## Healthcare Insurance Analysis

### Problem Statement

A significant public health concern is the rising cost of healthcare. Therefore, it's crucial to be able to predict future costs and gain a solid understanding of their causes. The insurance industry must also take this analysis seriously. This analysis may be used by healthcare insurance providers to make a variety of strategic and tactical decisions.

## Objective

The objective of this project is to predict patients' healthcare costs and to identify factors contributing to this prediction. It will also be useful to learn the interdependencies of different factors and comprehend the significance of various tools at various stages of the healthcare cost prediction process.

## **Install Important Libraries**

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings("ignore")
%matplotlib inline
```

### >> DATA SCIENCE

### 1. Collate the files so that all the information is in one place

```
df1=pd.read csv('Hospitalisation details.csv')
df2=pd.read csv('Medical Examinations.csv')
df3=pd.read csv('Names.csv')
df=pd.merge(df1, df2,on='Customer ID')
df=pd.merge(df, df3, on='Customer ID')
df.head()
  Customer ID year month date
                                 children
                                            charges Hospital tier City
tier
       Id2335
              1992
                      Jul
                              9
                                             563.84
                                                         tier - 2 tier
- 3
1
       Id2334 1992
                      Nov
                             30
                                             570,62
                                                         tier - 2 tier
- 1
2
       Id2333 1993
                      Jun
                             30
                                         0
                                             600.00
                                                         tier - 2 tier
- 1
3
       Id2332 1992
                      Sep
                             13
                                             604.54
                                                         tier - 3 tier
```

- 3										
4 - 3	Id23	31 199	8 Jul	27		0	637.26	tier	- 3	tier
3										
	tate ID tory \	BMI	HBA1C He	eart Is	ssues	Any	Transplants	Cancer		
0	R1013	17.58	4.51		No		No			No
1	R1013	17.60	4.39		No		No			No
2	R1013	16.47	6.35		No		No		,	Yes
3	R1013	17.70	6.28		No		No			No
4	R1013	22.34	5.57		No		No			No
	umber0fM	ajorSur	geries sn				Camaa	M.a. A	na	
0 1			1	No No			Rosendah	, Mr.  A l, Mr.		
2			1	No			Albai	no, Ms.	Jul	ie
3 4			1 1	No No	River	os (	Gonzalez, Mr Brietzke		D. S Jord	

### 2. Check for missing values in the dataset

```
df.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 2335 entries, 0 to 2334
Data columns (total 17 columns):
#
     Column
                              Non-Null Count
                                               Dtype
0
     Customer ID
                              2335 non-null
                                               object
1
                              2335 non-null
                                               object
     year
 2
     month
                              2335 non-null
                                               object
 3
     date
                              2335 non-null
                                               int64
 4
                              2335 non-null
     children
                                               int64
 5
                              2335 non-null
                                               float64
     charges
 6
     Hospital tier
                              2335 non-null
                                               object
 7
     City tier
                              2335 non-null
                                               object
 8
                              2335 non-null
     State ID
                                               object
 9
     BMI
                              2335 non-null
                                               float64
10
     HBA1C
                                               float64
                              2335 non-null
11
     Heart Issues
                              2335 non-null
                                               object
 12
     Any Transplants
                              2335 non-null
                                               object
 13
     Cancer history
                              2335 non-null
                                               object
     NumberOfMajorSurgeries
 14
                              2335 non-null
                                               object
 15
                              2335 non-null
     smoker
                                               object
 16
     name
                              2335 non-null
                                               object
```

```
dtypes: float64(3), int64(2), object(12)
memory usage: 310.2+ KB
```

Upon initial inspection, it looks like there are no null values. But the dataset contains '?' at some places. For Example -

```
df.loc[df['State ID']=='?']
     Customer ID year month date children
                                               charges Hospital
tier \
542
          Id1793 1995
                         Dec
                                               4827.90
                                                            tier - 1
2165
           Id170 2000
                         Sep
                                           1 37165.16
                                                            tier - 1
     City tier State ID
                            BMI
                                 HBA1C Heart Issues Any Transplants \
    tier - 2
542
                      ?
                         18.905
                                  4.91
                                                                 No
                                                ves
2165 tier - 3
                      ?
                        37.620
                                  6.32
                                                yes
                                                                yes
     Cancer history NumberOfMajorSurgeries smoker
name
542
                                                   Capriolo, Mr.
                 No
                                         1
                                               No
Michael
2165
                 No
                                              yes
                                                       Torphy, Mr.
Bobby
```

3. Find the percentage of rows that have trivial value (for example, ?), and delete such rows if they do not contain significant information.

```
trivial_counts=df.eq('?').sum().sum()
trivial_percentage=(trivial_counts/len(df))*100
print('Trivial Counts:',trivial_counts)
print('Trivial Percentgae: %.2f' % trivial_percentage + '%')
Trivial Counts: 11
Trivial Percentgae: 0.47%
```

### Detailed columnwise Trivial counts

```
trivial_counts=df.eq('?').sum()
trivial_percentage=(trivial_counts/len(df))*100 ##
Here,len(df)==2335

# Create a table with trivial counts and percentages
table = [['Features', 'Trivial Counts', 'Trivial Percentage (%)']]
for feat, count in trivial_counts.items():
    table.append([feat, count, trivial_percentage[feat]])

# Print the table using tabulate
```

<pre>from tabulate import tabu print(tabulate(table, hea +</pre>	late ders='first		efmt='psql'))
+   Features 	+   Trivial	Counts	Trivial Percentage (%)
 +		0	0
   year		2	0.0856531
   month		3	0.12848
   date		0	0
   children		0	0
   charges		0	0
   Hospital tier		1	0.0428266
   City tier		1	0.0428266
   State ID		2	0.0856531
   BMI		0	0
   HBA1C		0	0
   Heart Issues		0	0
   Any Transplants		0	0
   Cancer history		0	0
NumberOfMajorSurgeries	I	0	0
smoker	I	2	0.0856531
name	I	0	Θ
 +	+		
+	+		

We can see that those trivial values contains around 0.1% data in each feature. So, we can eliminate them.

```
# Replace '?' with NaN values
df.replace('?',np.nan, inplace=True)
#Check Null Values
df.dropna(inplace=True)
df.isnull().sum()
Customer ID
                           0
                           0
vear
month
                           0
date
                           0
children
                           0
                           0
charges
Hospital tier
                           0
                           0
City tier
State ID
                           0
BMI
                           0
HBA1C
                           0
Heart Issues
                           0
Any Transplants
                           0
Cancer history
                           0
NumberOfMajorSurgeries
                           0
smoker
                           0
                           0
name
dtype: int64
#check Duplicate Values
df['Customer ID'].duplicated().sum()
np.int64(0)
df.info()
<class 'pandas.core.frame.DataFrame'>
Index: 2325 entries, 0 to 2334
Data columns (total 17 columns):
 #
     Column
                              Non-Null Count
                                               Dtype
     -----
 0
     Customer ID
                              2325 non-null
                                               object
 1
     year
                              2325 non-null
                                               obiect
 2
     month
                              2325 non-null
                                               object
 3
     date
                              2325 non-null
                                               int64
 4
     children
                              2325 non-null
                                               int64
 5
     charges
                              2325 non-null
                                               float64
                              2325 non-null
 6
     Hospital tier
                                               object
 7
                              2325 non-null
     City tier
                                               object
 8
     State ID
                              2325 non-null
                                               object
 9
     BMI
                              2325 non-null
                                               float64
```

```
10 HBA1C
                           2325 non-null
                                           float64
11 Heart Issues
                           2325 non-null
                                           object
12 Any Transplants
                           2325 non-null
                                           object
13 Cancer history
                           2325 non-null
                                           object
    NumberOfMajorSurgeries 2325 non-null
                                           object
15
   smoker
                           2325 non-null
                                           object
16 name
                           2325 non-null
                                           object
dtypes: float64(3), int64(2), object(12)
memory usage: 327.0+ KB
```

So, total 10 data have been dropped, which is 0.43% of the total data.

4. Use the necessary transformation methods to deal with the nominal and ordinal categorical variables in the dataset.

df.hea	ad()	<b>-</b>							
Cust tier	tomer ID	year ı	month	date	children	charges	Hospital <sup>-</sup>	tier	City
0 - 3	`Id2335	1992	Jul	9	0	563.84	tier	- 2	tier
1 - 1	Id2334	1992	Nov	30	0	570.62	tier	- 2	tier
2 - 1	Id2333	1993	Jun	30	0	600.00	tier	- 2	tier
3 - 3	Id2332	1992	Sep	13	0	604.54	tier	- 3	tier
4 - 3	Id2331	1998	Jul	27	0	637.26	tier	- 3	tier
histor 0 F	ry \ R1013 17 R1013 17	.58	BA1C F 4.51 4.39 6.35	leart I	No No No No	Transplar	nts Cancer No No No		No No Yes
			6.28		No		No		No
			5.57		No		No		No
Numb 0 1 2 3	oerOfMajo	rSurge	ries s 1 1 1 1	moker No No No No No	Riveros	Roseno Al Gonzalez,	man, Mr. / dahl, Mr. lbano, Ms. Mr. Juan tzke, Mr.	Aaron Evan Jul	n P .ie Sr.

We observe that 'Hospital tier' and 'City tier' columns have alphanumeric values. We aim to convert them into numerical values, where 'tier-1' becomes '1', 'tier-2' becomes '2', etc., to enable more efficient data handling.

```
df['Hospital tier num'] = df['Hospital tier'].str.extract('tier - (\)
d+)').astype(int)
df['City tier num'] = df['City tier'].str.extract('tier - (\)
d+)').astype(int)
df.head()
  Customer ID year month date
                                  children
                                             charges Hospital tier City
tier \
       Id2335 1992
                      Jul
                                              563.84
0
                               9
                                          0
                                                          tier - 2 tier
- 3
1
                                          0
                                              570.62
                                                          tier - 2 tier
       Id2334 1992
                       Nov
                              30
- 1
2
       Id2333 1993
                       Jun
                              30
                                          0
                                              600.00
                                                          tier - 2 tier
- 1
3
       Id2332 1992
                              13
                                          0
                                              604.54
                                                          tier - 3 tier
                       Sep
- 3
       Id2331 1998
                       Jul
                              27
                                              637.26
                                                          tier - 3 tier
- 3
  State ID
              BMI
                   HBA1C Heart Issues Any Transplants Cancer
history \
     R1013
            17.58
                     4.51
                                    No
                                                     No
                                                                     No
     R1013
           17.60
                     4.39
                                    No
                                                     No
                                                                     No
1
2
     R1013 16.47
                    6.35
                                    No
                                                     No
                                                                    Yes
3
     R1013 17.70
                     6.28
                                    No
                                                     No
                                                                     No
     R1013 22.34
                     5.57
                                    No
                                                     No
                                                                     No
  NumberOfMajorSurgeries smoker
                                                                  name \
0
                              No
                                                 German, Mr.
                                                               Aaron K
                        1
                                               Rosendahl, Mr.
1
                        1
                              No
                                                                Evan P
2
                        1
                              No
                                                   Albano, Ms.
                                                                 Julie
3
                        1
                              No
                                  Riveros Gonzalez, Mr. Juan D. Sr.
4
                        1
                                                Brietzke, Mr.
                              No
   Hospital tier num
                      City tier num
0
                    2
                                   3
                    2
1
                                   1
                    2
2
                                   1
3
                    3
                                   3
4
                    3
                                   3
```

5. The dataset has State ID, which has around 16 states. All states are not represented in equal proportions in the data. Creating dummy variables for all regions may also result in too many insignificant predictors. Nevertheless, only R1011, R1012, and R1013 are worth investigating further. Design a suitable strategy to create dummy variables with these restraints.

```
df['State ID'].value counts()
State ID
         609
R1013
R1011
         574
R1012
         572
         159
R1024
R1026
          84
          70
R1021
R1016
          64
          40
R1025
          38
R1023
R1017
          36
R1019
          26
R1022
          14
R1014
          13
R1015
          11
R1018
           9
           6
R1020
Name: count, dtype: int64
# Create a new column called 'stateflag' that categorizes state IDs.
df['State Flag'] = np.where((df['State ID'] == 'R1011') | (df['State
ID'] == 'R1012') | (df['State ID'] == 'R1013'), df['State ID'],
'other')
df['State_Flag'].replace('R1011',1,inplace=True)
df['State Flag'].replace('R1012',2,inplace=True)
df['State_Flag'].replace('R1013',3,inplace=True)
df['State Flag'].replace('other',0,inplace=True)
df.head()
  Customer ID year month date children charges Hospital tier City
tier
       Id2335
              1992
                      Jul
                                             563.84
                                                          tier - 2 tier
0
- 3
       Id2334 1992
                      Nov
                              30
                                         0
                                             570.62
                                                          tier - 2 tier
1
- 1
2
       Id2333 1993
                      Jun
                              30
                                             600.00
                                                          tier - 2 tier
- 1
3
       Id2332 1992
                      Sep
                              13
                                         0
                                             604.54
                                                          tier - 3 tier
- 3
```

4 - 3		31 1998	3 Jul	27	0	637.26	tier -	3 tier
_	State ID Story \	BMI	HBA1C He	art Issue	s Any	Transplants	Cancer	
0	R1013	17.58	4.51	N	lo	No		No
1	R1013	17.60	4.39	N	lo	No		No
2	R1013	16.47	6.35	N	lo	No		Yes
3	R1013	17.70	6.28	N	lo	No		No
4	R1013	22.34	5.57	N	lo	No		No
0 1 2 3 4	lumber0fM	ajorSurq	geries sm 1 1 1 1 1	No No No	eros (	Rosendah <sup>-</sup>	no, Ms. . Juan D	Evan P Julie
	Hospital	_tier_n	um City_	tier_num	State	e_Flag		
0 1 2 3			2 2 2 3 3	- 3 1 1 3 3		3 3 3 3		
4			3	3		3		

6. The variable NumberOfMajorSurgeries also appears to have string values. Apply a suitable method to clean up this variable.

```
# Replace 'No major surgery' with '0'.
df['NumberOfMajorSurgeries'].replace('No major
surgery','0',inplace=True)
df['NumberOfMajorSurgeries'] =
df['NumberOfMajorSurgeries'].astype(int)
df.head()
                                            charges Hospital tier City
  Customer ID year month date children
tier
       Id2335
              1992
                      Jul
                                             563.84
                                                         tier - 2 tier
0
- 3
       Id2334
              1992
                      Nov
                             30
                                         0
                                             570.62
                                                         tier - 2 tier
1
- 1
2
       Id2333
              1993
                      Jun
                             30
                                         0
                                             600.00
                                                         tier - 2 tier
- 1
3
       Id2332 1992
                                                         tier - 3 tier
                      Sep
                             13
                                         0
                                             604.54
- 3
```

4 - :		31 1998	3 Jul	27	0	637.26	tier -	3 tier
	State ID story \	BMI	HBA1C Hea	art Issı	ues Any	Transplants	Cancer	
0	R1013	17.58	4.51		No	No		No
1	R1013	17.60	4.39		No	No		No
2	R1013	16.47	6.35		No	No		Yes
3	R1013	17.70	6.28		No	No		No
4	R1013	22.34	5.57		No	No		No
	N b 0. £	M C						
naı		MajorSu	rgeries s					
0			1	No		German	, Mr. A	aron K
1			1	No		Rosendah	l, Mr.	Evan P
2			1	No		Alba	no, Ms.	Julie
3			1	No F	Riveros	Gonzalez, Mr	. Juan	D. Sr.
4			1	No		Brietzk	e, Mr.	Jordan
0	Hospital	_tier_n	um City_ <sup>.</sup> 2		n State 3	_		
1			2		l	3 3 3 3		
2			2 3	3	l 3	3		
4			3	3	3	3		

7. Age appears to be a significant factor in this analysis. Calculate the patients' ages based on their dates of birth.

```
#Covert 'year' column into int datatype.
df['year']=df['year'].astype('int')

# create a dictionary to map month names to integers
month_no = {'Jun': 6, 'Jul': 7, 'Aug': 8, 'Sep': 9, 'Oct': 10, 'Nov':
11, 'Dec': 12}

# convert month column to integer using the dictionary
df['month'] = df['month'].map(month_no)

# Calculate Date of Birth
```

```
import datetime
df['D.O.B']=pd.to datetime({'year': df['year'], 'month': df['month'],
'day': df['date']})
#Age Calculation
def age calculation(DOB):
    today = datetime.datetime.today()
    age = today.year - DOB.year - ((today.month, today.day) <</pre>
(DOB.month, DOB.day))
    return (age)
df['Age'] = df['D.O.B'].apply(age calculation)
df.head()
                     month date children charges Hospital tier City
  Customer ID year
tier \
                         7
                               9
                                         0
                                             563.84
       Id2335 1992
                                                          tier - 2
tier - 3
       Id2334 1992
                        11
                              30
                                         0
                                             570.62
                                                          tier - 2
1
tier - 1
       Id2333 1993
                         6
                              30
                                         0
                                             600.00
                                                          tier - 2
tier - 1
       Id2332 1992
                         9
                                         0
                                                          tier - 3
                              13
                                             604.54
tier - 3
       Id2331 1998
                         7
                              27
                                             637.26
                                                          tier - 3
tier - 3
  State ID
                        Any Transplants Cancer history
              BMI
NumberOfMajorSurgeries
0
     R1013 17.58
                                     No
                                                     No
1
1
     R1013 17.60
                                     No
                                                     No
1
2
     R1013 16.47
                                     No
                                                    Yes
1
3
     R1013 17.70
                                     No
                                                     No
1
4
     R1013 22.34
                                     No
                                                     No
1
  smoker
                                        name Hospital tier num
City_tier_num \
                        German, Mr. Aaron K
                                                              2
      No
3
1
                      Rosendahl, Mr. Evan P
      No
                                                              2
1
2
      No
                          Albano, Ms. Julie
                                                              2
1
3
          Riveros Gonzalez, Mr. Juan D. Sr.
                                                              3
      No
```

```
3
4
      No
                         Brietzke, Mr. Jordan
                                                                   3
3
   State_Flag
                            Age
                     D.O.B
0
             3 1992-07-09
                             32
1
             3 1992-11-30
                             31
2
             3 1993-06-30
                             31
3
             3 1992-09-13
                             31
4
             3 1998-07-27
                             26
[5 rows x 22 columns]
```

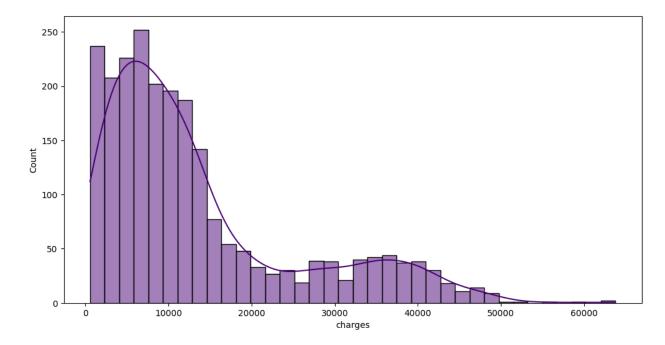
8. The gender of the patient may be an important factor in determining the cost of hospitalization. The salutations in a beneficiary's name can be used to determine their gender. Make a new field for the beneficiary's gender.

```
#Create a new column called Gender where define Male as 1 and Female
as 0.
gender = [1 if 'Mr.' in name else 0 for name in df['name']]
df['Gender']=gender
df.head()
 Customer ID year
                     month date children charges Hospital tier City
tier
       Id2335 1992
                                9
                                              563.84
                                                          tier - 2
0
tier - 3
       Id2334 1992
                        11
                              30
                                          0
                                              570.62
                                                          tier - 2
tier - 1
       Id2333 1993
                         6
                              30
                                              600.00
                                                          tier - 2
tier - 1
       Id2332 1992
                         9
                                          0
                                              604.54
                                                          tier - 3
                              13
tier - 3
       Id2331
              1998
                              27
                                          0
                                              637,26
                                                          tier - 3
tier - 3
  State ID
              BMI
                        Cancer history NumberOfMajorSurgeries
smoker \
     R1013
           17.58
                                     No
                                                             1
                                                                    No
     R1013
           17.60
                                     No
                                                                    No
     R1013
           16.47
                                    Yes
                                                                    No
     R1013
           17.70
                                     No
                                                                    No
     R1013 22.34
                                     No
                                                                    No
```

,		name	<pre>Hospital_tier_num</pre>	<pre>City_tier_num</pre>
0 (	German, Mr.	Naron K	2	3
· ·	Jerman, m	Adi Oli IX	2	J
1 Ros	sendahl, Mr.	Evan P	2	1
2	Albano, Ms.	Julie	2	1
3 Riveros Gonzale	ez, Mr. Juan	D. Sr.	3	3
4 B	rietzke, Mr.	Jordan	3	3
1 3 1992 2 3 1993 3 3 1992	D.O.B Age -07-09 32 -11-30 31 -06-30 31 -09-13 31 -07-27 26	Gender 1 1 0 1		
[5 rows x 23 colur	mns]			

9. You should also visualize the distribution of costs using a histogram, box and whisker plot, and swarm plot.

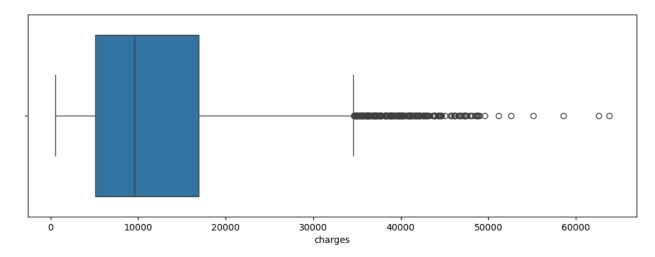
```
plt.figure(figsize=(12, 6))
sns.histplot(df['charges'], legend=True, kde=True, color="#490275")
plt.show()
```



#### Observation:

- The distribution is right-skewed (positively skewed), with a longer tail on the right side. This indicates that while most 'charges' values are lower, there are some significantly higher values.
- The majority of 'charges' values fall below 20,000. Beyond this point, the frequency of 'charges' decreases, but there are still some occurrences up to 60,000.
- The highest peak in the histogram occurs between 5,000 and 10,000.

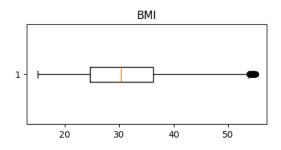
```
plt.figure(figsize=(12, 4))
sns.boxplot(x=df['charges'])
plt.show()
```

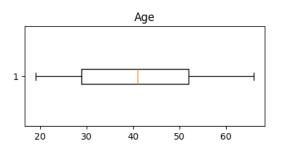


#### Observation:

- The median is around 9,000, which indicates that half of the 'charges' values are below this point.
- 50% of the central data is contained in the box, ranging approximately from 5,000 to 17,000.
- The lower whisker ends near 0, while the upper whisker ends around 35,000.
- The outliers extend up to approximately 63,000.

```
#Check Outliers in other features such as Age and BMI.
cols=['BMI','Age']
fig,ax=plt.subplots(1,2,figsize=(12,2))
for i,col in enumerate(cols):
    ax[i].boxplot(x=df[col].values,vert=False)
    ax[i].set_title(col)
plt.subplots_adjust(wspace=0.5)
plt.show()
```

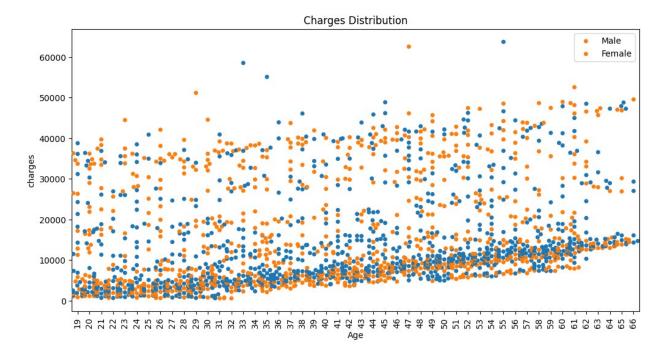




### Observation:

- The median BMI is around 30 and age is around 40.
- For BMI, there are several outliers beyond the upper whisker, with values approximately exceeding 55.
- For age, the whiskers range from around 18 to 64.

```
plt.figure(figsize=(12,6))
sns.swarmplot(data=df, y='charges', x='Age', hue='Gender')
plt.title('Charges Distribution')
plt.legend(labels=['Male','Female'])
plt.xticks(rotation=90)
plt.show()
```

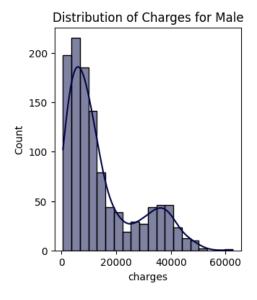


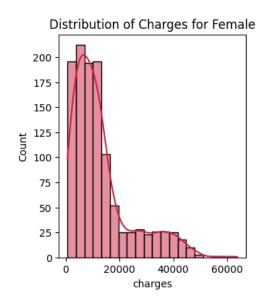
# 10. State how the distribution is different across gender and tiers of hospitals.

```
# Plot histogram for male and Female charges

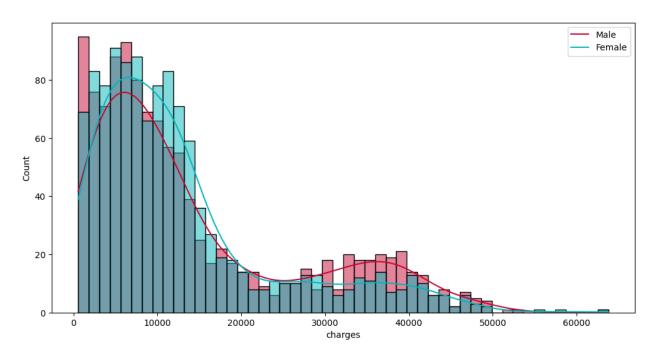
fig, ax = plt.subplots(1, 2, figsize=(10, 4))
for i, (gender, label, color) in enumerate(zip([1, 0], ['Male', 'Female'], ['#020647', '#cf2340'])):
    sns.histplot(df[df['Gender'] == gender]['charges'], ax=ax[i],
kde=True, bins=20, color=color)
    ax[i].set_title(f'Distribution of Charges for {label}')

fig.subplots_adjust(wspace=1)
plt.show()
```





```
# Plot histogram for combined male and Female charges.
plt.figure(figsize=(12, 6))
custom_palette = ["#06b7ba", "#c70a36"]
sns.histplot(data=df, x='charges', hue='Gender', alpha=0.5, bins=50,
palette=custom_palette,kde=True)
plt.legend(labels=['Male', 'Female'])
plt.show()
```

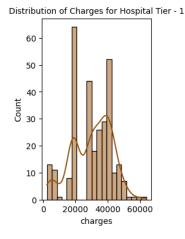


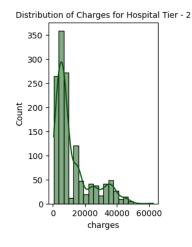
```
# Plot histogram for Different Tier of hospital charges

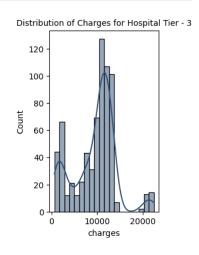
fig, ax = plt.subplots(1,3,figsize=(12,4))
hospital_tier = [1,2,3]
color_pal = ['#9c5209','#045908','#315378']

for i, hospital_tier in enumerate(hospital_tier):
    sns.histplot(df[df['Hospital_tier_num'] == hospital_tier]
['charges'], ax=ax[i], kde=True, bins=20,color=color_pal[i])
    ax[i].set_title(f'Distribution of Charges for Hospital Tier -
{hospital_tier}', fontsize=10)

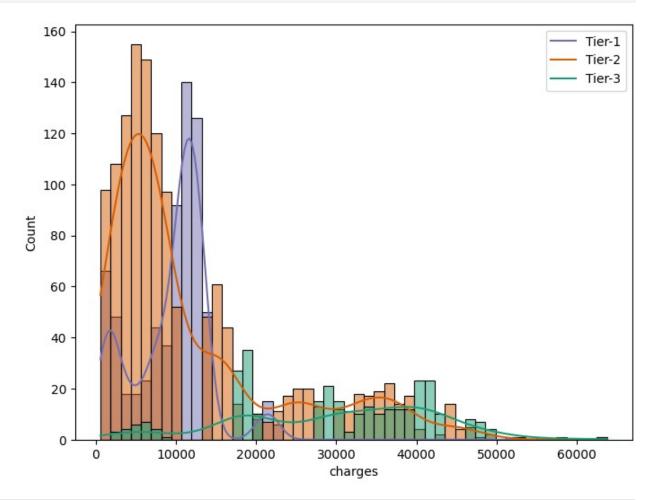
fig.subplots_adjust(wspace=1)
plt.show()
```



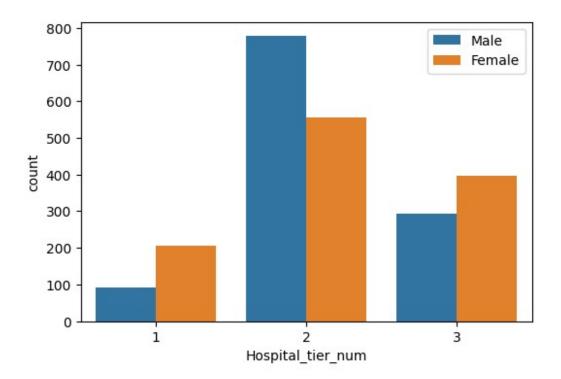




```
# Plot histogram for combined different Tier of hospital charges
plt.figure(figsize=(8, 6))
sns.histplot(data=df, x='charges', hue='Hospital_tier_num', alpha=0.5,
bins=50, palette='Dark2', kde=True)
plt.legend(labels=['Tier-1', 'Tier-2', 'Tier-3'])
plt.show()
```



```
# Plot countplot for different tiers of hospitals across gender.
plt.figure(figsize=(6, 4))
sns.countplot(data=df, x='Hospital_tier_num', hue='Gender')
plt.legend(labels=['Male', 'Female'])
plt.show()
```



## 11. Create a radar chart to showcase the median hospitalization cost for each tier of hospitals.

```
# Calculate the median hospitalization cost for each tier of hospitals
tier medians = df.groupby('Hospital tier num')
['charges'].median().reset index()
#covert tier medians into dataframe to plot radar plot
r_theta = {'r': tier_medians['charges'].tolist(), 'theta':['Tier-
1, 'Tier-2, 'Tier-3]
radar data = pd.DataFrame(r theta)
radar data
               theta
   32097.435
              Tier-1
   7168.760
              Tier-2
1
             Tier-3
  10676.830
import plotly.express as px
radar plot=px.line polar(radar data,
r='r',theta='theta',line close=True)
radar_plot.update_traces(fill='toself')
radar plot.show()
{"config":{"plotlyServerURL":"https://plot.ly"},"data":
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```
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```

```
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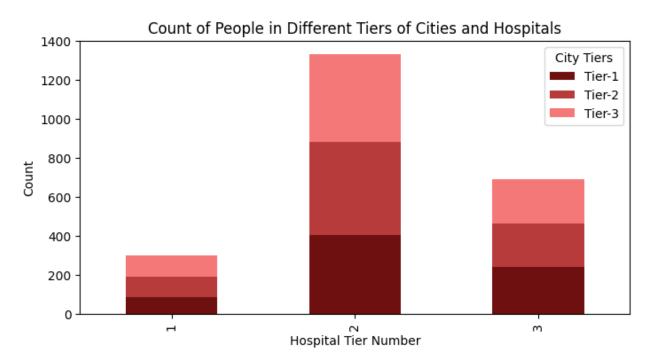
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```

12. Create a frequency table and a stacked bar chart to visualize the count of people in the different tiers of cities and hospitals.

```
#Create crosstab frequency table
freq_table = pd.crosstab(df['Hospital_tier_num'],df['City_tier_num'])
freq_table

#Calculating hospital tier wise and City tier wise total population
frequency_table=freq_table.copy()
total_city_population = frequency_table.sum(axis=1)
total_hospital_population = frequency_table.sum(axis=0)
frequency_table['City_Total_Population'] = total_city_population
```

```
frequency table.loc['Hosp Total Population'] =
total hospital population
#Calculating total polulation
total population = total hospital population.sum()
frequency table.loc['Hosp Total Population','City Total Population'] =
total_population
frequency table
City tier num
                                           City Total Population
Hospital tier num
                        85.0
                              106.0
                                     109.0
                                                            300.0
2
                       403.0
                             479.0
                                     452.0
                                                           1334.0
3
                       241.0 222.0
                                    228.0
                                                            691.0
Hosp Total Population 729.0 807.0 789.0
                                                           2325.0
# Create a stacked bar chart
fig, ax = plt.subplots(figsize=(8, 4))
freq table.plot(kind='bar', stacked=True, ax=ax, color=['#6e1010',
'#b83b3b', '#f57878'])
plt.title('Count of People in Different Tiers of Cities and
Hospitals')
plt.xlabel('Hospital Tier Number')
plt.ylabel('Count')
plt.legend(title='City Tiers', labels=['Tier-1', 'Tier-2', 'Tier-3'])
plt.show()
```



- 13. Test the following null hypotheses:
- a. The average hospitalization costs for the three types of hospitals are not significantly different.
- b. The average hospitalization costs for the three types of cities are not significantly different.
- c. The average hospitalization cost for smokers is not significantly different from the average cost for nonsmokers.
- d. Smoking and heart issues are independent.

Question	X-Variable	Y-Variable	X-Variable Datatype	Y-Variable Datatype	Statistical Test
Q.13(a)	Hospital Tier	Avg charges	Categorical	Continuous	ANOVA
Q.13(b)	City Tier	Avg charges	Categorical	Continuous	ANOVA
Q.13(c)	Smoker	Avg charges	Categorical	Continuous	ANOVA
Q.13(d)	Smoker	Heart Issues	Categorical	Categorical	Chi Square

H0: Equal Means (or no relationship between variables)

Ha: Unequal Means (or effect or relationship between variables)

### Q. 13(a)

```
# Checking the Avg. value of different Hospital Tiers
avg tier1 charges=df[df['Hospital tier num'] == 1]['charges'].mean()
avg_tier2_charges=df[df['Hospital_tier_num'] == 2]['charges'].mean()
avg tier3_charges=df[df['Hospital_tier_num'] == 3]['charges'].mean()
print('Avg. Tier-1 Hospital Charges: ',avg_tier1_charges,'\n', 'Avg.
Tier-2 Hospital Charges: ',avg tier2 charges,'\n', 'Avg. Tier-3
Hospital Charges: ', avg tier3 charges)
#Perform the ANOVA test for the average hospitalization costs for the
three types of hospitals.
Tier1_charges=df[df['Hospital_tier_num'] == 1]['charges']
Tier2 charges=df[df['Hospital tier num'] == 2]['charges']
Tier3 charges=df[df['Hospital tier num'] == 3]['charges']
from scipy.stats import f_oneway
f stat, p val = f oneway(\overline{\text{Tier1}} charges,\overline{\text{Tier2}} charges,\overline{\text{Tier3}} charges)
print('F-statistic: %.3f' % f stat)
print('p-value: %.3f' % p_val)
# Interpret the results
if p val < 0.05:
```

```
print('Reject the null hypothesis:','\n', 'Conclusion: The average
hospitalization costs for the three types of hospitals are
significantly different.')
else:
    print('Fail to reject the null hypothesis:','\n', 'Conclusion: The
average hospitalization costs for the three types of hospitals are not
significantly different.')

Avg. Tier-1 Hospital Charges: 30131.995899999998
Avg. Tier-2 Hospital Charges: 11875.883860569715
Avg. Tier-3 Hospital Charges: 9487.456222865412
F-statistic: 493.990
p-value: 0.000
Reject the null hypothesis:
Conclusion: The average hospitalization costs for the three types of
hospitals are significantly different.
```

### Q. 13(b)

```
# Checking the Avg. value of different City Tiers
avg_tier1_charges=df[df['City_tier_num'] == 1]['charges'].mean()
avg_tier2_charges=df[df['City_tier_num'] == 2]['charges'].mean()
avg_tier3_charges=df[df['City_tier_num'] == 3]['charges'].mean()
print('Avg. Tier-1 Hospital Charges: ',avg_tier1_charges,'\n', 'Avg.
Tier-2 Hospital Charges: ',avg tier2 charges,'\n', 'Avg. Tier-3
Hospital Charges: ', avg_tier3_charges)
#Perform the ANOVA test for the average hospitalization costs for the
three types of cities.
Tier1 charges=df[df['City tier num'] == 1]['charges']
Tier2_charges=df[df['City_tier_num'] == 2]['charges']
Tier3 charges=df[df['City tier num'] == 3]['charges']
f stat, p val = f oneway(Tier1 charges, Tier2 charges, Tier3 charges)
print('F-statistic: %.3f' % f stat)
print('p-value: %.3f' % p val)
if p val < 0.05:
    print('Reject the null hypothesis:','\n', 'Conclusion: The average
hospitalization costs for the three types of cities are significantly
different.')
else:
    print('Fail to reject the null hypothesis:','\n', 'Conclusion: The
average hospitalization costs for the three types of cities are not
significantly different.')
Avg. Tier-1 Hospital Charges: 13009.972578875171
Avg. Tier-2 Hospital Charges: 13471.919281288725
 Avg. Tier-3 Hospital Charges: 14045.312065906212
```

```
F-statistic: 1.454
p-value: 0.234
Fail to reject the null hypothesis:
Conclusion: The average hospitalization costs for the three types of cities are not significantly different.
```

### Q. 13(c)

```
# Checking the Avg. value of Smokers and Non-smokers
avg_Smoker_charges=df[df['smoker'] == 'yes']['charges'].mean()
avg_NonSmoker_charges=df[df['smoker'] == 'No']['charges'].mean()
print('Avg. Smoker Charges: ',avg_Smoker_charges,'\n', 'Avg. Non-
Smoker Charges: ',avg NonSmoker charges)
#Perform the ANOVA test for the average hospitalization costs for the
smokers.
Smoker charges=df[df['smoker'] == 'yes']['charges']
NonSmoker charges=df[df['smoker'] == 'No']['charges']
f stat, p val = f oneway(Smoker charges, NonSmoker charges)
print('F-statistic: %.3f' % f stat)
print('p-value: %.3f' % p val)
if p val < 0.05:
    print('Reject the null hypothesis:','\n', 'Conclusion: The average
hospitalization costs for the smokers and Non-smokers are
significantly different.')
else:
    print('Fail to reject the null hypothesis:','\n', 'Conclusion: The
average hospitalization costs for the smokers and Non-smokers are not
significantly different.')
Avg. Smoker Charges: 32866.96022633745
Avg. Non-Smoker Charges: 8409.19924959217
F-statistic: 5499.054
p-value: 0.000
Reject the null hypothesis:
 Conclusion: The average hospitalization costs for the smokers and
Non-smokers are significantly different.
```

### Q. 13(d)

```
# Create a contingency table
contingency_table = pd.crosstab(df['smoker'], df['Heart Issues'])
contingency_table

Heart Issues No yes
smoker
```

```
No
              1108 731
               297 189
ves
# Perform the Chi-Square test
from scipy.stats import chi2 contingency
chi2, p, dof, expected = chi2 contingency(contingency table)
print('Chi-Square Statistic: %.3f' % chi2)
print('p-value: %.3f' % p)
# Interpret the results
if p < 0.05:
    print('Reject the null hypothesis:','\n', 'Conclusion: Smoking and
heart issues are not independent.')
else:
    print('Fail to reject the null hypothesis:','\n', 'Conclusion:
Smoking and heart issues are independent.')
Chi-Square Statistic: 0.086
p-value: 0.769
Fail to reject the null hypothesis:
Conclusion: Smoking and heart issues are independent.
```

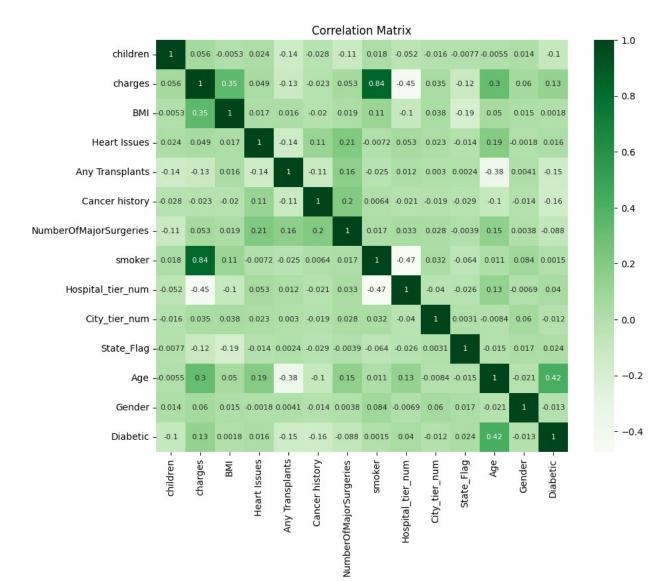
### >> MACHINE LEARNING

1. Examine the correlation between predictors to identify highly correlated predictors.

```
#Creting a new Dataframe with relevant features only
df1=df.copv()
df.drop(['Customer ID', 'year', 'month', 'date', 'Hospital tier', 'City
tier','State ID','name','D.O.B'], axis=1, inplace=True)
df.head()
   children
                              HBA1C Heart Issues Any Transplants \
             charges
                         BMI
0
          0
              563.84 17.58
                               4.51
                                               No
                                                                No
1
          0
              570.62 17.60
                               4.39
                                               No
                                                                No
2
          0
              600.00
                      16.47
                               6.35
                                               No
                                                                No
3
                      17.70
          0
              604.54
                               6.28
                                               No
                                                                No
              637.26 22.34
                               5.57
                                               No
                                                                No
  Cancer history
                   NumberOfMajorSurgeries smoker
                                                   Hospital tier num \
0
              No
                                               No
                                                                    2
                                         1
1
              No
                                               No
2
                                         1
                                                                    2
             Yes
                                               No
3
                                         1
                                                                    3
              No
                                               No
4
                                                                    3
              No
                                               No
   City tier num State Flag Age Gender
```

```
0
                3
                            3
                                32
                                          1
1
                1
                            3
                                31
                                          1
2
                            3
                1
                                31
                                          0
3
                3
                            3
                                          1
                                31
                3
                            3
4
                                          1
                                26
# Consider HBA1C>=8 as a Diabetic customer and assign diabetic
customer as 1 and non-diabetic as 0.
df['Diabetic'] = df['HBA1C'].apply(lambda x: '1' if x >= 8 else '0')
df.drop(['HBA1C'], axis=1, inplace=True)
df.head()
                         BMI Heart Issues Any Transplants Cancer
   children charges
history
          0
              563.84
                      17.58
                                        No
                                                         No
0
No
              570.62 17.60
                                        No
                                                         No
1
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2
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3
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#Transform categorical columns into numerical using Label Encoding
df2=df.copy()
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
categorical col=['Heart Issues','Any Transplants','Cancer
history','smoker']
for i in categorical col:
    df[i] = le.fit transform(df[i])
df3=df.copy()
df.head()
```

```
children charges
                         BMI
                              Heart Issues Any Transplants Cancer
history \
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# Correction between predictors using Heatmap
plt.figure(figsize=(10, 8))
sns.heatmap(df.corr(), annot=True, cmap='Greens',
square=True,annot_kws={'fontsize': 8})
plt.title('Correlation Matrix')
plt.show()
```



#### Observation:

The heatmap reveals some notable correlations:

- Charges are strongly correlated with smoking, and also show a significant correlation with BMI and age.
- Additionally, age is correlated with diabetes, indicating a possible relationship between these two factors.
- 2. Develop a regression model Linear or Ridge. Evaluate the model with k-fold cross validation.

Also, ensure that you apply all the following suggestions:

- Implement the stratified 5-fold cross validation technique for both model building and validation.
- Utilize effective standardization techniques and hyperparameter tuning.
- Incorporate sklearn-pipelines to streamline the workflow.
- Apply appropriate regularization techniques to address the biasvariance trade-off.
- Create five folds in the data, and introduce a variable to identify the folds.
- Develop Gradient Boost model and determine the variable importance scores, and identify the redundant variables.

```
#Define X and Y variable for the Regression Model
y=df['charges']
df.pop('charges')
X=df
```

## Regression Model with Ridge Regression using K-Fold Cross Validation

```
from sklearn.model_selection import KFold
from sklearn.linear_model import Ridge
from sklearn.model_selection import GridSearchCV
from sklearn.preprocessing import StandardScaler
from sklearn.pipeline import Pipeline
from sklearn.metrics import mean_squared_error, mean_absolute_error,
explained_variance_score, r2_score
```

```
# Define the k-fold cross validation object
ridge kfold = KFold(n splits=5, shuffle=True, random state=34)
# Define the pipeline for Ridge Regression
ridge pipeline = Pipeline([
    ('scaler', StandardScaler()),
    ('ridge', Ridge())
])
# Define the hyperparameter tuning space for Ridge Regression
ridge param grid = {'ridge alpha': [0.001, 0.01, 0.1, 1, 10]}
# Perform k-fold cross validation for Ridge Regression with
hyperparameter tuning
ridge maes = []
ridge mses = []
ridge rmses = []
ridge accuracies = []
ridge_r2s = []
for i, (ridge train idx, ridge val idx) in
enumerate(ridge kfold.split(X)):
    ridge_X_train, ridge_X_val = X.iloc[ridge_train_idx],
X.iloc[ridge val idx]
    ridge_y_train, ridge_y_val = y.iloc[ridge_train_idx],
y.iloc[ridge_val_idx]
    print(f"Fold {i}:")
    print(f" Train: index={ridge train idx}")
    print(f" Test: index={ridge_val_idx}")
    ridge grid search = GridSearchCV(ridge pipeline, ridge param grid,
cv=5, scoring='neg mean squared error')
    ridge grid search.fit(ridge X train, ridge y train)
    ridge y pred = ridge grid search.predict(ridge X val)
    ridge maes.append(mean absolute error(ridge y val, ridge y pred))
    ridge mses.append(mean squared_error(ridge_y_val, ridge_y_pred))
    ridge_rmses.append(mean_squared_error(ridge_y_val,
ridge y pred)**(1/2.0))
    ridge accuracies.append(explained variance score(ridge y val,
ridge y pred))
    ridge r2s.append(r2 score(ridge y val, ridge y pred))
# Get the best hyperparameters and the corresponding model
ridge best params = ridge grid search.best params
ridge best model = ridge grid search.best estimator
print(f'Best Hyperparameters: {ridge best params}')
```

```
# Calculate the average of the metrics
avg ridge mae = sum(ridge maes) / len(ridge maes)
avg_ridge_mse = sum(ridge_mses) / len(ridge_mses)
avg ridge rmse = sum(ridge rmses) / len(ridge rmses)
avg ridge accuracy = sum(ridge accuracies) / len(ridge accuracies) *
avg ridge r2 = sum(ridge r2s) / len(ridge r2s)
print(f'Average MAE: {avg_ridge_mae}')
print(f'Average MSE: {avg_ridge_mse}')
print(f'Average RMSE: {avg_ridge_rmse}')
print(f'Average Accuracy: {avg ridge accuracy:.2f}%')
print(f'Average R-squared: {avg ridge r2:.2f}')
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224	234	236	244	249	252	253	254	256	261	267	268	272	281
286	292	295	299	301	309	311	318	320	321	323	329	334	335
338	341	342	346	349	356	357	368	377	378	387	392	394	406
416	420	421	423	424	431	434	456	457	466	473	479	483	490
492	496	504	507	512	519	522	527	530	533	537	540	545	546
547	553	556	560	563	565	568	575	578	580	581	583	584	589
590	594	596	603	605	610	621	636	641	643	646	647	660	662
664	665	668	677	678	681	683	688	690	692	709	714	718	720
721	725	731	738	744	748	750	751	754	756	763	768	770	773
788	804	807	809	813	817	825	826	831	833	837	842	848	855
857	860	868	877	880	885	886	890	898	899	903	908	914	925
934	936	947	948	955	956	962	963	968	969	971	972	976	977
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193	198	202	208	216	219	220	225	232	257	259	263	269	271
275	276	277	280	283	285	289	290	293	294	303	313	315	316
317	322	331	337	339	340	343	345	355	359	366	369	370	374
375	385	389	397	402	404	405	411		417	422	445	447	448
451	460	462	470	471	472	476	487	491	494	503	508	511	518

520	521	523	525	526	529	531	532	535	539	549	550	551	569
582	591	597	599	604	620	623	625	626	627	633	645	650	651
653	658	674	679	693	697	698	703	707	708	717	719	722	724
728	729	741	747	753	757	758	764	782	793	795	796	798	801
805	810	815	822	823	834	836	844	846	847	850	853	854	856
859	865	866	871	872	875	878	893	896	904	905	906	907	911
915	917	921	923	927	928	933	937	943	949	954	958	964	965
982	986	987	995	1005	1012		1020	1021		1040	1043	1046	1056
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1404 1462	1415	1417	1410		1424		1433			1509	1513	1455 1521	1522
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                                                                    861
 863
      867
            883
                 889
                      894
                           916
                                929
                                     938
                                          945
                                               950
                                                     952
                                                          959
                                                               960
 967
      974
            975
                 981
                      984
                           993 1006 1008 1009 1013 1016 1018 1022 1024
 1027 1032 1038 1041 1042 1044 1047 1048 1055 1060 1061 1070 1073 1079
 1082 1091 1092 1094 1096 1101 1106 1108 1116 1120 1123 1124 1125 1126
 1129 1130 1131 1146 1150 1157 1159 1166 1170 1171 1179 1190 1196 1208
 1209 1210 1212 1218 1227 1230 1232 1239 1243 1248 1254 1263 1267 1271
 1280 1282 1284 1285 1301 1303 1305 1317 1320 1324 1332 1346 1348 1350
 1352 1358 1362 1368 1371 1372 1381 1382 1389 1399 1405 1412 1413 1414
 1422 1429 1434 1436 1441 1447 1448 1463 1468 1470 1472 1499 1507 1508
 1514 1516 1518 1520 1526 1537 1542 1549 1556 1557 1559 1561 1565 1571
 1588 1606 1613 1623 1629 1634 1654 1656 1668 1675 1677 1680 1682 1691
 1696 1701 1705 1708 1724 1733 1737 1743 1744 1752 1753 1754 1756 1757
 1758 1763 1770 1772 1774 1783 1795 1799 1800 1802 1810 1812 1818 1832
 1834 1840 1841 1862 1863 1866 1868 1869 1872 1891 1897 1898 1900 1914
 1933 1936 1937 1948 1950 1954 1956 1957 1959 1961 1968 1971 1972 1978
 1979 1980 1984 1986 1988 1991 1992 1998 2002 2005 2010 2015 2020 2038
 2042 2043 2056 2059 2065 2066 2070 2091 2092 2094 2095 2105 2107 2112
 2133 2145 2147 2151 2152 2154 2165 2166 2167 2168 2170 2174 2183 2187
 2191 2200 2212 2215 2216 2217 2220 2222 2224 2230 2231 2239 2241 2245
```

```
2248 2254 2258 2268 2274 2279 2281 2282 2283 2285 2290 2291 2292 2295 2300 2313 2316]
Best Hyperparameters: {'ridge__alpha': 0.001}
Average MAE: 2753.121926443369
Average MSE: 20010465.494363222
Average RMSE: 4461.915143040959
Average Accuracy: 85.67%
Average R-squared: 0.86
```

The Ridge Regression model with an alpha of 0.001 demonstrates strong predictive performance, with a high R-squared value and accuracy, along with reasonably low error metrics (MAE, MSE, RMSE). This suggests that the model is well-regularized and effectively captures the underlying patterns in the data while maintaining generalizability across different folds of the cross-validation.

### Gradient Boosting Model with Variable Importance Scores

```
from sklearn.ensemble import GradientBoostingRegressor
# Define the pipeline for Gradient Boosting
gb pipeline = Pipeline([
    ('scaler', StandardScaler()),
    ('gb', GradientBoostingRegressor())
])
# Define the hyperparameter tuning space for Gradient Boosting
gb_param_grid = {
    'gb__n_estimators': [100, 200, 300, 400, 500, 1000],
    #'gb learning rate': [0.3, 0.5, 1],
    #'gb max depth': [1,2],
    #'gb min samples split': [3,4,5],
    #'gb min samples leaf': [1,2]
}
# Define the k-fold cross validation object
gb kfold = KFold(n splits=5, shuffle=True, random state=42)
# Perform k-fold cross validation for Gradient Boosting with
hyperparameter tuning
qb maes = []
gb mses = []
gb rmses = []
gb accuracies = []
gb r2s = []
for i, (qb train idx, qb val idx) in enumerate(qb kfold.split(X)):
    gb_X_train, gb_X_val = X.iloc[gb_train_idx], X.iloc[gb_val_idx]
    gb y train, gb y val = y.iloc[gb train idx], y.iloc[gb val idx]
    print(f"Fold {i+1}:")
```

```
#print(f" Train: index={gb train idx}")
   #print(f" Test: index={gb val idx}")
   gb grid search = GridSearchCV(gb pipeline, gb param grid, cv=5,
scoring='neg mean squared error', verbose=2)
   gb grid search.fit(gb X train, gb y train)
   gb y pred = gb grid search.predict(gb X val)
   gb maes.append(mean absolute error(gb y val, gb y pred))
   gb mses.append(mean_squared_error(gb_y_val, gb_y_pred))
   gb_rmses.append(mean_squared_error(gb_y_val, gb_y_pred)**(1/2.0))
   gb accuracies.append(explained variance score(gb y val,
gb y pred))
   gb_r2s.append(r2_score(gb_y_val, gb_y_pred))
# Get the best hyperparameters and the corresponding model
gb best params = gb grid search.best params
gb best model = gb grid search.best estimator
print(f'Best Hyperparameters: {gb best params}')
Fold 1:
Fitting 5 folds for each of 5 candidates, totalling 25 fits
[CV] END .....gb n estimators=50; total
time=
      0.05
[CV] END .....gb_n_estimators=50; total
time=
      0.0s
[CV] END .....gb_n_estimators=50; total
time=
      0.0s
[CV] END .....gb n estimators=50; total
time= 0.0s
[CV] END .....gb n estimators=50; total
      0.0s
time=
[CV] END .....gb n estimators=100; total
time=
      0.1s
[CV] END .....gb__n_estimators=100; total
time=
      0.1s
[CV] END .....gb n estimators=100; total
time=
      0.1s
[CV] END .....gb__n_estimators=100; total
      0.1s
time=
[CV] END .....gb__n_estimators=100; total
time=
      0.1s
[CV] END .....gb__n_estimators=200; total
time= 0.4s
[CV] END .....gb n estimators=200; total
      0.3s
time=
[CV] END .....gb__n_estimators=200; total
time=0.4s
[CV] END .....gb n estimators=200; total
```

```
time=
     0.4s
[CV] END .....gb n estimators=200; total
time=
     0.3s
[CV] END .....gb__n_estimators=300; total
time=
     0.6s
[CV] END .....gb n estimators=300; total
     0.6s
time=
[CV] END .....gb n estimators=300; total
     0.6s
time=
[CV] END .....gb__n_estimators=300; total
     0.6s
time=
[CV] END .....gb_n_estimators=300; total
     0.6s
time=
[CV] END .....gb n estimators=400; total
time=
     0.8s
[CV] END .....gb n estimators=400; total
time=
     0.8s
[CV] END .....gb_n_estimators=400; total
time=
     1.0s
[CV] END .....gb n estimators=400; total
time=
     0.9s
[CV] END .....gb n estimators=400; total
     0.8s
time=
Fold 2:
Fitting 5 folds for each of 5 candidates, totalling 25 fits
[CV] END .....gb_n_estimators=50; total
time=
     0.0s
[CV] END .....gb n estimators=50; total
time=
     0.0s
[CV] END .....gb_n_estimators=50; total
     0.0s
time=
[CV] END .....gb n estimators=50; total
time=
     0.0s
[CV] END .....gb n estimators=50; total
     0.1s
time=
[CV] END .....gb__n_estimators=100; total
     0.2s
time=
[CV] END .....gb__n_estimators=100; total
     0.2s
time=
[CV] END .....gb n estimators=100; total
     0.2s
time=
[CV] END .....gb n estimators=100; total
time=
     0.1s
[CV] END .....gb n estimators=100; total
     0.1s
time=
[CV] END .....gb_n_estimators=200; total
     0.4s
time=
[CV] END .....gb n estimators=200; total
time=
     0.4s
```

```
[CV] END .....gb_n_estimators=200; total
time=
     0.3s
[CV] END .....gb_n_estimators=200; total
     0.4s
time=
[CV] END .....gb__n_estimators=200; total
     0.4s
time=
[CV] END .....gb n estimators=300; total
     0.6s
time=
[CV] END .....gb n estimators=300; total
time=
     0.7s
[CV] END .....gb n estimators=300; total
     0.7s
[CV] END .....gb n estimators=300; total
time=
     0.6s
[CV] END .....gb n estimators=300; total
time=
     0.8s
[CV] END .....gb__n_estimators=400; total
     1.3s
time=
[CV] END .....gb n estimators=400; total
time=
     1.0s
[CV] END .....gb__n_estimators=400; total
time=
     0.9s
[CV] END .....gb__n_estimators=400; total
time= 1.3s
[CV] END .....gb n estimators=400; total
time=
     1.0s
Fold 3:
Fitting 5 folds for each of 5 candidates, totalling 25 fits
[CV] END .....gb n estimators=50; total
     0.0s
time=
[CV] END .....gb n estimators=50; total
     0.1s
time=
[CV] END .....gb n estimators=50; total
time=
     0.0s
[CV] END .....gb n estimators=50; total
time=
     0.0s
[CV] END .....gb n estimators=50; total
     0.1s
[CV] END .....gb__n_estimators=100; total
     0.2s
time=
[CV] END .....gb__n_estimators=100; total
time=
     0.2s
[CV] END .....gb_n_estimators=100; total
     0.2s
time=
[CV] END .....gb_n_estimators=100; total
     0.2s
time=
[CV] END .....gb n estimators=100; total
     0.2s
time=
[CV] END .....gb n estimators=200; total
```

```
time=
     0.6s
[CV] END .....gb n estimators=200; total
time=
     0.4s
[CV] END .....gb__n_estimators=200; total
     0.4s
[CV] END .....gb n estimators=200; total
     0.3s
time=
[CV] END .....gb n estimators=200; total
time=
     0.3s
[CV] END .....gb__n_estimators=300; total
     0.6s
time=
[CV] END .....gb_n_estimators=300; total
     0.6s
time=
[CV] END .....gb n estimators=300; total
time=
     0.6s
[CV] END .....gb n estimators=300; total
time=
     0.5s
[CV] END .....gb_n_estimators=300; total
     0.6s
[CV] END .....gb__n_estimators=400; total
time=
     0.8s
[CV] END .....gb n estimators=400; total
     0.8s
time=
[CV] END .....gb__n_estimators=400; total
     0.8s
time=
[CV] END .....gb__n_estimators=400; total
     0.9s
time=
[CV] END .....gb_n_estimators=400; total
time= 0.8s
Fold 4:
Fitting 5 folds for each of 5 candidates, totalling 25 fits
[CV] END .....gb_n_estimators=50; total
     0.0s
time=
[CV] END .....gb n estimators=50; total
     0.0s
time=
[CV] END .....gb_n_estimators=50; total
     0.0s
time=
[CV] END .....gb _ n_estimators=50; total
     0.0s
time=
[CV] END .....gb n estimators=50; total
     0.0s
time=
[CV] END .....gb__n_estimators=100; total
time=
     0.2s
[CV] END .....gb n estimators=100; total
     0.1s
time=
[CV] END .....gb_n_estimators=100; total
     0.2s
time=
[CV] END .....gb n estimators=100; total
time=
     0.2s
```

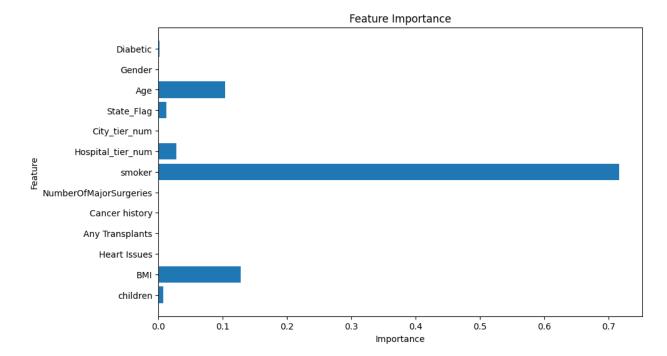
```
[CV] END .....gb_n_estimators=100; total
time=
     0.2s
[CV] END .....gb_n_estimators=200; total
     0.4s
time=
[CV] END .....gb n estimators=200; total
     0.4s
time=
[CV] END .....gb n estimators=200; total
     0.4s
time=
[CV] END .....gb n estimators=200; total
time=
     0.4s
[CV] END .....gb n estimators=200; total
     0.4s
[CV] END .....gb n estimators=300; total
time=
     0.6s
[CV] END .....gb n estimators=300; total
time=
     0.6s
[CV] END .....gb__n_estimators=300; total
time=
     0.6s
[CV] END .....gb__n_estimators=300; total
time=
     0.6s
[CV] END .....gb__n_estimators=300; total
time=
     0.6s
[CV] END .....gb__n_estimators=400; total
time=
     0.9s
[CV] END .....gb__n_estimators=400; total
     0.8s
time=
[CV] END .....gb__n_estimators=400; total
time=
     0.8s
[CV] END .....gb n estimators=400; total
time=
     0.8s
[CV] END .....gb n estimators=400; total
     0.8s
time=
Fold 5:
Fitting 5 folds for each of 5 candidates, totalling 25 fits
[CV] END .....gb n estimators=50; total
time=
     0.0s
[CV] END .....gb_n_estimators=50; total
     0.0s
[CV] END .....gb_n_estimators=50; total
     0.0s
time=
[CV] END .....gb__n_estimators=50; total
     0.0s
time=
[CV] END .....gb_n_estimators=50; total
time=
     0.0s
[CV] END .....gb_n_estimators=100; total
time=
     0.1s
[CV] END .....gb n estimators=100; total
     0.1s
time=
[CV] END .....gb n estimators=100; total
```

```
time=
      0.1s
[CV] END .....gb n estimators=100; total
time=
      0.1s
[CV] END .....gb n estimators=100; total
      0.1s
[CV] END .....gb n estimators=200; total
      0.5s
time=
[CV] END ......gb n estimators=200; total
time=
      0.3s
[CV] END .....gb__n_estimators=200; total
time=
      0.4s
[CV] END .....gb_n_estimators=200; total
      0.3s
time=
[CV] END .....gb n_estimators=200; total
time=
      0.4s
[CV] END .....gb__n_estimators=300; total
time=
      0.6s
[CV] END .....gb__n_estimators=300; total
time=0.6s
[CV] END .....gb__n_estimators=300; total
time=
      0.6s
[CV] END .....gb n estimators=300; total
      0.65
time=
[CV] END .....gb__n_estimators=300; total
      0.6s
time=
[CV] END .....gb__n_estimators=400; total
      0.8s
time=
[CV] END .....gb n estimators=400; total
time= 0.8s
[CV] END .....gb_n_estimators=400; total
      0.9s
time=
[CV] END .....gb n estimators=400; total
time= 1.3s
[CV] END .....gb n estimators=400; total
time=
      1.3s
Best Hyperparameters: {'gb n estimators': 400}
# Calculate the average of the metrics
avq qb mae = sum(gb maes) / len(gb_maes)
avg_gb_mse = sum(gb_mses) / len(gb_mses)
avg gb rmse = sum(gb rmses) / len(gb rmses)
avg gb accuracy = sum(gb accuracies) / len(gb accuracies) * 100
avg gb r2 = sum(gb \ r2s) / len(gb \ r2s)
print(f'Average MAE: {avg_gb_mae}')
print(f'Average MSE: {avg gb mse}')
print(f'Average RMSE: {avg gb rmse}')
print(f'Average Accuracy: {avg_gb_accuracy:.2f}%')
print(f'Average R-squared: {avg_gb r2:.2f}')
```

```
Average MAE: 2125.803116928094
Average MSE: 13082270.633482825
Average RMSE: 3584.6096743740395
Average Accuracy: 90.80%
Average R-squared: 0.91
```

The Gradient Boosting model outperforms the Ridge Regression model in terms of error metrics (MAE, MSE, RMSE), accuracy, and R-squared value. This suggests that Gradient Boosting is better suited for this regression task, providing more accurate and reliable predictions compared to Ridge Regression.

```
# Get the feature importance scores
feature importances =
gb best model.named steps['gb'].feature importances
# Create a DataFrame to store the feature importance scores
df feature importances = pd.DataFrame({'Feature': X.columns,
'Importance': feature importances})
# Sort the DataFrame by importance in descending order
df feature importances sort =
df feature importances.sort values(by='Importance', ascending=False)
print(df feature importances sort)
                   Feature Importance
6
                    smoker
                              0.717429
1
                       BMI
                              0.127649
10
                       Age
                              0.103964
7
         Hospital tier num
                              0.027680
                State Flag
9
                              0.011800
0
                  children
                              0.007196
12
                  Diabetic
                              0.001128
11
                    Gender
                              0.001030
8
             City tier num
                              0.000821
2
              Heart Issues
                              0.000607
5
    NumberOfMajorSurgeries
                              0.000324
3
           Any Transplants
                              0.000237
4
            Cancer history
                              0.000136
# Plot feature importances
plt.figure(figsize=(10, 6))
plt.barh(range(len(df feature importances)),
df feature importances['Importance'])
plt.yticks(range(len(df feature importances)),
df_feature_importances['Feature'])
plt.xlabel('Importance')
plt.ylabel('Feature')
plt.title('Feature Importance')
plt.show()
```



The feature importance scores show that 'smoker' is the most significant predictor, while features like 'Cancer history' and 'Any Transplants' have minimal impact.

```
# Identify redundant variables by selecting features with importance <
0.01
redundant_variables =
df_feature_importances[df_feature_importances['Importance'] < 0.01]
['Feature'].tolist()
print(f'Redundant Variables: {redundant_variables}')

Redundant Variables: ['children', 'Heart Issues', 'Any Transplants',
'Cancer history', 'NumberOfMajorSurgeries', 'City_tier_num', 'Gender',
'Diabetic']</pre>
```

### 3. Case scenario:

Estimate the cost of hospitalization for Christopher, Ms. Jayna (Date of birth 12/28/1988; height 170 cm; and weight 85 kgs). She lives with her partner and two children in a tier-1 city, and her state's State ID is R1011. She was found to be nondiabetic (HbA1c = 5.8). She smokes but is otherwise healthy. She has had no transplants or major surgeries. Her father died of lung cancer. Hospitalization costs will be estimated using tier-1 hospitals.

```
#Age Calculation
jayna_dob = '12/28/1988'
jayna_dob = pd.to_datetime(jayna_dob, format='%m/%d/%Y')
```

```
jayna_age = age_calculation(jayna dob.date())
print(jayna age)
35
#BMI Calculation
iavna height = 170
jayna weight = 85
jayna bmi = round(jayna weight / ((jayna height/100) ** 2),2)
Convert height from cm to m
print(jayna bmi)
29.41
#Make Dataframe
jayna df =pd.DataFrame({'children':[2], 'BMI':[jayna bmi],
       'Heart Issues':[0], 'Any Transplants':[0], 'Cancer history':
[1],
       'NumberOfMajorSurgeries':[0], 'smoker':[1],
'Hospital tier num':[1],
       'City tier num':[1], 'State Flag':[1], 'Age':[jayna age],
'Gender': [0], 'Diabetic':[0]})
jayna df
   children
               BMI Heart Issues Any Transplants
                                                   Cancer history
0
     2
            29.41
  NumberOfMajorSurgeries
                                   Hospital tier num City tier num \
                           smoker
0
                        0
                                1
   State Flag Age Gender
                            Diabetic
0
           1 35
#Hospital cost Calculation
ridge hospitalization cost = ridge grid search.predict(jayna df)
print(f"Predicted Hospitalization Cost:
{round(ridge hospitalization cost[0], 3)}")
Predicted Hospitalization Cost: 31879.883
```

## 4. Find the predicted hospitalization cost using the best models (Gradient Boosting Model)

```
gb_hospitalization_cost= gb_grid_search.predict(jayna_df)
print(f"Predicted Hospitalization Cost:
{round(gb_hospitalization_cost[0],3)}")
Predicted Hospitalization Cost: 27831.229
```