

In [1]: import numpy as np import pandas as pd import matplotlib.pyplot as plt import seaborn as sns import warnings warnings.simplefilter("ignore")

Out[2]:

| | Gene One | Gene Two | Cancer Present |
|---|----------|----------|----------------|
| 0 | 4.3 | 3.9 | 1 |
| 1 | 2.5 | 6.3 | 0 |
| 2 | 5.7 | 3.9 | 1 |
| 3 | 6.1 | 6.2 | 0 |
| 4 | 7.4 | 3.4 | 1 |

In [3]: ► df.info()

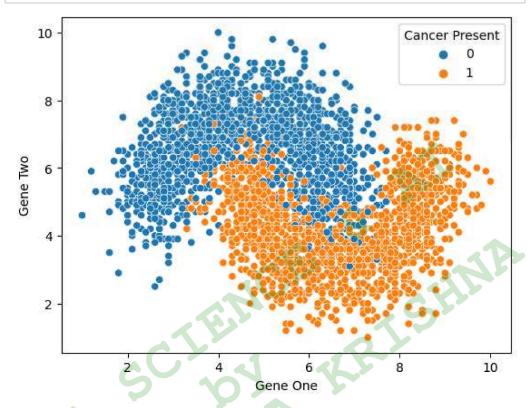
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3000 entries, 0 to 2999
Data columns (total 3 columns):

dtypes: float64(2), int64(1)

memory usage: 70.4 KB

In [4]: In sns.scatterplot(x='Gene One',y='Gene Two',hue='Cancer Present',data=di
plt.show()





```
In [5]: M df.isnull().sum()
```

Out[5]: Gene One 0
Gene Two 0
Cancer Present 0
dtype: int64

X & y

```
In [6]:  X = df.drop('Cancer Present',axis=1)
y = df['Cancer Present']
```

Train|Test Split

Scaling Data

Hyperparameter Tuning for KNN Classifier

KNN model with best hyper parameters

```
#Modelling
In [10]:
            from sklearn.neighbors import KNeighborsClassifier
            knn = KNeighborsClassifier(n_neighbors=19)
            knn.fit(X train,y train)
            # Prediction
            y_pred_test = knn.predict(X_test)
            y_pred_train = knn.predict(X_train)
            #Evaluation
            from sklearn.metrics import accuracy score
            print("Train accuracy:",accuracy_score(y_train,y_pred_train))
            print("Test accuracy:",accuracy_score(y_test,y_pred_test))
            from sklearn.model selection import cross val score
            print("Cross Validation Score:",cross_val_score(knn,X,y,cv=5).mean())
            Train accuracy: 0.94083333333333333
            Cross Validation Score: 0.9313333333333333
          ▶ | from sklearn.metrics import confusion matrix
In [11]:
            confusion_matrix(y_test,y_pred_test)
   Out[11]: array([[264, 19],
                   [ 21, 296]], dtype=int64)
```




| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 0.93 | 0.93 | 0.93 | 283 |
| 1 | 0.94 | 0.93 | 0.94 | 317 |
| | | | | |
| accuracy | | | 0.93 | 600 |
| macro avg | 0.93 | 0.93 | 0.93 | 600 |
| weighted avg | 0.93 | 0.93 | 0.93 | 600 |

Prediction on New data

New Data

Out[13]:

| | Gene One | Gene Two |
|---|----------|----------|
| 0 | 4.9 | 3.9 |

preprocess the new data

use KNN model to predict on new data

```
In [15]: H knn.predict(df_scaled)
```

Out[15]: array([1], dtype=int64)