```
matplotlib.pyplot for plotting
       #and chi2_contingency from scipy.stats for hypothesis testing
       import pandas as pd
       import matplotlib.pyplot as plt
       import seaborn as sns
       import numpy as np
       #Setting the option to display all columns for preview
       pd.set_option("display.max.columns", None)
       #Open the dataset and print the variables info
       inpatient = pd.read_csv("Hospital_Inpatient_Discharges__SPARCS_De-
Identified___2021_20231012.csv", low_memory=False)
       #Examine the dataset
       inpatient.info()
       #fill null values
       for col in inpatient.columns:
         inpatient[col] = inpatient[col].fillna(inpatient[col].mode()[0])
       #Selected needed columns only
```

#import the librarys needed include the Panda for opening the file and seaborn and

```
inpatient = inpatient[['Facility Name', 'Age Group', 'Gender', 'Race', 'Ethnicity', 'Length
of Stay', 'Type of Admission',
```

'Patient Disposition', 'CCSR Diagnosis Description', 'APR DRG

Description',

'APR Severity of Illness Description', 'APR Medical Surgical Description', 'Payment Typology 1',

'Payment Typology 2', 'Payment Typology 3']]

#create a new column called Insured

inpatient["Insured"] = inpatient['Payment Typology 1'] == "Self-Pay"

#display the first 20 records

inpatient.head(20)

#made the values of 120+ 121 and change the datatype of Length of Stay to numbers inpatient["Length of Stay"] = inpatient["Length of Stay"].replace('120 +', 121) inpatient["Length of Stay"] = pd.to_numeric(inpatient["Length of Stay"])

#remove all records of death people

inpatient = inpatient[inpatient['Patient Disposition'] != "Expired"]

#show the summary descriptive values of the column Length of stay

```
inpatient['Length of Stay'].describe()
       # check the distribution of the Length of stay variable
       plt.figure(figsize=(10, 6))
       sns.histplot(data=inpatient, x='Length of Stay', bins = 50)
       plt.title('Distribution of Length of Stay')
       plt.ylabel('Count')
       plt.grid(True)
       plt.show()
       #Distribution of the length of stay by payment typology
       ax=sns.barplot(x="Payment Typology 1", y="Length of Stay", data=inpatient,
errorbar=None)
       ax.bar_label(ax.containers[0], fontsize=10);
       plt.xticks(rotation=90)
       plt.title('Length of Stay vs. Primary Payment')
       plt.show()
       #creating a crosstab table
       inpatient_tab = pd.crosstab(inpatient['Length of Stay'], inpatient['Insured'], margins =
True)
       inpatient_tab
```

```
# label encoding
from sklearn.preprocessing import LabelEncoder
for col in inpatient.columns:
  inpatient_le = LabelEncoder()
  inpatient[col] = inpatient_le.fit_transform(inpatient[col])
inpatient.head()
from sklearn.feature_selection import chi2
x = inpatient.drop(columns=['Length of Stay'], axis=1)
y = inpatient['Length of Stay']
chi\_scores = chi2(x, y)
chi_scores
pd.options.plotting.backend = "plotly"
#a plot of the chi_scores, higher the chi value, higher the importance
chi_values = pd.Series(chi_scores[0], index=x.columns)
chi_values.sort_values(ascending=False, inplace=True)
fig = chi_values.plot.bar()
fig.update_yaxes(tickformat=".0s").show() # show number as is
import pandas as pd
#pd.options.plotting.backend = "plotly"
\# if p-value > 0.05, lower the importance
```

```
p_values = pd.Series(chi_scores[1], index=x.columns)
p_values.sort_values(ascending=False, inplace=True)
p_values.plot.bar()
```

fig.update_yaxes(tickformat=".0s").show() # show number as is