# **Problem Statement:-**

# Which model is suitable for Insurance Dataset

# Import all the required packages

#### In [1]:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
```

# **Step-1:- Data Collection**

# In [2]:

```
df=pd.read_csv(r"C:\Users\thara\Downloads\insurance.csv")
df
```

#### Out[2]:

	age	sex	bmi	children	smoker	region	charges
0	19	female	27.900	0	yes	southwest	16884.92400
1	18	male	33.770	1	no	southeast	1725.55230
2	28	male	33.000	3	no	southeast	4449.46200
3	33	male	22.705	0	no	northwest	21984.47061
4	32	male	28.880	0	no	northwest	3866.85520
1333	50	male	30.970	3	no	northwest	10600.54830
1334	18	female	31.920	0	no	northeast	2205.98080
1335	18	female	36.850	0	no	southeast	1629.83350
1336	21	female	25.800	0	no	southwest	2007.94500
1337	61	female	29.070	0	yes	northwest	29141.36030

<sup>1338</sup> rows × 7 columns

# **Step-2:- Data Preprocessing**

## In [3]:

# The shape of a DataFrame is a tuple of array dimensions that tells the number of rows and columns of a given DataFrame df.shape

# Out[3]:

(1338, 7)

#### In [4]:

```
# The head() returns the first n rows for the object based on position.
df.head()
```

#### Out[4]:

charges	region	smoker	children	bmi	sex	age	
16884.92400	southwest	yes	0	27.900	female	19	0
1725.55230	southeast	no	1	33.770	male	18	1
4449.46200	southeast	no	3	33.000	male	28	2
21984.47061	northwest	no	0	22.705	male	33	3
3866.85520	northwest	no	0	28.880	male	32	4

## In [5]:

```
# The tail function in Python displays the last five rows of the dataframe by default.
df.tail()
```

#### Out[5]:

	age	sex	bmi	children	smoker	region	charges
133	<b>33</b> 50	male	30.97	3	no	northwest	10600.5483
133	<b>34</b> 18	female	31.92	0	no	northeast	2205.9808
133	<b>35</b> 18	female	36.85	0	no	southeast	1629.8335
133	<b>36</b> 21	female	25.80	0	no	southwest	2007.9450
133	<b>37</b> 61	female	29.07	0	yes	northwest	29141.3603

#### In [6]:

```
# To check the null values.
df.isnull().sum()
```

## Out[6]:

age 0
sex 0
bmi 0
children 0
smoker 0
region 0
charges 0
dtype: int64

There Are No Null Values In The Above DataFrame

## In [7]:

```
# Check the data types for each column.
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1338 entries, 0 to 1337
Data columns (total 7 columns):
# Column
              Non-Null Count Dtype
              1338 non-null
 0
                             int64
    age
1
    sex
              1338 non-null
                              object
              1338 non-null
                              float64
    bmi
 3
    children 1338 non-null
                             int64
 4
    smoker
              1338 non-null
                              object
              1338 non-null
    region
                              object
   charges
             1338 non-null
                              float64
dtypes: float64(2), int64(2), object(3)
memory usage: 73.3+ KB
```

#### In [8]:

```
# Explore numerical columns stats.
# The describe function only displays the numerical columns present in the Dataset.

df.describe()
```

#### Out[8]:

	age	bmi	children	charges
count	1338.000000	1338.000000	1338.000000	1338.000000
mean	39.207025	30.663397	1.094918	13270.422265
std	14.049960	6.098187	1.205493	12110.011237
min	18.000000	15.960000	0.000000	1121.873900
25%	27.000000	26.296250	0.000000	4740.287150
50%	39.000000	30.400000	1.000000	9382.033000
75%	51.000000	34.693750	2.000000	16639.912515
max	64.000000	53.130000	5.000000	63770.428010

#### In [9]:

```
# Explore our age variable.
df.age.describe()
```

## Out[9]:

```
1338.000000
count
          39.207025
mean
           14.049960
std
          18.000000
min
25%
           27.000000
50%
           39.000000
75%
           51.000000
           64.000000
max
Name: age, dtype: float64
```

## In [10]:

```
df.duplicated().sum()
```

## Out[10]:

1

## In [11]:

```
df=df.drop_duplicates()
df
```

## Out[11]:

	age	sex	bmi	children	smoker	region	charges
0	19	female	27.900	0	yes	southwest	16884.92400
1	18	male	33.770	1	no	southeast	1725.55230
2	28	male	33.000	3	no	southeast	4449.46200
3	33	male	22.705	0	no	northwest	21984.47061
4	32	male	28.880	0	no	northwest	3866.85520
1333	50	male	30.970	3	no	northwest	10600.54830
1334	18	female	31.920	0	no	northeast	2205.98080
1335	18	female	36.850	0	no	southeast	1629.83350
1336	21	female	25.800	0	no	southwest	2007.94500
1337	61	female	29.070	0	yes	northwest	29141.36030

1337 rows × 7 columns

```
In [12]:
```

```
J={"sex":{"female":1,"male":0}}
df=df.replace(J)
print(df)

age sex bmi children smoker region charges
```

```
0
               27.900
                                        southwest 16884.92400
       19
            1
                               0
                                   yes
1
       18
             0
               33.770
                               1
                                    no
                                         southeast
                                                    1725.55230
             0 33.000
                                                     4449.46200
2
       28
                               3
                                    no
                                        southeast
             0 22.705
                               0
                                        northwest 21984.47061
3
       33
                                     no
4
       32
               28.880
                               0
                                    no
                                        northwest
                                                    3866.85520
1333
       50
            0
               30.970
                                        northwest 10600.54830
                               3
                                    no
            1 31.920
       18
                               0
                                    no northeast
                                                     2205.98080
1334
1335
       18
             1
               36.850
                               0
                                    no
                                         southeast
                                                     1629.83350
1336
       21
            1 25.800
                               0
                                    no southwest
                                                    2007.94500
1337
       61
             1 29.070
                               0
                                   yes northwest 29141.36030
```

[1337 rows x 7 columns]

```
In [13]:
```

```
J={"smoker":{"yes":1,"no":0}}
df=df.replace(J)
df
```

#### Out[13]:

	age	sex	bmi	children	smoker	region	charges
0	19	1	27.900	0	1	southwest	16884.92400
1	18	0	33.770	1	0	southeast	1725.55230
2	28	0	33.000	3	0	southeast	4449.46200
3	33	0	22.705	0	0	northwest	21984.47061
4	32	0	28.880	0	0	northwest	3866.85520
1333	50	0	30.970	3	0	northwest	10600.54830
1334	18	1	31.920	0	0	northeast	2205.98080
1335	18	1	36.850	0	0	southeast	1629.83350
1336	21	1	25.800	0	0	southwest	2007.94500
1337	61	1	29.070	0	1	northwest	29141.36030

1337 rows × 7 columns

# Feature Scaling:

Feature Scaling is done to normalize the features in the dataset into a finite range.

## In [14]:

```
#Training the model

X=df[['age', 'sex', 'bmi', 'children', 'smoker']]
y=df['charges']
```

## In [15]:

```
from sklearn.model_selection import train_test_split
X_train,X_test,y_train,y_test=train_test_split(X,y,test_size=0.3,random_state=100)
```

# **Step-3:- Data Visualization**

## In [16]:

```
# Explore expenses relationships.

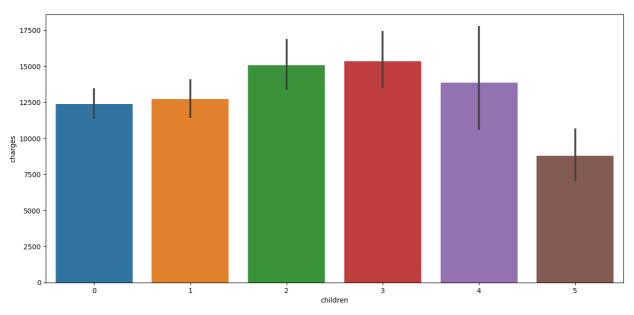
# A bar plot or bar chart is a graph that represents the category of data with rectangular bars with lengths and heights

plt.figure(figsize=(15,7))

sns.barplot(x=df.children, y=df.charges)
```

## Out[16]:

<Axes: xlabel='children', ylabel='charges'>

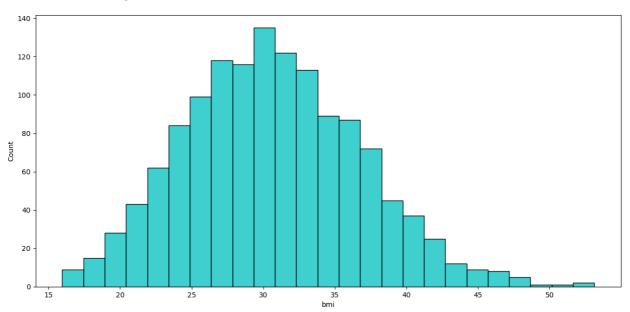


## In [17]:

```
# Visualize our BMI column.
# A histogram is a graph showing frequency distributions.
# It is a graph showing the number of observations within each given interval.
plt.figure(figsize=(15,7))
sns.histplot(data=df, x='bmi',color='c')
```

#### Out[17]:

<Axes: xlabel='bmi', ylabel='Count'>



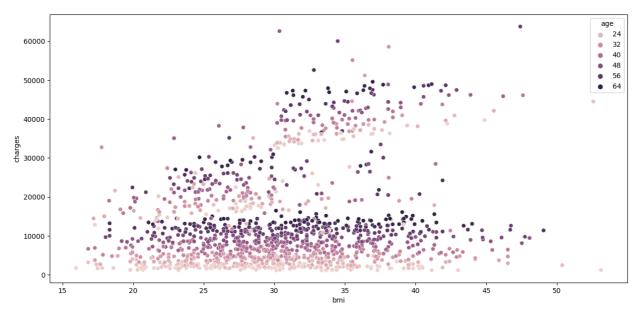
#### In [18]:

```
# With Pyplot, you can use the scatter() function to draw a scatter plot.

# The scatter() function plots one dot for each observation. It needs two arrays of the same length, one for the values plt.figure(figsize=(15,7))
sns.scatterplot(x=df.bmi, y=df.charges,hue=df.age)
```

## Out[18]:

<Axes: xlabel='bmi', ylabel='charges'>



## In [19]:

df.charges.corr(df.bmi)

# Out[19]:

0.19840083122624932

#### In [20]:

df.charges.corr(df.children)

#### Out[20]:

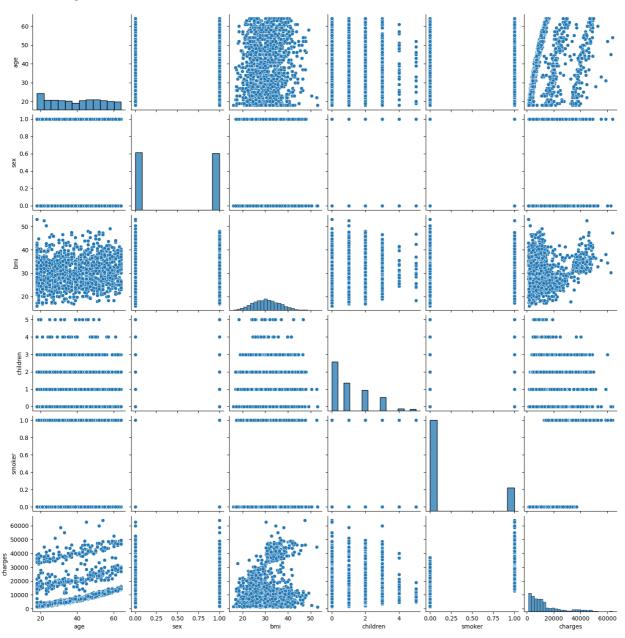
0.06738935083963239

#### In [21]:

sns.pairplot(df)

## Out[21]:

<seaborn.axisgrid.PairGrid at 0x269335f2ad0>



# Step-4:- Data Modelling

# **Applying LinearRegression**

```
In [22]:
```

```
# Fit Model to the training set
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_train, y_train)
```

# Out[22]:

```
LinearRegression
LinearRegression()
```

```
In [23]:
# Predict the test set result
y_pred = regressor.predict(X_test)
```

```
In [24]:
```

```
from sklearn.metrics import r2_score
score = r2_score(y_test, y_pred)
```

#### In [25]:

score

## Out[25]:

0.757820140931517

We have a R^2 score of 0.75 which tells us that our model is accurate.

# **Applying LogisticRegression**

#### In [26]:

```
#Logistic Regression
x=np.array(df['charges']).reshape(-1,1)
y=np.array(df['smoker']).reshape(-1,1)
df.dropna(inplace=True)
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.3,random_state=1)
from sklearn.linear_model import LogisticRegression
lr=LogisticRegression(max_iter=10000)
```

#### In [27]:

```
lr.fit(x_train,y_train)
```

C:\Users\thara\AppData\Local\Programs\Python\Python310\lib\site-packages\sklearn\utils\validation.py:1143: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n\_samples, ), for example using ravel().  $y = \text{column\_or\_1d}(y, \text{warn=True})$ 

#### Out[27]:

```
LogisticRegression
LogisticRegression(max_iter=10000)
```

## In [28]:

```
score=lr.score(x_test,y_test)
print(score)
```

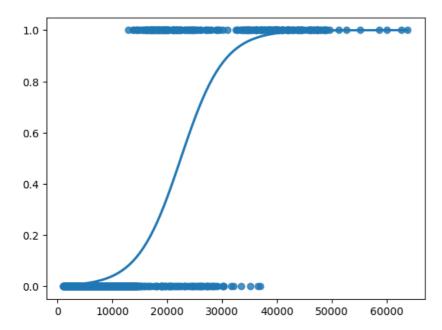
0.9253731343283582

#### In [29]:

sns.regplot(x=x,y=y,data=df,logistic=True,ci=None)

# Out[29]:

<Axes: >



We got the best fit score for Logistic Regression.

# **Applying Decision Tree**

we are going to check that if we may get better accuracy by implementing Decision Tree and RandomForest

### In [43]:

```
#Decision tree

from sklearn.tree import DecisionTreeClassifier
clf=DecisionTreeClassifier(random_state=0)
clf.fit(x_train,y_train)
```

# Out[43]:

```
DecisionTreeClassifier
DecisionTreeClassifier(random_state=0)
```

#### In [44]:

```
score=clf.score(x_test,y_test)
print(score)
```

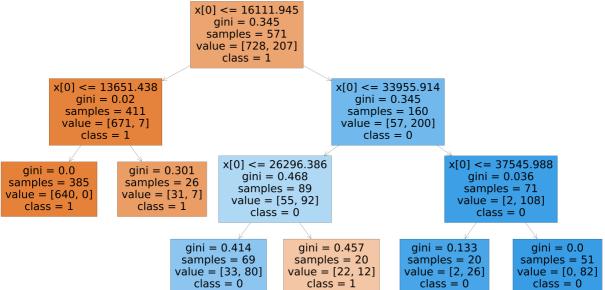
0.900497512437811

### **Random Forest**

```
In [45]:
#Random forest classifier
from sklearn.ensemble import RandomForestClassifier
rfc=RandomForestClassifier()
rfc.fit(X_train,y_train)
C:\Users\thara\AppData\Local\Temp\ipykernel_25876\1232785509.py:4: DataConversionWarning: A column-vector
y was passed when a 1d array was expected. Please change the shape of y to (n_samples,), for example using
ravel().
   rfc.fit(X_train,y_train)
Out[45]:
 ▼ RandomForestClassifier
RandomForestClassifier()
In [49]:
rf = RandomForestClassifier()
In [50]:
params={\'max depth':[2,3,5,10,20],\'min samples leaf':[5,10,20,50,100,200],\'n estimators':[10,25,30,50,100,200]}
In [51]:
from sklearn.model_selection import GridSearchCV
grid_search = GridSearchCV(estimator=rf,param_grid=params,cv = 2, scoring='accuracy')
grid_search.fit(x_train,y_train)
ation.py:686: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please
change the shape of y to (n_samples,), for example using ravel().
   estimator.fit(X_train, y_train, **fit_params)
ation.py:686: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please
change the shape of y to (n_samples,), for example using ravel().
    estimator.fit(X_train, y_train, **fit_params)
ation.py:686: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please
change the shape of y to (n_samples,), for example using ravel().
    estimator.fit(X_train, y_train, **fit_params)
 \verb| C:\Users \land AppData \land Programs \land Python \land Python 310 \land Site-packages \land Sklearn \land validation \land Validation
ation.py:686: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please
change the shape of y to (n_samples,), for example using ravel().
    estimator.fit(X_train, y_train, **fit_params)
ation.py:686: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please
change the shape of y to (n_samples,), for example using ravel().
In [52]:
grid_search.best_score_
Out[52]:
0.9219193250242501
In [53]:
rf_best=grid_search.best_estimator_
rf_best
Out[53]:
                                                  RandomForestClassifier
RandomForestClassifier(max_depth=3, min_samples_leaf=20, n_estimators=50)
```

#### In [61]:

```
from sklearn.tree import plot_tree
plt.figure(figsize=(80,40))
plot_tree(rf_best.estimators_[5],class_names=['1','0'],filled=True);
```



#### In [63]:

```
score=rfc.score(X_test,y_test)
print(score)
```

0.7786069651741293

## **Conclusion:**

Based on accuracy scores of all models that were implemented we can conclude that "Logistic Regression" is the best model for the given DataSet

# In [ ]: