

```
In [221... # ! pip install -q -U keras-tuner
```

```
In [222... import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
import matplotlib.image as mpimg
from datetime import time
# preprocessing and pipeline
from sklearn.model_selection import train_test_split, cross_val_score, GridSearchCV
from sklearn.compose import ColumnTransformer
from sklearn.preprocessing import OneHotEncoder, LabelEncoder, StandardScaler
from sklearn.pipeline import Pipeline
from outlier_cleaner import OutlierCleaner

# sklearn libraries
from sklearn.linear_model import LinearRegression, Ridge, Lasso
from sklearn.svm import SVC, SVR
from sklearn.tree import DecisionTreeRegressor, DecisionTreeClassifier
from sklearn.neighbors import KNeighborsRegressor, KNeighborsClassifier
from sklearn.ensemble import (
    RandomForestRegressor, GradientBoostingRegressor, AdaBoostRegressor,
    VotingRegressor, StackingRegressor, RandomForestClassifier, VotingClassifier
)

# sklearn metrics
from sklearn.metrics import r2_score, mean_squared_error, accuracy_score, confusion_matrix
from extended_sklearn_metrics import evaluate_model_with_cross_validation

# extra model
import xgboost as xgb

import joblib

# deep learning
from imblearn.over_sampling import SMOTE
import tensorflow as tf
from tensorflow.keras import layers, models
import keras_tuner as kt
from tensorflow import keras
```

```
In [223... df = pd.read_csv('heart_attack_prediction_indonesia.csv', nrows=30000)
df1 = df.copy()
```

Feature Engineering (higher values = more risk)

```
In [224... df['alcohol_consumption'] = df['alcohol_consumption'].fillna('None')
stress_mapping = {
    'Low': 1,
    'Moderate': 2,
    'High': 3
}
df['stress_level'] = df['stress_level'].map(stress_mapping)
smoking_mapping = {
```

```

    'Never': 1,
    'Past': 2,
    'Current': 3
}
df['smoking_status'] = df['smoking_status'].map(smoking_mapping)
alcohol_consumption_mapping = {
    'None': 1,
    'Moderate': 2,
    'High': 3
}
df['alcohol_consumption'] = df['alcohol_consumption'].map(alcohol_consumption_ma
physical_activity_mapping = {
    'Low': 1,
    'Moderate': 2,
    'High': 3
}
df['physical_activity'] = df['physical_activity'].map(physical_activity_mapping)
dietary_habits_mapping = {
    'Unhealthy': 1,
    'Healthy': 2,
}
df['dietary_habits'] = df['dietary_habits'].map(dietary_habits_mapping)
air_pollution_mapping = {
    'Low': 1,
    'Moderate': 2,
    'High': 3
}
df['air_pollution_exposure'] = df['air_pollution_exposure'].map(air_pollution_ma
income_level_mapping = {
    'Low': 3,
    'Middle': 2,
    'High': 1
}
df['income_level'] = df['income_level'].map(income_level_mapping)
df['age_group'] = np.where(
    df['age'] < 40, 1, np.where(
        df['age'] < 55, 2, np.where(
            df['age'] < 65, 3, 4
        )
    )
)
df.age_group.unique()

```

Out[224... array([3, 2, 4, 1])

In [225... df.columns.tolist()

```
Out[225... ['age',
             'gender',
             'region',
             'income_level',
             'hypertension',
             'diabetes',
             'cholesterol_level',
             'obesity',
             'waist_circumference',
             'family_history',
             'smoking_status',
             'alcohol_consumption',
             'physical_activity',
             'dietary_habits',
             'air_pollution_exposure',
             'stress_level',
             'sleep_hours',
             'blood_pressure_systolic',
             'blood_pressure_diastolic',
             'fasting_blood_sugar',
             'cholesterol_hdl',
             'cholesterol_ldl',
             'triglycerides',
             'EKG_results',
             'previous_heart_disease',
             'medication_usage',
             'participated_in_free_screening',
             'heart_attack',
             'age_group']
```

```
In [226... df['health_risk_score'] = (
    df['hypertension'] +
    df['diabetes'] +
    df['obesity'] +
    df['family_history'] +
    df['smoking_status'] +
    df['alcohol_consumption'] +
    (1 - df['physical_activity']) +
    (1 - df['dietary_habits']) +
    df['air_pollution_exposure'] +
    df['stress_level'] +
    df.income_level
)

# df['cholesterol_ratio'] = df['cholesterol_hdl']/df['cholesterol_ldl']
# df = df.drop(labels=['cholesterol_hdl', 'cholesterol_ldl'], axis=1)
df['obesity_risk_score'] = np.where(df['gender'] == 'Male' , df['obesity'] * 1 +
# df = df.drop(labels=['obesity', 'waist_circumference'], axis=1)
# df['pulse_pressure'] = df['blood_pressure_systolic'] - df['blood_pressure_dias
# df = df.drop(labels=['blood_pressure_systolic', 'blood_pressure_diastolic'], a
df['stress_to_sleep_ratio'] = df['stress_level'] / df['sleep_hours']
# df = df.drop(columns=['stress_level', 'sleep_hours'])
df['mean_artierial_pressure'] = (2* df['blood_pressure_systolic'] + df['blood_pre
# df = df.drop(columns=['blood_pressure_systolic', 'blood_pressure_diastolic'])
df['triglyceride-hdl-ratio'] = df['triglycerides'] / df['cholesterol_hdl']
# df = df.drop(columns=['triglycerides', 'cholesterol_hdl'])
```

```
In [227... # df_risk_score = df[['health_risk_score', 'hypertension', 'diabetes', 'obesity_r
# df_risk_score.corr().sort_values(by='health_risk_score', ascending=False).styl
```

```
In [228... # df = df.drop(columns=['physical_activity', 'dietary_habits'])
```

```
In [229... df.isna().any()
```

```
Out[229... age                False
gender            False
region           False
income_level     False
hypertension     False
diabetes         False
cholesterol_level False
obesity          False
waist_circumference False
family_history   False
smoking_status   False
alcohol_consumption False
physical_activity False
dietary_habits   False
air_pollution_exposure False
stress_level     False
sleep_hours      False
blood_pressure_systolic False
blood_pressure_diastolic False
fasting_blood_sugar False
cholesterol_hdl  False
cholesterol_ldl  False
triglycerides    False
EKG_results      False
previous_heart_disease False
medication_usage False
participated_in_free_screening False
heart_attack      False
age_group         False
health_risk_score False
obesity_risk_score False
stress_to_sleep_ratio False
mean_arterial_pressure False
triglyceride-hdl-ratio False
dtype: bool
```

```
In [230... # df.health_risk_score.unique()
```

Encoding

```
In [231... ## Encoding
cat_df = df.select_dtypes(include='object')
num_df = df.select_dtypes(exclude='object')
encoder = LabelEncoder()
for cols in cat_df:
    cat_df[cols+'_encoded'] = encoder.fit_transform(cat_df[cols])
cat_df = cat_df.select_dtypes(exclude='object')
df = pd.concat([cat_df, num_df], axis=1)
```

Cleaning Outliers

```
In [232... cleaner = OutlierCleaner(df, preserve_index=True)
cleaned_df, info = cleaner.clean_columns(
    method='zscore',
    show_progress=True
)
df = cleaned_df
```

Cleaning columns: 100%|██████████| 34/34 [00:00<00:00, 188.79it/s]

Visualization of correlation of features with heart attack

```
In [233... df.waist_circumference.min(), df.waist_circumference.max()
# df.waist_circumference.unique()
```

Out[233... (45, 142)

```
In [234... df.columns
```

```
Out[234... Index(['gender_encoded', 'region_encoded', 'EKG_results_encoded', 'age',
      'income_level', 'hypertension', 'diabetes', 'cholesterol_level',
      'obesity', 'waist_circumference', 'family_history', 'smoking_status',
      'alcohol_consumption', 'physical_activity', 'dietary_habits',
      'air_pollution_exposure', 'stress_level', 'sleep_hours',
      'blood_pressure_systolic', 'blood_pressure_diastolic',
      'fasting_blood_sugar', 'cholesterol_hdl', 'cholesterol_ldl',
      'triglycerides', 'previous_heart_disease', 'medication_usage',
      'participated_in_free_screening', 'heart_attack', 'age_group',
      'health_risk_score', 'obesity_risk_score', 'stress_to_sleep_ratio',
      'mean_arterial_pressure', 'triglyceride-hdl-ratio'],
      dtype='object')
```

```
In [235... results = []
for i in df1.columns:
    results.append({
        f'{i}': df1[i].unique()
    })
results
```

```

Out[235... [{ 'age': array([60, 53, 62, 73, 52, 64, 49, 61, 57, 32, 34, 48, 42, 58, 44, 38,
72,
55, 37, 56, 41, 59, 47, 51, 77, 54, 40, 31, 39, 63, 46, 67, 33, 50,
66, 71, 25, 45, 65, 84, 68, 81, 43, 36, 70, 35, 87, 90, 82, 80, 30,
76, 74, 29, 69, 79, 78, 27, 75, 28, 85, 86, 83, 26, 88, 89]),
dtype=int64)},
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{ 'region': array(['Rural', 'Urban'], dtype=object)},
{ 'income_level': array(['Middle', 'Low', 'High'], dtype=object)},
{ 'hypertension': array([0, 1], dtype=int64)},
{ 'diabetes': array([1, 0], dtype=int64)},
{ 'cholesterol_level': array([211, 208, 231, 202, 232, 238, 165, 186, 121, 196,
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228, 265, 177, 192, 176, 207, 174, 225, 170, 130, 251, 201, 159,
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133, 168, 188, 285, 252, 216, 220, 212, 116, 182, 250, 175, 246,
215, 226, 240, 210, 147, 254, 227, 243, 223, 146, 241, 173, 256,
244, 198, 247, 187, 164, 217, 218, 152, 161, 303, 203, 206, 245,
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195, 239, 235, 118, 113, 229, 123, 117, 197, 233, 273, 224, 272,
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260, 135, 261, 279, 140, 137, 274, 114, 115, 122, 107, 100, 278,
124, 283, 318, 143, 306, 288, 287, 120, 276, 299, 269, 129, 325,
262, 106, 319, 112, 293, 284, 301, 281, 295, 302, 296, 292, 126,
289, 108, 282, 109, 267, 102, 290, 300, 286, 308, 312, 326, 309,
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{ 'alcohol_consumption': array([nan, 'Moderate', 'High'], dtype=object)},
{ 'physical_activity': array(['High', 'Moderate', 'Low'], dtype=object)},
{ 'dietary_habits': array(['Unhealthy', 'Healthy'], dtype=object)},
{ 'air_pollution_exposure': array(['Moderate', 'High', 'Low'], dtype=object)},
{ 'stress_level': array(['Moderate', 'High', 'Low'], dtype=object)},
{ 'sleep_hours': array([5.97060316, 5.64381314, 6.33619667, ..., 6.71533855, 6.
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87, 156, 201, 159, 172, 200, 140, 84, 102, 126, 239, 85, 197,
222, 169, 223, 107, 275, 254, 165, 80, 179, 111, 244, 127, 144,
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{'previous_heart_disease': array([0, 1], dtype=int64)},
{'medication_usage': array([0, 1], dtype=int64)},
{'participated_in_free_screening': array([0, 1], dtype=int64)},
{'heart_attack': array([0, 1], dtype=int64)}]

```

In [236...

```

results = []
for i in df.columns:
    results.append({
        f'{i}': df[i].unique()
    })
results

```



```

Out[236... [{"gender_encoded": array([1, 0])},
             {"region_encoded": array([0, 1])},
             {"EKG_results_encoded": array([1, 0])},
             {"age": array([60, 53, 62, 73, 52, 64, 49, 61, 57, 32, 34, 48, 42, 58, 38, 72,
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                    76, 74, 29, 69, 79, 78, 27, 75, 28, 86, 85, 83, 26, 88, 89],
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             {"hypertension": array([0, 1], dtype=int64)},
             {"diabetes": array([1, 0], dtype=int64)},
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                    138, 141, 140, 137, 139, 142,  47], dtype=int64)},
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             {"smoking_status": array([1, 2, 3], dtype=int64)},
             {"alcohol_consumption": array([1, 2, 3], dtype=int64)},
             {"physical_activity": array([3, 2, 1], dtype=int64)},
             {"dietary_habits": array([1, 2], dtype=int64)},
             {"air_pollution_exposure": array([2, 3, 1], dtype=int64)},
             {"stress_level": array([2, 3, 1], dtype=int64)},
             {"sleep_hours": array([5.97060316, 5.64381314, 6.33619667, ..., 6.71533855, 6.
44848813,
                    5.8914506 ])},
             {"blood_pressure_systolic": array([113, 132, 116, 136, 127, 131, 128, 109, 15
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```

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```

```

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{'medication_usage': array([0, 1], dtype=int64)},
{'participated_in_free_screening': array([0, 1], dtype=int64)},
{'heart_attack': array([0, 1], dtype=int64)},
{'age_group': array([3, 2, 4, 1])},
{'health_risk_score': array([ 7, 11,  9, 12, 10, 14,  8, 13,  6,  5,  4, 15],
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{'obesity_risk_score': array([0. , 0.5, 1.5, 1. ])},
{'stress_to_sleep_ratio': array([0.33497453, 0.53155551, 0.15782338, ..., 0.29
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```

```
46875 , 1.703125 ,  
      1.10869565]]}]
```

In [237...

```
corr_df = df.corr()[['heart_attack']].sort_values(by='heart_attack', ascending=F  
corr_df.style.background_gradient(cmap='coolwarm', axis=None)
```

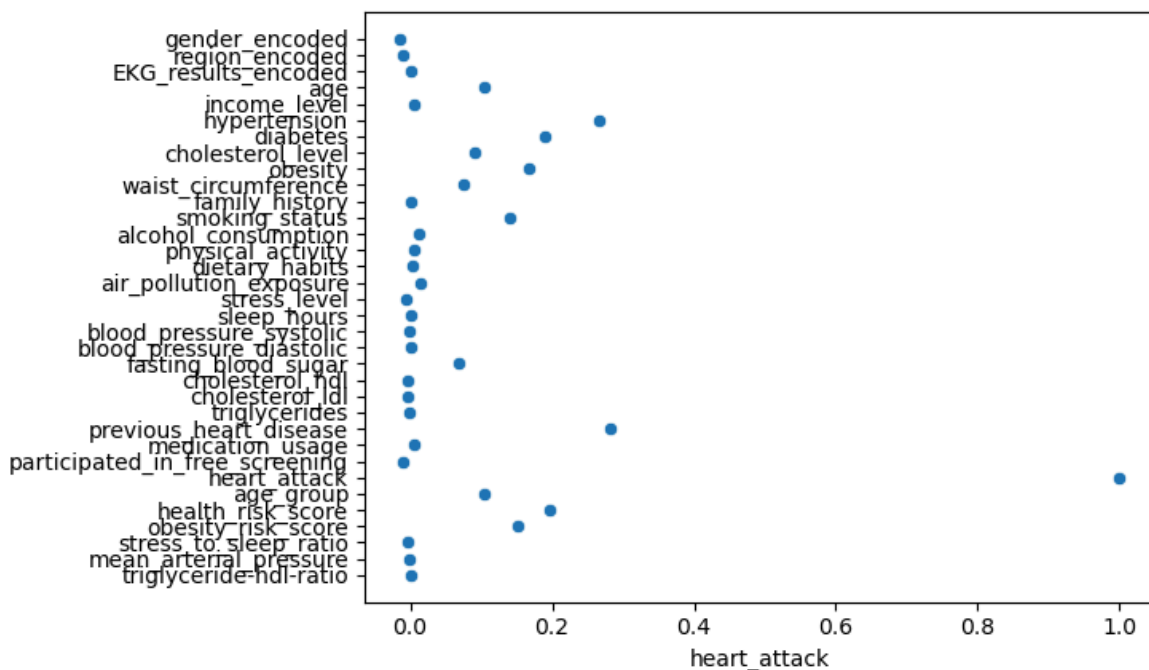
Out[237...

heart_attack	
heart_attack	1.000000
previous_heart_disease	0.281527
hypertension	0.266837
health_risk_score	0.197037
diabetes	0.188859
obesity	0.166300
obesity_risk_score	0.152269
smoking_status	0.141086
age	0.103794
age_group	0.103686
cholesterol_level	0.090968
waist_circumference	0.074709
fasting_blood_sugar	0.068222
air_pollution_exposure	0.013178
alcohol_consumption	0.011258
income_level	0.005869
medication_usage	0.004564
physical_activity	0.004182
dietary_habits	0.003435
triglyceride-hdl-ratio	0.001366
family_history	0.001147
blood_pressure_diastolic	0.000394
EKG_results_encoded	0.000035
sleep_hours	-0.000019
mean_arterial_pressure	-0.001625
blood_pressure_systolic	-0.001841
triglycerides	-0.002392
cholesterol_ldl	-0.004101
cholesterol_hdl	-0.004503
stress_to_sleep_ratio	-0.004751
stress_level	-0.006581
participated_in_free_screening	-0.009895
region_encoded	-0.010467

heart_attack

gender_encoded -0.014083

```
In [238... sns.scatterplot(data=df.corr(), x='heart_attack', y=df.columns.tolist())
plt.show()
```

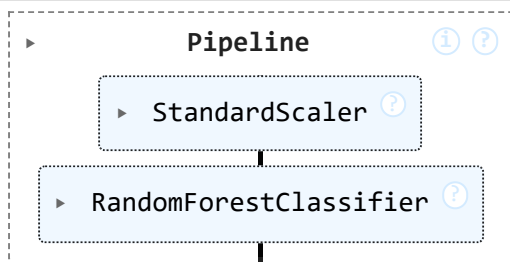


Modelling

```
In [239... x = df.drop('heart_attack', axis=1)
y = df['heart_attack']
X_train, X_test, y_train, y_test = train_test_split(x, y, test_size=0.3, random_
# scaler = StandardScaler()
# X_train_scaled = scaler.fit_transform(X_train)
# X_test_scaled = scaler.fit_transform(X_test)
# X_train_scaled, X_train_scaled[0]
SEED = 42
rf = RandomForestClassifier(
    criterion='entropy',
    max_depth=5,
    max_features='sqrt',
    min_samples_leaf=1,
    min_samples_split=3,
)
# param_grid = {
#     'classifier__criterion': ['gini', 'entropy', 'log_loss'],
#     'classifier__max_depth': [2, 4, 5],
#     'classifier__min_samples_split': [2, 3, 4],
#     'classifier__min_samples_leaf': [1, 2, 3],
#     'classifier__max_features': ['sqrt', 'log2'],
# }
rf_pipeline = Pipeline([
    ('scaler', StandardScaler()),
    ('classifier', rf),
])
# grid_search = GridSearchCV(
```

```
# estimator = rf_pipeline,
# param_grid = param_grid,
# scoring = 'accuracy',
# n_jobs = None,
# cv = 5,
# verbose = 2,
# )
# grid_search.fit(X_train, y_train)
# scaler = StandardScaler()
# X_train = scaler.fit_transform(X_train)
# X_test = scaler.transform(X_test)
# rf.fit(X_train, y_train)
rf_pipeline.fit(X_train, y_train)
```

Out[239...



In [240...

```
x.columns
```

Out[240...

```
Index(['gender_encoded', 'region_encoded', 'EKG_results_encoded', 'age',
      'income_level', 'hypertension', 'diabetes', 'cholesterol_level',
      'obesity', 'waist_circumference', 'family_history', 'smoking_status',
      'alcohol_consumption', 'physical_activity', 'dietary_habits',
      'air_pollution_exposure', 'stress_level', 'sleep_hours',
      'blood_pressure_systolic', 'blood_pressure_diastolic',
      'fasting_blood_sugar', 'cholesterol_hdl', 'cholesterol_ldl',
      'triglycerides', 'previous_heart_disease', 'medication_usage',
      'participated_in_free_screening', 'age_group', 'health_risk_score',
      'obesity_risk_score', 'stress_to_sleep_ratio', 'mean_arterial_pressure',
      'triglyceride-hdl-ratio'],
      dtype='object')
```

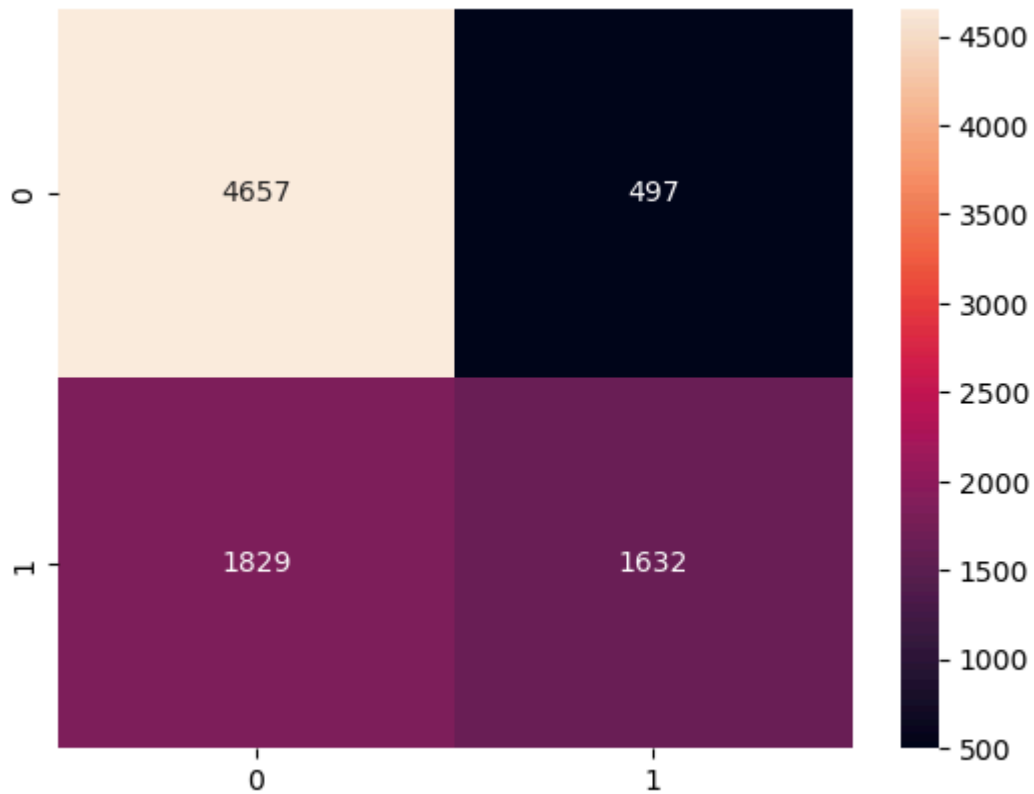
In [241...

```
y_pred = rf_pipeline.predict(X_test)
print(classification_report(y_test, y_pred))
```

	precision	recall	f1-score	support
0	0.72	0.90	0.80	5154
1	0.77	0.47	0.58	3461
accuracy			0.73	8615
macro avg	0.74	0.69	0.69	8615
weighted avg	0.74	0.73	0.71	8615

In [242...

```
sns.heatmap(confusion_matrix(y_test, y_pred), annot=True, fmt='d')
plt.show()
```



Saving model to joblib file

```
In [243... # joblib.dump(rf_pipeline, 'heart_attack_prediction_model.joblib')
```

Save clean df as CSV file

```
In [244... df.to_csv('clean_hap.csv', index=False)
```

```
In [ ]:
```