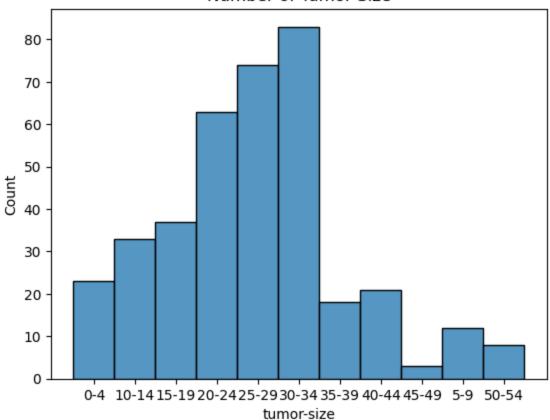
```
In [124... import pandas as pd
         import sklearn
         import seaborn as sns
         import matplotlib.pyplot as plt
         from sklearn.model_selection import train_test_split, GridSearchCV
         from sklearn import datasets
         from sklearn.linear_model import SGDClassifier
         from sklearn.metrics import accuracy_score, recall_score, precision_score, f
         from sklearn.neighbors import KNeighborsClassifier
         import numpy as np
         import warnings
         from sklearn.preprocessing import LabelEncoder
         from sklearn.metrics import classification report
In [125... | warnings.filterwarnings('ignore')
         # Read in our dataset
         cancer_data = pd.read_csv('project1.csv')
In [125... | # Identify the shape of raw data
         cancer_data.shape
Out[125... (386, 10)
In [125... # Identify the sie of raw data
         cancer_data.size
Out[125... 3860
In [125... # Get info about datatype
         cancer_data.info()
        <class 'pandas.core.frame.DataFrame'>
        RangeIndex: 386 entries, 0 to 385
        Data columns (total 10 columns):
             Column
                          Non-Null Count Dtype
         0
                          386 non-null
             class
                                          object
         1
             age
                          386 non-null
                                          object
                          386 non-null
         2
             menopause
                                          object
         3
            tumor-size 385 non-null
                                          object
            inv-nodes 385 non-null
                                          object
         5
             node-caps
                          386 non-null
                                          object
            deg-malig
                          386 non-null int64
         7
             breast
                          386 non-null
                                          object
         8
             breast-quad 386 non-null
                                          object
             irradiat
                          386 non-null
                                          object
        dtypes: int64(1), object(9)
        memory usage: 30.3+ KB
In [125... # cast column vals to type category
         cancer data['class'] = cancer data['class'].astype("category")
         cancer_data['age'] = cancer_data['age'].astype("category")
         cancer_data['menopause'] = cancer_data['menopause'].astype("category")
         cancer_data['tumor-size'] = cancer_data['tumor-size'].astype("category")
```

```
cancer_data['inv-nodes'] = cancer_data['inv-nodes'].astype("category")
         cancer_data['node-caps'] = cancer_data['node-caps'].astype("category")
         cancer data['breast'] = cancer data['breast'].astype("category")
         cancer_data['breast-quad'] = cancer_data['breast-quad'].astype("category")
         cancer_data['irradiat'] = cancer_data['irradiat'].astype("category")
In [125... # Check if any duplicate rows exist in the data
         cancer_data.duplicated().sum()
Out [125... 11
In [125... # Treat duplicated rows
         cancer_data = cancer_data.drop_duplicates()
In [125... # Check to see if any missing data exists in the dataset
         cancer data.isnull().sum()
Out[125... class
                         0
                         0
          age
          menopause
          tumor-size
                         1
          inv-nodes
                         1
          node-caps
                         0
          deg-malig
          breast
                         0
          breast-quad
          irradiat
          dtype: int64
In [125... # Treating missing values from 'tumor-size' and 'inv-nodes' columns with the
         cancer data['tumor-size'] = cancer data.groupby(['breast-quad'])['tumor-size']
         cancer data['inv-nodes'] = cancer data.groupby(['breast-quad'])['inv-nodes']
In [125... # Histogram Counting the tumor size
         sns.histplot(data=cancer_data['tumor-size'])
         plt.title("Number of Tumor Size")
Out[125... Text(0.5, 1.0, 'Number of Tumor Size')
```

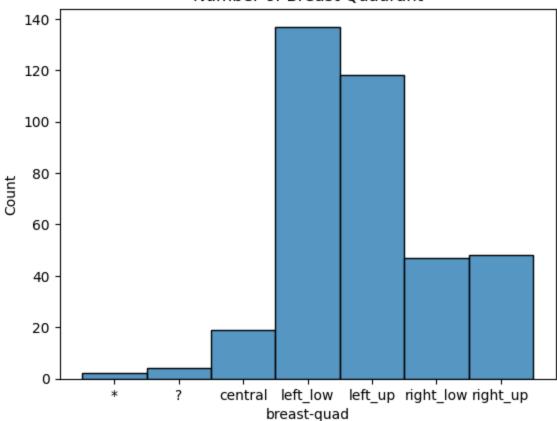
Number of Tumor Size



In [126... # Histogram Counting the location of the tumor
 sns.histplot(data=cancer_data['breast-quad'])
 plt.title("Number of Breast Quadrant")

Out[126... Text(0.5, 1.0, 'Number of Breast Quadrant')

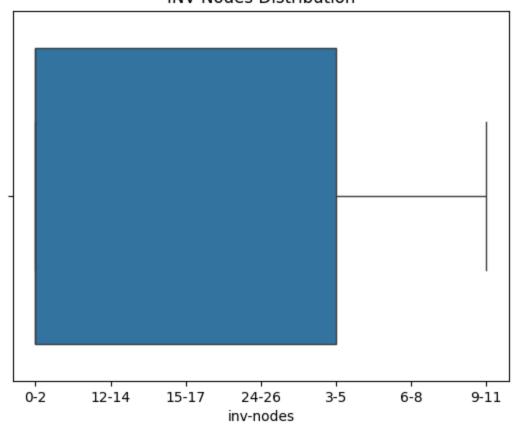
Number of Breast Quadrant



```
In [126... # Box Plot for Inv-Nodes Column
    sns.boxplot(data=cancer_data, x="inv-nodes")
    plt.title("INV Nodes Distribution")
```

Out[126... Text(0.5, 1.0, 'INV Nodes Distribution')

INV Nodes Distribution



In [126... # Performing One hot encoding on Categorical Columns
 cancer_data = pd.get_dummies(cancer_data, columns=["class", "menopause", "ac
 cancer_data.info()

> <class 'pandas.core.frame.DataFrame'> Index: 375 entries, 0 to 385 Data columns (total 36 columns):

```
Column
                             Non-Null Count Dtype
___
    _____
                             _____
                                             ____
0
                             375 non-null
                                             int64
    deg-malig
1
    class_recurrence-events 375 non-null
                                             bool
2
    menopause_lt40
                             375 non-null
                                             bool
 3
    menopause premeno
                             375 non-null
                                             bool
4
    age 30-39
                             375 non-null
                                             bool
5
    age_40-49
                             375 non-null
                                             bool
6
    age 50-59
                             375 non-null
                                             bool
7
                             375 non-null
    age 60-69
                                             bool
8
    age 70-79
                             375 non-null
                                             bool
9
    tumor-size 10-14
                             375 non-null
                                             bool
10 tumor-size 15-19
                             375 non-null
                                             bool
11 tumor-size_20-24
                             375 non-null
                                             bool
12 tumor-size_25-29
                             375 non-null
                                             bool
13 tumor-size 30-34
                             375 non-null
                                             bool
14 tumor-size 35-39
                             375 non-null
                                             bool
15 tumor-size_40-44
                             375 non-null
                                             bool
16 tumor-size 45-49
                             375 non-null
                                             bool
17 tumor-size 5-9
                             375 non-null
                                             bool
18 tumor-size_50-54
                             375 non-null
                                             bool
19 inv-nodes 12-14
                             375 non-null
                                             bool
20 inv-nodes 15-17
                             375 non-null
                                             bool
21 inv-nodes_24-26
                             375 non-null
                                             bool
22 inv-nodes 3-5
                             375 non-null
                                             bool
23 inv-nodes 6-8
                             375 non-null
                                             bool
24 inv-nodes_9-11
                             375 non-null
                                             bool
25 node-caps ?
                             375 non-null
                                             bool
26 node-caps no
                             375 non-null
                                             bool
27 node-caps_yes
                             375 non-null
                                             bool
28 breast right
                             375 non-null
                                             bool
29 breast-quad_?
                             375 non-null
                                             bool
30 breast-quad_central
                             375 non-null
                                             bool
31 breast-quad left low
                             375 non-null
                                             bool
32 breast-quad_left_up
                             375 non-null
                                             bool
33 breast-quad_right_low
                             375 non-null
                                             bool
34 breast-quad_right_up
                             375 non-null
                                             bool
35 irradiat yes
                             375 non-null
                                             bool
dtypes: bool(35), int64(1)
```

memory usage: 18.7 KB

```
In [126... # Assigning subset of data as X and y
         X = cancer_data.drop('class_recurrence-events', axis=1)
         y = cancer data["class recurrence-events"]
```

```
In [126... # Linear Classification
         X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.3, r
         clf = SGDClassifier(loss="perceptron", alpha=0.1)
         clf.fit(X_train, y_train)
```

```
Out [126...
```

SGDClassifier

SGDClassifier(alpha=0.1, loss='perceptron')

```
In [126... # Accuracy for Linear Classification
         accuracy_test = accuracy_score(y_test, clf.predict(X_test))
         accuracy_train = accuracy_score(y_train, clf.predict(X_train))
         print(f"Test Accuracy: {accuracy test:.4f}")
         print(f"Train Accuracy: {accuracy train:.4f}\n")
         # Recall for Linear Classification
         recall_test = recall_score(y_test, clf.predict(X_test), average='weighted')
         recall_train = recall_score(y_train, clf.predict(X_train), average='weighted
         print(f"Test Recall: {recall test:.4f}")
         print(f"Train Recall: {recall train:.4f}\n")
         # Precision for Linear Classification
         precision_test = precision_score(y_test, clf.predict(X_test), average='weighted")
         precision_train = precision_score(y_train, clf.predict(X_train), average='we
         print(f"Test Precision: {precision test:.4f}")
         print(f"Train Precision: {precision train:.4f}\n")
         # F-1 Score for Linear Classification
         f1_test = f1_score(y_test, clf.predict(X_test), average='weighted')
         f1_train = f1_score(y_train, clf.predict(X_train), average='weighted')
         print(f"Test F1 Score: {f1_test:.4f}")
         print(f"Train F1 Score: {f1 train:.4f}\n")
         print(classification_report(y_test, clf.predict(X_test)))
```

Test Accuracy: 0.5398
Train Accuracy: 0.6031

Test Recall: 0.5398
Train Recall: 0.6031

Test Precision: 0.6558
Train Precision: 0.7092

Test F1 Score: 0.5526 Train F1 Score: 0.6112

	precision	recall	f1-score	support
False	0.78	0.46	0.58	78
True	0.37	0.71	0.49	35
accuracy			0.54	113
macro avg	0.58	0.59	0.54	113
weighted avg	0.66	0.54	0.55	113

```
In [126... # Instantiate the model for K-Nearest Neighbor Classifier
knn = KNeighborsClassifier(n_neighbors=3)
knn.fit(X_train, y_train)
```

Out [126...

KNeighborsClassifier

KNeighborsClassifier(n_neighbors=3)

```
In [126... # Accuracy for K-Nearest Neighbor Classifier
         accuracy_test = accuracy_score(y_test, knn.predict(X_test))
         accuracy_train = accuracy_score(y_train, knn.predict(X_train))
         print(f"Test Accuracy: {accuracy test:.4f}")
         print(f"Train Accuracy: {accuracy_train:.4f}\n")
         # Recall for K-Nearest Neighbor Classifier
         recall_test = recall_score(y_test, knn.predict(X_test), average='weighted')
         recall_train = recall_score(y_train, knn.predict(X_train), average='weighted
         print(f"Test Recall: {recall test:.4f}")
         print(f"Train Recall: {recall_train:.4f}\n")
         # Precision for K-Nearest Neighbor Classifier
         precision_test = precision_score(y_test, knn.predict(X_test), average='weight

         precision train = precision score(y train, knn.predict(X train), average='we
         print(f"Test Precision: {precision_test:.4f}")
         print(f"Train Precision: {precision_train:.4f}\n")
         # F-1 Score for K-Nearest Neighbor Classifier
         f1_test = f1_score(y_test, knn.predict(X_test), average='weighted')
         f1 train = f1 score(y train, knn.predict(X train), average='weighted')
         print(f"Test F1 Score: {f1_test:.4f}")
         print(f"Train F1 Score: {f1 train:.4f}\n")
         print(classification_report(y_test, knn.predict(X_test)))
```

Test Accuracy: 0.6460 Train Accuracy: 0.7672

Test Recall: 0.6460 Train Recall: 0.7672

Test Precision: 0.5988
Train Precision: 0.7592

Test F1 Score: 0.6100 Train F1 Score: 0.7575

	precision	recall	f1-score	support
False	0.70	0.85	0.77	78
True	0.37	0.20	0.26	35
accuracy			0.65	113
macro avg	0.54	0.52	0.51	113
weighted avg	0.60	0.65	0.61	113

```
In [126... knn_search_accuracy = KNeighborsClassifier()
    param_grid = {"n_neighbors": np.arange(1, 100)}
    knn_gscv = GridSearchCV(knn_search_accuracy, param_grid, cv=5)
    knn_gscv.fit(X_train, y_train)
    best_knn = knn_gscv.best_estimator_
```

```
In [126... knn_search_recall = KNeighborsClassifier()
    knn_gscv2 = GridSearchCV(knn_search_recall, param_grid, cv=5, scoring="recall
    knn_gscv2.fit(X_train, y_train)
    best_recall = knn_gscv2.best_estimator_
```

```
In [127... # Recall for K-Nearest Neighbor Classifier using Grid search CV
models = [knn, best_knn, best_recall]
for m in models:

    recall_test=recall_score(y_test, m.predict(X_test), average='weighted')
    print(f'Recall of {m} on test data is : {recall_test}')
    recall_train=recall_score(y_train, m.predict(X_train), average='weighted
    print(f'Recall of {m} on train data is : {recall_train}\n')
```

```
Recall of KNeighborsClassifier(n_neighbors=3) on test data is: 0.6460176991 150443

Recall of KNeighborsClassifier(n_neighbors=3) on train data is: 0.767175572 519084

Recall of KNeighborsClassifier(n_neighbors=12) on test data is: 0.672566371 681416

Recall of KNeighborsClassifier(n_neighbors=12) on train data is: 0.68702290 07633588

Recall of KNeighborsClassifier(n_neighbors=1) on test data is: 0.5486725663 716814

Recall of KNeighborsClassifier(n_neighbors=1) on train data is: 0.988549618 3206107
```

```
In [127... # Accuracy for K-Nearest Neighbor Classifier using Grid search CV
         accuracy_test = accuracy_score(y_test, best_knn.predict(X_test))
         accuracy train = accuracy score(y train, best knn.predict(X train))
         print(f"Test Accuracy: {accuracy_test:.4f}")
         print(f"Train Accuracy: {accuracy train:.4f}\n")
         # Precision for K-Nearest Neighbor Classifier using Grid search CV
         precision test = precision score(y test, best knn.predict(X test), average='
         precision_train = precision_score(y_train, best_knn.predict(X_train), averag
         print(f"Test Precision: {precision test:.4f}")
         print(f"Train Precision: {precision train:.4f}\n")
         # F-1 Score for K-Nearest Neighbor Classifier using Grid search CV
         f1 test = f1 score(y test, best knn.predict(X test), average='weighted')
         f1_train = f1_score(y_train, best_knn.predict(X_train), average='weighted')
         print(f"Test F1 Score: {f1 test:.4f}")
         print(f"Train F1 Score: {f1 train:.4f}\n")
         print(classification report(y test, best knn.predict(X test)))
```

Test Accuracy: 0.6726 Train Accuracy: 0.6870

Test Precision: 0.5524
Train Precision: 0.6600

Test F1 Score: 0.5696 Train F1 Score: 0.6223

	precision	recall	f1-score	support
False True	0.69 0.25	0.96 0.03	0.80 0.05	78 35
accuracy macro avg weighted avg	0.47 0.55	0.50 0.67	0.67 0.43 0.57	113 113 113