```
File - models
```

C:\Users\chole\miniconda3\python.exe C:\Users/chole/Desktop/Cervical_cancer_behavior_risk/
models.py

Shape of complete dataset

X shape: (72, 19) y shape: (72,)

Shape of split dataset X_train shape: (54, 19) y_train shape: (54,) X_test shape: (18, 19) y_test shape: (18,)

Logistic Regression results

Best model parameters

{'model__C': 0.08858667904100823, 'model__class_weight': {0: 1, 1: 1.4736842105263157}}

Best model score (accuracy)

0.9466666666666667

Test data results

Classification report

	precision	recall	f1-score	support
0 1	0.92 0.67	0.85 0.80	0.88 0.73	13 5
accuracy			0.83	18
macro avg	0.79	0.82	0.80	18
weighted avg	0.85	0.83	0.84	18

AUC score

0.823076923076923

Accuracy score

0.8333333333333333

Test data results with changed threshold

Best Threshold=0.686739, G-Mean=0.894

Classification report

support	f1-score	recall	precision	
13	0.96	1.00	0.93	0
5	0.89	0.80	1.00	1
18	0.94			accuracy
18	0.93	0.90	0.96	macro avg
18	0.94	0.94	0.95	weighted avg

AUC score

0.9

Accuracy score 0.9444444444444444

categorical Naive Bayes results

Best model parameters

File - models

{'alpha': 0.0677966102677966}

Best model score (accuracy)

0.8699999999999999

Test data results

Classification report

support	f1-score	recall	precision	Classification
13	0.96	0.92	1.00	0
5	0.91	1.00	0.83	1
18	0.94			accuracy
18	0.93	0.96	0.92	macro avg
18	0.95	0.94	0.95	weighted avg

AUC score

0.9615384615384616

Accuracy score

0.944444444444444

Test data results with changed threshold

Best Threshold=0.999951, G-Mean=1.000

Classification report

	precision	recall	f1-score	support
0 1	1.00 1.00	1.00 1.00	1.00 1.00	13 5
accuracy			1.00	18
macro avg	1.00	1.00	1.00	18
weighted avg	1.00	1.00	1.00	18

AUC score

1.0

Accuracy score

1.0

Process finished with exit code 0