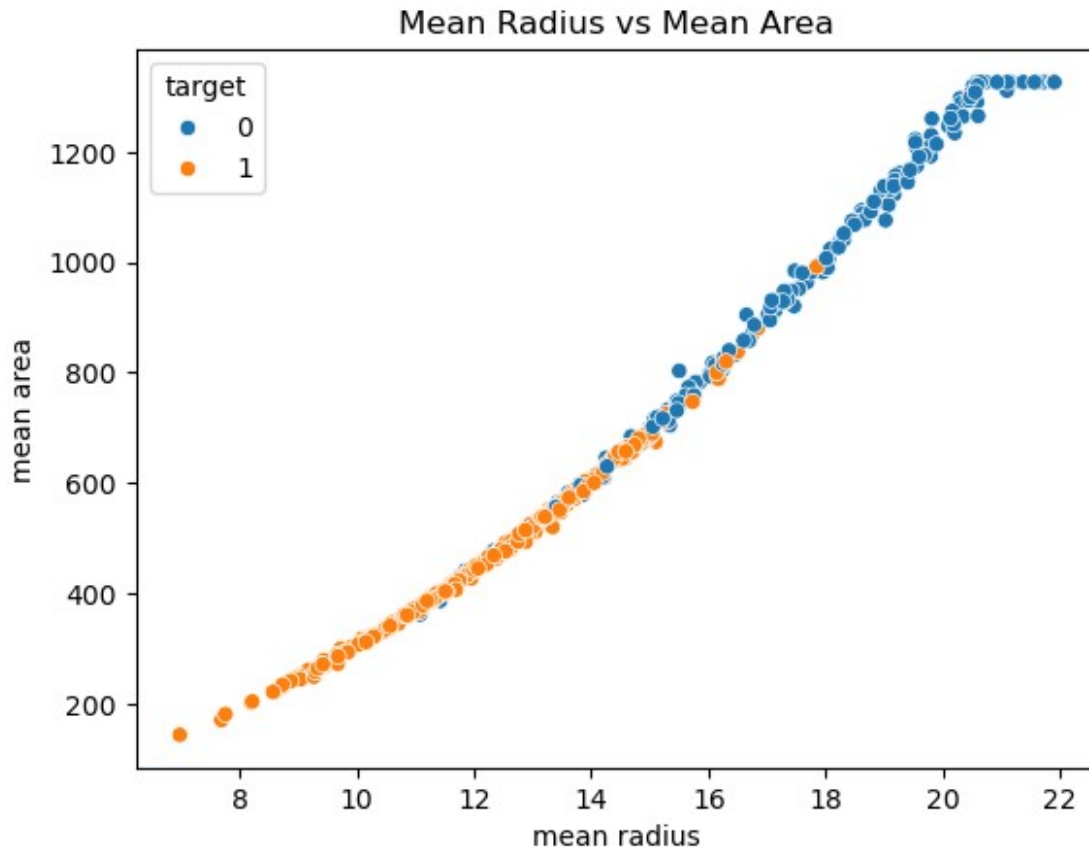
```
# Histogram  
df1.hist(bins=30, figsize=(15, 12))  
plt.tight_layout()  
plt.show()
```



```
# Compute correlation matrix  
corr_matrix = df1.corr()  
  
# Heatmap  
plt.figure(figsize=(20, 22))  
sns.heatmap(corr_matrix, annot=True, fmt=".2f", cmap="coolwarm")  
plt.title("Correlation Heatmap")  
plt.show()
```



```
# Scatterplot
sns.scatterplot(x="mean radius", y="mean area", hue="target", data=df)
plt.title("Mean Radius vs Mean Area")
plt.show()
```



## PREPROCESSING STEPS AND EXPLANATION

1. Fetched basic details of the dataset with `info()`, `describe()`, `shape` and `dtypes`
2. Checked for duplicate values and null values. The data don't have any null/ duplicated values
3. Added boxplot for every column to visualise if there is any outliers present also added a single box plot chart combining every feature to compare the outliers
4. Used IQR method to find the outliers of all features
5. Used capping method further to fix the outliers
6. Target feature doesn't have any outliers so dropped it for outlier fixing
7. After Capping added box plot again to see the changes after outlier fixation
8. Checked for the skew value for every feature . skew was in a good range.
9. Drawn a histogram for every feature
10. Added a correlation heatmap to see the relationship
11. Added a scatterplot for mean radius vs mean area

## FEATURE SELECTION

```
# Using correlation matrix

# Compute correlation matrix
corr_matrix = df1.corr().abs()

# Select upper triangle of correlation matrix
upper = corr_matrix.where(np.triu(np.ones(corr_matrix.shape),
k=1).astype(bool))

# Find features with correlation > 0.9
to_drop = [column for column in upper.columns if any(upper[column] >
0.9)]

print(f"Features to drop due to high correlation: {to_drop}")

# Drop features
df1_reduced = df1.drop(columns=to_drop)

Features to drop due to high correlation: ['mean perimeter', 'mean
area', 'mean concave points', 'perimeter error', 'area error', 'worst
radius', 'worst texture', 'worst perimeter', 'worst area', 'worst
concavity', 'worst concave points']

df1_reduced.head()
```

	mean radius	mean texture	mean smoothness	mean compactness	\
0	17.99	10.38	0.118400	0.22862	
1	20.57	17.77	0.084740	0.07864	
2	19.69	21.25	0.109600	0.15990	
3	11.42	20.38	0.133695	0.22862	
4	20.29	14.34	0.100300	0.13280	

	mean concavity	mean symmetry	mean fractal dimension	radius error	\
0	0.28241	0.2419	0.07871	0.84865	
1	0.08690	0.1812	0.05667	0.54350	
2	0.19740	0.2069	0.05999	0.74560	
3	0.24140	0.2464	0.07875	0.49560	
4	0.19800	0.1809	0.05883	0.75720	

	texture error	smoothness error	compactness error	concavity error	\
0	0.9053	0.006399	0.049040	0.05373	
1	0.7339	0.005225	0.013080	0.01860	

2	0.7869	0.006150	0.040060	0.03832
3	1.1560	0.009110	0.061505	0.05661
4	0.7813	0.011490	0.024610	0.05688

	concave points error	symmetry error	fractal dimension error	\
0	0.01587	0.03003		0.006193
1	0.01340	0.01389		0.003532
2	0.02058	0.02250		0.004571
3	0.01867	0.03596		0.008023
4	0.01885	0.01756		0.005115

	worst smoothness	worst compactness	worst symmetry	\
0	0.1622	0.62695	0.41915	
1	0.1238	0.18660	0.27500	
2	0.1444	0.42450	0.36130	
3	0.1901	0.62695	0.41915	
4	0.1374	0.20500	0.23640	

	worst fractal dimension	target
0	0.11890	0
1	0.08902	0
2	0.08758	0
3	0.12301	0
4	0.07678	0

## SETTING X AND Y

```
y = df1_reduced['target']
y
0      0
1      0
2      0
3      0
4      0
..
564    0
565    0
566    0
567    0
568    1
Name: target, Length: 569, dtype: int32

x = df1_reduced.drop('target',axis=1)
x
```



	mean radius	mean texture	mean smoothness	mean compactness	\
0	17.99	10.38	0.118400	0.22862	
1	20.57	17.77	0.084740	0.07864	
2	19.69	21.25	0.109600	0.15990	
3	11.42	20.38	0.133695	0.22862	
4	20.29	14.34	0.100300	0.13280	
..	...	...	...	...	
564	21.56	22.39	0.111000	0.11590	
565	20.13	28.25	0.097800	0.10340	
566	16.60	28.08	0.084550	0.10230	
567	20.60	29.33	0.117800	0.22862	
568	7.76	24.54	0.057975	0.04362	
	mean concavity	mean symmetry	mean fractal dimension	radius	error \
0	0.28241	0.2419		0.07871	0.84865
1	0.08690	0.1812		0.05667	0.54350
2	0.19740	0.2069		0.05999	0.74560
3	0.24140	0.2464		0.07875	0.49560
4	0.19800	0.1809		0.05883	0.75720
..	...	...		...	..
564	0.24390	0.1726		0.05623	0.84865
565	0.14400	0.1752		0.05533	0.76550
566	0.09251	0.1590		0.05648	0.45640
567	0.28241	0.2397		0.07016	0.72600
568	0.00000	0.1587		0.05884	0.38570
	texture error	smoothness error	compactness error	concavity	error \
0	0.90530	0.006399	0.049040		0.05373
1	0.73390	0.005225	0.013080		0.01860
2	0.78690	0.006150	0.040060		0.03832
3	1.15600	0.009110	0.061505		0.05661
4	0.78130	0.011490	0.024610		0.05688

```

..          ...          ...          ...
..
564          1.25600          0.010300          0.028910
0.05198
565          2.43415          0.005769          0.024230
0.03950
566          1.07500          0.005903          0.037310
0.04730
567          1.59500          0.006522          0.061505
0.07117
568          1.42800          0.007189          0.004660
0.00000

```

```

          concave points error  symmetry error  fractal dimension error \
0          0.01587          0.03003          0.006193
1          0.01340          0.01389          0.003532
2          0.02058          0.02250          0.004571
3          0.01867          0.03596          0.008023
4          0.01885          0.01756          0.005115
..          ...          ...          ...
564          0.02454          0.01114          0.004239
565          0.01678          0.01898          0.002498
566          0.01557          0.01318          0.003892
567          0.01664          0.02324          0.006185
568          0.00000          0.02676          0.002783

```

```

          worst smoothness  worst compactness  worst symmetry \
0          0.16220          0.62695          0.41915
1          0.12380          0.18660          0.27500
2          0.14440          0.42450          0.36130
3          0.19010          0.62695          0.41915
4          0.13740          0.20500          0.23640
..          ...          ...          ...
564          0.14100          0.21130          0.20600
565          0.11660          0.19220          0.25720
566          0.11390          0.30940          0.22180
567          0.16500          0.62695          0.40870
568          0.08996          0.06444          0.28710

```

```

          worst fractal dimension
0          0.11890
1          0.08902
2          0.08758
3          0.12301
4          0.07678
..          ...
564          0.07115
565          0.06637
566          0.07820
567          0.12301

```

568 0.07039

[569 rows x 19 columns]

## FEATURE SCALING

```
minmax_scaler = MinMaxScaler()
```

```
#Applying scaling
```

```
x_normalized = minmax_scaler.fit_transform(x)
```

```
# converting into dataframe
```

```
x_normalized = pd.DataFrame(x_normalized)
```

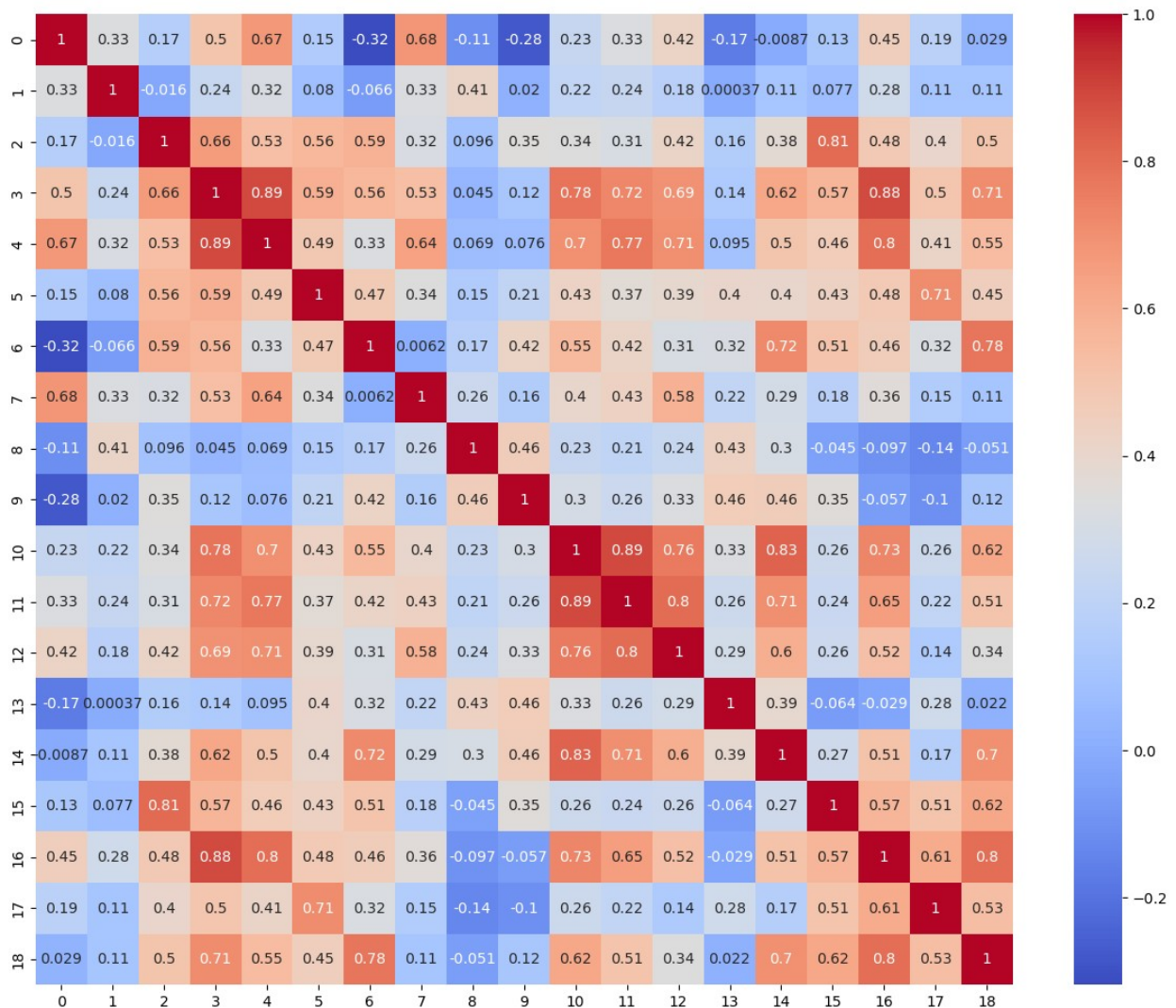
```
x_normalized.head()
```

	0	1	2	3	4	5	6
\							
0	0.737918	0.032627	0.798006	1.000000	1.000000	0.966716	
0.998611							
1	0.910852	0.392501	0.353473	0.283215	0.307709	0.517751	
0.233067							
2	0.851867	0.561967	0.681788	0.671573	0.698984	0.707840	
0.348385							
3	0.297540	0.519601	1.000000	1.000000	0.854786	1.000000	
1.000000							
4	0.892084	0.225469	0.558967	0.542057	0.701108	0.515533	
0.308093							
	7	8	9	10	11	12	
13 \							
0	1.000000	0.262832	0.429967	0.789631	0.651352	0.626827	
0.788803							
1	0.586041	0.180188	0.322246	0.182742	0.225482	0.529268	
0.213975							
2	0.860205	0.205743	0.407120	0.638077	0.464541	0.812860	
0.520621							
3	0.521061	0.383712	0.678717	1.000000	0.686265	0.737420	
1.000000							
4	0.875941	0.203043	0.897096	0.377331	0.689538	0.744530	
0.344683							
	14	15	16	17	18		
0	0.743273	0.762755	1.000000	1.000000	0.939532		
1	0.369967	0.436224	0.265667	0.451171	0.499926		
2	0.515726	0.611395	0.662392	0.779745	0.478741		
3	1.000000	1.000000	1.000000	1.000000	1.000000		
4	0.592043	0.551871	0.296351	0.304207	0.319847		

```
correlation = x_normalized.corr()
```

```
plt.figure(figsize=(15, 12))
```

```
sns.heatmap(correlation, annot=True, cmap='coolwarm')
plt.show()
```



## SPLITTING TRAIN AND TEST

```
x_train, x_test, y_train, y_test = train_test_split(x_normalized, y,
test_size=0.2, random_state=42)
```

## BUILDING MODELS

```
models = {
    "Logistic Regression": LogisticRegression(max_iter=500),
    "Decision Tree": DecisionTreeClassifier(max_depth=5),
    "Random Forest": RandomForestClassifier(n_estimators=100),
    "SVM": SVC(kernel='linear', C=1),
    "k-NN": KNeighborsClassifier(n_neighbors=5)
}
```

## LOGISTIC REGRESSION

A logistic (sigmoid) function is used in the linear model known as logistic regression to forecast probability. Based on a threshold (usually 0.5), it generates probabilities for binary classification and designates a label (e.g., malignant or non-cancerous). In the feature space, the decision boundary is linear.

Why its suitable: It works well with linearly separable datasets and is easy to understand. Datasets on breast cancer frequently exhibit distinct patterns that logistic regression is well suited to model. In medical analysis, the coefficients offer useful insights regarding feature relevance.

## DECISION TREE CLASSIFIER

Based on feature values, decision trees divide the data into subsets and produce branches until a leaf node makes a categorization. Criteria such as Gini Impurity or Information Gain are used to determine the divides.

Why its suitable: It efficiently manages both numerical and category features. The tree structure can be used to describe the classification process in medical circumstances because it is simple to see and understand. Even if they are non-linear, it captures intricate correlations in the data.

## RANDOM FOREST CLASSIFIER

Multiple decision trees are constructed during training using the Random Forest ensemble approach, which then aggregates the results (using majority vote for classification). By bootstrapping the data and using random feature subsets for splitting, it adds unpredictability.

By lessening overfitting, which can happen in individual decision trees, it increases accuracy. robust to outliers and noisy data, which makes it appropriate for datasets with different feature distributions, such as breast cancer. It can help identify the elements impacting forecasts by ranking the relevance of features.

## SVC

By maximizing the margin between data points of distinct classes, SVC seeks to identify the hyperplane that best divides them. For improved separability, it can convert data into higher dimensions using kernels (such as linear and RBF). It performs well in high-dimensional areas, such as the feature-rich breast cancer dataset. useful for datasets in which the initial feature space does not readily allow for the separation of classes. When there are more features than samples, it is resistant to overfitting.

## K- NEAREST NEIGHBOURS (K-NN)

A data point is categorized by k-NN using the majority class of its k nearest neighbors in the feature space. Euclidean and other distance metrics are used to find neighbors. It is a straightforward non-parametric approach that doesn't assume anything about the distribution of the underlying data. Ideal for datasets with distinct and well-clustered class distributions. Because it is sensitive to feature magnitudes, it works best when feature scaling (such as standardization) is used.

```

# Train and evaluate each model
results = {}
for model_name, model in models.items():
    print(f"Training {model_name}...")
    model.fit(x_train, y_train) # Train the model
    y_pred = model.predict(x_test) # Make predictions

    # Evaluate the model
    accuracy = accuracy_score(y_test, y_pred)
    results[model_name] = accuracy

    print(f"{model_name} Accuracy: {accuracy:.4f}")
    print(f"{model_name} Classification Report:\n{classification_report(y_test, y_pred)}\n")

    # Generate Confusion Matrix
    cm = confusion_matrix(y_test, y_pred)
    print(f"{model_name} Confusion Matrix:\n{cm}\n")

    # Visualize Confusion Matrix
    plt.figure(figsize=(6, 4))
    sns.heatmap(cm, annot=True, fmt="d", cmap="Blues",
xticklabels=["Class 0", "Class 1"], yticklabels=["Class 0", "Class 1"])
    plt.title(f"Confusion Matrix for {model_name}")
    plt.ylabel("Actual")
    plt.xlabel("y_pred")
    plt.show()

```

Training Logistic Regression...

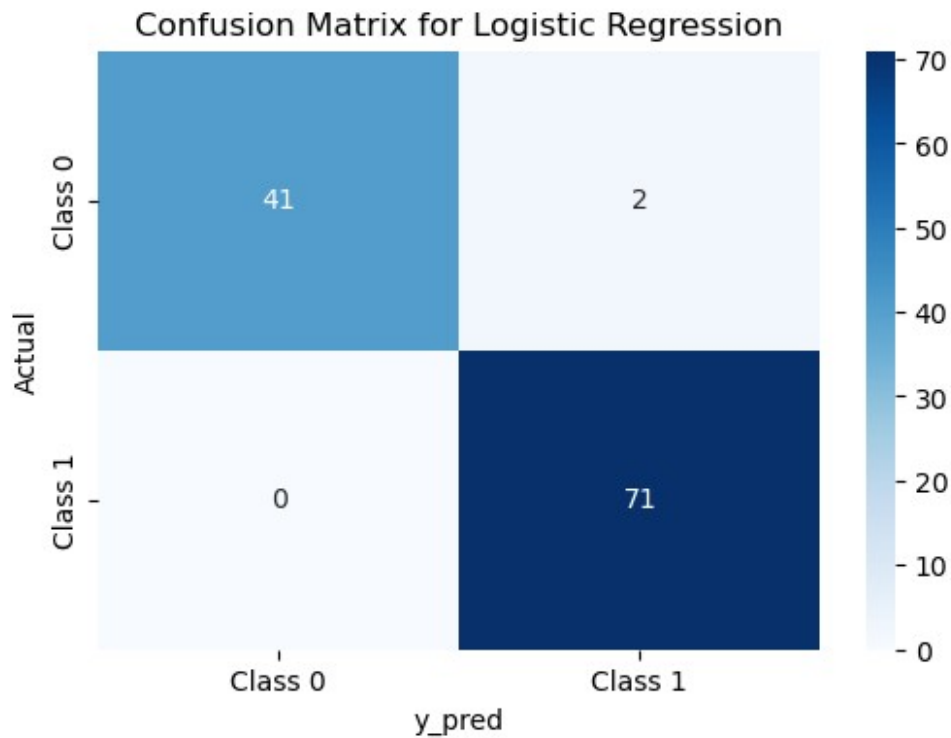
Logistic Regression Accuracy: 0.9825

Logistic Regression Classification Report:

	precision	recall	f1-score	support
0	1.00	0.95	0.98	43
1	0.97	1.00	0.99	71
accuracy			0.98	114
macro avg	0.99	0.98	0.98	114
weighted avg	0.98	0.98	0.98	114

Logistic Regression Confusion Matrix:

```
[[41  2]
 [ 0 71]]
```



Training Decision Tree...

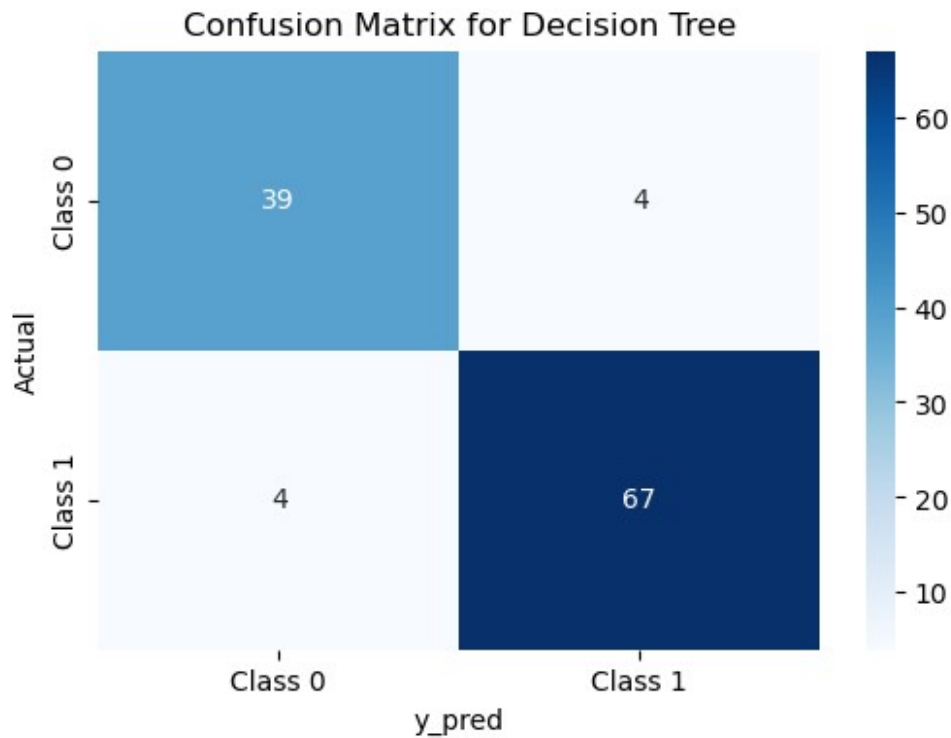
Decision Tree Accuracy: 0.9298

Decision Tree Classification Report:

	precision	recall	f1-score	support
0	0.91	0.91	0.91	43
1	0.94	0.94	0.94	71
accuracy			0.93	114
macro avg	0.93	0.93	0.93	114
weighted avg	0.93	0.93	0.93	114

Decision Tree Confusion Matrix:

```
[[39  4]
 [ 4 67]]
```



Training Random Forest...

Random Forest Accuracy: 0.9649

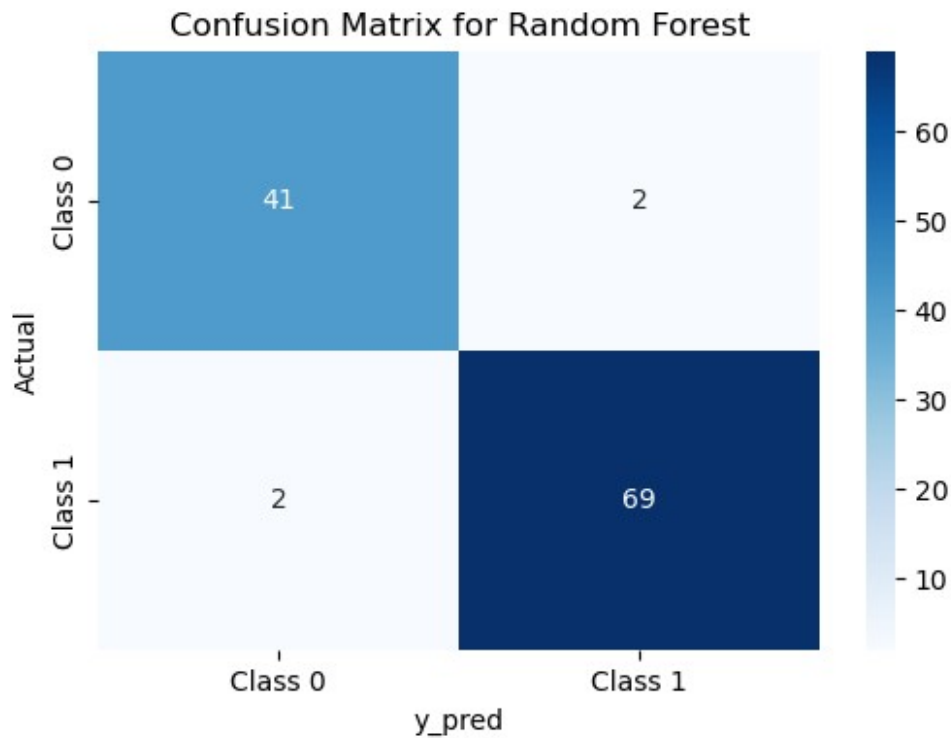
Random Forest Classification Report:

	precision	recall	f1-score	support
0	0.95	0.95	0.95	43
1	0.97	0.97	0.97	71
accuracy			0.96	114
macro avg	0.96	0.96	0.96	114
weighted avg	0.96	0.96	0.96	114

Random Forest Confusion Matrix:

```
[[41  2]
 [ 2 69]]
```





Training SVM...

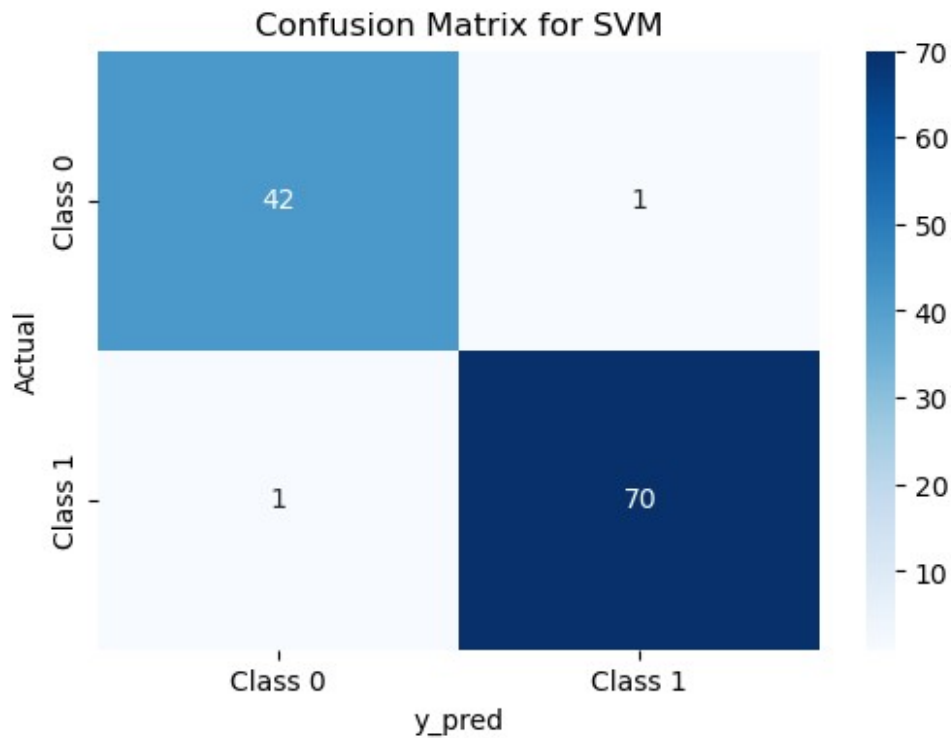
SVM Accuracy: 0.9825

SVM Classification Report:

	precision	recall	f1-score	support
0	0.98	0.98	0.98	43
1	0.99	0.99	0.99	71
accuracy			0.98	114
macro avg	0.98	0.98	0.98	114
weighted avg	0.98	0.98	0.98	114

SVM Confusion Matrix:

```
[[42  1]
 [ 1 70]]
```



Training k-NN...

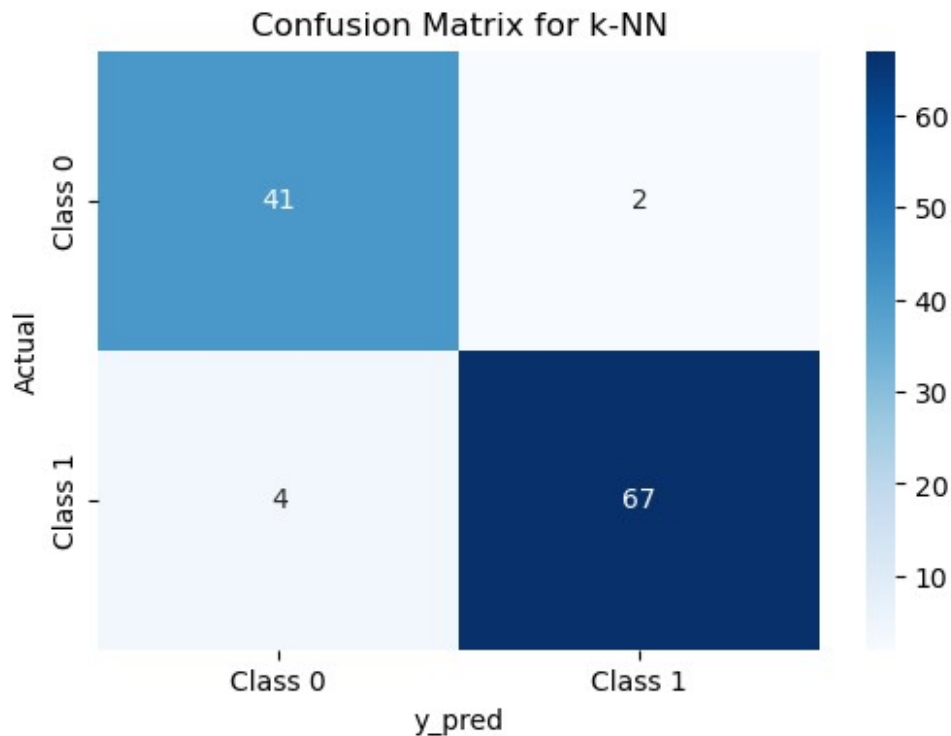
k-NN Accuracy: 0.9474

k-NN Classification Report:

	precision	recall	f1-score	support
0	0.91	0.95	0.93	43
1	0.97	0.94	0.96	71
accuracy			0.95	114
macro avg	0.94	0.95	0.94	114
weighted avg	0.95	0.95	0.95	114

k-NN Confusion Matrix:

```
[[41  2]
 [ 4 67]]
```



## MODEL EVALUATION

```
# Comparison of results
print("Model Accuracy Comparison:")
for model, acc in results.items():
    print(f"{model}: {acc:.4f}")

Model Accuracy Comparison:
Logistic Regression: 0.9825
Decision Tree: 0.9298
Random Forest: 0.9649
SVM: 0.9825
k-NN: 0.9474

## finding best model
# Get the best model from sorted_results
best_model_name = max(results, key=results.get)
best_model_accuracy = results[best_model_name]

print(f"\nThe Best Model is: {best_model_name}")
print(f"Accuracy: {best_model_accuracy:.4f}")

The Best Model is: Logistic Regression
Accuracy: 0.9825
```

## HYPERPARAMETER TUNING WITH GRIDSEARCH CV

```
# Define the Logistic Regression model
logreg = LogisticRegression(max_iter=1000)

# Define the parameter grid
param_grid = {
    'C': [0.01, 0.1, 1, 10, 100],          # Regularization strength
    'penalty': ['l1', 'l2', 'elasticnet'], # Regularization type
    'solver': ['liblinear', 'saga']        # Solver choice
}

# Perform GridSearchCV
grid_search = GridSearchCV(estimator=logreg, param_grid=param_grid,
                           scoring='accuracy', cv=5, verbose=2)
grid_search.fit(x_train, y_train)

# Best parameters and best score
print("Best Parameters:", grid_search.best_params_)
print("Best Cross-Validated Accuracy:", grid_search.best_score_)

Fitting 5 folds for each of 30 candidates, totalling 150 fits
[CV] END .....C=0.01, penalty=l1, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l1, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=liblinear; total
time= 0.0s
```

```
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=l2, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=elasticnet, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=elasticnet, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=elasticnet, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=elasticnet, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=elasticnet, solver=liblinear; total
time= 0.0s
[CV] END .....C=0.01, penalty=elasticnet, solver=saga; total
time= 0.0s
[CV] END .....C=0.01, penalty=elasticnet, solver=saga; total
time= 0.0s
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Best Parameters: {'C': 10, 'penalty': 'l2', 'solver': 'saga'}
Best Cross-Validated Accuracy: 0.9758241758241759

```

*# Evaluate the best model on the test set*

```

best_logreg_model = grid_search.best_estimator_
test_accuracy = best_logreg_model.score(x_test, y_test)
print("Test Accuracy of Best Logistic Regression Model:",
test_accuracy)

```

Test Accuracy of Best Logistic Regression Model: 0.9824561403508771

## SAVING THE MODEL

*# Save the tuned logistic regression model*

```

joblib.dump(best_logreg_model,
'classification_lr_model(breast_cancer).joblib')
print("Model saved as classification_lr_model(breast_cancer).joblib.")

```

Model saved as classification\_lr\_model(breast\_cancer).joblib.