



SRI RAMACHANDRA
INSTITUTE OF HIGHER EDUCATION AND RESEARCH
(Category - I Deemed to be University) Porur, Chennai
SRI RAMACHANDRA ENGINEERING AND TECHNOLOGY

LUNG CANCER PREDICTION

CA 4 REPORT

Quarter IV (Year 1)

Submitted by

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E0220201

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in

COMPUTER SCIENCE AND ENGINEERING

(Artificial Intelligence & Machine Learning)

Sri Ramachandra Engineering and Technology

Sri Ramachandra Institute of Higher Education and Research, Porur, Chennai -600116

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BONAFIDE CERTIFICATE

Certified that this project report “**Lung Cancer Prediction**” is the bonafide work of **Joselyn Diana Cindrella M, Keerthivasan S R** Reg No. **E0120017, E0220201** who carried out the internship work under my supervision.

Signature of Faculty Mentor Signature of Vice-Principal

Chennai-600116

Prof.Rajive Gandhi

Evaluation Date:

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I wish to thank my faculty Prof. Rajive Gandhi Department of Computer Science and Engineering, Sri Ramachandra Engineering and Technology for extending help and encouragement throughout the project. Without his/her continuous guidance and persistent help, this project would not have been a success for me.

I am grateful to the Department of Computer Science and Engineering, Sri Ramachandra Engineering and Technology, our beloved parents, and friends for extending support, who helped us to overcome obstacles in the study.

1. INTRODUCTION:

Lung cancer, also known as lung carcinoma, is a malignant lung tumor characterized by uncontrolled cell growth in tissues of the lung. This growth can spread beyond the lung by the process of metastasis into nearby tissue or other parts of the body.

In this study, we tried to predict Lung Cancer using 6 different algorithms:

- 1) Logistic regression classification
- 2) Naive Bayes classification
- 3) Decision tree classification
- 4) Random forest classification
- 5) K-Nearest Neighbor classification

2. ABOUT THE DATASET:

This Dataset is taken from Kaggle

<https://www.kaggle.com/datasets/yusufdede/lung-cancer-dataset>

The dataset has 7 columns and 59 rows

The columns include Name, Surname, Age, Smoking Status, Area Q, Alcohol, and Result.

3. CODE:

Importing the Required Libraries

```
In [10]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

from sklearn import preprocessing
from sklearn.preprocessing import MinMaxScaler

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

Reading the dataset

```
In [11]: data = pd.read_csv('C:/Users/Joselyn Cindrella M/Desktop/lung_cancer_examples.csv')
data.head()
```

```
Out[11]:
```

| | Name | Surname | Age | Smokes | AreaQ | Alkhol | Result |
|---|--------|-------------|-----|--------|-------|--------|--------|
| 0 | John | Wick | 35 | 3 | 5 | 4 | 1 |
| 1 | John | Constantine | 27 | 20 | 2 | 5 | 1 |
| 2 | Camela | Anderson | 30 | 0 | 5 | 2 | 0 |
| 3 | Alex | Telles | 28 | 0 | 8 | 1 | 0 |
| 4 | Diego | Maradona | 68 | 4 | 5 | 6 | 1 |

```
In [12]: print('Dataset :',data.shape)

Dataset : (59, 7)
```

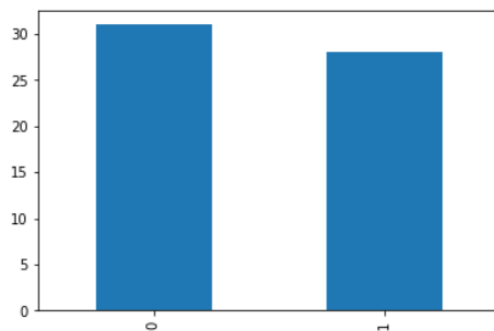
```
In [13]: data[0:10]
```

```
Out[13]:
```

| | Name | Surname | Age | Smokes | AreaQ | Alkhol | Result |
|---|-----------|-------------|-----|--------|-------|--------|--------|
| 0 | John | Wick | 35 | 3 | 5 | 4 | 1 |
| 1 | John | Constantine | 27 | 20 | 2 | 5 | 1 |
| 2 | Camela | Anderson | 30 | 0 | 5 | 2 | 0 |
| 3 | Alex | Telles | 28 | 0 | 8 | 1 | 0 |
| 4 | Diego | Maradona | 68 | 4 | 5 | 6 | 1 |
| 5 | Cristiano | Ronaldo | 34 | 0 | 10 | 0 | 0 |
| 6 | Mihail | Tal | 58 | 15 | 10 | 0 | 0 |
| 7 | Kathy | Bates | 22 | 12 | 5 | 2 | 0 |
| 8 | Nicole | Kidman | 45 | 2 | 6 | 0 | 0 |
| 9 | Ray | Milland | 52 | 18 | 4 | 5 | 1 |

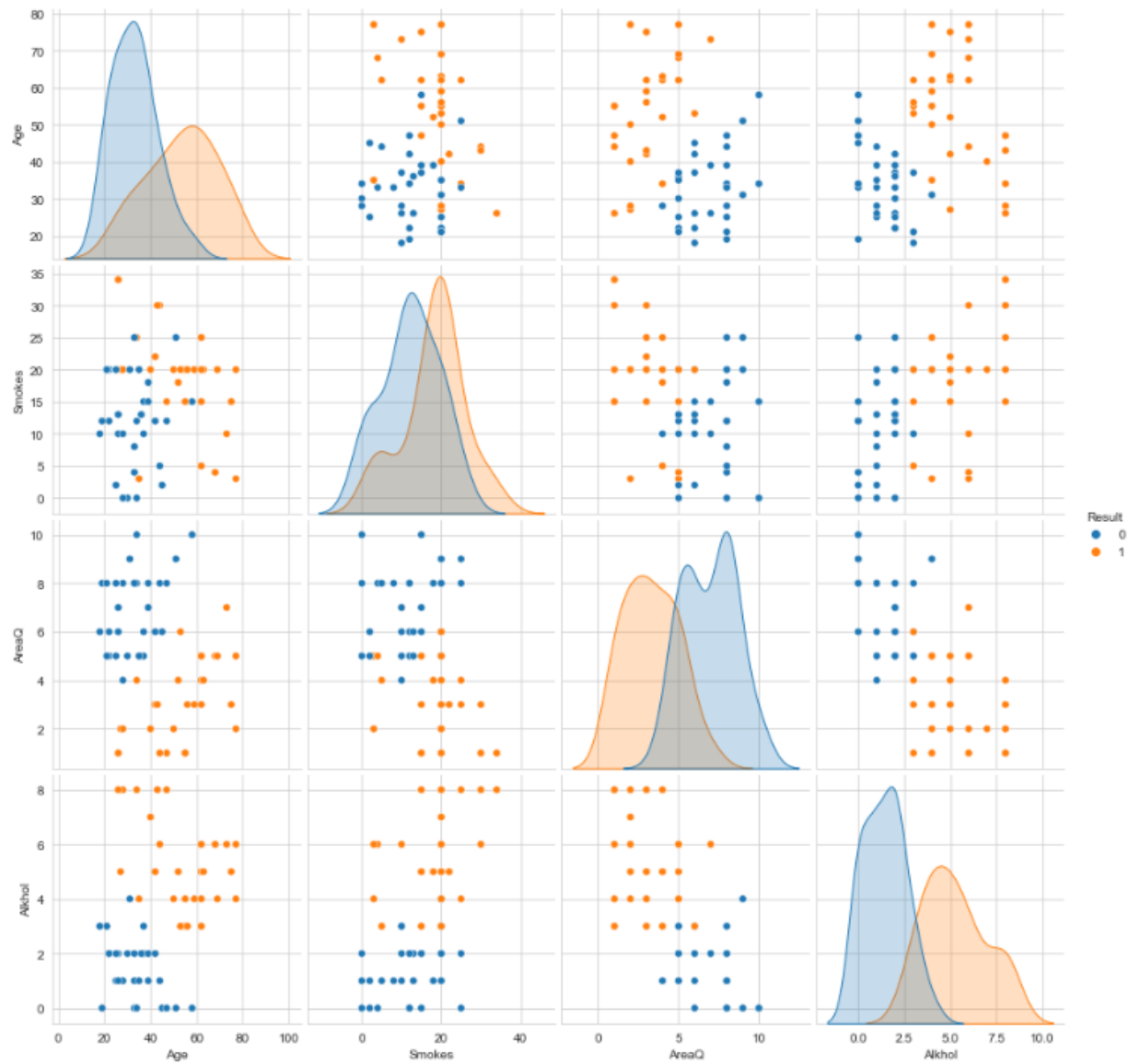
Distribution of Diagnosis

```
In [14]: data.Result.value_counts()[0:30].plot(kind='bar')
plt.show()
```



Visualizing the data

```
In [15]: sns.set_style("whitegrid")
sns.pairplot(data,hue="Result",size=3);
plt.show()
```



Removing irrelevant variables

```
In [17]: data1 = data.drop(columns=['Name', 'Surname'],  
                             axis=1)  
data1 = data1.dropna(how='any')  
print(data1.shape)  
data1.head()
```

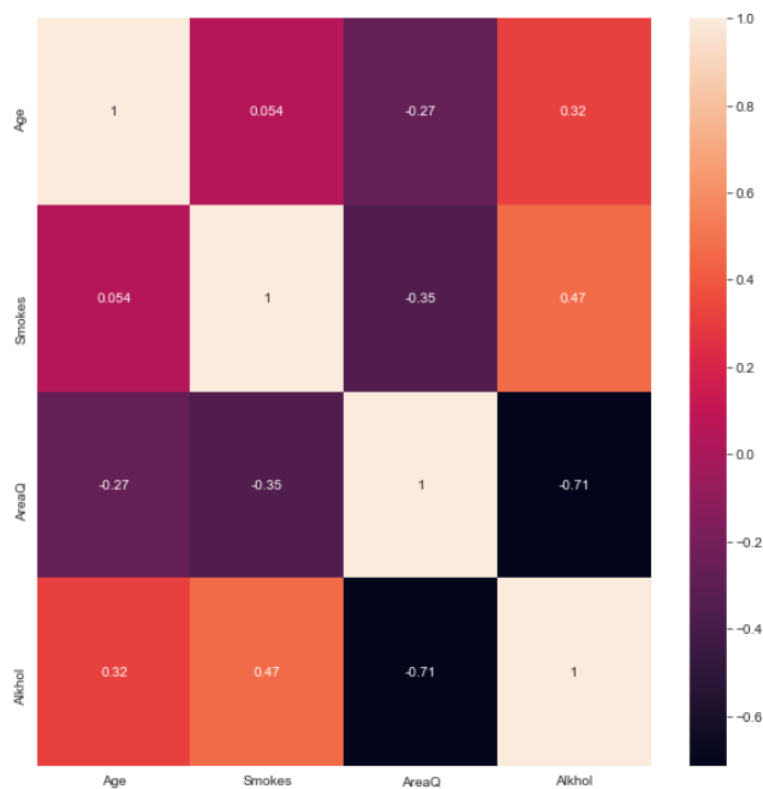
(59, 5)

Out[17]:

| | Age | Smokes | AreaQ | Alkhol | Result |
|---|-----|--------|-------|--------|--------|
| 0 | 35 | 3 | 5 | 4 | 1 |
| 1 | 27 | 20 | 2 | 5 | 1 |
| 2 | 30 | 0 | 5 | 2 | 0 |
| 3 | 28 | 0 | 8 | 1 | 0 |
| 4 | 68 | 4 | 5 | 6 | 1 |

Observations

```
In [18]: correlation = data1.drop(['Result'], axis=1).corr()  
plt.figure(figsize=(10, 10))  
sns.heatmap(correlation, annot=True)  
plt.show()
```



Data Training and Testing

```
In [32]: from sklearn.model_selection import train_test_split
Y = data1['Result']
X = data1.drop(columns=['Result'])
X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.50, random_state=9)
```

```
In [33]: print('X train shape: ', X_train.shape)
print('Y train shape: ', Y_train.shape)
print('X test shape: ', X_test.shape)
print('Y test shape: ', Y_test.shape)
```

```
X train shape: (29, 4)
Y train shape: (29,)
X test shape: (30, 4)
Y test shape: (30,)
```

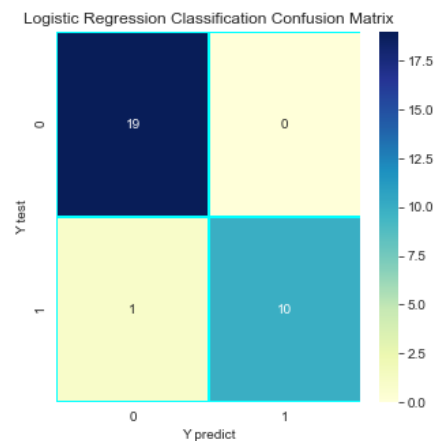
Logistic Regression Classification

```
In [34]: from sklearn.linear_model import LogisticRegression
logreg = LogisticRegression(C=10)
logreg.fit(X_train, Y_train)
Y_predict1 = logreg.predict(X_test)
```

```
In [35]: from sklearn.metrics import confusion_matrix
import seaborn as sns

logreg_cm = confusion_matrix(Y_test, Y_predict1)
f, ax = plt.subplots(figsize=(5,5))
sns.heatmap(logreg_cm, annot=True, linewidth=0.7, linecolor='cyan', fmt='g', ax=ax, cmap="YlGnBu")
plt.title('Logistic Regression Classification Confusion Matrix')
plt.xlabel('Y predict')
plt.ylabel('Y test')
plt.show()

score_logreg = logreg.score(X_test, Y_test)
print(score_logreg)
```



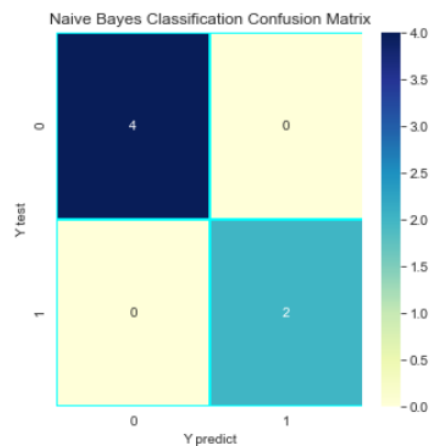
0.9666666666666667

Naive Bayes Classification

```
In [28]: from sklearn.naive_bayes import GaussianNB
nbcla = GaussianNB()
nbcla.fit(X_train, Y_train)
Y_predict3 = nbcla.predict(X_test)
```

```
In [29]: nbcla_cm = confusion_matrix(Y_test, Y_predict3)
f, ax = plt.subplots(figsize=(5,5))
sns.heatmap(nbcla_cm, annot=True, linewidth=0.7, linecolor='cyan', fmt='g', ax=ax, cmap="YlGnBu")
plt.title('Naive Bayes Classification Confusion Matrix')
plt.xlabel('Y predict')
plt.ylabel('Y test')
plt.show()

score_nbcla = nbcla.score(X_test, Y_test)
print(score_nbcla)
```



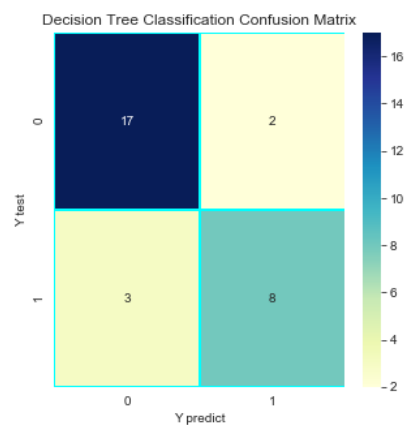
1.0

Decision Tree Classification

```
In [38]: from sklearn.tree import DecisionTreeClassifier
dtcla = DecisionTreeClassifier(random_state=9)
dtcla.fit(X_train, Y_train)
Y_predict4 = dtcla.predict(X_test)
```

```
In [39]: dtcla_cm = confusion_matrix(Y_test, Y_predict4)
f, ax = plt.subplots(figsize=(5,5))
sns.heatmap(dtcla_cm, annot=True, linewidth=0.7, linecolor='cyan', fmt='g', ax=ax, cmap="YlGnBu")
plt.title('Decision Tree Classification Confusion Matrix')
plt.xlabel('Y predict')
plt.ylabel('Y test')
plt.show()

score_dtcla = dtcla.score(X_test, Y_test)
print(score_dtcla)
```



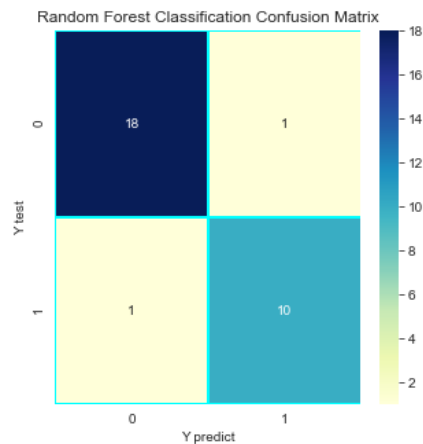
0.8333333333333334

Random forest Classification

```
In [40]: from sklearn.ensemble import RandomForestClassifier
rfcla = RandomForestClassifier(n_estimators=100, random_state=9, n_jobs=-1)
rfcla.fit(X_train, Y_train)
Y_predict5 = rfcla.predict(X_test)
```

```
In [41]: rfcla_cm = confusion_matrix(Y_test, Y_predict5)
f, ax = plt.subplots(figsize=(5,5))
sns.heatmap(rfcla_cm, annot=True, linewidth=0.7, linecolor='cyan', fmt='g', ax=ax, cmap="YlGnBu")
plt.title('Random Forest Classification Confusion Matrix')
plt.xlabel('Y predict')
plt.ylabel('Y test')
plt.show()

score_rfcla = rfcla.score(X_test, Y_test)
print(score_rfcla)
```



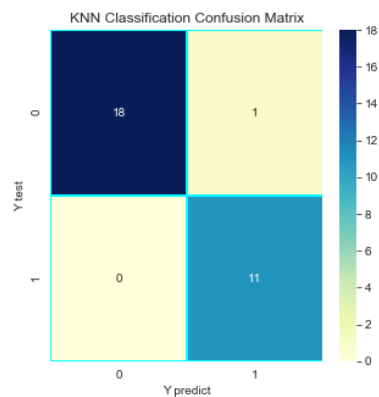
0.9333333333333333

K Nearest Neighbor Classification

```
In [42]: from sklearn.neighbors import KNeighborsClassifier
knncla = KNeighborsClassifier(n_neighbors=5, n_jobs=-1)
knncla.fit(X_train, Y_train)
Y_predict6 = knncla.predict(X_test)
```

```
In [43]: knncla_cm = confusion_matrix(Y_test, Y_predict6)
f, ax = plt.subplots(figsize=(5,5))
sns.heatmap(knncla_cm, annot=True, linewidth=0.7, linecolor='cyan', fmt='g', ax=ax, cmap="YlGnBu")
plt.title('KNN Classification Confusion Matrix')
plt.xlabel('Y predict')
plt.ylabel('Y test')
plt.show()

score_knncla = knncla.score(X_test, Y_test)
print(score_knncla)
```



0.9666666666666667

Test Score

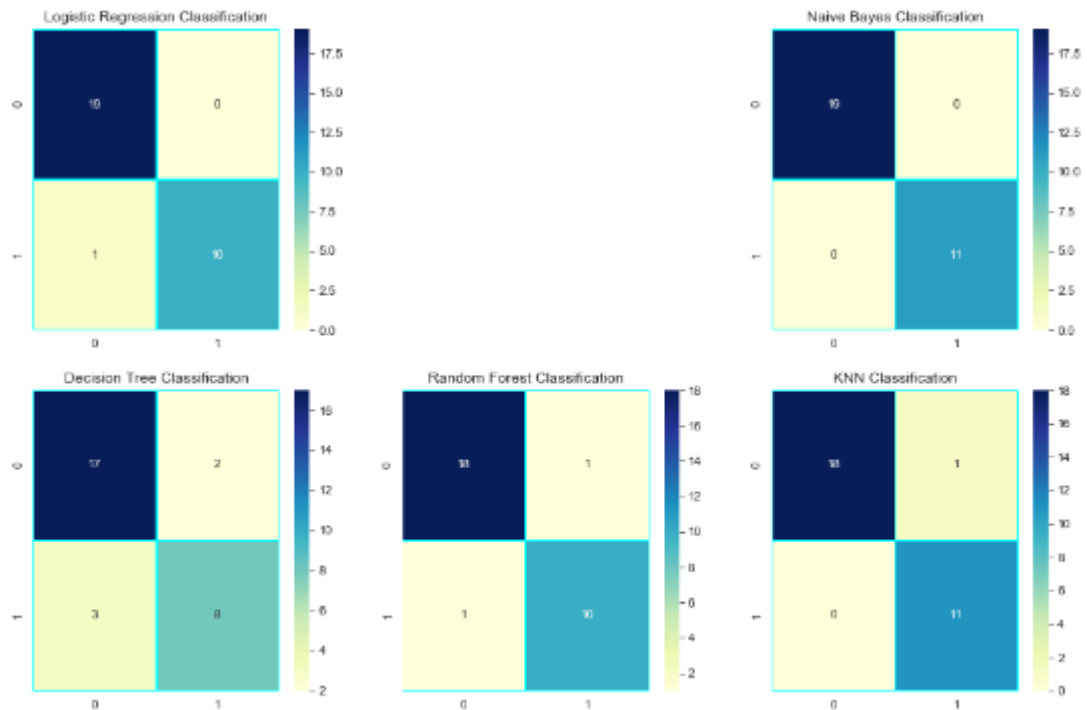
```
In [44]: Testscores = pd.Series([score_logreg, score_nbcla, score_dtcla, score_rfcla, score_knncla],
                                index=['Logistic Regression Score', 'Naive Bayes Score', 'Decision Tree Score', 'Random Forest Score', 'K-Nearest Neighbour Score'],
                                dtype=float64)

print(Testscores)
```

| | |
|---------------------------|----------|
| Logistic Regression Score | 0.966667 |
| Naive Bayes Score | 1.000000 |
| Decision Tree Score | 0.833333 |
| Random Forest Score | 0.933333 |
| K-Nearest Neighbour Score | 0.966667 |

The confusion matrix

```
In [45]: fig = plt.figure(figsize=(15,15))
ax1 = fig.add_subplot(3, 3, 1)
ax1.set_title('Logistic Regression Classification')
ax2 = fig.add_subplot(3, 3, 3)
ax2.set_title('Naive Bayes Classification')
ax3 = fig.add_subplot(3, 3, 4)
ax3.set_title('Decision Tree Classification')
ax4 = fig.add_subplot(3, 3, 5)
ax4.set_title('Random Forest Classification')
ax5 = fig.add_subplot(3, 3, 6)
ax5.set_title('KNN Classification')
sns.heatmap(data=logreg_cm, annot=True, linewidth=0.7, linecolor='cyan', cmap="YlGnBu", fmt='g', ax=ax1)
sns.heatmap(data=nbccla_cm, annot=True, linewidth=0.7, linecolor='cyan', cmap="YlGnBu", fmt='g', ax=ax2)
sns.heatmap(data=dtcla_cm, annot=True, linewidth=0.7, linecolor='cyan', cmap="YlGnBu", fmt='g', ax=ax3)
sns.heatmap(data=rfcla_cm, annot=True, linewidth=0.7, linecolor='cyan', cmap="YlGnBu", fmt='g', ax=ax4)
sns.heatmap(data=knncla_cm, annot=True, linewidth=0.7, linecolor='cyan', cmap="YlGnBu", fmt='g', ax=ax5)
plt.show()
```



ROC Curve

```
In [46]: from sklearn.metrics import roc_curve

# Logistic Regression Classification
Y_predict1_proba = logreg.predict_proba(X_test)
Y_predict1_proba = Y_predict1_proba[:, 1]
fpr, tpr, thresholds = roc_curve(Y_test, Y_predict1_proba)
plt.subplot(331)
plt.plot([0,1],[0,1], 'k--')
plt.plot(fpr, tpr, label='ANN')
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('ROC Curve Logistic Regression')
plt.grid(True)

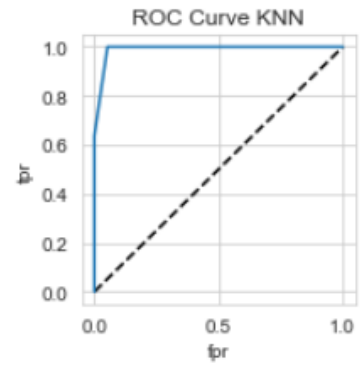
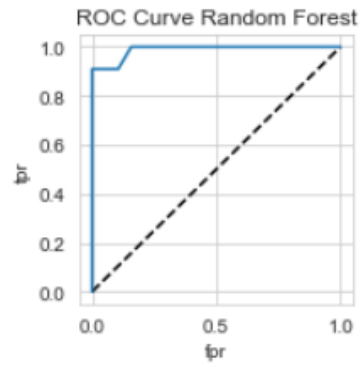
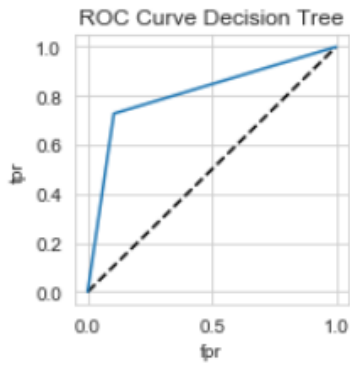
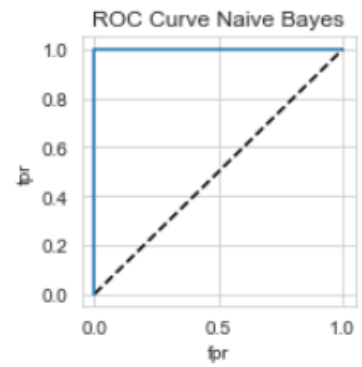
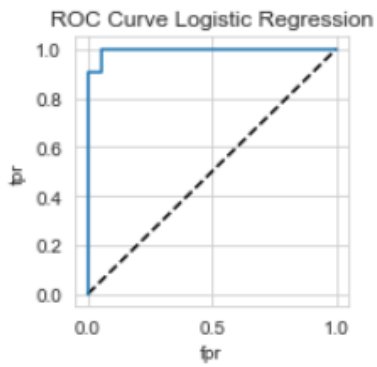
# Naive Bayes Classification
Y_predict3_proba = nbcla.predict_proba(X_test)
Y_predict3_proba = Y_predict3_proba[:, 1]
fpr, tpr, thresholds = roc_curve(Y_test, Y_predict3_proba)
plt.subplot(333)
plt.plot([0,1],[0,1], 'k--')
plt.plot(fpr, tpr, label='ANN')
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('ROC Curve Naive Bayes')
plt.grid(True)

# Decision Tree Classification
Y_predict4_proba = dtcla.predict_proba(X_test)
Y_predict4_proba = Y_predict4_proba[:, 1]
fpr, tpr, thresholds = roc_curve(Y_test, Y_predict4_proba)
plt.subplot(334)
plt.plot([0,1],[0,1], 'k--')
plt.plot(fpr, tpr, label='ANN')
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('ROC Curve Decision Tree')
plt.grid(True)

# Random Forest Classification
Y_predict5_proba = rfcla.predict_proba(X_test)
Y_predict5_proba = Y_predict5_proba[:, 1]
fpr, tpr, thresholds = roc_curve(Y_test, Y_predict5_proba)
plt.subplot(335)
plt.plot([0,1],[0,1], 'k--')
plt.plot(fpr, tpr, label='ANN')
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('ROC Curve Random Forest')
plt.grid(True)

# KNN Classification
Y_predict6_proba = knncla.predict_proba(X_test)
Y_predict6_proba = Y_predict6_proba[:, 1]
fpr, tpr, thresholds = roc_curve(Y_test, Y_predict6_proba)
plt.subplot(336)
plt.plot([0,1],[0,1], 'k--')
plt.plot(fpr, tpr, label='ANN')
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.title('ROC Curve KNN')
plt.grid(True)

plt.subplots_adjust(top=2, bottom=0.08, left=0.10, right=1.4, hspace=0.45, wspace=0.45)
plt.show()
```



The predicted Accuracy for each algorithm is

Logistic regression-96%

Naive Bayes-100%

Decision Tree-83%

Random Forest-93%

KNN-96%