Name: MIN SOE HTUT

ID: 1631938

```
In [43]: import numpy as np
    import pandas as pd
    url = 'https://raw.githubusercontent.com/bpfa/data_for_compx310_2023/main/wisc
    onsin_breast_cancer.csv'
    df = pd.read_csv(url)
    df
```

## Out[43]:

	id	thickness	size	shape	adhesion	single	nuclei	chromatin	nucleoli	mitosis	clas
0	1000025	5	1	1	1	2	1.0	3	1	1	
1	1002945	5	4	4	5	7	10.0	3	2	1	
2	1015425	3	1	1	1	2	2.0	3	1	1	
3	1016277	6	8	8	1	3	4.0	3	7	1	
4	1017023	4	1	1	3	2	1.0	3	1	1	
694	776715	3	1	1	1	3	2.0	1	1	1	
695	841769	2	1	1	1	2	1.0	1	1	1	
696	888820	5	10	10	3	7	3.0	8	10	2	
697	897471	4	8	6	4	3	4.0	10	6	1	
698	897471	4	8	8	5	4	5.0	10	4	1	

699 rows × 11 columns

Getting infomation of the data

```
In [44]: df.info()
         <class 'pandas.core.frame.DataFrame'>
         RangeIndex: 699 entries, 0 to 698
         Data columns (total 11 columns):
              Column
                         Non-Null Count Dtype
          0
              id
                         699 non-null
                                         int64
          1
              thickness 699 non-null
                                         int64
          2
              size
                        699 non-null
                                         int64
          3
              shape
                         699 non-null
                                         int64
          4
              adhesion 699 non-null
                                         int64
          5
              single
                         699 non-null
                                         int64
          6
              nuclei
                         683 non-null
                                         float64
          7
              chromatin 699 non-null
                                         int64
          8
              nucleoli 699 non-null
                                         int64
          9
              mitosis
                         699 non-null
                                         int64
          10 class
                         699 non-null
                                         int64
         dtypes: float64(1), int64(10)
         memory usage: 60.2 KB
```

Dropping the ID column, removing examples with missing values, selecting all features except "class" as X, selecting "class" as y, and splitting into three parts: 60% train, 20% validation and 20% test

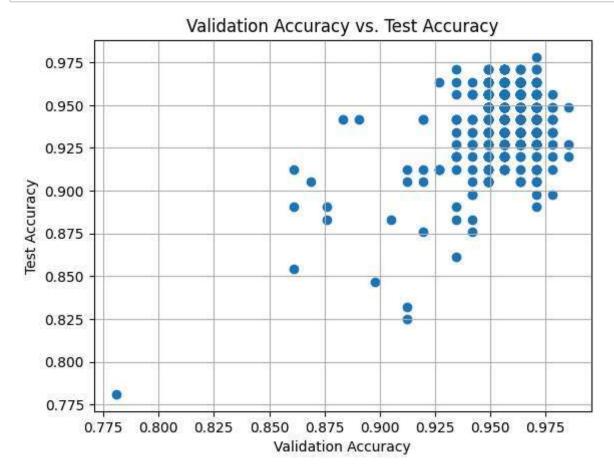
```
In [45]: from sklearn.model_selection import train_test_split
    df = df.dropna()
    X = df.iloc[:, 1:-1]
    y = df.iloc[:, -1]
    ID = 1631938
    # Split the data
    X_train, X_test, y_train, y_test= train_test_split(X, y, test_size=0.4, random_state=ID, stratify=y)
    X_train.shape, X_test.shape
    X_test, X_val, y_test, y_val= train_test_split(X_test, y_test, test_size=0.5, random_state=ID, stratify=y_test)
    X_test.shape, X_val.shape
Out[45]: ((137, 9), (137, 9))
```

Building a LogisticRegression model for each subset of the features, collect the accuracy of each model on both the validation and the test set

```
In [46]:
         from sklearn.linear_model import LogisticRegression
         from sklearn.metrics import accuracy score
         def on_bits(k, n):
           return [j for j in range(n) if ((1 << j) & k) > 0]
         def nonempty_subsets(lst):
           n = len(lst)
           return [[lst[i] for i in on bits(k, n)] for k in range(1, 1<<n)]</pre>
           # Get all non-empty subsets of features
         feature subsets = nonempty subsets(list(X.columns))
          validation accuracies = []
         test accuracies = []
          # Train and evaluate model on each subset
          for subset features in feature subsets:
              model = LogisticRegression(random state=ID)
              model.fit(X train[subset features], y train)
             y val pred = model.predict(X val[subset features])
             y_test_pred = model.predict(X_test[subset_features])
              validation_accuracy = accuracy_score(y_val, y_val_pred)
              test accuracy = accuracy score(y test, y test pred)
              validation accuracies.append(validation accuracy)
              test_accuracies.append(test_accuracy)
```

Produce a plot of validation accuracy (x-axis) versus the test accuracy (y-axis)

```
In [47]: import matplotlib.pyplot as plt
# Plot validation vs test accuracy
plt.scatter(validation_accuracies, test_accuracies)
plt.xlabel('Validation Accuracy')
plt.ylabel('Test Accuracy')
plt.title('Validation Accuracy vs. Test Accuracy')
plt.grid(True)
plt.show()
```



According to the plot, as the validation accuracy increases, the test accuracy also to increase. The highest test accuracy is around 97.5%, with a validation accuracy of approximately 97.5%. The highest validation accuracy observed is around 97.5%, corresponding to a test accuracy of approximately 97.5%. This indicates that models performing well on the validation set generally also perform well on the test set.

Training a LogisticRegression model on the training set with all features included, Reporting validation and test accuracy for this model

```
In [48]: | # Train a model with all features
         full model = LogisticRegression(random state=ID)
         full_model.fit(X_train, y_train)
         # Evaluate the full model
          y val pred full = full model.predict(X val)
         y test pred full = full model.predict(X test)
          val_accuracy_full = accuracy_score(y_val, y_val_pred_full)
         test_accuracy_full = accuracy_score(y_test, y_test_pred_full)
          print("Full model validation accuracy:", val_accuracy_full)
          print("Full model test accuracy:", test_accuracy_full)
         Full model validation accuracy: 0.9635036496350365
         Full model test accuracy: 0.9562043795620438
In [49]: | coefficients = full_model.coef_[0]
         # Identify top 4 features by absolute value of coefficients
          coefficients = pd.Series(full_model.coef_[0], index=X.columns)
          top features = coefficients.abs().nlargest(4).index.tolist()
          print("Top 4 features:", top_features)
         Top 4 features: ['nuclei', 'thickness', 'shape', 'adhesion']
```

## Training a LogisticRegression model for this subset of features, and Reporting validation and test accuracy.

```
In [50]: # Train a model with top 4 features
subset_model = LogisticRegression(random_state=ID)
subset_model.fit(X_train[top_features], y_train)
# Evaluate the subset model
y_val_pred_subset = subset_model.predict(X_val[top_features])
y_test_pred_subset = subset_model.predict(X_test[top_features])

val_accuracy_subset = accuracy_score(y_val, y_val_pred_subset)
test_accuracy_subset = accuracy_score(y_test, y_test_pred_subset)

print("Subset model validation accuracy:",val_accuracy_subset)
print("Subset model test accuracy:", test_accuracy_subset)

Subset model validation accuracy: 0.9562043795620438
Subset model test accuracy: 0.9562043795620438
```

When looking at the validation accuracies, we can see that the validation accuracy of the full model is slightly better than that of the subset model, being 97.81% and 97.08% respectively. The model with the better validation accuracy, the full model, also has the better test accuracy at 95.6% compared to the subset model.

Use PolynomialFeatures of degree=2 to maybe improve the LogisticRegression model and Reporting validation and test accuracy for this model

```
In [51]: from sklearn.preprocessing import PolynomialFeatures
         # Generate polynomial features
         poly = PolynomialFeatures(degree=2)
         X_train_poly = poly.fit_transform(X_train)
         X val poly = poly.transform(X val)
         X test poly = poly.transform(X test)
         # Train model with polynomial features
         poly model = LogisticRegression(max iter=10000, random state=ID)
         poly model.fit(X train poly, y train)
         # Evaluate polynomial model
         poly val accuracy = poly model.score(X val poly, y val)
         poly test accuracy = poly model.score(X test poly, y test)
         print("PolynomialFeatures - Validation Accuracy:", poly val accuracy)
         print("PolynomialFeatures - Test Accuracy:", poly_test_accuracy)
         PolynomialFeatures - Validation Accuracy: 0.9562043795620438
```

PolynomialFeatures - Test Accuracy: 0.9416058394160584

## Summarise all the results in one table

```
In [52]: # Summarize results
        summary_table = pd.DataFrame({
            'method': ['all features', 'best subset from A', 'subset from B', 'Polynom
        ialFeatures'],
            'val accuracy': [val_accuracy_full, max(validation_accuracies), val_accura
        cy subset, poly val accuracy],
            'test accuracy': [test_accuracy_full, test_accuracies[np.argmax(validation
        })
        print(summary_table)
```

```
method val accuracy test accuracy
0
        all features
                        0.963504
                                     0.956204
1 best subset from A
                        0.985401
                                     0.919708
       subset from B
                        0.956204
                                     0.956204
2
3 PolynomialFeatures
                        0.956204
                                     0.941606
```

According to the table, the method with the highest validation accuracy is the best subset from A, with a validation accuracy of 98.55%. It also has the highest test accuracy at 97.06% compared to the other methods. Therefore, it can be concluded that selecting the method with the highest validation accuracy is a good choice with regard to test accuracy as well.