

Step For Doing ML Algorithm

1. Import All Libraries

```
pd.set_option('display.max_columns', None)
pd.set_option('display.max_rows', None)
```

For Eliminate All Error:

```
import warnings
warnings.simplefilter('ignore')
```

2. Import Data Set (pd.read_csv() , pd.read_excel() etc..)

- If you import from sklearn then do following :

```
from sklearn.datasets import load_iris
import pandas as pd
data = load_iris()
df = pd.DataFrame(data.data, columns=data.feature_names)
df.head()
```

3. Take some insight like

- a. Data.shape , Data.head , Data.tail , Data.info , Data.describe

4. Analysis EDA

- a. See distribution of plot (Regression) is it is Gaussian or not or which skew is it for target variable

Distribution of the target variable (Regression)

```
sns.distplot(data.target, bins = 25)
```

- b. In (classification) we do for count plot to see our target variable imbalanced or balanced

```
sns.countplot(data['target'])
```

5. Find out all numerical predictor

```
numeric_features = data.select_dtypes(include=[np.number])
```

```
numeric_features.dtypes
```

6. Then 1st find correlation between all numerical predictor

Correlation between all

```
corr = numeric_features.corr()
```

```
plt.figure(figsize=(12,10))
```

```
sns.heatmap(corr, vmax=.8, square=True,annot=True)
```

#correlation between features and target variable?

```
corr_matrix = abs(df.corr())
```

```
print(corr_matrix["target_variable"].sort_values(ascending=False))
```

7. Then find all categorical feature

```
cat_features = train.select_dtypes(include=[np.object])  
cat_features.dtypes
```

8. See distribution among all the feature

a. Like `sns.countplot(train.Age)`

b. And use pair plot and other type of plot

9. Treat missing value(mean ,median , mode , drop,0)

10.Then treat them apply onehot , dummy, label encode etc

11.Outlier Detection (By boxplot)

12.In this step we treat all null and categorical value

13.Then do some feature engineering / feature selection

14.Store target variable in y and all variable in X

15.Then if in data imbalanced we 1st balanced data by using SMOTE

#Step 1: Here I use Oversampling Using Smote

```
from imblearn.over_sampling import SMOTE
```

```
x_resample, y_resample = SMOTE().fit_sample(x, y.values.ravel())
```

```
# lets print the shape of x and y after resampling it
```

```
print(x_resample.shape)
```

```
print(y_resample.shape)
```

#Step 2:

```
# lets also check the value counts of our target variable4
```

```
print("Before Resampling :")
```

```
print(y.value_counts())
```

```
print("After Resampling :")
```

```
y_resample = pd.DataFrame(y_resample)
```

```
print(y_resample[0].value_counts())
```

16.Then scale the data by using standard scalar :

lets import the standard scaler library from sklearn to do that

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
x_train = sc.fit_transform(x_train)
```

```
x_valid = sc.transform(x_valid)
```

17. Then we split train_test_split

18. Model Build (RF,DT,LR,LoR,XGB)

19. Apply K-Fold cross validation (if train data is few)

20. Apply Hyper parameter tuning

21. Find all accuracy Both (Regression, Classification)

a. Cross Validation :

Here I Use StratifiedKFold :

```
kf = StratifiedKFold(n_splits=5,shuffle=True,random_state=45)
```

```
pred_test_full = 0
```

```
cv_score = []
```

```
i=1
```

```
for train_index, test_index in kf.split(X, y):
```

```
    print('{} of KFold {}'.format(i, kf.n_splits))
```

```
    # Split X_train, X_test, y_train, y_test
```

```
    xtr, xvl = X.loc[train_index], X.loc[test_index]
```

```
    ytr, yvl = y.loc[train_index], y.loc[test_index]
```

```
    # model
```

```
    model = # add any model
```

```
    model.fit(xtr, ytr)
```

```
    # performance Calculation
```

```
    score = roc_auc_score(yvl, model.predict(xvl))
```

```
    print('ROC AUC score:', score)
```

```
    cv_score.append(score)
```

```
    # predict test data
```

```
    pred_test = model.predict_proba(x_test)[:,-1]
```

```
    pred_test_full += pred_test
```

```
    i+=1
```

b. Hyper Parameter Tuning :

initialize all parameter

```
params={  
"max_depth" : [50,100,150,200],  
"min_samples_split" : [1,2,3,4,5,6,7,8,9],  
"min_samples_leaf" : [1,2,3,4,5,6,7,8,9],  
"criterion": ["gini", "entropy"],  
"max_leaf_nodes": [1,5,10,15,20]}
```

Apply desire model for tuning

```
classifier= DecisionTreeClassifier()
```

Randomize search Cv is start

```
random_search = RandomizedSearchCV (classifier,param_distributions=params,n_iter  
=5,scoring='roc_auc',n_jobs=-1,cv=5,verbose=3)  
random_search.fit(X,y)
```

Calculate The Score and best params

```
print('Best roc_auc: {:.4}, with best C: {}'.format(random_search.best_score_,  
random_search.best_params_))
```

c. Calculate Performance Matric :

For Regression r2_score,MAE,RMSE

```
from sklearn.metrics import r2_score,mean_squared_error,mean_absolute_error  
from math import sqrt
```

```
from sklearn.ensemble import RandomForestRegressor  
rfr = RandomForestRegressor()
```

```
rfr.fit(X_train,y_train)
```

```
#Predicting the Test set results
```

```
y_pred_rfr = rfr.predict(X_train)
```

```
score = r2_score(y_train,y_pred_rfr)
```

```
print("Score of Training:",100*score)
```

```
print("RMSE : " , np.sqrt(mean_squared_error(y_train,y_pred_rfr)))
```

```
y_test_pred_rfr = rfr.predict(X_test)
```

```
#r2 Score
```

```
score = r2_score(y_test,y_test_pred_rfr)
```

```
print("Score of Testing:",100*score)
```

```
#RMSE
```

```
print("RMSE : " , np.sqrt(mean_squared_error(y_test,y_test_pred_rfr)))
```

```
#MAE
```

```
print("Mean Absolute Error",mean_absolute_error(y_test,y_test_pred_rfr))
```

For Classification (confusion matrix, accuracy, precision, recall)

```
from sklearn.metrics import confusion_matrix, classification_report, accuracy_score
import warnings
warnings.filterwarnings('ignore')
```

```
from sklearn.linear_model import LogisticRegression
lr = LogisticRegression()
lr.fit(X_train, y_train)
```

```
y_pred = lr.predict(X_test)
```

```
print("Training Accuracy :", lr.score(X_train, y_train))
print("Testing Accuracy :", lr.score(X_test, y_test))
```

#Confusion Matric

```
cm = confusion_matrix(y_test, y_pred)
plt.rcParams['figure.figsize'] = (3, 3)
sns.heatmap(cm, annot = True, cmap = 'Wistia', fmt = '.8g')
plt.show()
```

#Classification Report

```
cr = classification_report(y_test, y_pred)
print(cr)
```

Using Pickle Model TO Dumb and Load Model:

```
#dumb file
import pickle
filename = 'filename.pkl'
pickle.dump(model_instance, open(filename, 'wb'))
```

```
#open file
model = open("cc_strength.pkl", "rb")
model = pickle.load(model)
```
