

Project Development Phase Model Performance Test

| | |
|---------------|--|
| Date | 10 November 2022 |
| Team ID | PNT2022TMID08852 |
| Project Name | Project – University admit eligibility predictor |
| Maximum Marks | 10 Marks |

Model Performance Testing:

Project team shall fill the following information in model performance testing template.

| S.No. | Parameter | Values | Screenshot |
|-------|----------------|---|------------|
| 1. | Metrics | Regression Model: MAE - , MSE - , RMSE - , R2 score - Classification Model: Confusion Matrix - , Accuracy Score- & Classification Report - | See Below |
| 2. | Tune the Model | Hyperparameter Tuning - Validation Method - | See Below |

1. Metrics

In [150...

```
from sklearn.metrics import accuracy_score, recall_score, roc_auc_score, confusion_matrix

print('Accuracy Score:', accuracy_score(y_test, y_pred))
print('Recall Score:', recall_score(y_test, y_pred))
print('ROC AUC Score:', roc_auc_score(y_test, y_pred))
print('Confussion Matrix:\n', confusion_matrix(y_test, y_pred))
```

```
Accuracy Score: 0.9333333333333333
Recall Score: 1.0
ROC AUC Score: 0.6
Confussion Matrix:
[[ 1  4]
 [ 0 55]]
```

In [143...

```
from sklearn.metrics import mean_squared_error, r2_score, mean_absolute_error
import numpy as np
print('Mean Absolute Error:', mean_absolute_error(y_test, y_predict))
print('Mean Squared Error:', mean_squared_error(y_test, y_predict))
print('Root Mean Squared Error:', np.sqrt(mean_squared_error(y_test, y_predict)))
```

```
Mean Absolute Error: 0.04198977717341872
Mean Squared Error: 0.0033348539701851436
Root Mean Squared Error: 0.05774819451883447
```

2. Tune the Model

Hyperparameter Tuning:

- The number of features is important and should be tuned in random forest classification.
- Initially all parameters in the dataset are taken as independent values to arrive at the dependent decision of Chronic Kidney Disease or No Chronic Kidney Disease.
- But the result was not accurate so used only 8 more correlated values as independent values to arrive at the dependent decision of Chronic Kidney Disease or not.

Validation Method:

It involves **partitioning the training data set into subsets, where one subset is held out to test the performance of the model**. This data set is called the validation data set.

Cross validation is to use different models and identify the best:

Logistic Regression Model performance values:

```
In [147... from sklearn.linear_model._logistic import LogisticRegression
           lore = LogisticRegression(random_state=0, max_iter=1000)
           lr = lore.fit(X_train, y_train)

In [148... y_pred = lr.predict(X_test)

In [150... from sklearn.metrics import accuracy_score, recall_score, roc_auc_score, confusion_matrix
           print('Accuracy Score:', accuracy_score(y_test, y_pred))
           print('Recall Score:', recall_score(y_test, y_pred))
           print('ROC AUC Score:', roc_auc_score(y_test, y_pred))
           print('Confussion Matrix:\n', confusion_matrix(y_test, y_pred))

Accuracy Score: 0.9333333333333333
Recall Score: 1.0
ROC AUC Score: 0.6
Confussion Matrix:
[[ 1  4]
 [ 0 55]]
```

