

Heart Disease Diagnosis

Overview

In this article we will be closely working with the heart disease diagnosis and for that we will be looking into the heart disease dataset and from that dataset we will derieve various insights that helps us know the weightage of each feature and how they are interrelated to each other but this time our soul aim is to detect the probablity of person that will be effected by a saviour heart problem or not.

Takeaways from this article

1. Data insight: As mentioned here we will be working with the heart disease detection dataset and we will be putting out interesting inference from the data to derieve some meaningful results.
2. EDA: Exploratory data analysis is the key step for the getting the meaningful results.
3. Feature engineering: After getting the insights from the data we have to alter the features so that they can move forward for the model building phase.
4. Model building: In this phase we will be building our Machine learning model for heart disease detection.

Importing Necessary Libraries

```
In [1]: # Plotting Libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
#import cufflinks as cf
%matplotlib inline

# Metrics for Classification technique
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

# Scaler
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import RandomizedSearchCV, train_test_split

from xgboost import XGBClassifier
from catboost import CatBoostClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
```

Data Loading

Our first step is to extract train and test data.

In [2]: *# Importing Data*

```
data = pd.read_csv("heart.csv")  
data.head(6) # Mention no of rows to be displayed from the top in the argument
```

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
5	57	1	0	140	192	0	1	148	0	0.4	1	0	1	1

Exploratory Data Analysis

In [3]: *#Size of the dataset*
data.shape

Out[3]: (303, 14)

I have a dataset with 303 rows which indicates a smaller set of data.

In [4]: data.info()

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 303 entries, 0 to 302
Data columns (total 14 columns):
#   Column      Non-Null Count  Dtype
---  -
0   age         303 non-null    int64
1   sex         303 non-null    int64
2   cp          303 non-null    int64
3   trestbps    303 non-null    int64
4   chol        303 non-null    int64
5   fbs         303 non-null    int64
6   restecg     303 non-null    int64
7   thalach     303 non-null    int64
8   exang       303 non-null    int64
9   oldpeak     303 non-null    float64
10  slope       303 non-null    int64
11  ca          303 non-null    int64
12  thal        303 non-null    int64
13  target      303 non-null    int64
dtypes: float64(1), int64(13)
memory usage: 33.3 KB
```

- Out of 14 features, I have 13 int type and only one with float data type.
- Woah! We have no missing values in our dataset.

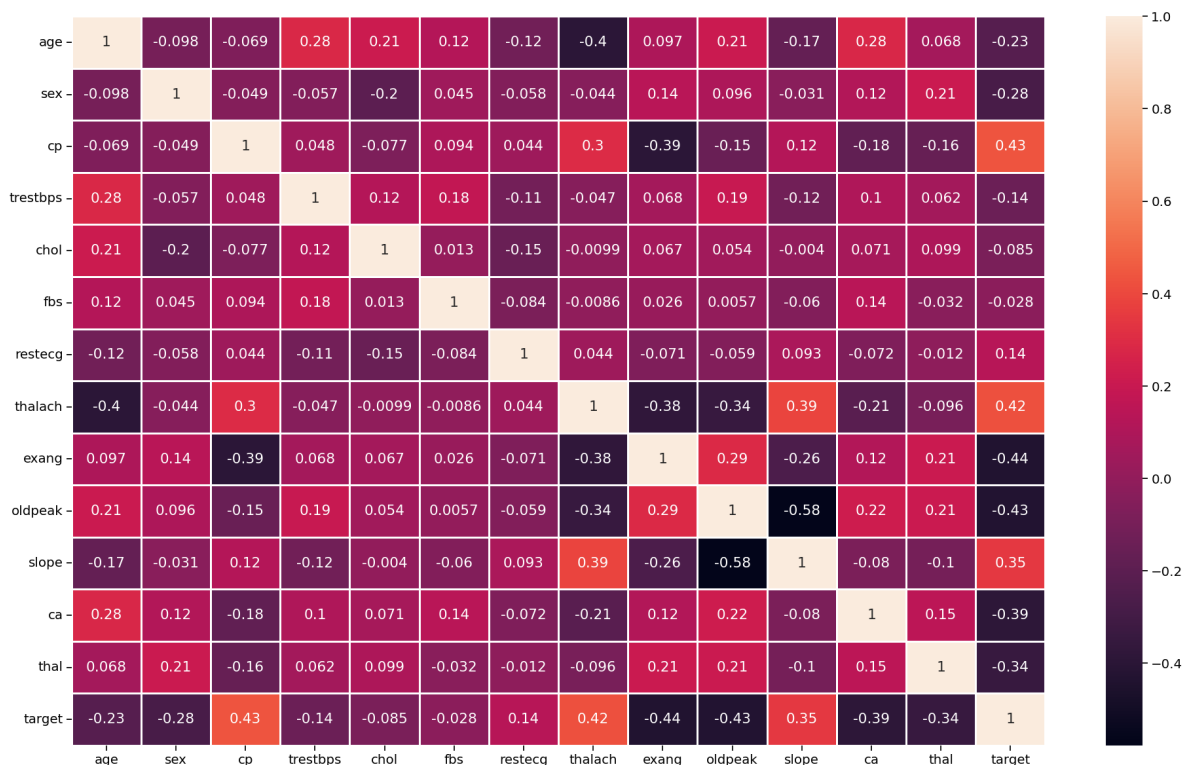
In [5]: data.describe()

Out[5]:

	age	sex	cp	trestbps	chol	fbs	restecg	
count	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303.000000	303
mean	54.366337	0.683168	0.966997	131.623762	246.264026	0.148515	0.528053	149
std	9.082101	0.466011	1.032052	17.538143	51.830751	0.356198	0.525860	22
min	29.000000	0.000000	0.000000	94.000000	126.000000	0.000000	0.000000	71
25%	47.500000	0.000000	0.000000	120.000000	211.000000	0.000000	0.000000	133
50%	55.000000	1.000000	1.000000	130.000000	240.000000	0.000000	1.000000	153
75%	61.000000	1.000000	2.000000	140.000000	274.500000	0.000000	1.000000	166
max	77.000000	1.000000	3.000000	200.000000	564.000000	1.000000	2.000000	202

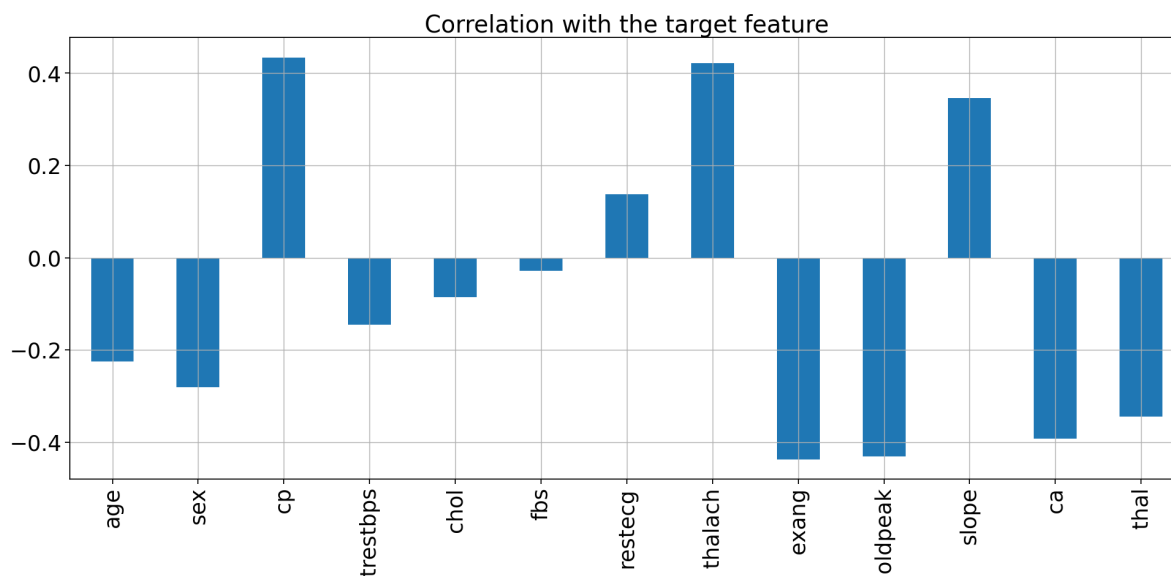
Let's check correleation between various features.

```
In [6]: plt.figure(figsize=(20,12))
sns.set_context('notebook',font_scale = 1.3)
sns.heatmap(data.corr(),annot=True,linewidth =2)
plt.tight_layout()
```



Let's check the correlation of various features with the target feature.

```
In [7]: # plt.figure(figsize=(20,12))
sns.set_context('notebook',font_scale = 2.3)
data.drop('target', axis=1).corrwith(data.target).plot(kind='bar', grid=True,
title="Correlation with the target feature")
plt.tight_layout()
```



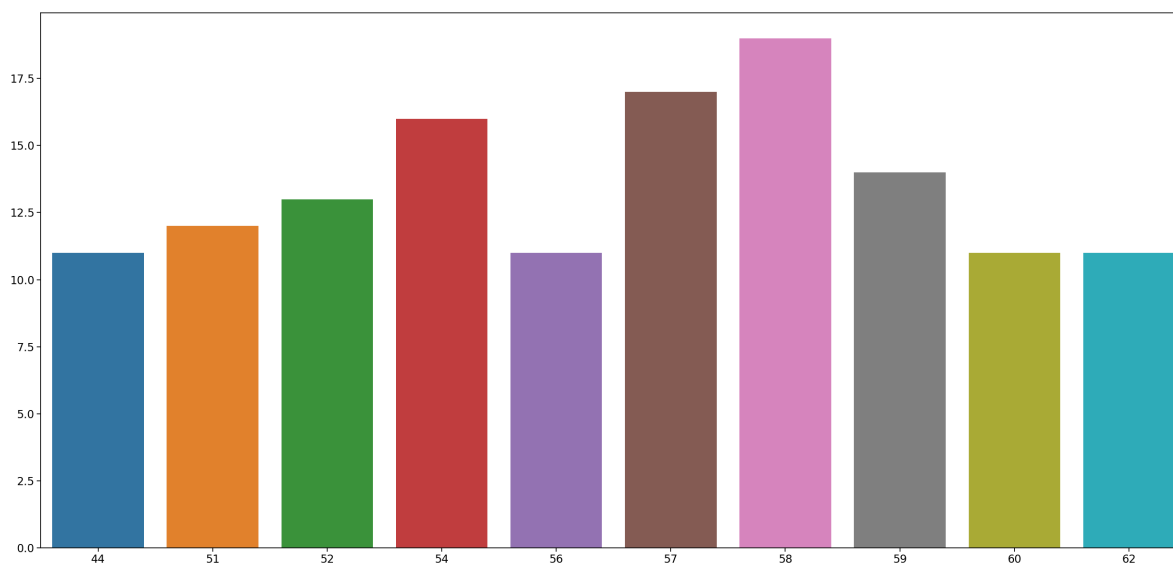
- Four feature("cp", "restecg", "thalach", "slope") are positively correlated with the target feature.
- Other features are negatively correlated with the target feature.

Individual Feature Analysis

Age("age") Analysis

In [8]: *# Let's check 10 ages and their count*

```
plt.figure(figsize=(25,12))
sns.set_context('notebook',font_scale = 1.5)
sns.barplot(x=data.age.value_counts()[:10].index,y=data.age.value_counts()[:10])
plt.tight_layout()
```



Let's check the range of age in the dataset.

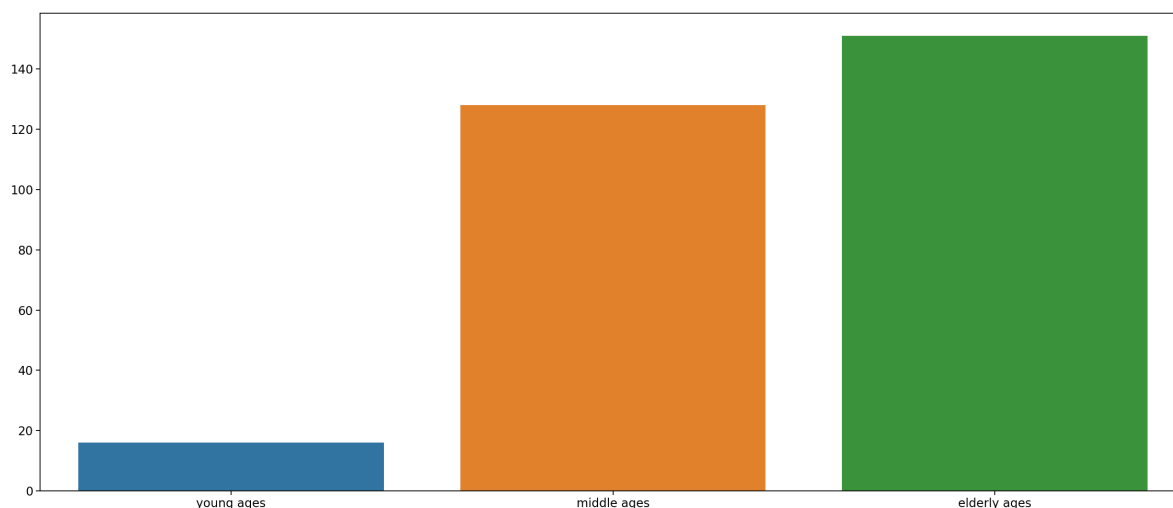
```
In [9]: minAge=min(data.age)
maxAge=max(data.age)
meanAge=data.age.mean()
print('Min Age :',minAge)
print('Max Age :',maxAge)
print('Mean Age :',meanAge)
```

```
Min Age : 29
Max Age : 77
Mean Age : 54.366336633663366
```

I should divide the Age feature into three parts - "Young", "Middle" and "Elder"

```
In [10]: Young = data[(data.age>=29)&(data.age<40)]  
Middle = data[(data.age>=40)&(data.age<55)]  
Elder = data[(data.age>55)]
```

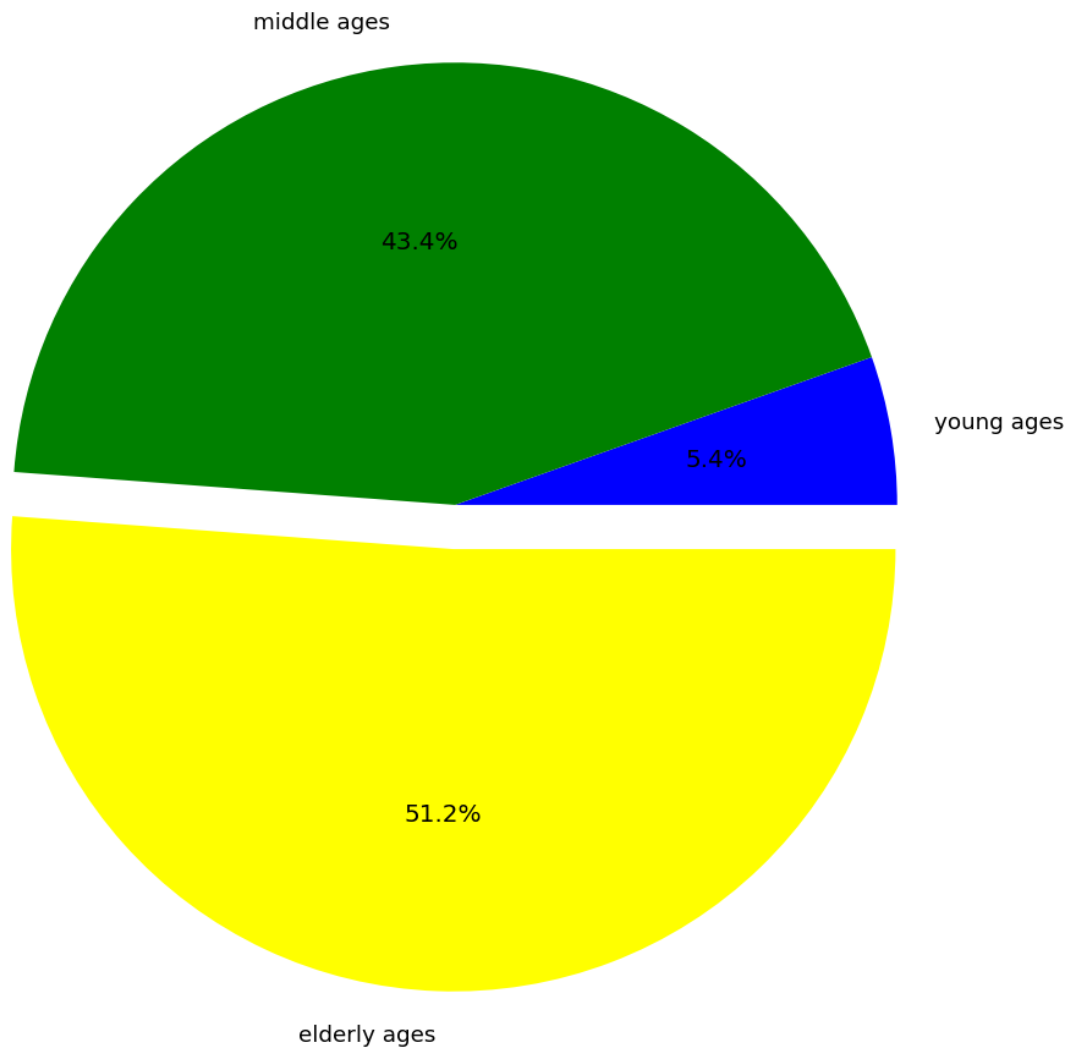
```
In [11]: plt.figure(figsize=(23,10))  
sns.set_context('notebook',font_scale = 1.5)  
sns.barplot(x=['young ages','middle ages','elderly ages'],y=[len(Young),len(Mi  
plt.tight_layout()
```



A large proportion of dataset contains Elder people.

Elderly people are more likely to suffer from heart disease.

```
In [12]: colors = ['blue', 'green', 'yellow']  
explode = [0,0,0.1]  
plt.figure(figsize=(10,10))  
sns.set_context('notebook', font_scale = 1.2)  
plt.pie([len(Young), len(Middle), len(Elder)], labels=['young ages', 'middle ages',  
plt.tight_layout()
```



Sex("sex") Feature Analysis

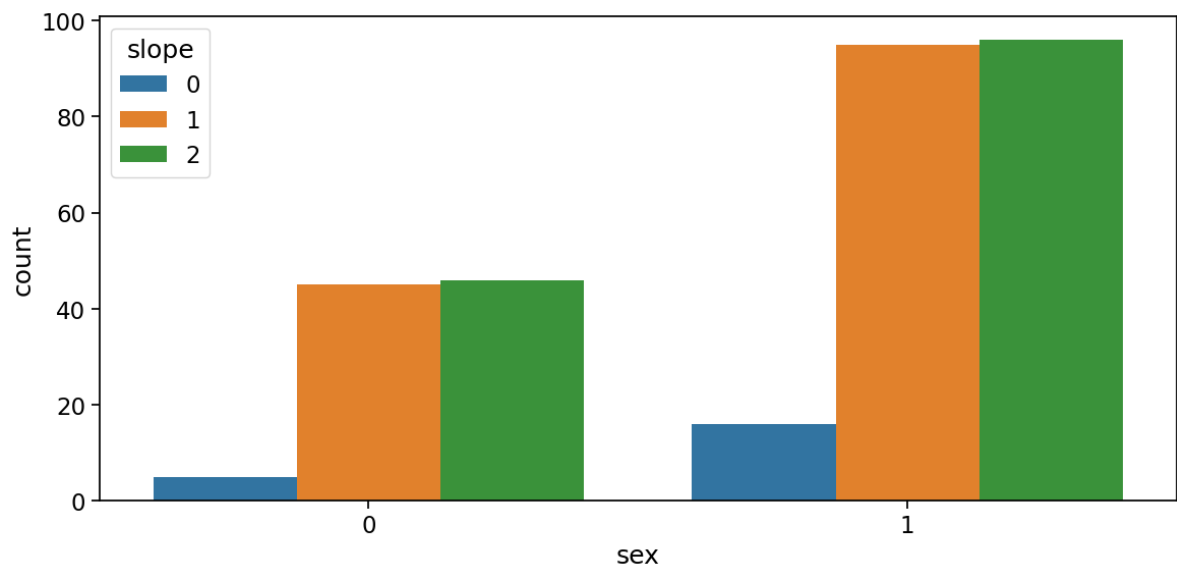
```
In [13]: plt.figure(figsize=(18,9))  
sns.set_context('notebook',font_scale = 1.5)  
sns.countplot(data['sex'])  
plt.tight_layout()
```



Ratio of Male to Female is approx 2:1


```
In [14]: # Let's plot the relation between sex and slope.
```

```
plt.figure(figsize=(12, 6))  
sns.set_context('notebook', font_scale=1.5)  
  
# Assuming 'sex' is on the x-axis and 'slope' is on the y-axis  
sns.countplot(x='sex', hue='slope', data=data)  
  
plt.tight_layout()  
plt.show()
```

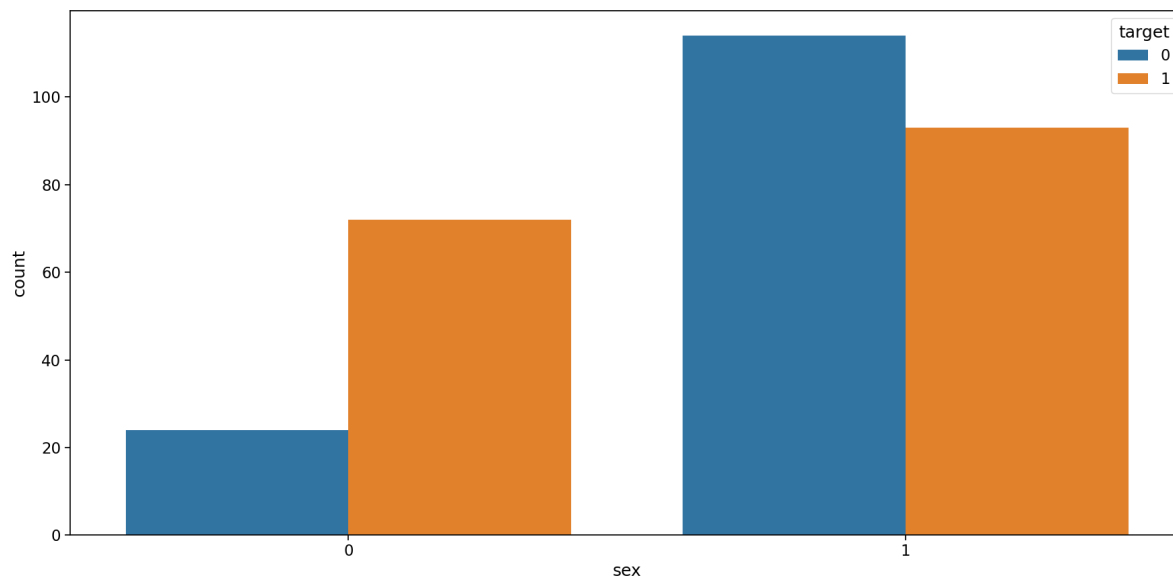


In [15]: *# Let's plot the relation between sex and target.*

```
plt.figure(figsize=(18, 9))
sns.set_context('notebook', font_scale=1.5)

# Assuming 'sex' is on the x-axis
sns.countplot(x='sex', hue='target', data=data)

plt.tight_layout()
plt.show()
```



Males are more likely to have heart disease than Female.

Chest Pain Type("cp") Analysis

```
In [16]: plt.figure(figsize=(18,9))  
sns.set_context('notebook',font_scale = 1.5)  
sns.countplot(data['cp'])  
plt.tight_layout()
```



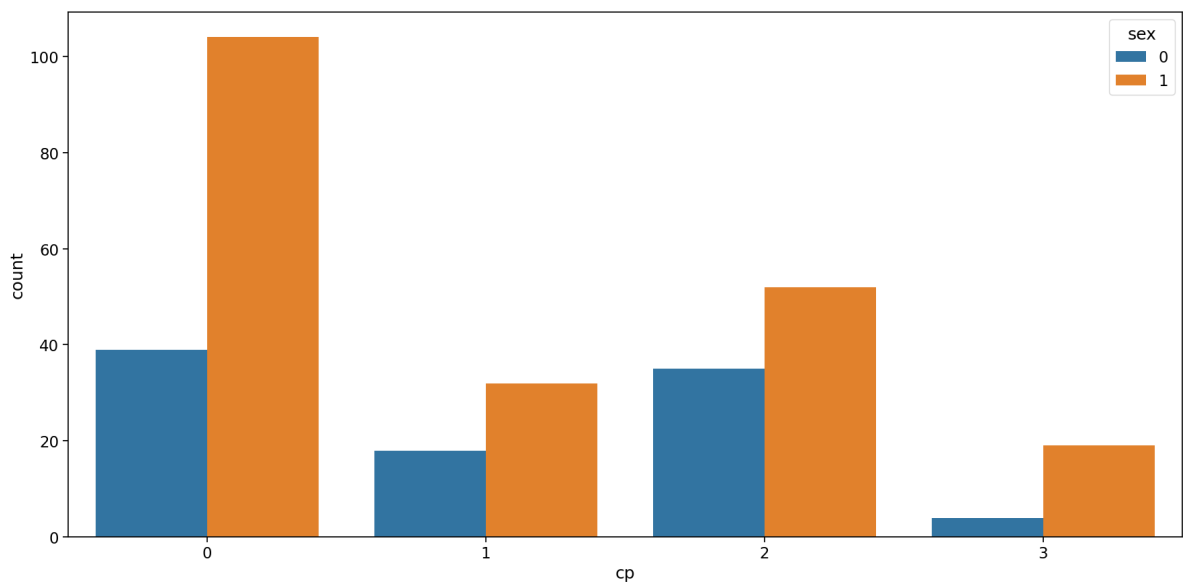
As seen, there are 4 types of chest pain

1. status at least
2. condition slightly distressed
3. condition medium problem
4. condition too bad

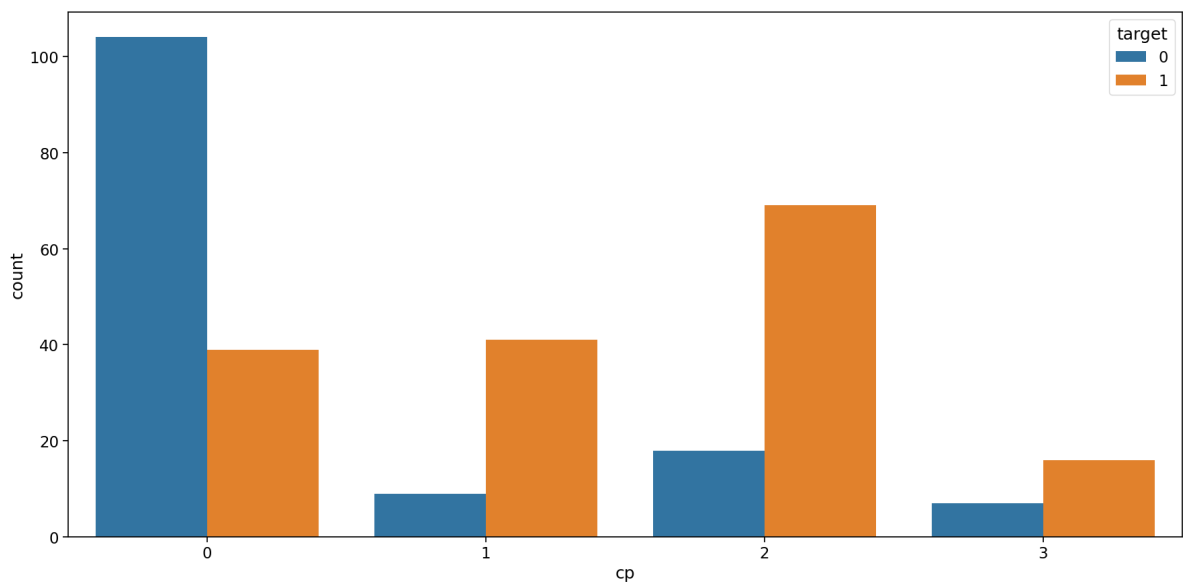
```
In [17]: plt.figure(figsize=(18, 9))
sns.set_context('notebook', font_scale=1.5)

# Assuming 'cp' is on the x-axis
sns.countplot(x='cp', hue='sex', data=data)

plt.tight_layout()
plt.show()
```



```
In [18]: plt.figure(figsize=(18,9))
sns.set_context('notebook', font_scale=1.5)
sns.countplot(x='cp', hue='target', data=data)
plt.tight_layout()
plt.show()
```



- People having least chest pain are not likely to heart disease.
- People having severe chest pain are likely to heart disease.

Elderly people are more likely to have chest pain.

Thal Analysis

```
In [19]: plt.figure(figsize=(18,9))
sns.set_context('notebook',font_scale = 1.5)
sns.countplot(data['thal'])
plt.tight_layout()
```



```
In [20]: data.head()
```

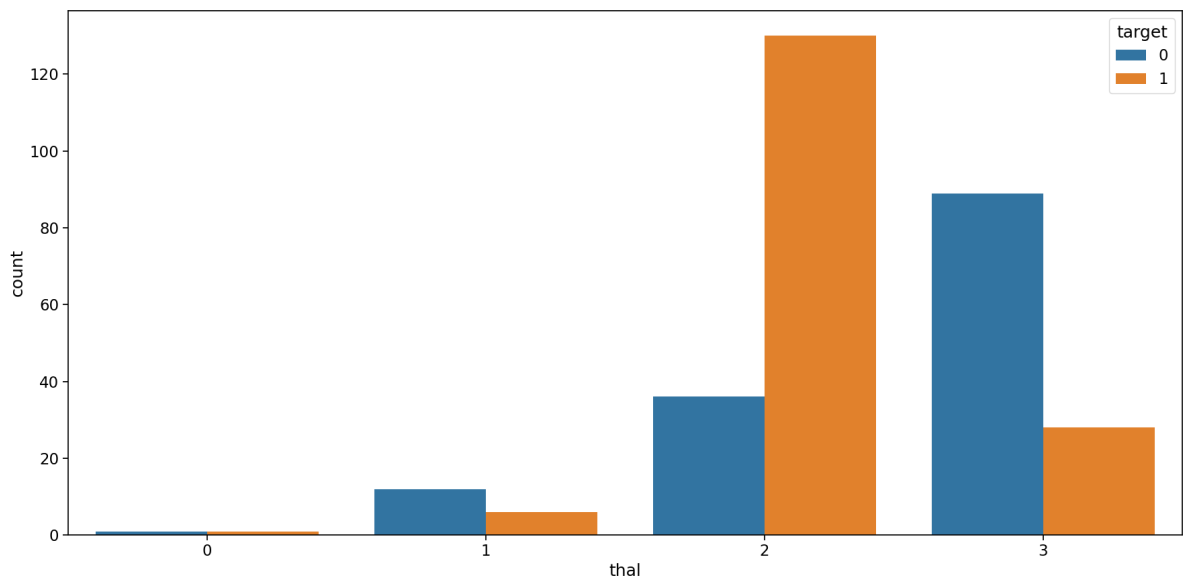
Out[20]:

	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

1. 3 = normal
2. 6 = fixed defect
3. 7 = reversable defect

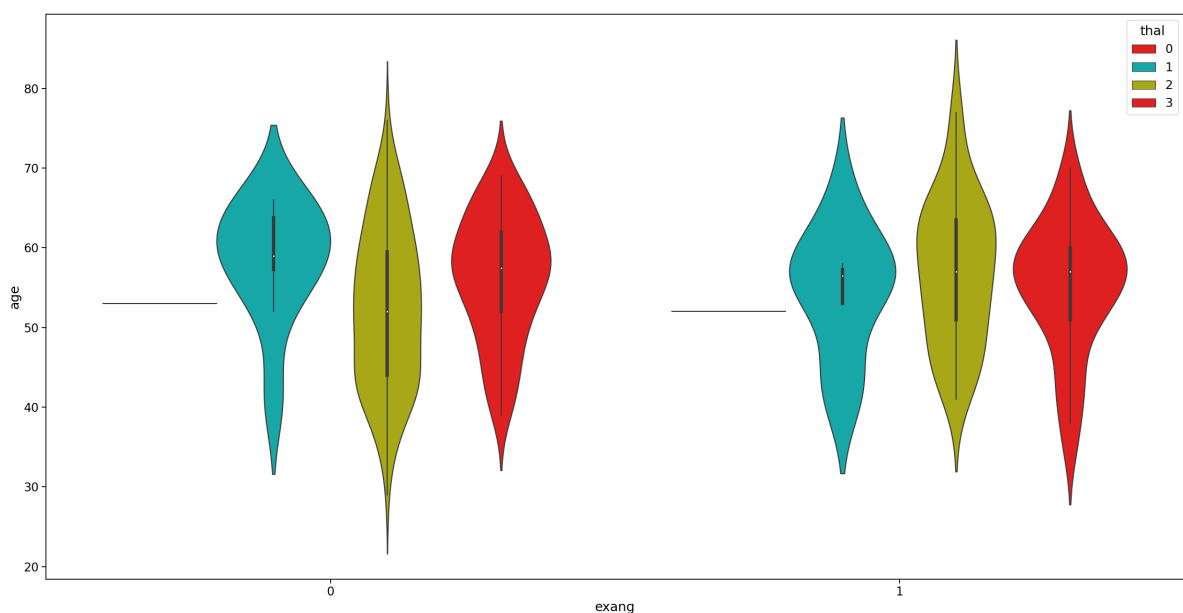
```
In [21]: plt.figure(figsize=(18,9))
plt.figure(figsize=(18, 9))
sns.set_context('notebook', font_scale=1.5)
sns.countplot(x='thal', hue='target', data=data)
plt.tight_layout()
plt.show()
```

<Figure size 1800x900 with 0 Axes>



People with fixed defect are more likely to have heart disease.

```
In [22]: plt.figure(figsize=(23,12))
sns.set_context('notebook', font_scale = 1.5)
sns.violinplot(x="exang", y="age", data=data, palette=["r", "c", "y"], hue="thal")
plt.tight_layout()
```



Target

```
In [23]: plt.figure(figsize=(18,9))
sns.set_context('notebook',font_scale = 1.5)
sns.countplot(data['target'])
plt.tight_layout()
```



Inference: The ratio between 1 and 0 is much less than 1.5 which indicates that target feature is not imbalanced. So for a balanced dataset, we can use accuracy_score as evaluation metrics for our model.

```
In [24]: data.head()
```

Out[24]:

	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

Feature Engineering

```
In [25]: categorical_val = []
continous_val = []
for column in data.columns:
    print("-----")
    print(f"{column} : {data[column].unique()}")
    if len(data[column].unique()) <= 10:
        categorical_val.append(column)
    else:
        continous_val.append(column)
```



```

-----
age : [63 37 41 56 57 44 52 54 48 49 64 58 50 66 43 69 59 42 61 40 71 51 65 5
3
46 45 39 47 62 34 35 29 55 60 67 68 74 76 70 38 77]
-----
sex : [1 0]
-----
cp : [3 2 1 0]
-----
trestbps : [145 130 120 140 172 150 110 135 160 105 125 142 155 104 138 128 1
08 134
122 115 118 100 124 94 112 102 152 101 132 148 178 129 180 136 126 106
156 170 146 117 200 165 174 192 144 123 154 114 164]
-----
chol : [233 250 204 236 354 192 294 263 199 168 239 275 266 211 283 219 340 2
26
247 234 243 302 212 175 417 197 198 177 273 213 304 232 269 360 308 245
208 264 321 325 235 257 216 256 231 141 252 201 222 260 182 303 265 309
186 203 183 220 209 258 227 261 221 205 240 318 298 564 277 214 248 255
207 223 288 160 394 315 246 244 270 195 196 254 126 313 262 215 193 271
268 267 210 295 306 178 242 180 228 149 278 253 342 157 286 229 284 224
206 167 230 335 276 353 225 330 290 172 305 188 282 185 326 274 164 307
249 341 407 217 174 281 289 322 299 300 293 184 409 259 200 327 237 218
319 166 311 169 187 176 241 131]
-----
fbs : [1 0]
-----
restecg : [0 1 2]
-----
thalach : [150 187 172 178 163 148 153 173 162 174 160 139 171 144 158 114 15
1 161
179 137 157 123 152 168 140 188 125 170 165 142 180 143 182 156 115 149
146 175 186 185 159 130 190 132 147 154 202 166 164 184 122 169 138 111
145 194 131 133 155 167 192 121 96 126 105 181 116 108 129 120 112 128
109 113 99 177 141 136 97 127 103 124 88 195 106 95 117 71 118 134
90]
-----
exang : [0 1]
-----
oldpeak : [2.3 3.5 1.4 0.8 0.6 0.4 1.3 0. 0.5 1.6 1.2 0.2 1.8 1. 2.6 1.5 3.
2.4
0.1 1.9 4.2 1.1 2. 0.7 0.3 0.9 3.6 3.1 3.2 2.5 2.2 2.8 3.4 6.2 4. 5.6
2.9 2.1 3.8 4.4]
-----
slope : [0 2 1]
-----
ca : [0 2 1 3 4]
-----
thal : [1 2 3 0]
-----
target : [1 0]

```

Now here first we will be removing the target column from our set of features then we will categorised all the categorical variables using get dummies method which will create the seperate column for each category suppose X variable contains 2 types of unique values then it

```
In [26]: categorical_val.remove('target')
dfs = pd.get_dummies(data, columns = categorical_val)
```

```
In [27]: dfs.head(6)
```

Out[27]:

	age	trestbps	chol	thalach	oldpeak	target	sex_0	sex_1	cp_0	cp_1	...	slope_2	ca_0	ca_1
0	63	145	233	150	2.3	1	0	1	0	0	...	0	1	0
1	37	130	250	187	3.5	1	0	1	0	0	...	0	1	0
2	41	130	204	172	1.4	1	1	0	0	1	...	1	1	0
3	56	120	236	178	0.8	1	0	1	0	1	...	1	1	0
4	57	120	354	163	0.6	1	1	0	1	0	...	1	1	0
5	57	140	192	148	0.4	1	0	1	1	0	...	0	1	0

6 rows × 31 columns



Now we will be using the standard scaler method to scale down the data so that it won't raise the outliers also dataset which is scaled to general units leads to have better accuracy.

```
In [28]: sc = StandardScaler()
col_to_scale = ['age', 'trestbps', 'chol', 'thalach', 'oldpeak']
dfs[col_to_scale] = sc.fit_transform(dfs[col_to_scale])
```

```
In [29]: dfs.head(6)
```

Out[29]:

	age	trestbps	chol	thalach	oldpeak	target	sex_0	sex_1	cp_0	cp_1	...	slope_2	ca_0	ca_1
0	0.952197	0.763956	-0.256334	0.015443	1.087338	1	0	1	0	0	...	0	1	0
1	-1.915313	-0.092738	0.072199	1.633471	2.122573	1	0	1	0	0	...	0	1	0
2	-1.474158	-0.092738	-0.816773	0.977514	0.310912	1	1	0	0	1	...	1	1	0
3	0.180175	-0.663867	-0.198357	1.239897	-0.206705	1	0	1	0	1	...	1	1	0
4	0.290464	-0.663867	2.082050	0.583939	-0.379244	1	1	0	1	0	...	1	1	0
5	0.290464	0.478391	-1.048678	-0.072018	-0.551783	1	0	1	1	0	...	0	1	0

6 rows × 31 columns



Modelling

Splitting our dataset

```
In [30]: X = dfs.drop('target', axis=1)
y = dfs.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
```

```
In [31]: X_train.head()
```

```
Out[31]:
```

	age	trestbps	chol	thalach	oldpeak	sex_0	sex_1	cp_0	cp_1	cp_2	...	s
124	-1.694735	-2.148802	-0.913400	1.283627	-0.896862	1	0	0	0	1	...	s
72	-2.797624	-0.092738	-0.816773	2.289429	-0.896862	0	1	0	1	0	...	s
15	-0.481558	-0.663867	-0.526890	0.365287	0.483451	1	0	0	0	1	...	s
10	-0.040403	0.478391	-0.140381	0.452748	0.138373	0	1	1	0	0	...	s
163	-1.805024	0.364165	-1.377212	1.021244	-0.896862	0	1	0	0	1	...	s

5 rows × 30 columns



Next I will work on following algorithms -

- KNN
- Random Forest Classifier
- XGBoost
- CatBoost

KNN

```
In [32]: knn = KNeighborsClassifier(n_neighbors = 10)
```

```
In [33]: knn.fit(X_train,y_train)
```

```
Out[33]:
```

▼

KNeighborsClassifier

KNeighborsClassifier(n_neighbors=10)

```
In [34]: print(X_test.flags)

<Flags(allows_duplicate_labels=True)>
```

```
In [35]: X_test = np.ascontiguousarray(X_test)
```

```
In [36]: y_pred1 = knn.predict(X_test)
```

```
D:\User\Yamuna\Lib\site-packages\sklearn\base.py:464: UserWarning: X does not  
have valid feature names, but KNeighborsClassifier was fitted with feature na  
mes  
  warnings.warn(
```

```
In [37]: print(accuracy_score(y_test,y_pred1))
```

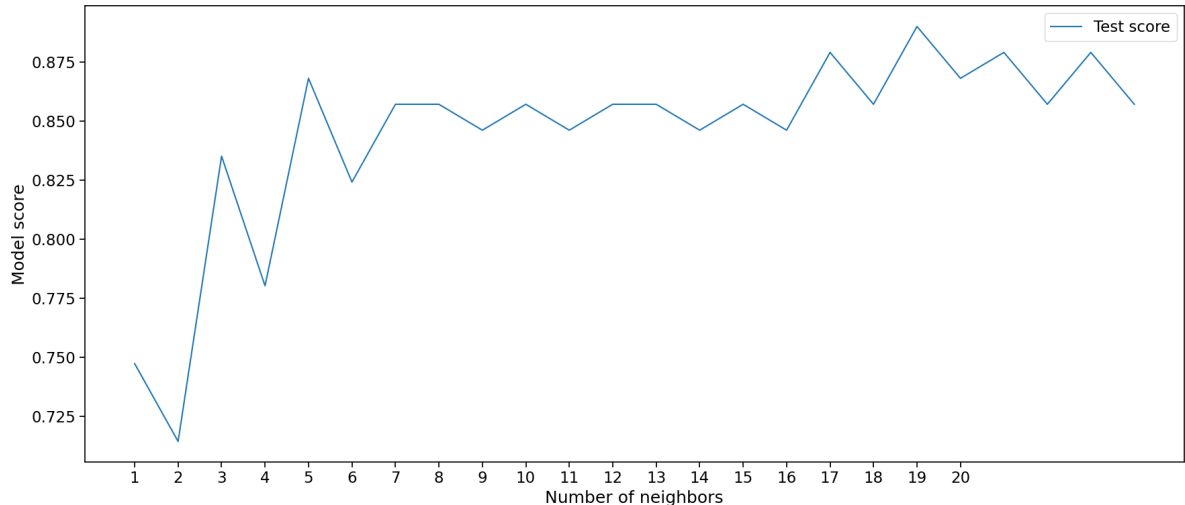
```
0.8571428571428571
```

In [38]: *# Hyperparameter Optimization*

```
test_score = []  
neighbors = range(1, 25)  
  
for k in neighbors:  
    model = KNeighborsClassifier(n_neighbors=k)  
    model.fit(X_train, y_train)  
    test_score.append(accuracy_score(y_test, model.predict(X_test)))
```

[illegible]


```
In [39]: plt.figure(figsize=(18, 8))
plt.plot(neighbors, test_score, label="Test score")
plt.xticks(np.arange(1, 21, 1))
plt.xlabel("Number of neighbors")
plt.ylabel("Model score")
plt.legend()
plt.tight_layout()
```



At K=19, i am getting highest test accuracy.

```
In [40]: knn = KNeighborsClassifier(n_neighbors = 19)
```

```
In [41]: knn.fit(X_train,y_train)
```

```
Out[41]: KNeighborsClassifier
KNeighborsClassifier(n_neighbors=19)
```

```
In [42]: y_pred1 = knn.predict(X_test)
```

```
D:\User\Yamuna\Lib\site-packages\sklearn\base.py:464: UserWarning: X does not
have valid feature names, but KNeighborsClassifier was fitted with feature na
mes
  warnings.warn(
```

```
In [43]: print(accuracy_score(y_test,y_pred1))
```

```
0.8901098901098901
```

I achieved accuracy 89% with KNN Model after Hyperparameter Optimization.

Random Forest Classifier

```
In [44]: rfc = RandomForestClassifier()
         rfc.fit(X_train,y_train)
         y_pred2 = rfc.predict(X_test)
```

```
D:\User\Yamuna\Lib\site-packages\sklearn\base.py:464: UserWarning: X does not
have valid feature names, but RandomForestClassifier was fitted with feature
names
  warnings.warn(
```

```
In [45]: print(accuracy_score(y_test,y_pred2))
```

0.8241758241758241

```
In [46]: ## Hyperparameter Optimization
```

```
max_depth = [int(x) for x in np.linspace(10, 110, num=11)]
max_depth.append(None)
```

```
params2 = {
```

```
'n_estimators': [int(x) for x in np.linspace(start=200, stop=2000, num=10)]
'max_features': ['auto', 'sqrt'],
'max_depth': max_depth,
'min_samples_split': [2, 5, 10],
'min_samples_leaf': [1, 2, 4],
'bootstrap': [True, False]
```

```
In [47]: rfc = RandomForestClassifier(random_state=42)
```

```
rfcs = RandomizedSearchCV(estimator=rfc, param_distributions=params2, n_iter=1
```

```
In [48]: rfcs.fit(X_train,y_train)
```

Fitting 5 folds for each of 100 candidates, totalling 500 fits

```
D:\User\Yamuna\Lib\site-packages\sklearn\model_selection\_validation.py:425:
FitFailedWarning:
205 fits failed out of a total of 500.
The score on these train-test partitions for these parameters will be set to
nan.
If these failures are not expected, you can try to debug them by setting erro
r_score='raise'.
```

Below are more details about the failures:

```
-----
---
104 fits failed with the following error:
Traceback (most recent call last):
  File "D:\User\Yamuna\Lib\site-packages\sklearn\model_selection\_validation.
py", line 732, in _fit_and_score
    estimator.fit(X_train, y_train, **fit_params)
  File "D:\User\Yamuna\Lib\site-packages\sklearn\base.py", line 1144, in wrap
per
    estimator._validate_params()
  File "D:\User\Yamuna\Lib\site-packages\sklearn\base.py", line 637, in _vali
date_params
    validate_parameter_constraints(
  File "D:\User\Yamuna\Lib\site-packages\sklearn\utils\_param_validation.py",
line 95, in validate_parameter_constraints
    raise InvalidParameterError(
sklearn.utils._param_validation.InvalidParameterError: The 'max_features' par
ameter of RandomForestClassifier must be an int in the range [1, inf), a floa
t in the range (0.0, 1.0], a str among {'sqrt', 'log2'} or None. Got 'auto' i
nstead.

-----
---
101 fits failed with the following error:
Traceback (most recent call last):
  File "D:\User\Yamuna\Lib\site-packages\sklearn\model_selection\_validation.
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ameter of RandomForestClassifier must be an int in the range [1, inf), a floa
t in the range (0.0, 1.0], a str among {'log2', 'sqrt'} or None. Got 'auto' i
nstead.

    warnings.warn(some_fits_failed_message, FitFailedWarning)
D:\User\Yamuna\Lib\site-packages\sklearn\model_selection\_search.py:976: User
Warning: One or more of the test scores are non-finite: [0.82513843 0.8157253
6 0.82059801          nan 0.82990033 0.81572536
0.82059801 0.80177187          nan          nan          nan 0.80642303
0.8158361 0.82059801 0.82535991          nan          nan 0.8158361
```

```

nan nan 0.8158361 0.81118494 0.82513843 nan
0.81118494 0.81129568 nan 0.82502769 nan 0.82502769
0.81572536 0.8158361 0.80642303 0.82990033 0.82990033 0.82535991
0.82502769 nan nan 0.82037652 nan 0.8345515
0.82990033 0.8345515 nan 0.8158361 nan 0.80631229
0.81572536 0.82048726 nan nan nan nan
0.82059801 0.81572536 nan nan nan 0.82978959
0.82524917 0.82513843 nan nan 0.81572536 nan
nan nan 0.81118494 nan 0.82048726 0.81572536
0.82502769 nan nan 0.82059801 0.81572536 0.81572536
0.82502769 nan nan nan nan nan
0.81572536 0.81118494 0.82513843 0.82990033 0.82990033 0.82037652
0.82990033 0.81572536 0.82037652 nan 0.81572536 nan
nan nan 0.81572536 nan]
warnings.warn(

```

Out[48]:

```

RandomizedSearchCV
  estimator: RandomForestClassifier
    RandomForestClassifier

```

In [49]: rfcs.best_estimator_

Out[49]:

```

RandomForestClassifier
RandomForestClassifier(max_depth=60, min_samples_leaf=2, min_samples_split=1
0,
n_estimators=600, random_state=42)

```

```

In [50]: y_pred2 = rfcs.predict(X_test)
print(accuracy_score(y_test,y_pred2))

```

0.8351648351648352

D:\User\Yamuna\Lib\site-packages\sklearn\base.py:464: UserWarning: X does not have valid feature names, but RandomForestClassifier was fitted with feature names

```

warnings.warn(

```

I achieved accuracy 83% approx with Random Forest Classifier Model. There is no improvement after Hyperparameter Optimization.

XGBoost

```

In [51]: xgb = XGBClassifier(random_state = 42)
xgb.fit(X_train,y_train)
y_pred3 = xgb.predict(X_test)

```

```
In [52]: print(accuracy_score(y_test,y_pred3))
```

```
0.8241758241758241
```

I achieved accuracy 83% approx with XGBoost Classifier Model.

CatBoost

```
In [53]: model4 = CatBoostClassifier(random_state=42)
```

```
In [54]: model4.fit(X_train,y_train)
y_pred4 = model4.predict(X_test)
```

```
51:   learn: 0.5158074      total: 203ms   remaining: 3.65s
52:   learn: 0.5164422      total: 204ms   remaining: 3.65s
53:   learn: 0.5141062      total: 206ms   remaining: 3.61s
54:   learn: 0.5117230      total: 208ms   remaining: 3.57s
55:   learn: 0.5094912      total: 210ms   remaining: 3.54s
56:   learn: 0.5079803      total: 211ms   remaining: 3.49s
57:   learn: 0.5056247      total: 213ms   remaining: 3.46s
58:   learn: 0.5034384      total: 215ms   remaining: 3.43s
59:   learn: 0.5012510      total: 217ms   remaining: 3.4s
60:   learn: 0.4984309      total: 219ms   remaining: 3.36s
61:   learn: 0.4955346      total: 220ms   remaining: 3.33s
62:   learn: 0.4930701      total: 222ms   remaining: 3.3s
63:   learn: 0.4908856      total: 224ms   remaining: 3.28s
64:   learn: 0.4887024      total: 226ms   remaining: 3.25s
65:   learn: 0.4867662      total: 228ms   remaining: 3.22s
66:   learn: 0.4846121      total: 230ms   remaining: 3.2s
67:   learn: 0.4821562      total: 231ms   remaining: 3.17s
68:   learn: 0.4800911      total: 233ms   remaining: 3.14s
69:   learn: 0.4794304      total: 234ms   remaining: 3.11s
70:   learn: 0.4772768      total: 235ms   remaining: 3.08s
71:   learn: 0.4750000      total: 237ms   remaining: 3.06s
```

```
In [55]: print(accuracy_score(y_test,y_pred4))
```

```
0.8131868131868132
```

I achieved accuracy 81% approx with CatBoost Classifier Model.

Conclusion

From the above models KNN is giving us the best accuracy which is 89%.