

SCREENSHOT – HEALTH CARE PROJECT

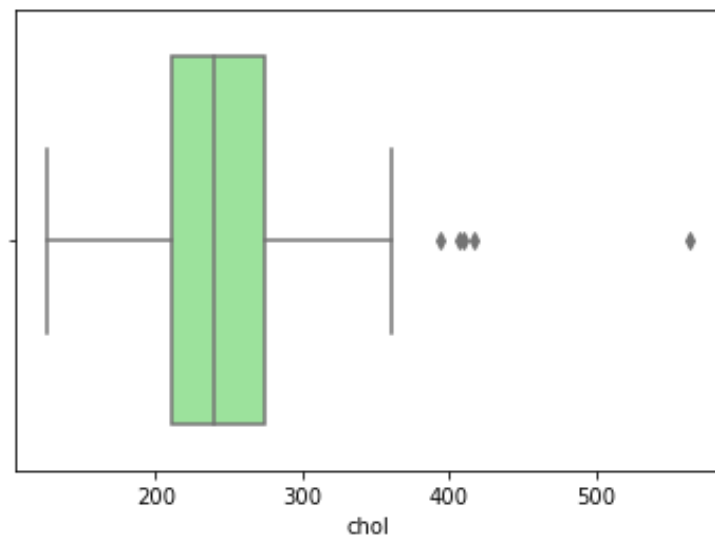
GIVEN DATA:

Out[2]:

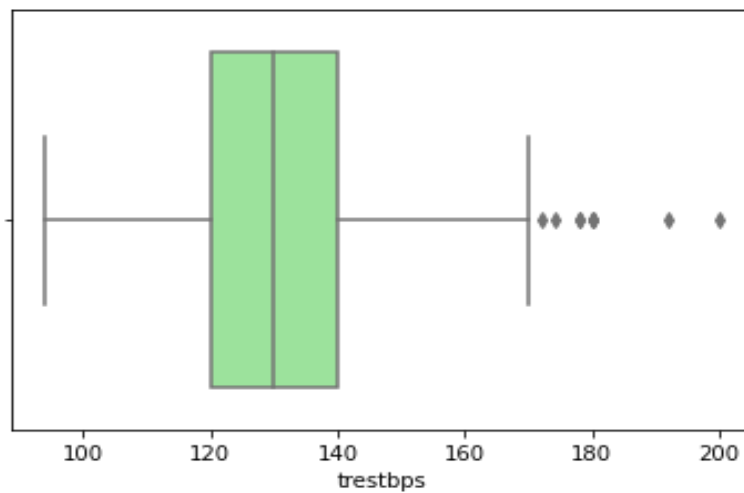
	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	target
0	63	1	3	145	233	1	0	150	0	2.3	0	0	1	1
1	37	1	2	130	250	0	1	187	0	3.5	0	0	2	1
2	41	0	1	130	204	0	0	172	0	1.4	2	0	2	1
3	56	1	1	120	236	0	1	178	0	0.8	2	0	2	1
4	57	0	0	120	354	0	1	163	1	0.6	2	0	2	1

DATA PREPROCESSING

Outliers in Cholesterol:

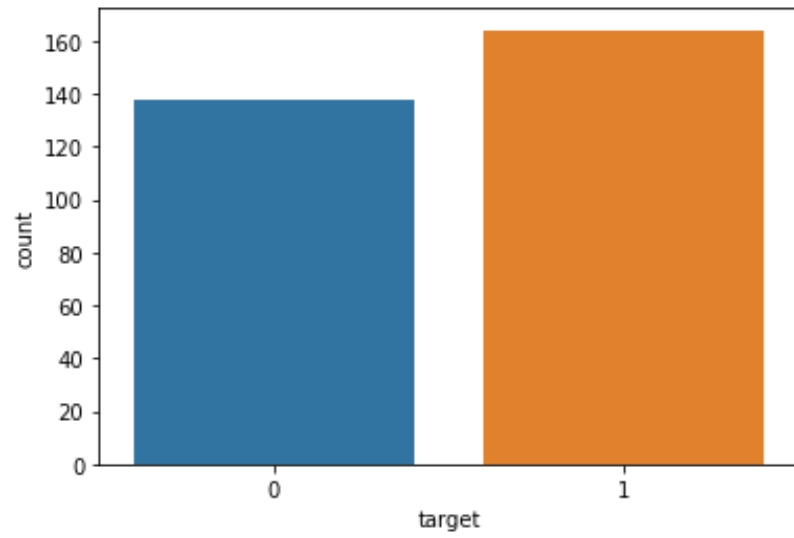


Outliers in Blood Pressure:

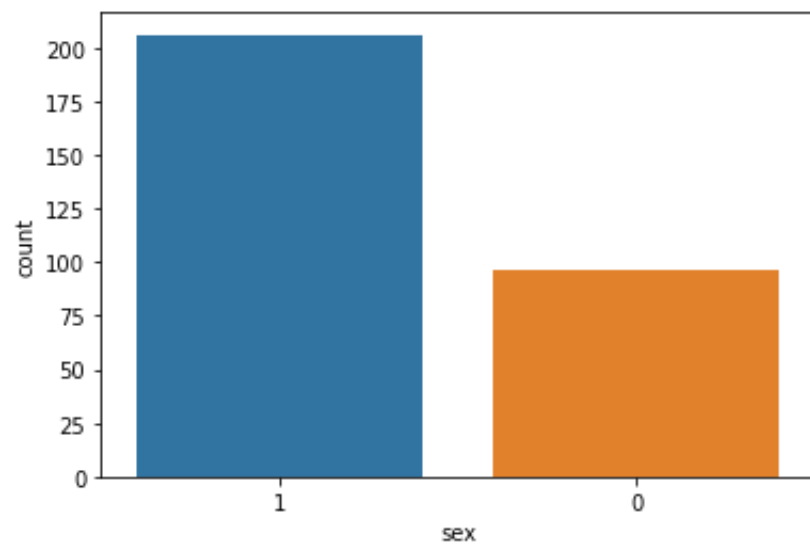


COUNTPLOT ANALYSIS

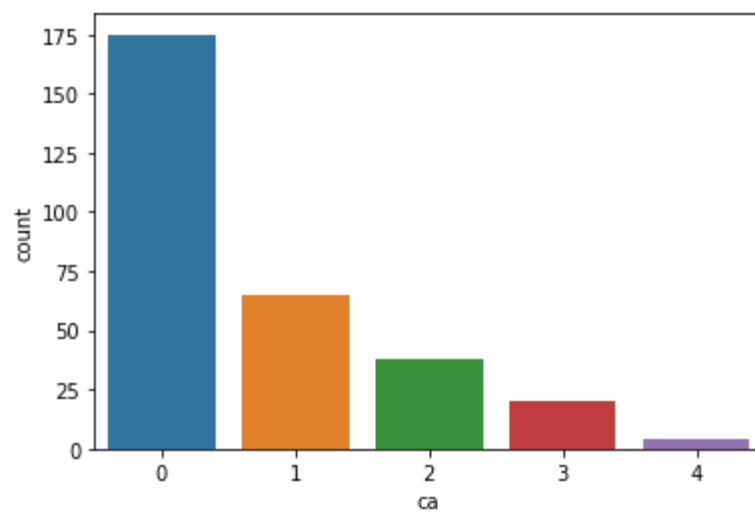
a) Target Variable:



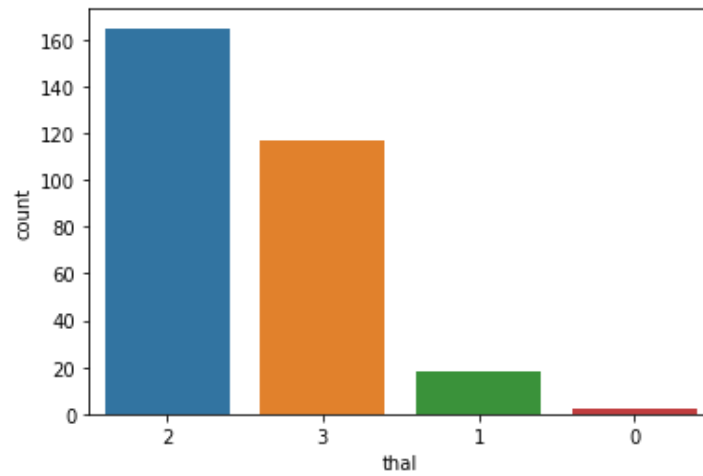
b) Sex



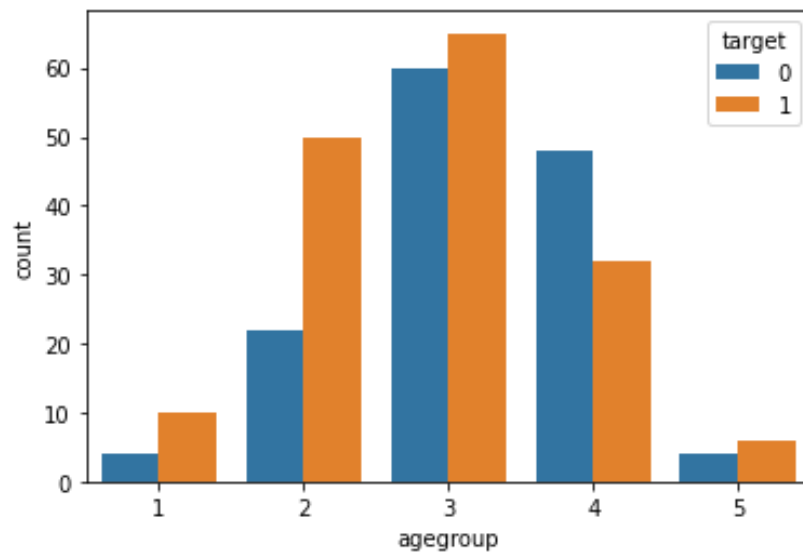
c) Ca



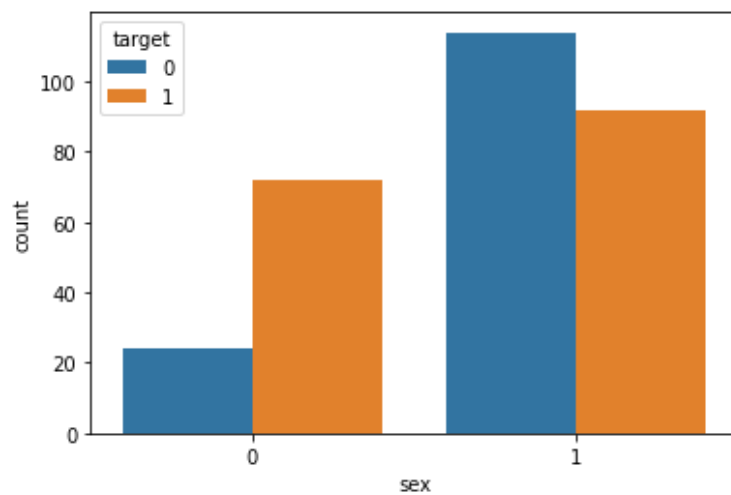
d) Thal



Analysis of Age group:

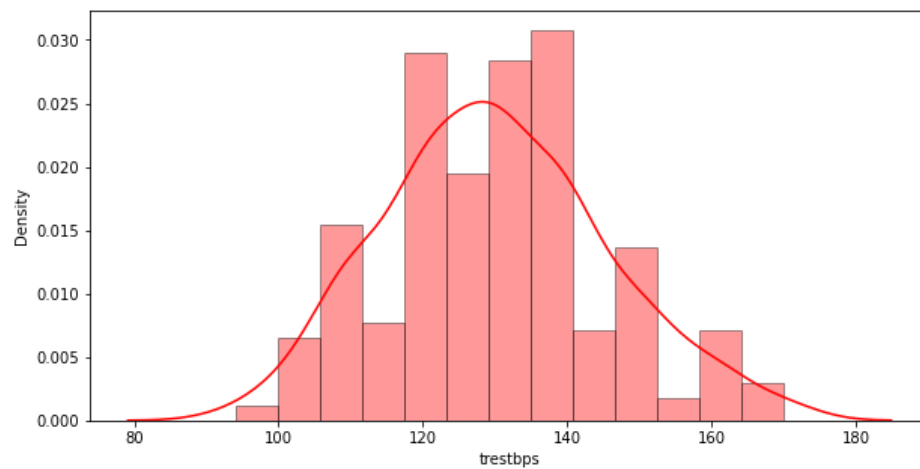


Gender Vs Target variable

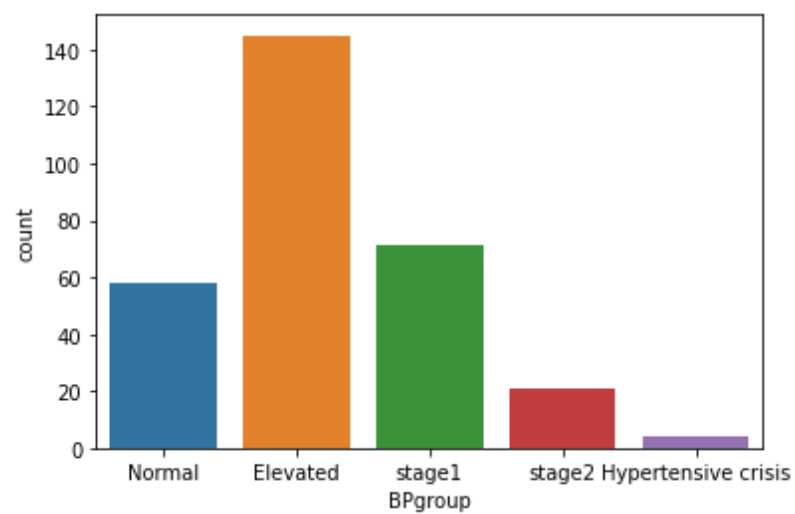


Analysis of Blood Pressure:

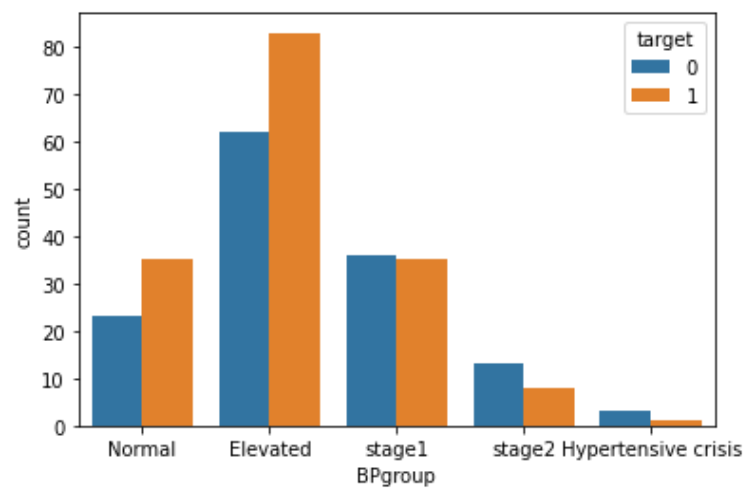
a) Distribution of BP:



b) Grouping of BP:

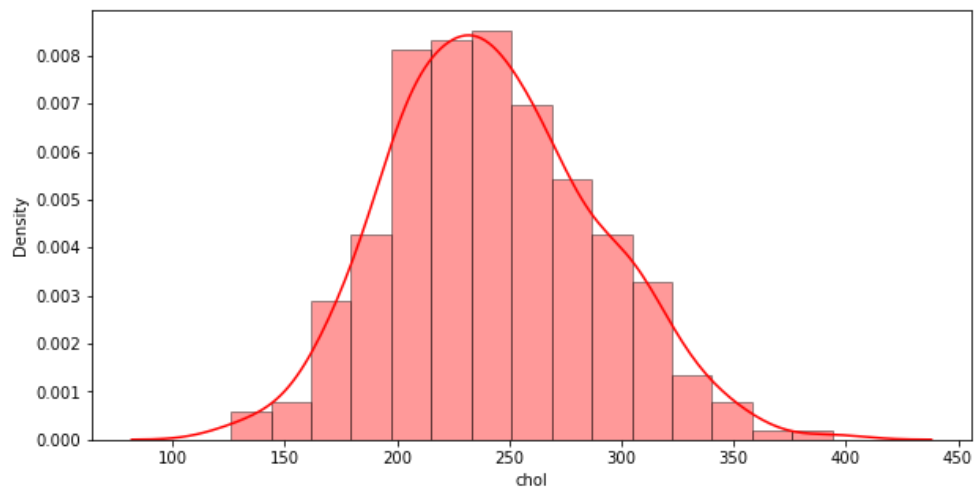


c) BP vs target variable:

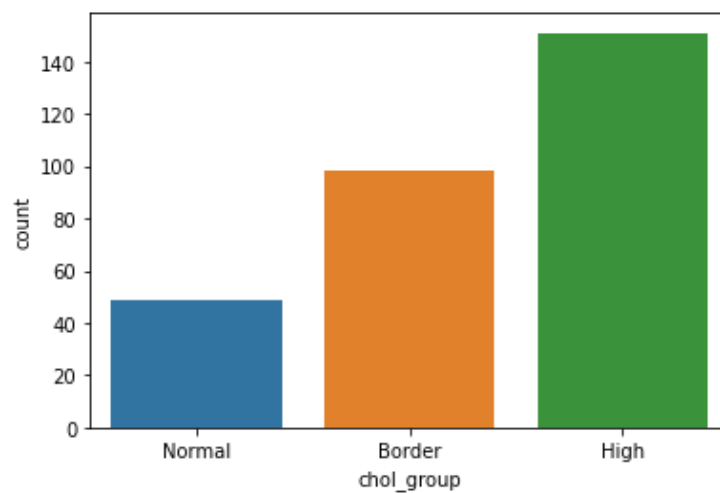


Analysis of Cholesterol:

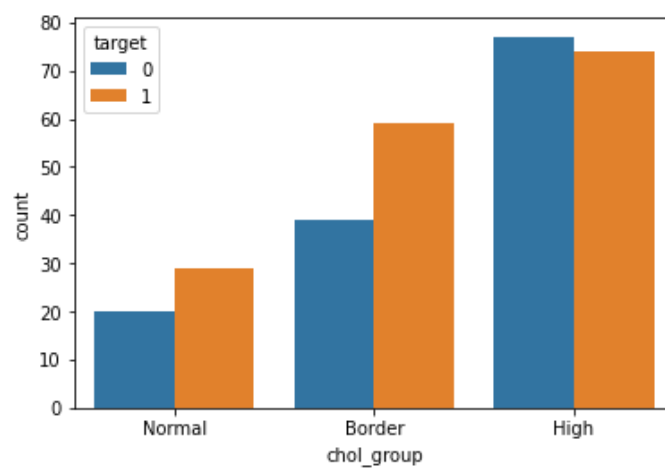
a) Distribution of data



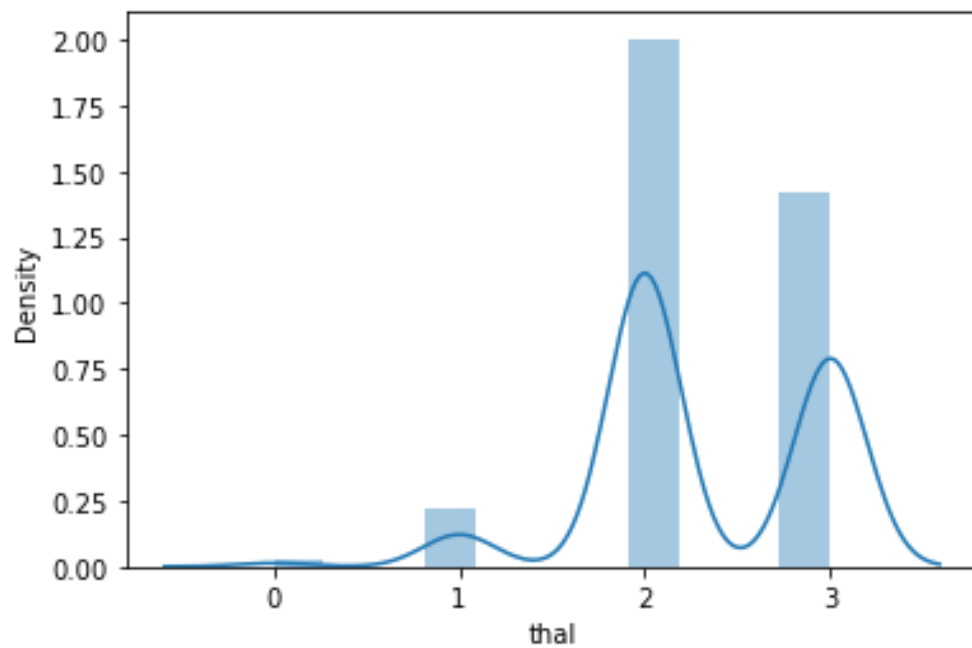
b) Grouping of cholesterol



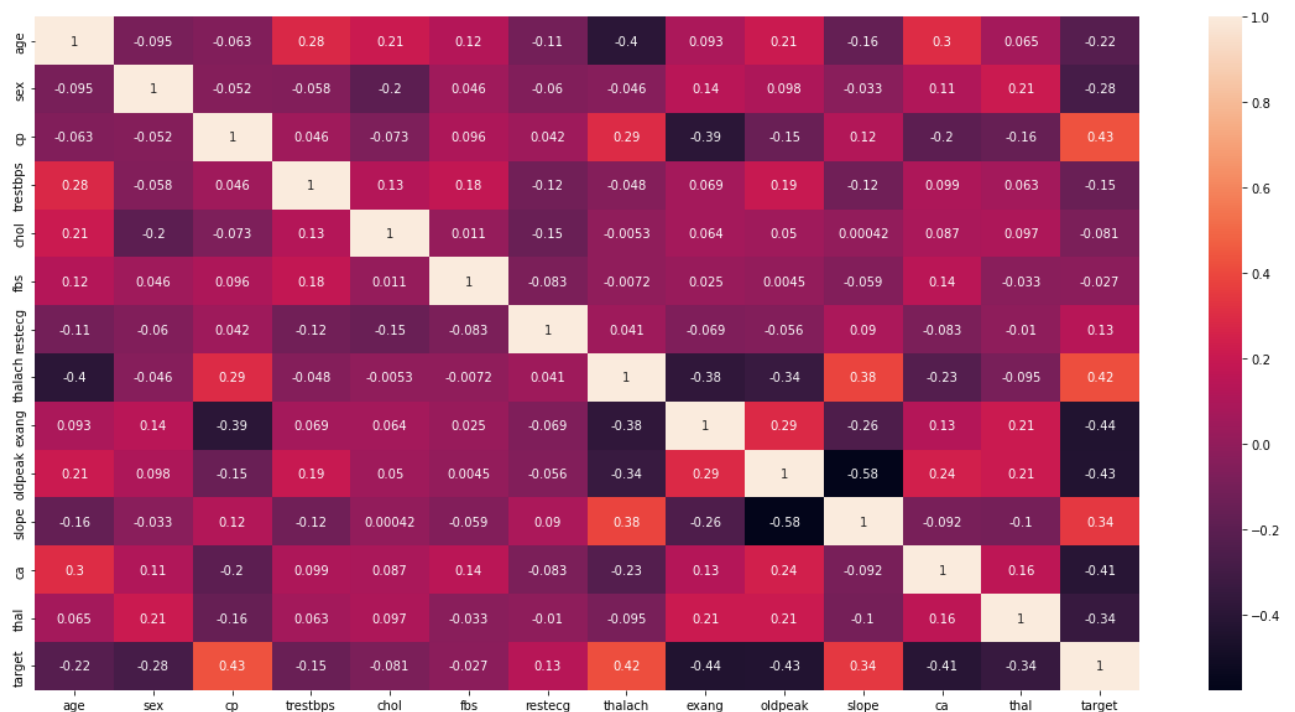
c) Cholesterol vs target variable



Study on Thalassemia



Correlation analysis between variables using Heat Map:



LOGISTIC REGRESSION:

Logistic Regression

```
In [101]: > classifier = LogisticRegression(solver='lbfgs',max_iter=10000)

In [102]: > classifier.fit(X_train, y_train)
Out[102]: LogisticRegression(max_iter=10000)

In [103]: > y_pred=lr.predict(X_test)

In [104]: > accuracy_score(y_test,y_pred)
Out[104]: 0.8131868131868132
```

RANDOM FOREST:

```
Out[86]: RandomForestClassifier(max_depth=50, n_estimators=900, random_state=0)

In [87]: > y_pred=rfc.predict(X_test)

In [91]: > accuracy_score(y_test,y_pred)
Out[91]: 0.8571428571428571

In [92]: > #Confusion Matrix
confusion_matrix(y_test,y_pred)
Out[92]: array([[36,  4],
               [ 9, 42]], dtype=int64)

In [95]: > print(classification_report(y_test,y_pred))
```

	precision	recall	f1-score	support
0	0.80	0.90	0.85	40
1	0.91	0.82	0.87	51
accuracy			0.86	91
macro avg	0.86	0.86	0.86	91
weighted avg	0.86	0.86	0.86	91