```
import warnings
warnings.filterwarnings('ignore')
# Import the numpy and pandas package
import numpy as np
import pandas as pd
# Read the given CSV file, and view some sample records
medical = pd.read_csv('../content/Medical Price Dataset.csv')
medical.head()
                       bmi children smoker
                                                                     1
        age
               sex
                                               region
                                                           charges
     0
        19 female 27.900
                                        yes southwest 16884.92400
              male 33.770
                                                        1725.55230
                                  1
                                         no southeast
     1
         18
      2
         28
              male 33.000
                                  3
                                         no southeast
                                                        4449.46200
                                  0
                                         no northwest 21984.47061
     3
         33
              male 22.705
              male 28.880
                                         no northwest
                                                        3866.85520
#Determining the number of rows and columns
medical.shape
     (1338, 7)
medical.describe() #summary of all the numeric columns in the dataset
                                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
              18.000000
                           15.960000
                                        0.000000
                                                   1121.873900
      min
      25%
              27.000000
                           26.296250
                                        0.000000
                                                  4740.287150
                          30.400000
      50%
              39 000000
                                        1.000000
                                                  9382 033000
      75%
              51.000000
                           34.693750
                                        2.000000 16639.912515
      max
              64 000000
                           53 130000
                                        5 000000 63770 428010
medical.info() #Datatypes of each column
     <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
                   1338 non-null
                                   object
         sex
                  1338 non-null
         bmi
                                   float64
     3
         children 1338 non-null
                                   int64
         smoker 1338 non-null
     4
                                   object
                   1338 non-null
     5
         region
                                   object
         charges 1338 non-null
     6
                                   float64
     dtypes: float64(2), int64(2), object(3)
     memory usage: 73.3+ KB
#Checking missing values
medical.isnull().sum()
                0
     age
     sex
                0
     hmi
                a
     children
                0
     smoker
                0
     region
                0
     charges
    dtype: int64
medical['sex'] = medical['sex'].map({'male': 0, 'female': 1})
medical['smoker'] = medical['smoker'].map({'yes': 1, 'no': 0})
medical.head()
```

	age	sex	bmi	children	smoker	region	charges	1
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	

#Import necessary libraies import matplotlib.pyplot as plt import seaborn as sns

#Binning the age column.
bins = [17,35,55,1000]

slots = ['Young adult','Senior Adult','Elder']

medical['Age\_range']=pd.cut(medical['age'],bins=bins,labels=slots)

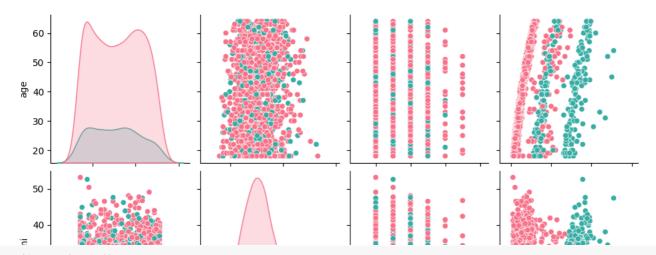
## medical.head()

	age	sex	bmi	children	smoker	region	charges	Age_range	7
0	19	1	27.900	0	1	southwest	16884.92400	Young adult	
1	18	0	33.770	1	0	southeast	1725.55230	Young adult	
2	28	0	33.000	3	0	southeast	4449.46200	Young adult	
3	33	0	22.705	0	0	northwest	21984.47061	Young adult	
4	32	0	28.880	0	0	northwest	3866.85520	Young adult	

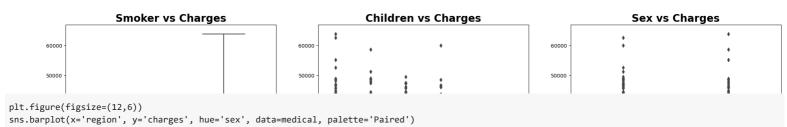
## medical.nunique().sort\_values()

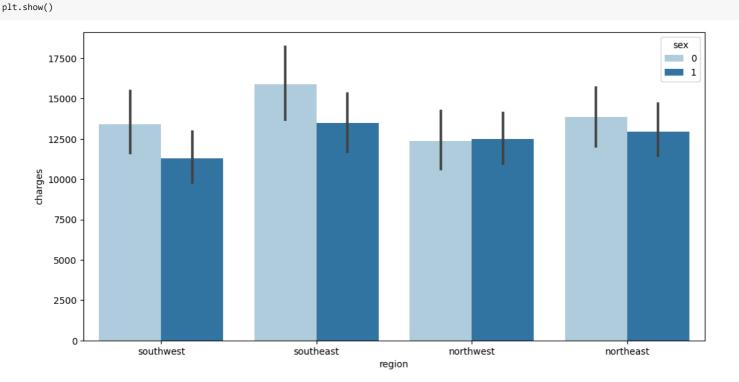
sex 2
smoker 2
Age\_range 3
region 4
children 6
age 47
bmi 548
charges 1337
dtype: int64

#Pairplot of all numerical variables
sns.pairplot(medical, vars=["age", 'bmi','children','charges'],hue='smoker',palette="husl")
plt.show()

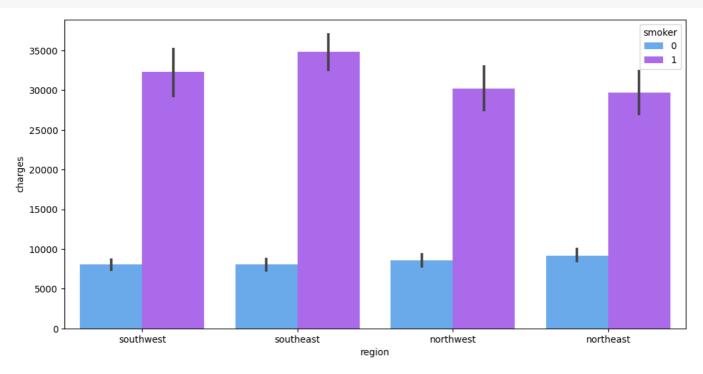


```
plt.figure(figsize=(25, 16))
plt.subplot(2,3,1)
sns.boxplot(x = 'smoker', y = 'charges', data = medical)
plt.title('Smoker vs Charges',fontweight="bold", size=20)
plt.subplot(2,3,2)
sns.boxplot(x = 'children', y = 'charges', data = medical,palette="husl")
plt.title('Children vs Charges',fontweight="bold", size=20)
plt.subplot(2,3,3)
sns.boxplot(x = 'sex', y = 'charges', data = medical, palette= 'husl')
plt.title('Sex vs Charges',fontweight="bold", size=20)
plt.subplot(2,3,4)
sns.boxplot(x = 'region', y = 'charges', data = medical,palette="bright")
plt.title('Region vs Charges',fontweight="bold", size=20)
plt.subplot(2,3,5)
sns.boxplot(x = 'Age_range', y = 'charges', data = medical, palette= 'husl')
plt.title('Age vs Charges',fontweight="bold", size=20)
plt.show()
```

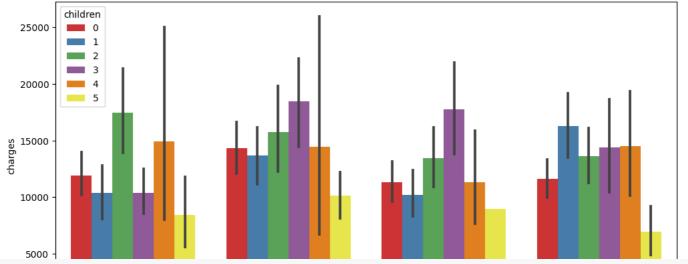




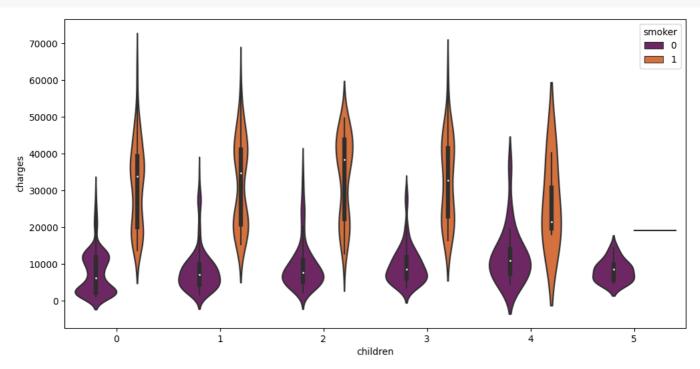
```
plt.figure(figsize=(12,6))
sns.barplot(x = 'region', y = 'charges',hue='smoker', data=medical, palette='cool')
plt.show()
```



```
plt.figure(figsize=(12,6))
sns.barplot(x='region', y='charges', hue='children', data=medical, palette='Set1')
plt.show()
```



plt.figure(figsize=(12,6))
sns.violinplot(x = 'children', y = 'charges', data=medical, hue='smoker', palette='inferno')
plt.show()



#Heatmap to see correlation between variables
plt.figure(figsize=(12, 8))
sns.heatmap(medical.corr(), cmap='RdYlGn', annot = True)
plt.title("Correlation between Variables")
plt.show()



medical.head()

	age	sex	bmi	children	smoker	region	charges	Age_range	10:
0	19	1	27.900	0	1	southwest	16884.92400	Young adult	
1	18	0	33.770	1	0	southeast	1725.55230	Young adult	
2	28	0	33.000	3	0	southeast	4449.46200	Young adult	
3	33	0	22.705	0	0	northwest	21984.47061	Young adult	
4	32	0	28.880	0	0	northwest	3866.85520	Young adult	
ē									

region=pd.get\_dummies(medical.region,drop\_first=True)

Age\_range=pd.get\_dummies(medical.Age\_range,drop\_first=True)

children= pd.get\_dummies(medical.children,drop\_first=True,prefix='children')

medical=pd.concat([region,Age\_range,children,medical],axis=1)
medical.head()

	northwest	southeast	southwest	Senior Adult	Elder	children_1	children_2	children_3	children_4	children_5	age	sex	bmi	children	smoker
0	0	0	1	0	0	0	0	0	0	0	19	1	27.900	0	1 s
1	0	1	0	0	0	1	0	0	0	0	18	0	33.770	1	0 s
2	0	1	0	0	0	0	0	1	0	0	28	0	33.000	3	0 s

medical.drop(['region', 'Age\_range', 'age','children'], axis = 1, inplace = True)
medical.head()

	northwest	southeast	southwest	Adult	Elder	children_1	children_2	children_3	children_4	children_5	sex	bmi	smoker	charges
(	0	0	1	0	0	0	0	0	0	0	1	27.900	1	16884.92400
	0	1	0	0	0	1	0	0	0	0	0	33.770	0	1725.55230
2	2 0	1	0	0	0	0	0	1	0	0	0	33.000	0	4449.46200
;	1	0	0	0	0	0	0	0	0	0	0	22.705	0	21984.47061
4	1	0	0	0	0	0	0	0	0	0	0	28.880	0	3866.85520

medical.shape

(1338, 14)

plt.figure(figsize=(15, 10))

sns.heatmap(medical.corr(), cmap='YlGnBu', annot = True)

plt.show()



from sklearn.model\_selection import train\_test\_split

# We specify this so that the train and test data set always have the same rows, respectively
#np.random.seed(0)
medical\_train, medical\_test = train\_test\_split(medical, train\_size = 0.7, random\_state = 100)

print(medical\_train.shape)
print(medical\_test.shape)

(936, 14) (402, 14)

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from sklearn.preprocessing import MinMaxScaler

medical.head()

₽		northwest	southeast	southwest	Senior Adult	Elder	children_1	children_2	children_3	children
	0	0	0	1	0	0	0	0	0	
	1	0	1	0	0	0	1	0	0	
	2	0	1	0	0	0	0	0	1	
	3	1	0	0	0	0	0	0	0	
	4							- Code —	+ Text	<b>&gt;</b>

#Instantiate an object
scaler = MinMaxScaler()

#Create a list of numeric variables
num\_vars=['bmi','charges']

#Fit on data

medical\_train[num\_vars] = scaler.fit\_transform(medical\_train[num\_vars])

medical\_train.head()

	northwest	southeast	southwest	Senior Adult	Elder	children_1	children_2	children_3	children_4	children_5	sex	bmi	smoker	charges
966	1	0	0	1	0	0	1	0	0	0	0	0.237692	1	0.364661
522	0	0	0	1	0	0	0	0	0	0	1	0.483051	0	0.139579
155	1	0	0	1	0	0	0	0	0	0	0	0.633844	0	0.093008
671	0	0	0	0	0	0	0	0	0	0	1	0.408932	0	0.045040
1173	1	0	0	1	0	0	1	0	0	0	0	0.357815	0	0.085173