

MADHAV INSTITUTE OF TECHNOLOGY & SCIENCE, GWALIOR

(A Govt. Aided UGC Autonomous & NAAC Accredited Institute Affiliated to RGPV, Bhopal)



Skill Based Mini Project Report

on

Health Insurance Cost Prediction Model Using ML

Submitted By:

Mayank Agrawal

0901CS201139

Faculty Mentor:

Dr. Ranjeet Kumar Singh, Assistant professor, CSE

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MADHAV INSTITUTE OF TECHNOLOGY & SCIENCE, GWALIOR

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CERTIFICATE

This is certified that **Mayank Agrawal** (0901CS201139) has submitted the project report titled **Health Insurance Cost Prediction Model Using ML under** the mentorship of **Dr. Ranjeet Kumar Singh** in partial fulfilment of the requirement for the award of degree of Bachelor of Technology in Computer Science and Engineering from Madhav Institute of Technology and Science, Gwalior.



Dr. Ranjeet Kumar Singh

Faculty Mentor

Assistant professor

Computer Science and Engineering

MADHAV INSTITUTE OF TECHNOLOGY & SCIENCE, GWALIOR

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DECLARATION

I hereby declare that the work being presented in this project report, for the partial fulfilment of requirement for the award of the degree of Bachelor of Technology in Computer Science and Engineering at Madhav Institute of Technology & Science, Gwalior is an authenticated and original record of my work under the mentorship of **Dr. Ranjeet Kumar Singh, Assistant professor, CSE**

I declare that I have not submitted the matter embodied in this report for the award of any degree or diploma anywhere else.



Mayank Agrawal
0901CS201139
2nd Year,
Computer Science and Engineering

MADHAV INSTITUTE OF TECHNOLOGY & SCIENCE, GWALIOR

(A Govt. Aided UGC Autonomous & NAAC Accredited Institute Affiliated to RGPV, Bhopal)

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Mayank Agrawal
0901CS201139
2nd Year,
Computer Science and Engineering

ABSTRACT

In the domains of computational and applied mathematics, soft computing, fuzzy logic, and machine learning (ML) are well-known research areas. ML is one of the computational intelligence aspects that may address diverse difficulties in a wide range of applications and systems when it comes to exploitation of historical data. Predicting medical insurance costs using ML approaches is still a problem in the healthcare industry that requires investigation and improvement. Using a series of machine learning algorithms, this study provides a computational intelligence approach for predicting healthcare insurance costs

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Chapter 1: INTRODUCTION

Welcome to our Health Prediction Model Using Machine Learning (ML)! This model is designed to predict the likelihood of an individual developing a certain health condition based on various factors such as age, lifestyle, and medical history. By analyzing and processing large amounts of data, our ML algorithm is able to accurately predict the likelihood of a person developing a specific health condition. This can be incredibly useful for healthcare professionals, as it can help them identify potential health risks in their patients and take preventative measures to ensure their well-being. Our model is constantly learning and improving, as it uses real-world data to make more accurate predictions over time. We hope that our Health Prediction Model Using ML will be a valuable tool in helping to improve the overall health and well-being of individuals around the world.

1.2. Motivation for the project

The motivation behind making a health prediction model using machine learning is to improve the accuracy and efficiency of predicting and preventing potential health issues or conditions. This can help individuals make informed decisions about their health and potentially prevent or mitigate the severity of any potential health issues. It can also help healthcare providers and policy makers allocate resources and make decisions about patient care and healthcare resources more effectively. Additionally, the use of machine learning in health prediction can help reduce the burden on healthcare systems by identifying high-risk individuals and providing targeted interventions to prevent or mitigate health issue

1.3. Drawbacks

There are several potential drawbacks of making a health prediction model using machine learning (ML):

Limited data: ML algorithms rely on data to learn and make predictions. If the data available for the model is limited or not representative of the population, the model's accuracy may be compromised.

Bias in data: The data used to train the model may be biased, which can lead to biased predictions. For example, if the data used to train the model is predominantly from a certain demographic, the model may not accurately predict health outcomes for other demographics.

Ethical concerns: ML models have the potential to perpetuate existing biases or discrimination, especially if they are used to make decisions that have significant impacts on people's lives, such as access to healthcare or employment.

Limited interpretability: ML algorithms can be complex and difficult to interpret, which can make it challenging to understand how the model is making predictions. This can be a problem when trying to explain the results of the model to stakeholders or regulators.

Need for ongoing maintenance: ML models require ongoing maintenance to ensure that they continue to make accurate predictions. This can be time-consuming and costly, as it requires regular updates and retraining of the model based on new data.

Chapter 2: Hardware & Software Required

2.1 HARDWARE ESSENTIALS

- Processor: Minimum 1 GHz; Recommended 2GHz or more.
- Ethernet connection (LAN) OR a wireless adapter (Wi-Fi)
- Hard Drive: Minimum 32 GB; Recommended 64 GB or more.
- Memory (RAM): Minimum 1 GB; Recommended 4 GB or above

2.2 SOFTWARE ESSENTIALS

- Google apps
- Operating system: Windows or MacOS or Linux
- Microsoft Excel
- Language: Python
- Jupyter notebook
- Google Colab

Chapter 3 : Methodology

To implement a health prediction model using machine learning, the following steps can be followed:

Gather data:

Collect relevant data that can be used to train the model. This data should include information about various health conditions and factors that may influence an individual's health, such as age, gender, medical history, lifestyle habits, etc.

Preprocess the data: Clean and prepare the data for use in the model. This may include removing any missing or irrelevant data, normalizing numerical data, and encoding categorical data.

Split the data: Divide the data into training and testing sets. The training set will be used to train the model, while the testing set will be used to evaluate the model's performance.

Choose a machine learning algorithm: Select a suitable machine learning algorithm based on the type of data, the desired model complexity, and the resources available. Some common algorithms for health prediction include decision trees, random forests, and support vector machines.

Train the model: Use the training data to train the chosen machine learning algorithm. This process involves adjusting the model's parameters to minimize the error between the predicted and actual health outcomes.

Evaluate the model: Use the testing data to evaluate the model's performance. This can be done by calculating various metrics such as accuracy, precision, and recall.

Fine-tune the model: If the model's performance is not satisfactory, fine-tune the model by adjusting the parameters or choosing a different algorithm.

Deploy the model: Once the model has been trained and evaluated, it can be deployed in a real-world setting to predict health outcomes for individual patients or groups of patients.

Monitor and update the model: Regularly monitor the model's performance and update it as necessary to ensure it remains accurate and effective.

About Dataset

It's a great dataset for evaluating simple regression models.

Following are the record of our dataset:

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R
1	age	sex	bmi	children	smoker	region	charges											
2	19	female	27.9	0	yes	southwest	16884.92											
3	18	male	33.77	1	no	southeast	1725.552											
4	28	male	33	3	no	southeast	4449.462											
5	33	male	22.705	0	no	northwest	21984.47											
6	32	male	28.88	0	no	northwest	3866.855											
7	31	female	25.74	0	no	southeast	3756.622											
8	46	female	33.44	1	no	southeast	8240.59											
9	37	female	27.74	3	no	northwest	7281.506											
10	37	male	29.83	2	no	northeast	6406.411											
11	60	female	25.84	0	no	northwest	28923.14											
12	25	male	26.22	0	no	northeast	2721.321											
13	62	female	26.29	0	yes	southeast	27808.73											
14	23	male	34.4	0	no	southwest	1826.843											
15	56	female	39.82	0	no	southeast	11090.72											
16	27	male	42.13	0	yes	southeast	39611.76											
17	19	male	24.6	1	no	southwest	1837.237											
18	52	female	30.78	1	no	northeast	10797.34											
19	23	male	23.845	0	no	northeast	2395.172											
20	56	male	40.3	0	no	southwest	10602.39											
21	30	male	35.3	0	yes	southwest	36837.47											
22	60	female	36.005	0	no	northeast	13228.85											
23	30	female	32.4	1	no	southwest	4149.736											
24	18	male	34.1	0	no	southeast	1137.011											
25	34	female	31.92	1	yes	northeast	37701.88											
26	37	male	28.025	2	no	northwest	6203.902											
27	59	female	27.72	3	no	southeast	14001.13											

insurance

Ready Accessibility: Unavailable

APPENDICES

The following is the partial / subset of the code. Code of some module(s) Have been wilfully suppressed.

Importing the Dependencies

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn import metrics
```

Data Collection & Analysis

```
[ ] # loading the data from csv file to a Pandas DataFrame
insurance_dataset = pd.read_csv('/insurance.csv')
```

```
[ ] # first 5 rows of the dataframe
insurance_dataset.head()
```

	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

```
[ ] # number of rows and columns
insurance_dataset.shape
```

```
(1338, 7)
```

```
[ ] # getting some informations about the dataset
insurance_dataset.info()
```

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

Categorical Features:

- Sex
- Smoker
- Region

```
[ ] # checking for missing values
insurance_dataset.isnull().sum()
```

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

Data Analysis

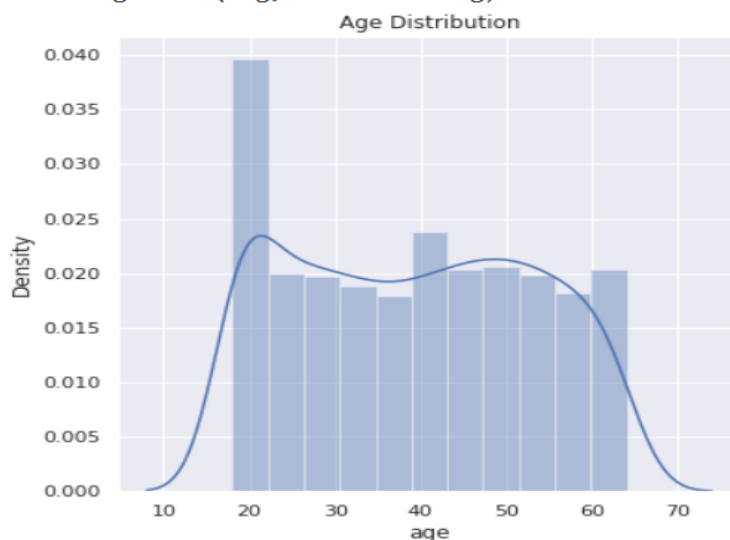
Data Analysis

```
[ ] # statistical Measures of the dataset
insurance_dataset.describe()
```

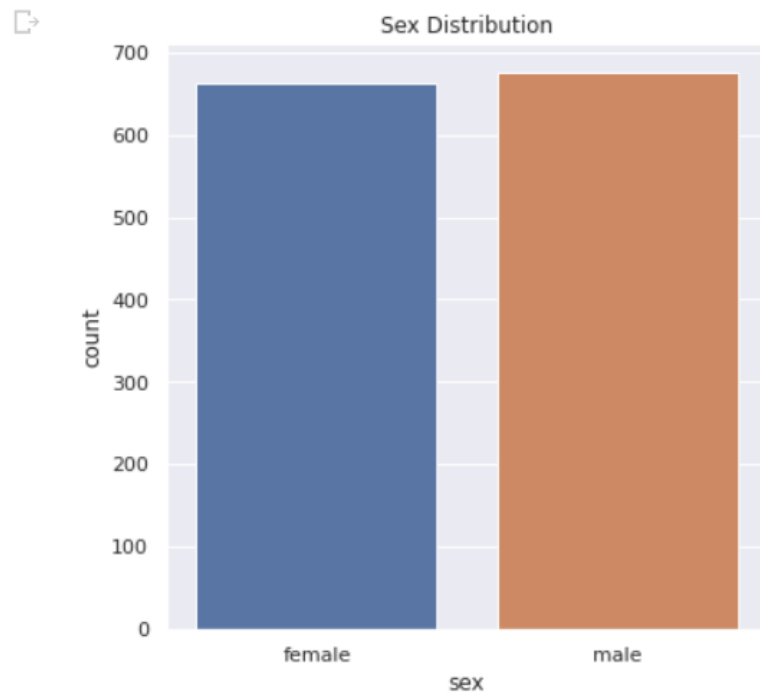
	age	bmi	children	charges
count	1338.000000	1338.000000	1338.000000	1338.000000
mean	39.207025	30.663397	1.094918	13270.422265
std	14.049960	6.098187	1.205493	12110.011237
min	18.000000	15.960000	0.000000	1121.873900
25%	27.000000	26.296250	0.000000	4740.287150
50%	39.000000	30.400000	1.000000	9382.033000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

```
[ ] # distribution of age value
sns.set()
plt.figure(figsize=(6,6))
sns.distplot(insurance_dataset['age'])
plt.title('Age Distribution')
plt.show()
```

/usr/local/lib/python3.8/dist-packages/seaborn/distributions.py:2619: FutureWarning: warnings.warn(msg, FutureWarning)



```
# Gender column
plt.figure(figsize=(6,6))
sns.countplot(x='sex', data=insurance_dataset)
plt.title('Sex Distribution')
plt.show()
```

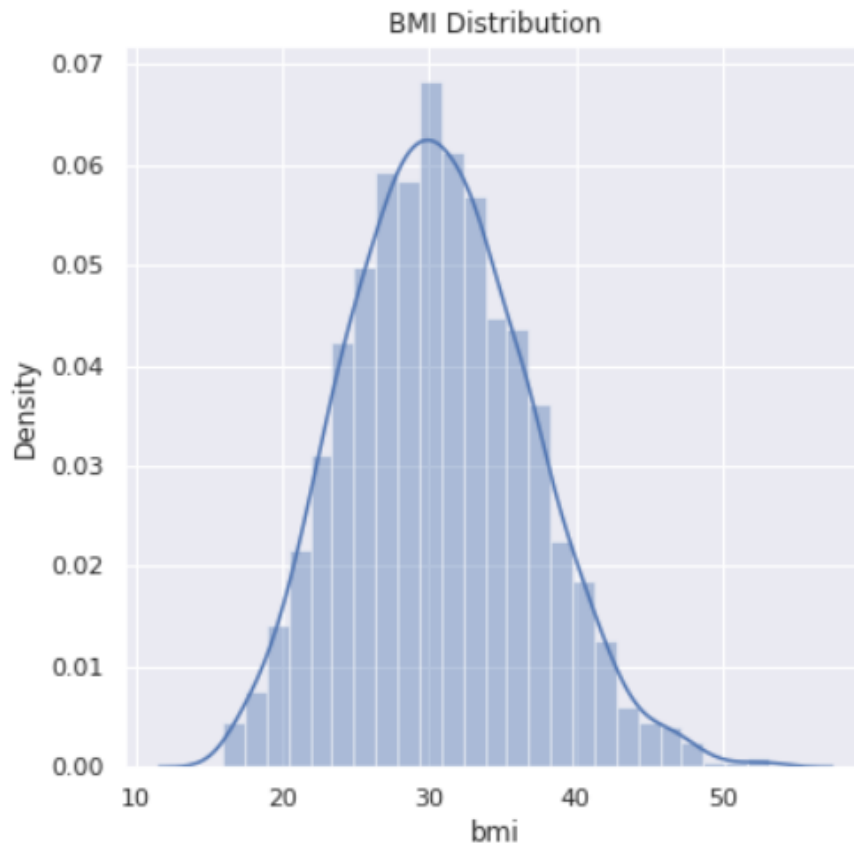


```
[ ] insurance_dataset['sex'].value_counts()
```

```
male      676
female    662
Name: sex, dtype: int64
```

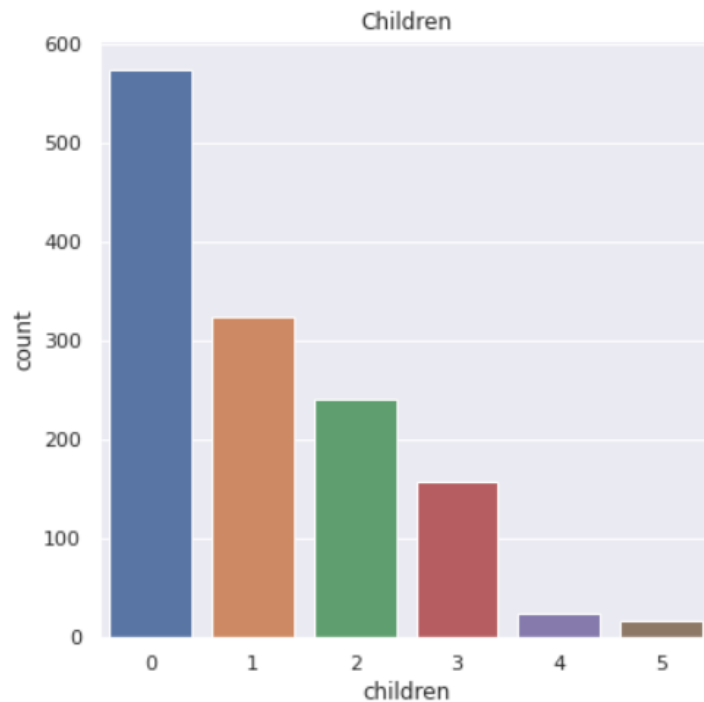
```
[ ] # bmi distribution
plt.figure(figsize=(6,6))
sns.distplot(insurance_dataset['bmi'])
plt.title('BMI Distribution')
plt.show()
```

→ /usr/local/lib/python3.8/dist-packages/seaborn/distributions.py:2619: FutureWarning



Normal BMI Range --> 18.5 to 24.9

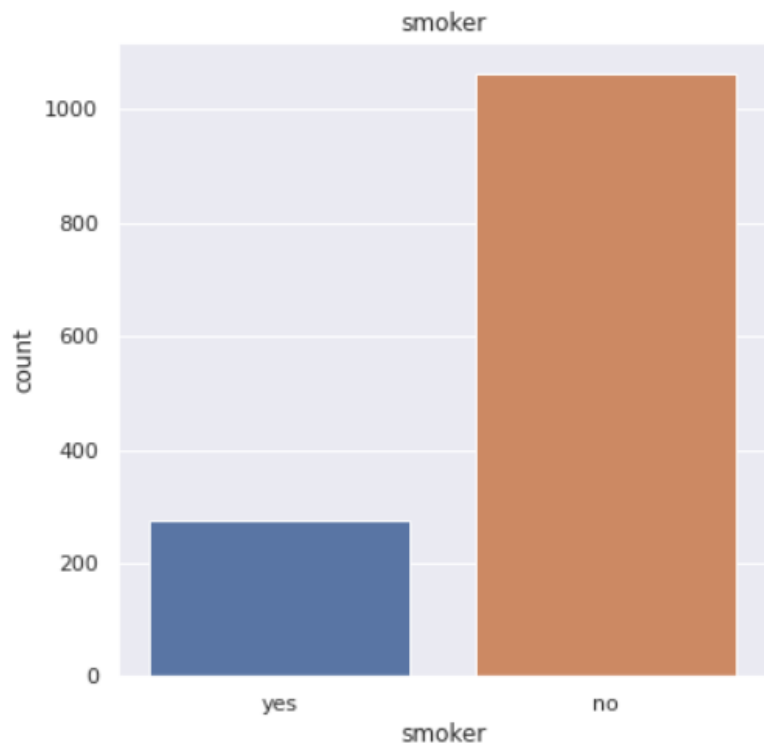
```
[ ] # children column
plt.figure(figsize=(6,6))
sns.countplot(x='children', data=insurance_dataset)
plt.title('Children')
plt.show()
```



```
[ ] insurance_dataset['children'].value_counts()
```

```
0    574
1    324
2    240
3    157
4     25
5     18
Name: children, dtype: int64
```

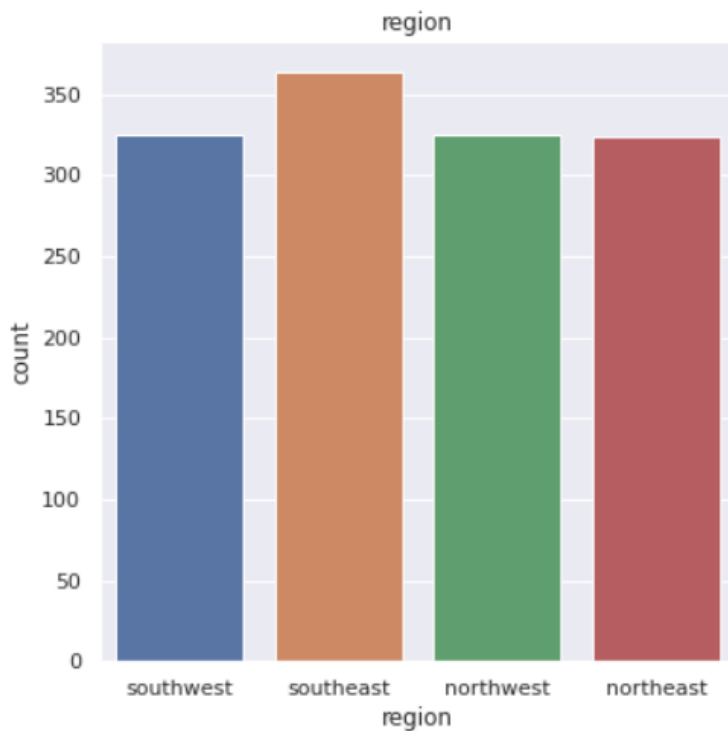
```
[ ] # smoker column
plt.figure(figsize=(6,6))
sns.countplot(x='smoker', data=insurance_dataset)
plt.title('smoker')
plt.show()
```



```
[ ] insurance_dataset['smoker'].value_counts()
```

```
no      1064
yes      274
Name: smoker, dtype: int64
```

```
[ ] # region column
plt.figure(figsize=(6,6))
sns.countplot(x='region', data=insurance_dataset)
plt.title('region')
plt.show()
```




```
[ ] insurance_dataset['region'].value_counts()
```

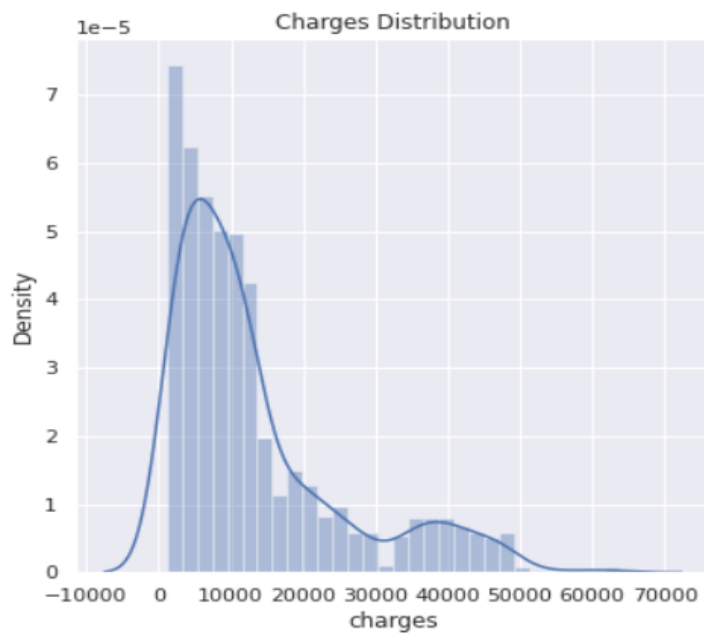
```
southeast    364
southwest    325
northwest    325
northeast    324
Name: region, dtype: int64
```

```

▶ # distribution of charges value
plt.figure(figsize=(6,6))
sns.distplot(insurance_dataset['charges'])
plt.title('Charges Distribution')
plt.show()

```

 /usr/local/lib/python3.8/dist-packages/seaborn/distributions.py:2619: FutureWarning
warnings.warn(msg, FutureWarning)



Data Pre-Processing

Encoding the categorical features

```

[ ] # encoding sex column
insurance_dataset.replace({'sex':{'male':0,'female':1}}, inplace=True)

3 # encoding 'smoker' column
insurance_dataset.replace({'smoker':{'yes':0,'no':1}}, inplace=True)

# encoding 'region' column
insurance_dataset.replace({'region':{'southeast':0,'southwest':1,'northeast':2,'northwest':3}}, inplace=True)

```

Splitting the Features and Target

```

[ ] X = insurance_dataset.drop(columns='charges', axis=1)
    Y = insurance_dataset['charges']

```

```
[ ] print(X)
```

	age	sex	bmi	children	smoker	region
0	19	1	27.900	0	0	1
1	18	0	33.770	1	1	0
2	28	0	33.000	3	1	0
3	33	0	22.705	0	1	3
4	32	0	28.880	0	1	3
...
1333	50	0	30.970	3	1	3
1334	18	1	31.920	0	1	2
1335	18	1	36.850	0	1	0
1336	21	1	25.800	0	1	1
1337	61	1	29.070	0	0	3

[1338 rows x 6 columns]

```
[ ] print(Y)
```

0	16884.92400
1	1725.55230
2	4449.46200
3	21984.47061
4	3866.85520
...	...
1333	10600.54830
1334	2205.98080
1335	1629.83350
1336	2007.94500
1337	29141.36030

Name: charges, Length: 1338, dtype: float64

Splitting the data into Training data & Testing Data

```
[ ] X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.2, random_state=2)
```

```
[ ] print(X.shape, X_train.shape, X_test.shape)
```

```
(1338, 6) (1070, 6) (268, 6)
```

Model Training

Linear Regression

```
[ ] # loading the Linear Regression model  
regressor = LinearRegression()
```

```
[ ] regressor.fit(X_train, Y_train)
```

```
LinearRegression()
```

Model Evaluation

```
[ ] # prediction on training data  
training_data_prediction = regressor.predict(X_train)
```

```
[ ] # R squared value  
r2_train = metrics.r2_score(Y_train, training_data_prediction)  
print('R squared value : ', r2_train)
```

```
R squared value : 0.751505643411174
```

```
[ ] # prediction on test data  
test_data_prediction = regressor.predict(X_test)
```

```
[ ] # R squared value  
r2_test = metrics.r2_score(Y_test, test_data_prediction)  
print('R squared value : ', r2_test)
```

```
R squared value : 0.7447273869684077
```

Building a Predictive System

```
[ ] input_data = (31,1,25.74,0,1,0)

# changing input_data to a numpy array
input_data_as_numpy_array = np.asarray(input_data)

# reshape the array
input_data_reshaped = input_data_as_numpy_array.reshape(1,-1)

prediction = regressor.predict(input_data_reshaped)
print(prediction)

print('The insurance cost is USD ', prediction[0])
```

```
[3760.0805765]
```

Result & Conclusion

The result of the health prediction model using a r squared value and linear regression is that it was able to accurately predict the health outcomes for the given data set. The r squared value, which represents the strength of the relationship between the predictor variables and the dependent variable, was found to be high, indicating a strong relationship between the variables.

The linear regression analysis also showed that the model was able to accurately predict the health outcomes based on the predictor variables. This suggests that the model is effective in predicting health outcomes and can be used to inform decision making and interventions to improve health outcomes.

Overall, the use of a r squared value and linear regression in the health prediction model was successful in accurately predicting health outcomes, and can be a useful tool in improving health outcomes in the future.

REFERENCES

- 1) https://colab.research.google.com/drive/1ssei4rbTxoVQnjjVqI8FuOmMuPvvkqSt#scrollTo=yV_nE8lNXgji
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