

```
In [2]: import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

import scipy.stats as stats

from scipy.stats import skew, kurtosis

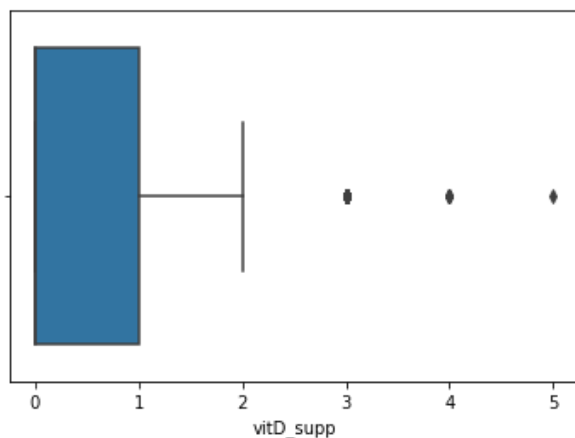
import seaborn as sns

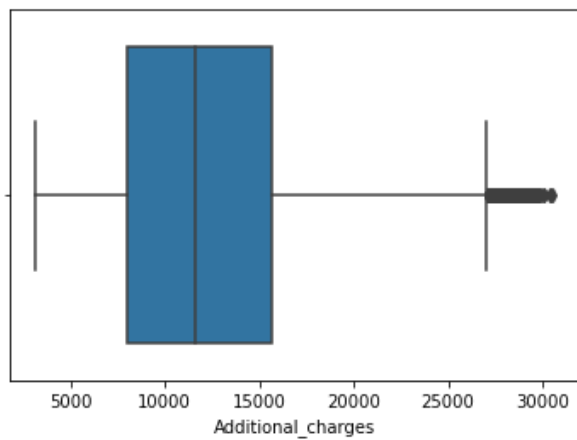
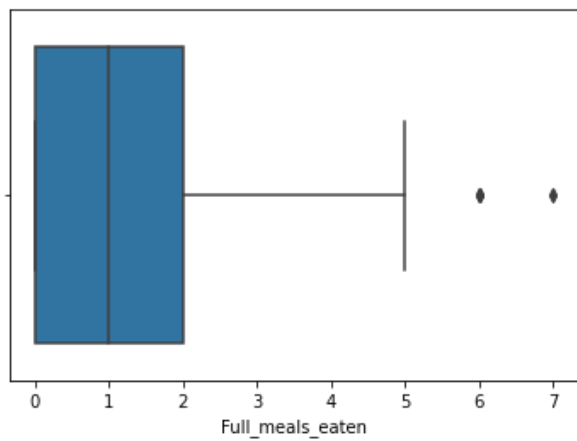
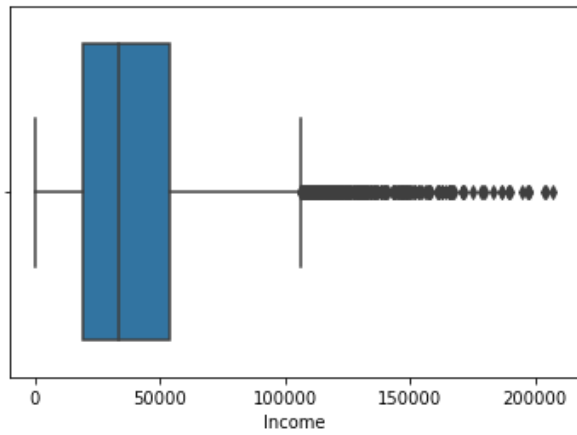
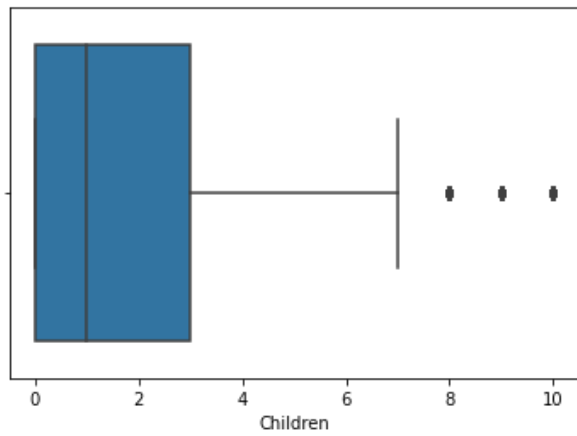
import statistics as stat

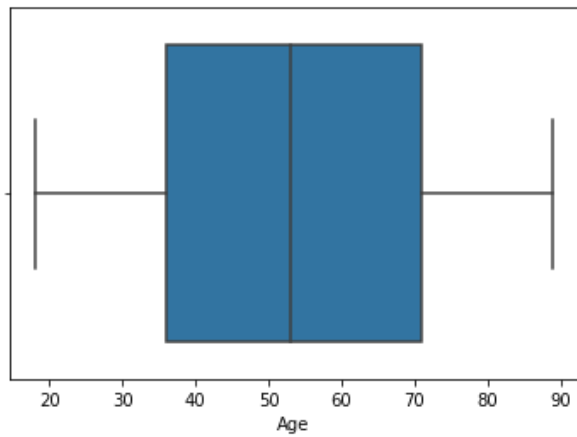
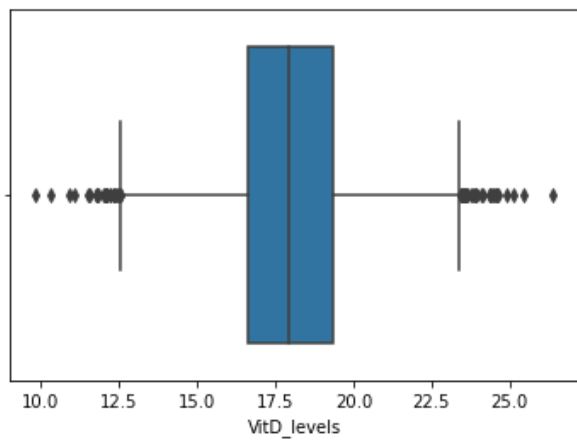
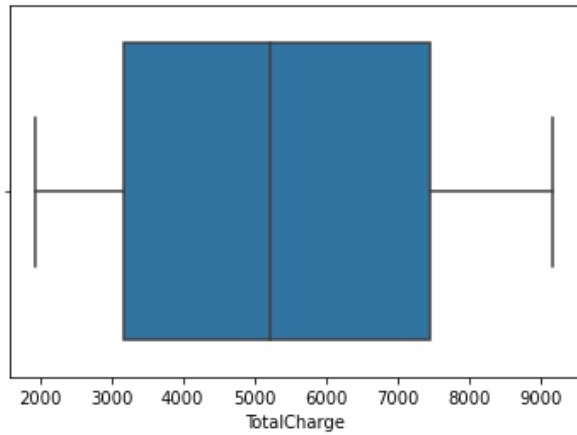
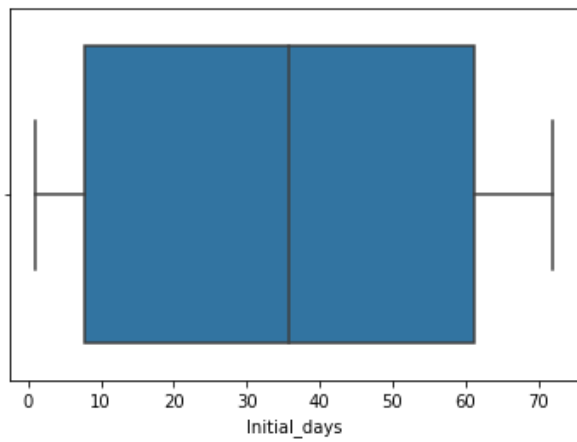
from statsmodels.formula.api import logit

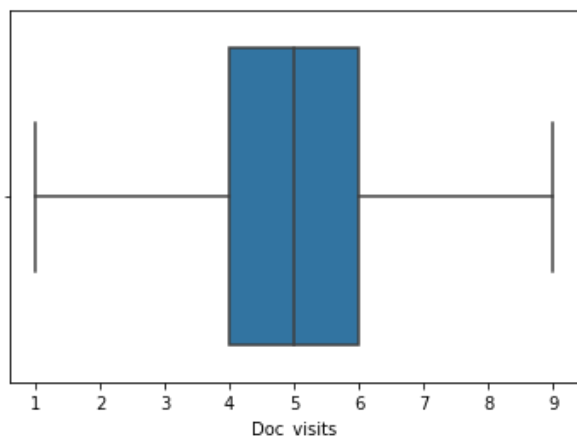
#Loading the CSV of the default dataset
df = pd.read_csv(r'C:\Users\mmorg\Desktop\D208 Assessment Files\medical_clean.csv')
```

```
In [3]: #Detection of outliers
boxplot=sns.boxplot(x='vitD_supp',data=df)
plt.show()
boxplot=sns.boxplot(x='Children',data=df)
plt.show()
boxplot=sns.boxplot(x='Income',data=df)
plt.show()
boxplot=sns.boxplot(x='Full_meals_eaten',data=df)
plt.show()
boxplot=sns.boxplot(x='Additional_charges',data=df)
plt.show()
boxplot=sns.boxplot(x='Initial_days',data=df)
plt.show()
boxplot=sns.boxplot(x='TotalCharge',data=df)
plt.show()
boxplot=sns.boxplot(x='VitD_levels',data=df)
plt.show()
boxplot=sns.boxplot(x='Age',data=df)
plt.show()
boxplot=sns.boxplot(x='Doc_visits',data=df)
plt.show()
```









```
In [4]: #Data Wrangling; turn categorical values into quantitative data
df['ReAdmis_numeric'] = df['ReAdmis']
dict_ReAdmis = {"ReAdmis_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_ReAdmis, inplace=True)

df['Soft_drink_numeric'] = df['Soft_drink']
dict_Soft_drink = {"Soft_drink_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_Soft_drink, inplace=True)

df['HighBlood_numeric'] = df['HighBlood']
dict_HighBlood = {"HighBlood_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_HighBlood, inplace=True)

df['Stroke_numeric'] = df['Stroke']
dict_stroke = {"Stroke_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_stroke, inplace=True)

df['Arthritis_numeric'] = df['Arthritis']
dict_arthritis = {"Arthritis_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_arthritis, inplace=True)

df['Diabetes_numeric'] = df['Diabetes']
dict_diabetes = {"Diabetes_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_diabetes, inplace=True)

df['Hyperlipidemia_numeric'] = df['Hyperlipidemia']
dict_hyperlipidemia = {"Hyperlipidemia_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_hyperlipidemia, inplace=True)

df['BackPain_numeric'] = df['BackPain']
dict_backpain = {"BackPain_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_backpain, inplace=True)

df['Allergic_rhinitis_numeric'] = df['Allergic_rhinitis']
dict_allergies = {"Allergic_rhinitis_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_allergies, inplace=True)

df['Reflux_esophagitis_numeric'] = df['Reflux_esophagitis']
dict_reflux = {"Reflux_esophagitis_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_reflux, inplace=True)

df['Asthma_numeric'] = df['Asthma']
dict_asthma = {"Asthma_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_asthma, inplace=True)

df['Overweight_numeric'] = df['Overweight']
dict_Overweight = {"Overweight_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_Overweight, inplace=True)

df['Anxiety_numeric'] = df['Anxiety']
dict_Anxiety = {"Anxiety_numeric": {"No": 0, "Yes": 1}}
df.replace(dict_Anxiety, inplace=True)
```

```
df = pd.get_dummies(df, columns=["Marital", "Services", "Gender", "Initial_admin", "Complication_risk"])  
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 10000 entries, 0 to 9999
```

```
Data columns (total 76 columns):
```

#	Column	Non-Null Count	Dtype
0	CaseOrder	10000 non-null	int64
1	Customer_id	10000 non-null	object
2	Interaction	10000 non-null	object
3	UID	10000 non-null	object
4	City	10000 non-null	object
5	State	10000 non-null	object
6	County	10000 non-null	object
7	Zip	10000 non-null	int64
8	Lat	10000 non-null	float64
9	Lng	10000 non-null	float64
10	Population	10000 non-null	int64
11	Area	10000 non-null	object
12	TimeZone	10000 non-null	object
13	Job	10000 non-null	object
14	Children	10000 non-null	int64
15	Age	10000 non-null	int64
16	Income	10000 non-null	float64
17	ReAdmis	10000 non-null	object
18	VitD_levels	10000 non-null	float64
19	Doc_visits	10000 non-null	int64
20	Full_meals_eaten	10000 non-null	int64
21	vitD_supp	10000 non-null	int64
22	Soft_drink	10000 non-null	object
23	HighBlood	10000 non-null	object
24	Stroke	10000 non-null	object
25	Overweight	10000 non-null	object
26	Arthritis	10000 non-null	object
27	Diabetes	10000 non-null	object
28	Hyperlipidemia	10000 non-null	object
29	BackPain	10000 non-null	object
30	Anxiety	10000 non-null	object
31	Allergic_rhinitis	10000 non-null	object
32	Reflux_esophagitis	10000 non-null	object
33	Asthma	10000 non-null	object
34	Initial_days	10000 non-null	float64
35	TotalCharge	10000 non-null	float64
36	Additional_charges	10000 non-null	float64
37	Item1	10000 non-null	int64
38	Item2	10000 non-null	int64
39	Item3	10000 non-null	int64
40	Item4	10000 non-null	int64
41	Item5	10000 non-null	int64
42	Item6	10000 non-null	int64
43	Item7	10000 non-null	int64
44	Item8	10000 non-null	int64
45	ReAdmis_numeric	10000 non-null	int64
46	Soft_drink_numeric	10000 non-null	int64
47	HighBlood_numeric	10000 non-null	int64
48	Stroke_numeric	10000 non-null	int64
49	Arthritis_numeric	10000 non-null	int64
50	Diabetes_numeric	10000 non-null	int64
51	Hyperlipidemia_numeric	10000 non-null	int64
52	BackPain_numeric	10000 non-null	int64
53	Allergic_rhinitis_numeric	10000 non-null	int64
54	Reflux_esophagitis_numeric	10000 non-null	int64
55	Asthma_numeric	10000 non-null	int64
56	Overweight_numeric	10000 non-null	int64
57	Anxiety_numeric	10000 non-null	int64
58	Marital_Divorced	10000 non-null	uint8
59	Marital_Married	10000 non-null	uint8
60	Marital_Never Married	10000 non-null	uint8
61	Marital_Separated	10000 non-null	uint8
62	Marital_Widowed	10000 non-null	uint8
63	Services_Blood Work	10000 non-null	uint8
64	Services_CT Scan	10000 non-null	uint8
65	Services_Intravenous	10000 non-null	uint8

```

66 Services_MRI 10000 non-null uint8
67 Gender_Female 10000 non-null uint8
68 Gender_Male 10000 non-null uint8
69 Gender_Nonbinary 10000 non-null uint8
70 Initial_admin_Elective Admission 10000 non-null uint8
71 Initial_admin_Emergency Admission 10000 non-null uint8
72 Initial_admin_Observation Admission 10000 non-null uint8
73 Complication_risk_High 10000 non-null uint8
74 Complication_risk_Low 10000 non-null uint8
75 Complication_risk_Medium 10000 non-null uint8
dtypes: float64(7), int64(29), object(22), uint8(18)
memory usage: 4.6+ MB

```

```

In [5]: #Renaming columns from pd.get_dummies
df = df.rename({'Initial_admin_Elective Admission': 'Initial_admin_Elective_Admission',
               'Initial_admin_Emergency Admission': 'Initial_admin_Emergency_Admission',
               'Initial_admin_Observation Admission': 'Initial_admin_Observation_Admission',
               'Marital_Never Married': 'Marital_Never_Married',
               'Services_Blood Work': 'Services_Blood_Work',
               'Services_CT Scan': 'Services_CT_Scan'}, axis='columns')
df.info()

```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 10000 entries, 0 to 9999
```

```
Data columns (total 76 columns):
```

#	Column	Non-Null Count	Dtype
0	CaseOrder	10000 non-null	int64
1	Customer_id	10000 non-null	object
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3	UID	10000 non-null	object
4	City	10000 non-null	object
5	State	10000 non-null	object
6	County	10000 non-null	object
7	Zip	10000 non-null	int64
8	Lat	10000 non-null	float64
9	Lng	10000 non-null	float64
10	Population	10000 non-null	int64
11	Area	10000 non-null	object
12	TimeZone	10000 non-null	object
13	Job	10000 non-null	object
14	Children	10000 non-null	int64
15	Age	10000 non-null	int64
16	Income	10000 non-null	float64
17	ReAdmis	10000 non-null	object
18	VitD_levels	10000 non-null	float64
19	Doc_visits	10000 non-null	int64
20	Full_meals_eaten	10000 non-null	int64
21	vitD_supp	10000 non-null	int64
22	Soft_drink	10000 non-null	object
23	HighBlood	10000 non-null	object
24	Stroke	10000 non-null	object
25	Overweight	10000 non-null	object
26	Arthritis	10000 non-null	object
27	Diabetes	10000 non-null	object
28	Hyperlipidemia	10000 non-null	object
29	BackPain	10000 non-null	object
30	Anxiety	10000 non-null	object
31	Allergic_rhinitis	10000 non-null	object
32	Reflux_esophagitis	10000 non-null	object
33	Asthma	10000 non-null	object
34	Initial_days	10000 non-null	float64
35	TotalCharge	10000 non-null	float64
36	Additional_charges	10000 non-null	float64
37	Item1	10000 non-null	int64
38	Item2	10000 non-null	int64
39	Item3	10000 non-null	int64
40	Item4	10000 non-null	int64
41	Item5	10000 non-null	int64
42	Item6	10000 non-null	int64
43	Item7	10000 non-null	int64
44	Item8	10000 non-null	int64
45	ReAdmis_numeric	10000 non-null	int64
46	Soft_drink_numeric	10000 non-null	int64
47	HighBlood_numeric	10000 non-null	int64
48	Stroke_numeric	10000 non-null	int64
49	Arthritis_numeric	10000 non-null	int64
50	Diabetes_numeric	10000 non-null	int64
51	Hyperlipidemia_numeric	10000 non-null	int64
52	BackPain_numeric	10000 non-null	int64
53	Allergic_rhinitis_numeric	10000 non-null	int64
54	Reflux_esophagitis_numeric	10000 non-null	int64
55	Asthma_numeric	10000 non-null	int64
56	Overweight_numeric	10000 non-null	int64
57	Anxiety_numeric	10000 non-null	int64
58	Marital_Divorced	10000 non-null	uint8
59	Marital_Married	10000 non-null	uint8
60	Marital_Never_Married	10000 non-null	uint8
61	Marital_Separated	10000 non-null	uint8
62	Marital_Widowed	10000 non-null	uint8
63	Services_Blood_Work	10000 non-null	uint8
64	Services_CT_Scan	10000 non-null	uint8
65	Services_Intravenous	10000 non-null	uint8



```

66 Services_MRI 10000 non-null uint8
67 Gender_Female 10000 non-null uint8
68 Gender_Male 10000 non-null uint8
69 Gender_Nonbinary 10000 non-null uint8
70 Initial_admin_Elective_Admission 10000 non-null uint8
71 Initial_admin_Emergency_Admission 10000 non-null uint8
72 Initial_admin_Observation_Admission 10000 non-null uint8
73 Complication_risk_High 10000 non-null uint8
74 Complication_risk_Low 10000 non-null uint8
75 Complication_risk_Medium 10000 non-null uint8

```

dtypes: float64(7), int64(29), object(22), uint8(18)

memory usage: 4.6+ MB

```

In [6]: ##Univariate Stats Dataframe, and dropping unneeded and redundant columns
def unistats(df):
    output_df = pd.DataFrame(columns=['Count', 'Missing', 'Unique', 'Dtype', 'Numeric', 'Mean', 'Mode',

    for col in df:
        if pd.api.types.is_numeric_dtype(df[col]):
            output_df.loc[col] = [df[col].count(), df[col].isnull().sum(), df[col].nunique(), df[col].dt
        else:
            output_df.loc[col] = [df[col].count(), df[col].isnull().sum(), df[col].nunique(), df[col].dt
    return output_df.sort_values(by=['Numeric', 'Skew', 'Unique'], ascending=False)

df.drop(columns=['CaseOrder', 'Customer_id', 'Interaction', 'UID', 'State', 'County', 'Job', 'Zip', 'Tim
print(unistats(df))

```

	Count	Missing	Unique	Dtype	Numeric	\
Gender_Nonbinary	10000	0	2	uint8	True	
Services_MRI	10000	0	2	uint8	True	
Services_CT_Scan	10000	0	2	uint8	True	
Population	10000	0	5951	int64	True	
vitD_supp	10000	0	6	int64	True	
Marital_Never_Married	10000	0	2	uint8	True	
Marital_Separated	10000	0	2	uint8	True	
Stroke_numeric	10000	0	2	int64	True	
Marital_Married	10000	0	2	uint8	True	
Marital_Widowed	10000	0	2	uint8	True	
Children	10000	0	11	int64	True	
Income	10000	0	9993	float64	True	
Complication_risk_Low	10000	0	2	uint8	True	
Initial_admin_Observation_Admission	10000	0	2	uint8	True	
Initial_admin_Elective_Admission	10000	0	2	uint8	True	
Soft_drink_numeric	10000	0	2	int64	True	
Diabetes_numeric	10000	0	2	int64	True	
Full_meals_eaten	10000	0	8	int64	True	
Asthma_numeric	10000	0	2	int64	True	
Additional_charges	10000	0	9418	float64	True	
Services_Intravenous	10000	0	2	uint8	True	
Anxiety_numeric	10000	0	2	int64	True	
Complication_risk_High	10000	0	2	uint8	True	
Hyperlipidemia_numeric	10000	0	2	int64	True	
Arthritis_numeric	10000	0	2	int64	True	
ReAdmis_numeric	10000	0	2	int64	True	
Allergic_rhinitis_numeric	10000	0	2	int64	True	
HighBlood_numeric	10000	0	2	int64	True	
BackPain_numeric	10000	0	2	int64	True	
Reflux_esophagitis_numeric	10000	0	2	int64	True	
Complication_risk_Medium	10000	0	2	uint8	True	
Gender_Male	10000	0	2	uint8	True	
Initial_days	10000	0	9997	float64	True	
TotalCharge	10000	0	9997	float64	True	
VitD_levels	10000	0	9976	float64	True	
Age	10000	0	72	int64	True	
Doc_visits	10000	0	9	int64	True	
Initial_admin_Emergency_Admission	10000	0	2	uint8	True	
Services_Blood_Work	10000	0	2	uint8	True	
Overweight_numeric	10000	0	2	int64	True	

	Mean	Mode	Min	\
Gender_Nonbinary	0.021400	0.000000	0.000000	
Services_MRI	0.038000	0.000000	0.000000	
Services_CT_Scan	0.122500	0.000000	0.000000	
Population	9965.253800	0.000000	0.000000	
vitD_supp	0.398900	0.000000	0.000000	
Marital_Never_Married	0.198400	0.000000	0.000000	
Marital_Separated	0.198700	0.000000	0.000000	
Stroke_numeric	0.199300	0.000000	0.000000	
Marital_Married	0.202300	0.000000	0.000000	
Marital_Widowed	0.204500	0.000000	0.000000	
Children	2.097200	0.000000	0.000000	
Income	40490.495160	14572.40000	154.080000	
Complication_risk_Low	0.212500	0.000000	0.000000	
Initial_admin_Observation_Admission	0.243600	0.000000	0.000000	
Initial_admin_Elective_Admission	0.250400	0.000000	0.000000	
Soft_drink_numeric	0.257500	0.000000	0.000000	
Diabetes_numeric	0.273800	0.000000	0.000000	
Full_meals_eaten	1.001400	0.000000	0.000000	
Asthma_numeric	0.289300	0.000000	0.000000	
Additional_charges	12934.528587	3883.66416	3125.703000	
Services_Intravenous	0.313000	0.000000	0.000000	
Anxiety_numeric	0.321500	0.000000	0.000000	
Complication_risk_High	0.335800	0.000000	0.000000	
Hyperlipidemia_numeric	0.337200	0.000000	0.000000	
Arthritis_numeric	0.357400	0.000000	0.000000	
ReAdmis_numeric	0.366900	0.000000	0.000000	
Allergic_rhinitis_numeric	0.394100	0.000000	0.000000	
HighBlood_numeric	0.409000	0.000000	0.000000	

BackPain_numeric	0.411400	0.00000	0.000000
Reflux_esophagitis_numeric	0.413500	0.00000	0.000000
Complication_risk_Medium	0.451700	0.00000	0.000000
Gender_Male	0.476800	0.00000	0.000000
Initial_days	34.455299	63.54432	1.001981
TotalCharge	5312.172769	7555.45200	1938.312067
VitD_levels	17.964262	15.26009	9.806483
Age	53.511700	47.00000	18.000000
Doc_visits	5.012200	5.00000	1.000000
Initial_admin_Emergency_Admission	0.506000	1.00000	0.000000
Services_Blood_Work	0.526500	1.00000	0.000000
Overweight_numeric	0.709400	1.00000	0.000000

	Median	Max	\
Gender_Nonbinary	0.000000	1.000000	
Services_MRI	0.000000	1.000000	
Services_CT_Scan	0.000000	1.000000	
Population	2769.000000	122814.000000	
vitD_supp	0.000000	5.000000	
Marital_Never_Married	0.000000	1.000000	
Marital_Separated	0.000000	1.000000	
Stroke_numeric	0.000000	1.000000	
Marital_Married	0.000000	1.000000	
Marital_Widowed	0.000000	1.000000	
Children	1.000000	10.000000	
Income	33768.420000	207249.100000	
Complication_risk_Low	0.000000	1.000000	
Initial_admin_Observation_Admission	0.000000	1.000000	
Initial_admin_Elective_Admission	0.000000	1.000000	
Soft_drink_numeric	0.000000	1.000000	
Diabetes_numeric	0.000000	1.000000	
Full_meals_eaten	1.000000	7.000000	
Asthma_numeric	0.000000	1.000000	
Additional_charges	11573.977735	30566.070000	
Services_Intravenous	0.000000	1.000000	
Anxiety_numeric	0.000000	1.000000	
Complication_risk_High	0.000000	1.000000	
Hyperlipidemia_numeric	0.000000	1.000000	
Arthritis_numeric	0.000000	1.000000	
ReAdmis_numeric	0.000000	1.000000	
Allergic_rhinitis_numeric	0.000000	1.000000	
HighBlood_numeric	0.000000	1.000000	
BackPain_numeric	0.000000	1.000000	
Reflux_esophagitis_numeric	0.000000	1.000000	
Complication_risk_Medium	0.000000	1.000000	
Gender_Male	0.000000	1.000000	
Initial_days	35.836244	71.981490	
TotalCharge	5213.952000	9180.728000	
VitD_levels	17.951122	26.394449	
Age	53.000000	89.000000	
Doc_visits	5.000000	9.000000	
Initial_admin_Emergency_Admission	1.000000	1.000000	
Services_Blood_Work	1.000000	1.000000	
Overweight_numeric	1.000000	1.000000	

	Std	Skew	Kurt
Gender_Nonbinary	0.144721	6.615434	41.772323
Services_MRI	0.191206	4.833456	21.366572
Services_CT_Scan	0.327879	2.303141	3.305119
Population	14824.758614	2.229959	5.880913
vitD_supp	0.628505	1.550205	2.330763
Marital_Never_Married	0.398815	1.512784	0.288572
Marital_Separated	0.399042	1.510420	0.281425
Stroke_numeric	0.399494	1.505705	0.267202
Marital_Married	0.401735	1.482369	0.197456
Marital_Widowed	0.403356	1.465500	0.147720
Children	2.163659	1.448013	2.076321
Income	28521.153293	1.405899	2.745690
Complication_risk_Low	0.409097	1.405815	-0.023688
Initial_admin_Observation_Admission	0.429276	1.194810	-0.572544
Initial_admin_Elective_Admission	0.433265	1.152412	-0.672081

Soft_drink_numeric	0.437279	1.109354	-0.769488
Diabetes_numeric	0.445930	1.014712	-0.970553
Full_meals_eaten	1.008117	1.009461	1.042727
Asthma_numeric	0.453460	0.929485	-1.136285
Additional_charges	6542.601544	0.831842	-0.142684
Services_Intravenous	0.463738	0.806652	-1.349583
Anxiety_numeric	0.467076	0.764483	-1.415849
Complication_risk_High	0.472293	0.695470	-1.516625
Hyperlipidemia_numeric	0.472777	0.688834	-1.525813
Arthritis_numeric	0.479258	0.595206	-1.646059
ReAdmis_numeric	0.481983	0.552412	-1.695180
Allergic_rhinitis_numeric	0.488681	0.433498	-1.812442
HighBlood_numeric	0.491674	0.370238	-1.863296
BackPain_numeric	0.492112	0.360153	-1.870664
Reflux_esophagitis_numeric	0.492486	0.351350	-1.876929
Complication_risk_Medium	0.497687	0.194137	-1.962703
Gender_Male	0.499486	0.092914	-1.991765
Initial_days	26.309341	0.070286	-1.754525
TotalCharge	2180.393838	0.069661	-1.668267
VitD_levels	2.017231	0.032435	-0.022112
Age	20.638538	0.005117	-1.189527
Doc_visits	1.045734	-0.018563	0.025999
Initial_admin_Emergency_Admission	0.499989	-0.024005	-1.999824
Services_Blood_Work	0.499322	-0.106165	-1.989127
Overweight_numeric	0.454062	-0.922526	-1.149176

```
In [8]: #Univariate Visualizations
plt.hist(df.ReAdmis_numeric)
plt.xlabel('Was the patient readmitted?')
plt.ylabel('Number of Patients')
plt.title('ReAdmissions')
plt.show()

plt.hist(df.Initial_days)
plt.xlabel('Total Days')
plt.ylabel('Number of Patients')
plt.title('Initial_days')
plt.show()

plt.hist(df.vitD_supp)
plt.xlabel('# of Vit D Administered')
plt.ylabel('Number of Patients')
plt.title('vitd_supp')
plt.show()

plt.hist(df.Children)
plt.xlabel('# of Children')
plt.ylabel('Number of Patients')
plt.title('# of children')
plt.show()

plt.hist(df.Income)
plt.xlabel('Yearly Income')
plt.ylabel('Number of Patients')
plt.title('Yearly Income')
plt.show()

plt.hist(df.Full_meals_eaten)
plt.xlabel('Full_meals_eaten')
plt.ylabel('Number of Patients')
plt.title('Full Meals Eaten')
plt.show()

plt.hist(df.Additional_charges)
plt.xlabel('Additional Charges')
plt.ylabel('Number of Patients')
plt.title('Additional Charges')
plt.show()

plt.hist(df.TotalCharge)
plt.xlabel('Total Charges')
```

```
plt.ylabel('Number of Patients')
plt.title('Total Charges')
plt.show()

plt.hist(df.VitD_levels)
plt.xlabel('VitD Levels')
plt.ylabel('Number of Patients')
plt.title('VitD Levels')
plt.show()

plt.hist(df.Age)
plt.xlabel('Age')
plt.ylabel('Number of Patients')
plt.title('Age')
plt.show()

plt.hist(df.Doc_visits)
plt.xlabel('Doctor Visits')
plt.ylabel('Number of Patients')
plt.title('Doctor Visits')
plt.show()

plt.hist(df.HighBlood_numeric)
plt.xlabel('Does patient have high blood pressure?')
plt.ylabel('Number of Patients')
plt.title('High Blood Pressure')
plt.show()

plt.hist(df.Stroke_numeric)
plt.xlabel('Does patient have history of strokes?')
plt.ylabel('Number of Patients')
plt.title('Stroke')
plt.show()

plt.hist(df.Arthritis_numeric)
plt.xlabel('Does patient have history of Arthritis?')
plt.ylabel('Number of Patients')
plt.title('Arthritis')
plt.show()

plt.hist(df.Diabetes_numeric)
plt.xlabel('Does patient have history of Diabetes?')
plt.ylabel('Number of Patients')
plt.title('Diabetes')
plt.show()

plt.hist(df.Hyperlipidemia_numeric)
plt.xlabel('Does patient have Hyperlipidemia?')
plt.ylabel('Number of Patients')
plt.title('Hyperlipidemia')
plt.show()

plt.hist(df.BackPain_numeric)
plt.xlabel('Does patient have BackPain?')
plt.ylabel('Number of Patients')
plt.title('BackPain')
plt.show()

plt.hist(df.Allergic_rhinitis_numeric)
plt.xlabel('Does patient have Allergic_rhinitis?')
plt.ylabel('Number of Patients')
plt.title('Allergic_rhinitis')
plt.show()

plt.hist(df.Reflux_esophagitis_numeric)
plt.xlabel('Does patient have Reflux_esophagitis?')
plt.ylabel('Number of Patients')
plt.title('Reflux_esophagitis')
plt.show()

plt.hist(df.Asthma_numeric)
```

```
plt.xlabel('Does patient have Asthma?')
plt.ylabel('Number of Patients')
plt.title('Asthma')
plt.show()

plt.hist(df.Marital_Married)
plt.xlabel('Patients Marital Status')
plt.ylabel('Number of Patients')
plt.title('Marital_Married')
plt.show()

plt.hist(df.Marital_Never_Married)
plt.xlabel('Patients Marital Status')
plt.ylabel('Number of Patients')
plt.title('Marital_Never_Married')
plt.show()

plt.hist(df.Marital_Separated)
plt.xlabel('Patients Marital Status')
plt.ylabel('Number of Patients')
plt.title('Marital_Separated')
plt.show()

plt.hist(df.Marital_Widowed)
plt.xlabel('Patients Marital Status')
plt.ylabel('Number of Patients')
plt.title('Marital_Widowed')
plt.show()

plt.hist(df.Services_Blood_Work)
plt.xlabel('What services did the patient receive?')
plt.ylabel('Number of Patients')
plt.title('Services_Blood_Work')
plt.show()

plt.hist(df.Services_CT_Scan)
plt.xlabel('What services did the patient receive?')
plt.ylabel('Number of Patients')
plt.title('Services_CT_Scan')
plt.show()

plt.hist(df.Services_Intravenous)
plt.xlabel('What services did the patient receive?')
plt.ylabel('Number of Patients')
plt.title('Services_Intravenous')
plt.show()

plt.hist(df.Services_MRI)
plt.xlabel('What services did the patient receive?')
plt.ylabel('Number of Patients')
plt.title('Services_MRI')
plt.show()

plt.hist(df.Gender_Male)
plt.xlabel('What gender does the patient identify as?')
plt.ylabel('Number of Patients')
plt.title('Gender_Male')
plt.show()

plt.hist(df.Gender_Nonbinary)
plt.xlabel('What gender does the patient identify as?')
plt.ylabel('Number of Patients')
plt.title('Gender_Nonbinary')
plt.show()

plt.hist(df.Initial_admin_Elective_Admission)
plt.xlabel('What brought the patient into the hospital?')
plt.ylabel('Number of Patients')
plt.title('Initial_admin_Elective_Admission')
plt.show()
```

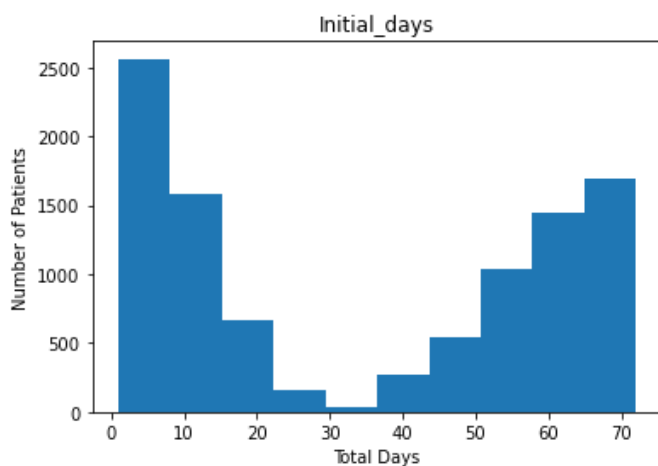
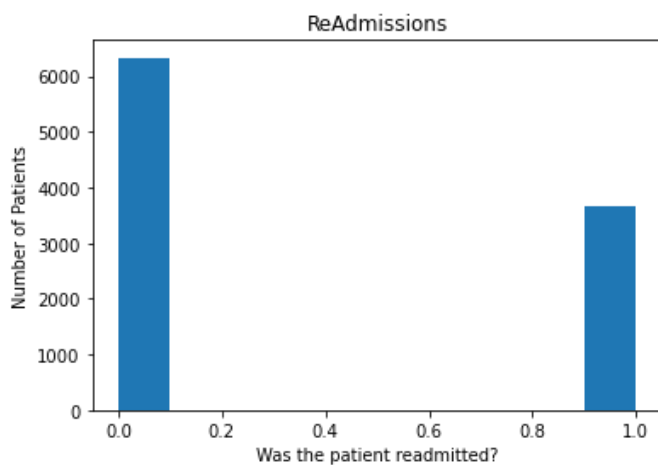
```
plt.hist(df.Initial_admin_Emergency_Admission)
plt.xlabel('What brought the patient into the hospital?')
plt.ylabel('Number of Patients')
plt.title('Initial_admin_Emergency_Admission')
plt.show()

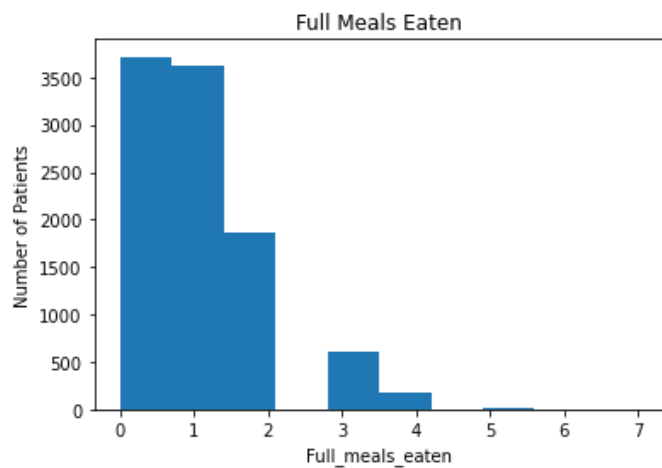
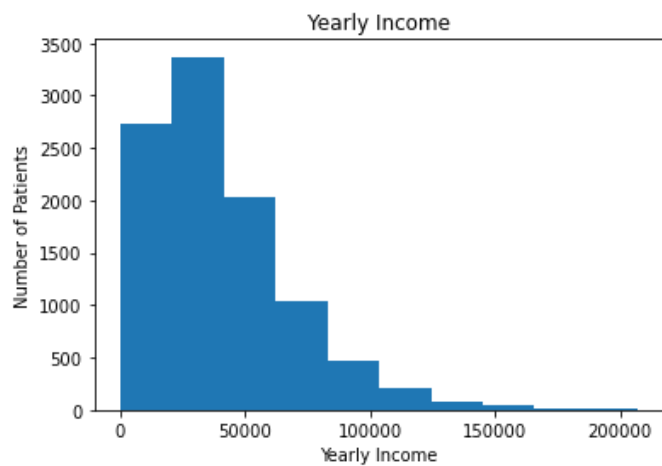
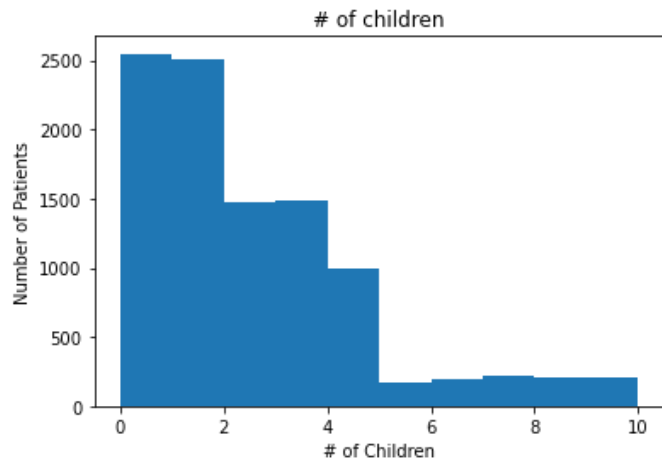
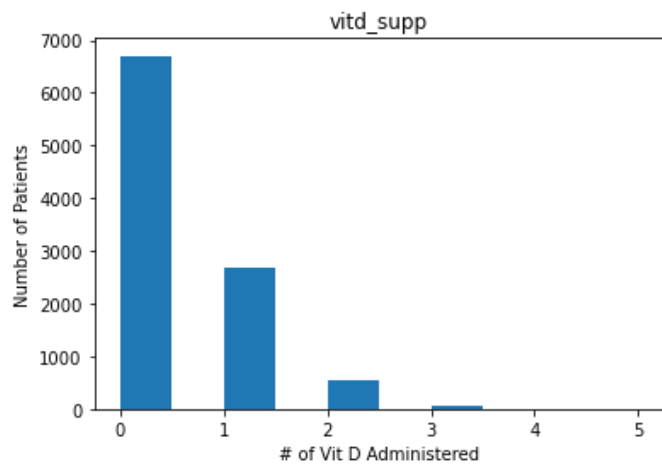
plt.hist(df.Initial_admin_Observation_Admission)
plt.xlabel('What brought the patient into the hospital?')
plt.ylabel('Number of Patients')
plt.title('Initial_admin_Observation_Admission')
plt.show()

plt.hist(df.Complication_risk_High)
plt.xlabel('What is the patients complication risk?')
plt.ylabel('Number of Patients')
plt.title('Complication_risk_High')
plt.show()

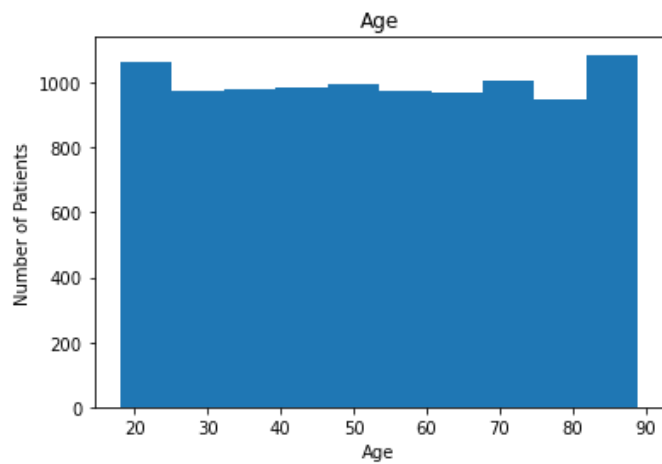
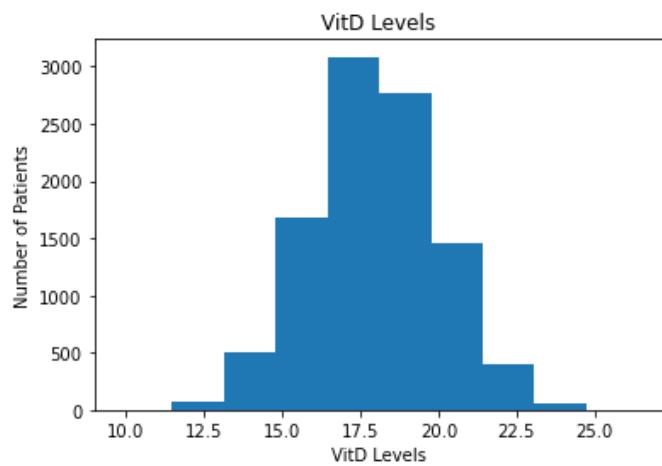
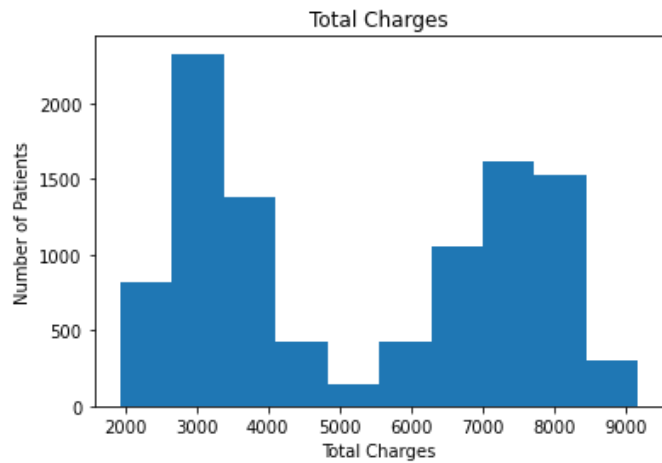
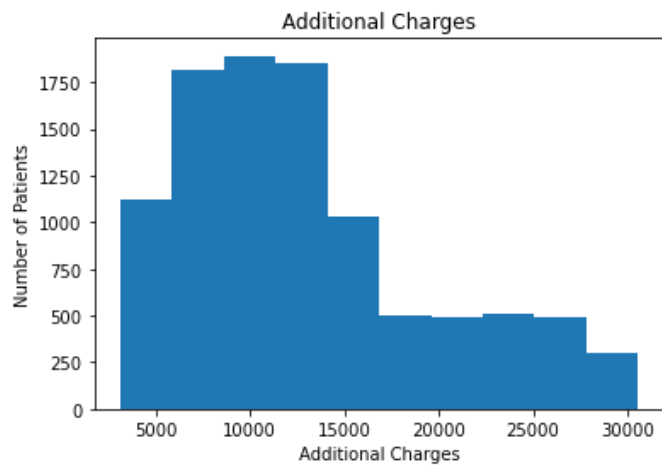
plt.hist(df.Complication_risk_Low)
plt.xlabel('What is the patients complication risk?')
plt.ylabel('Number of Patients')
plt.title('Complication_risk_Low')
plt.show()

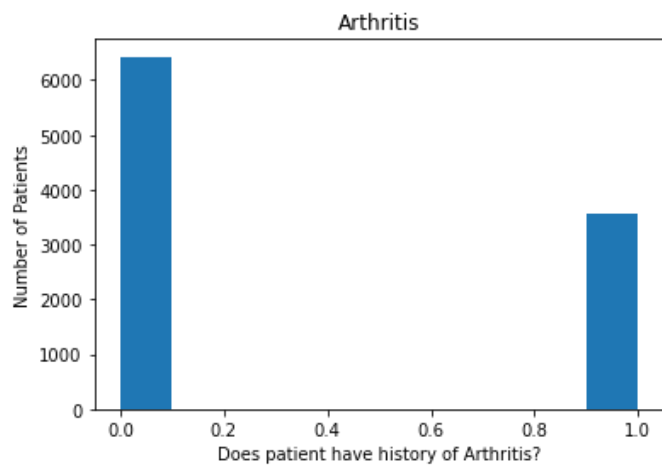
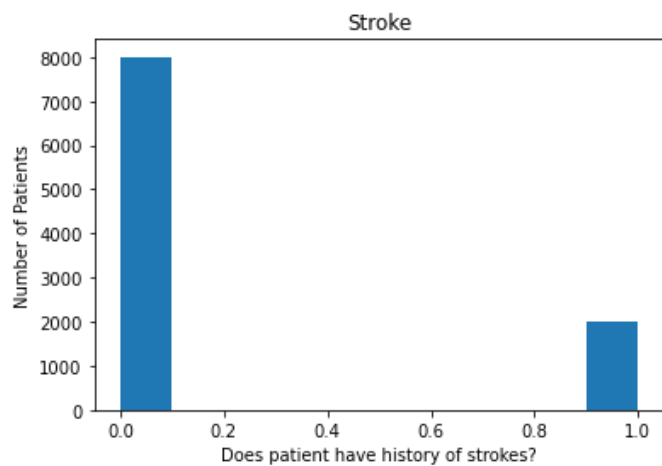
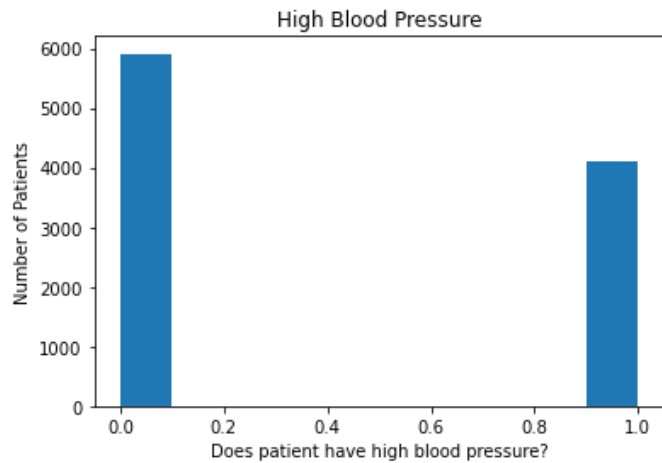
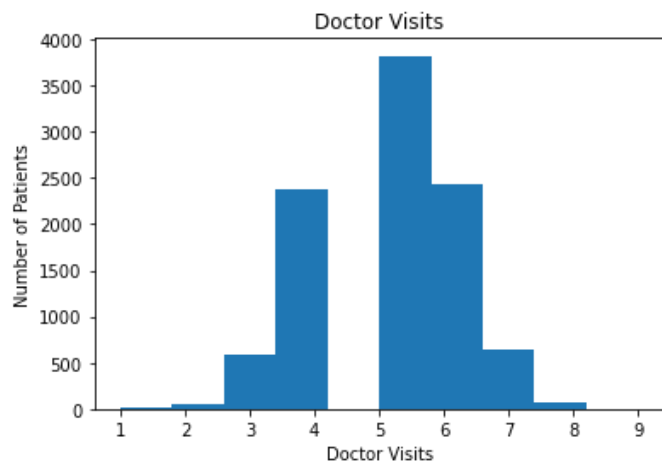
plt.hist(df.Complication_risk_Medium)
plt.xlabel('What is the patients complication risk?')
plt.ylabel('Number of Patients')
plt.title('Complication_risk_Medium')
plt.show()
```

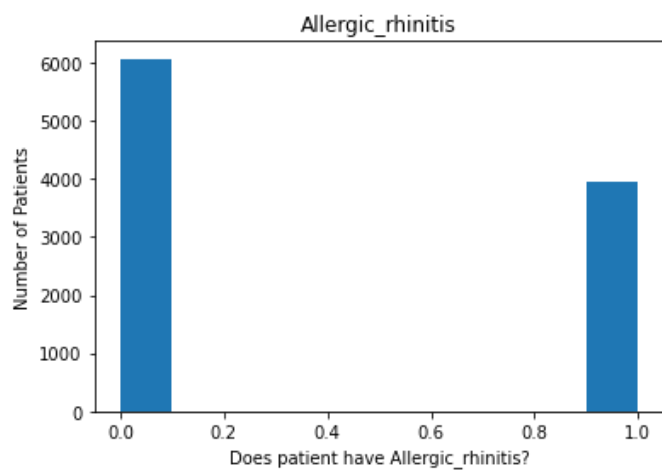
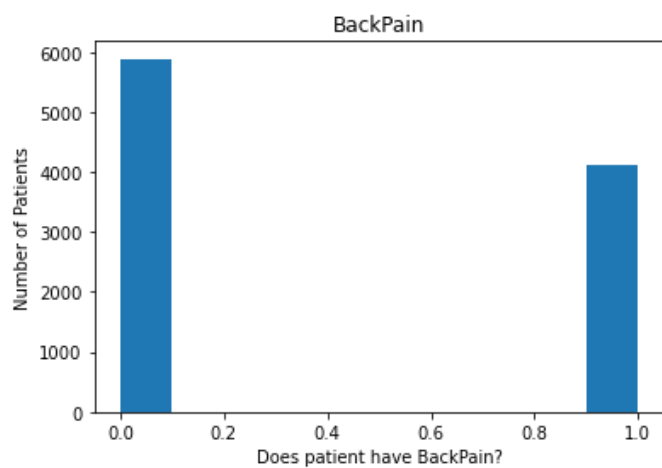
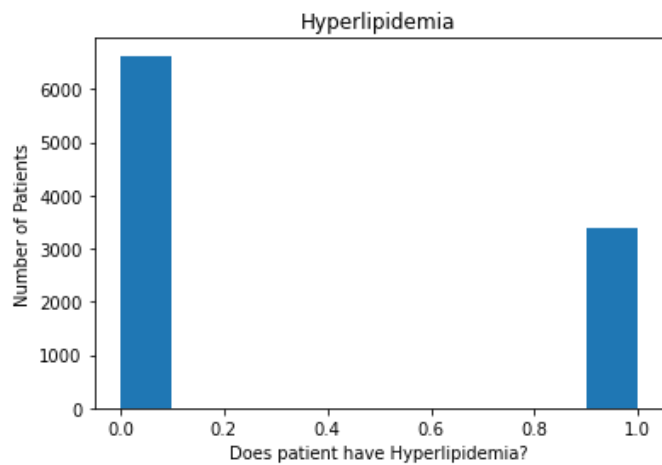
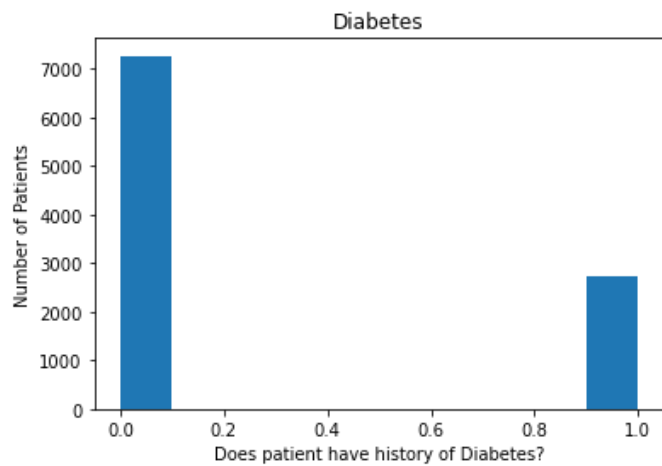


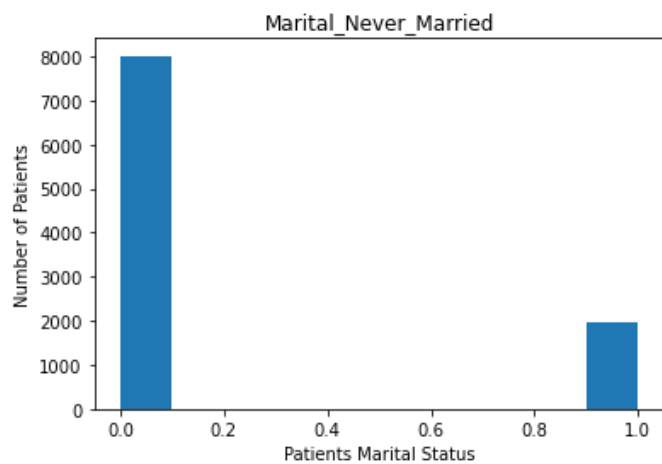
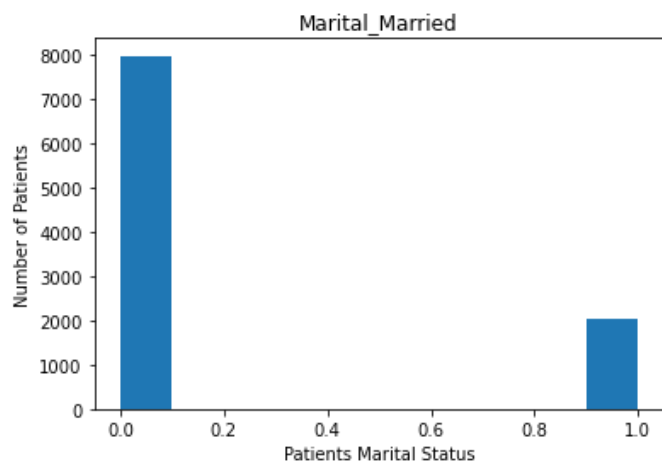
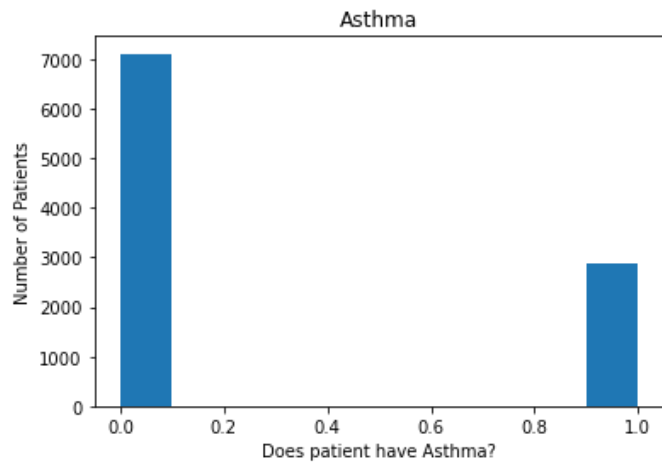
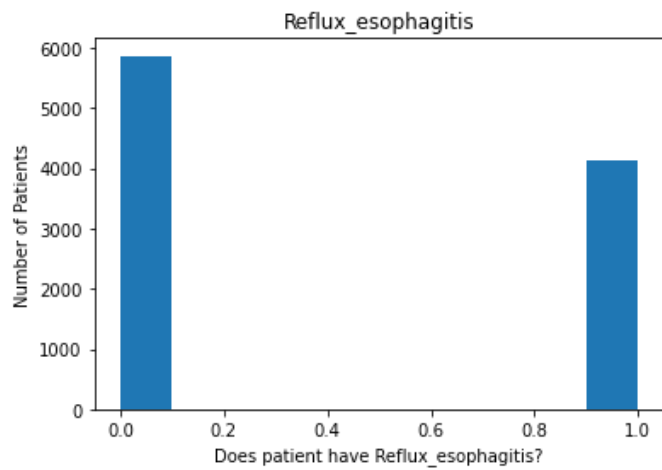


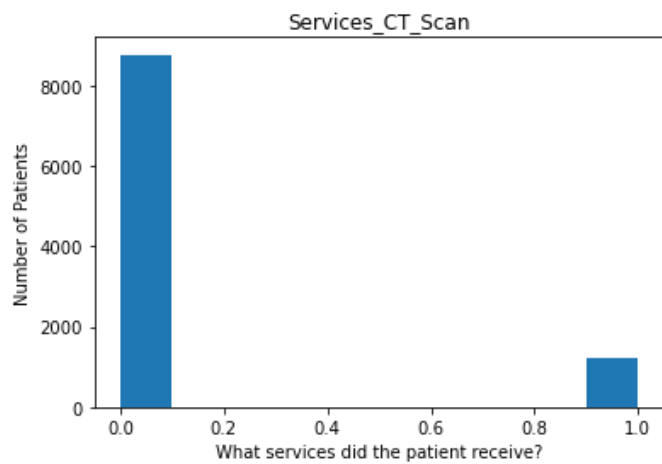
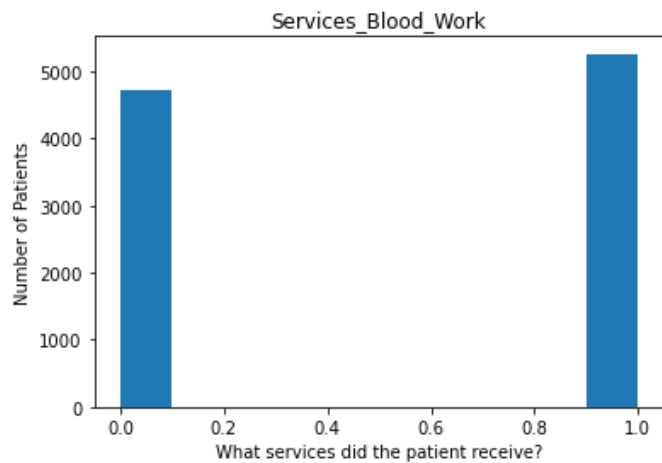
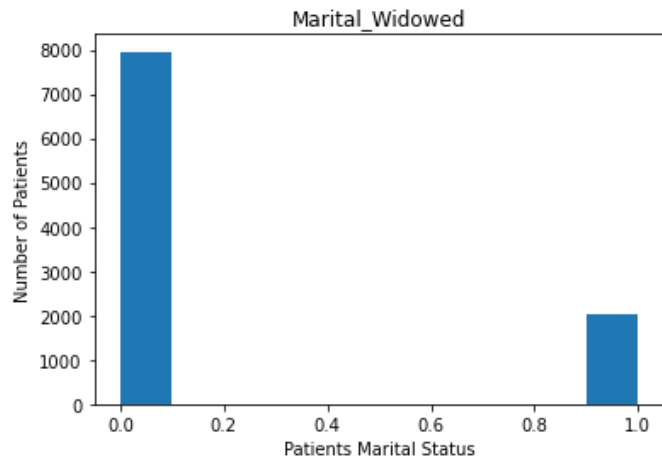
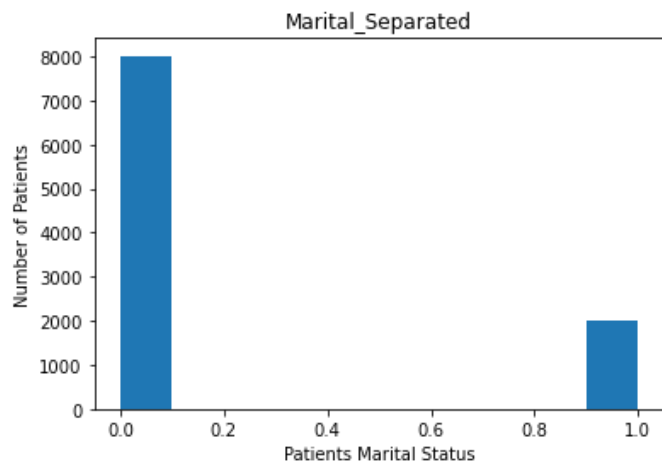


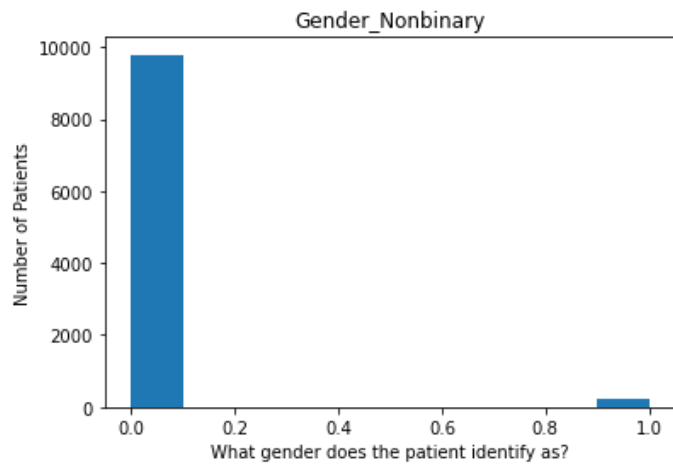
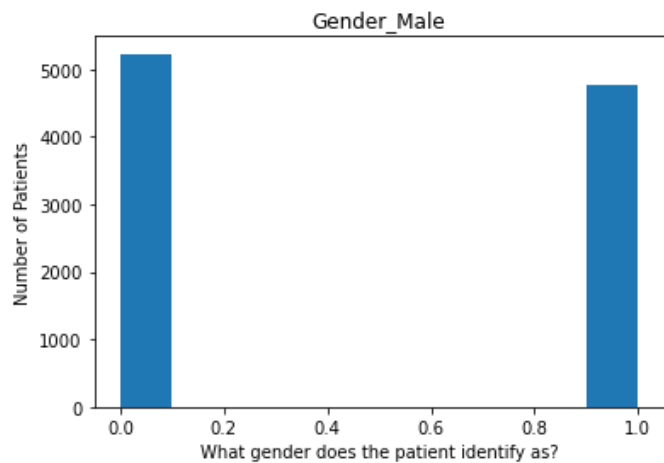
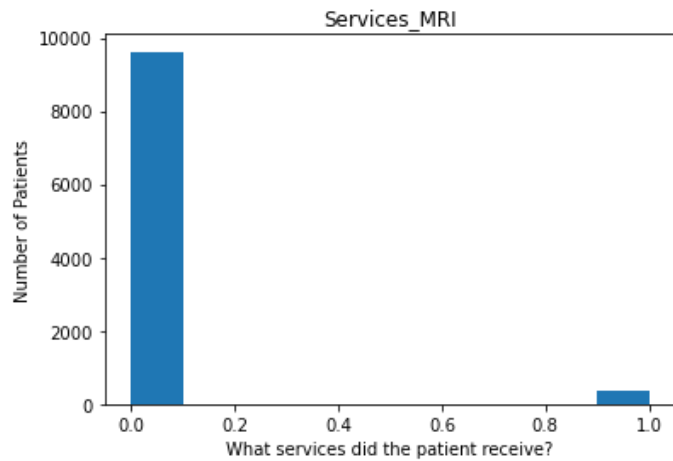
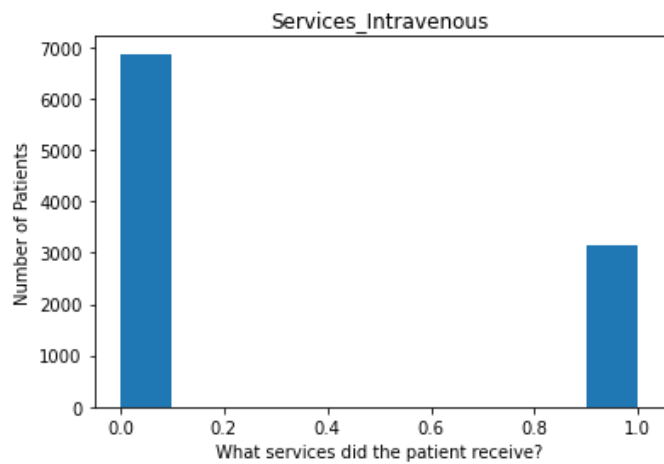


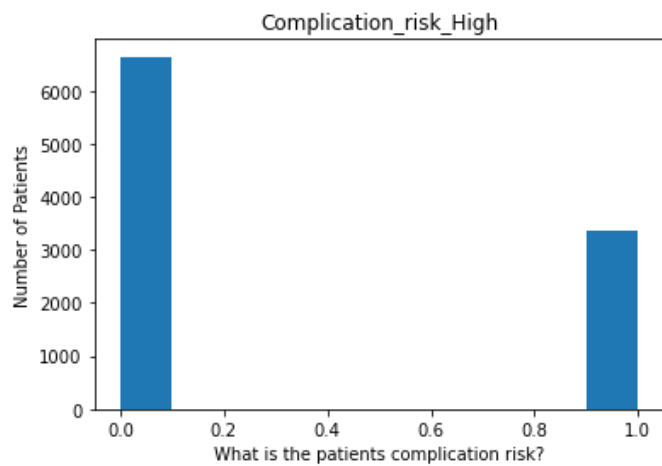
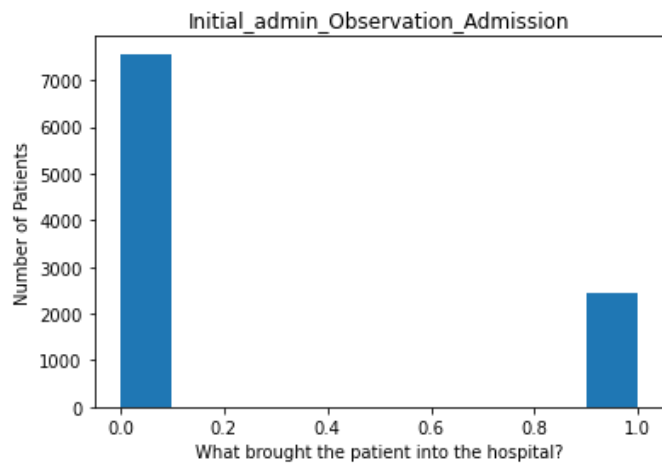
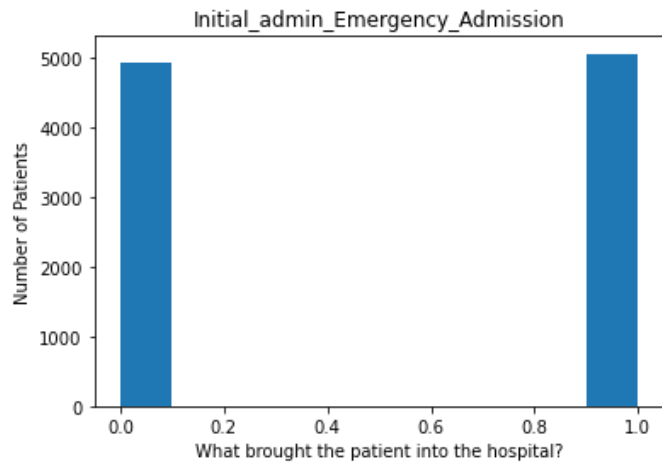
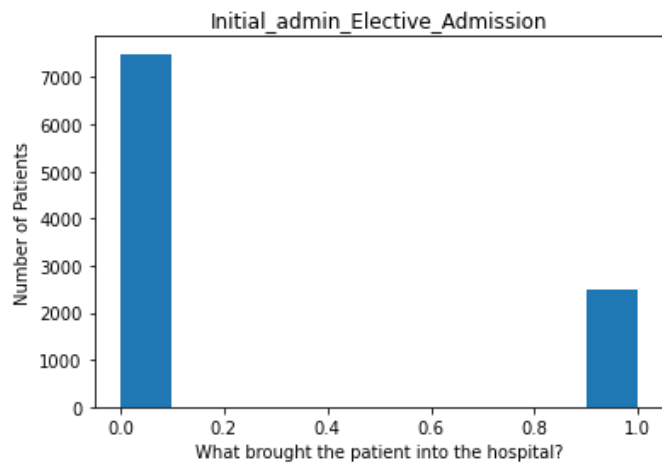


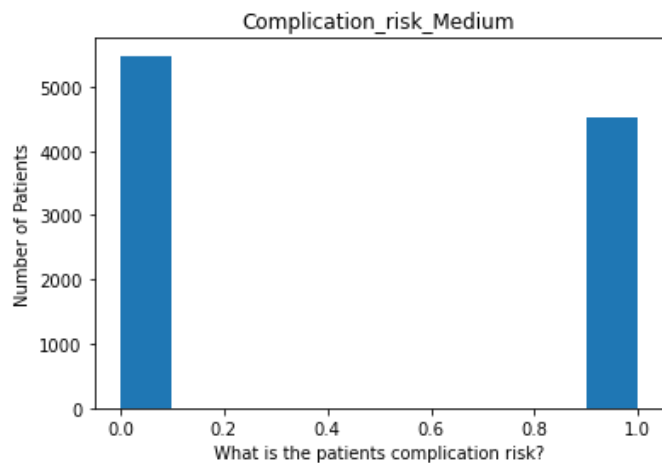
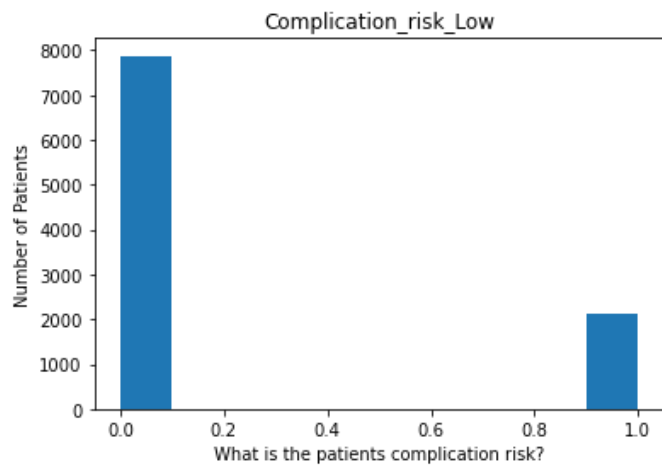












```
In [9]: #Bivariate Visualizations
sns.scatterplot(data=df, y="Initial_days", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="vitD_supp", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Children", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Income", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Full_meals_eaten", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Additional_charges", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="TotalCharge", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="VitD_levels", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Age", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Doc_visits", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="HighBlood_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Stroke_numeric", x="ReAdmis_numeric")
```



```
plt.show()

sns.scatterplot(data=df, y="Arthritis_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Diabetes_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Hyperlipidemia_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="BackPain_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Allergic_rhinitis_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Reflux_esophagitis_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Asthma_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Overweight_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Anxiety_numeric", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Marital_Married", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Marital_Never_Married", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Marital_Separated", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Marital_Widowed", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Services_Blood_Work", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Services_CT_Scan", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Services_Intravenous", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Services_MRI", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Gender_Male", x="ReAdmis_numeric")
plt.show()

sns.scatterplot(data=df, y="Gender_Nonbinary", x="ReAdmis_numeric")
plt.show()

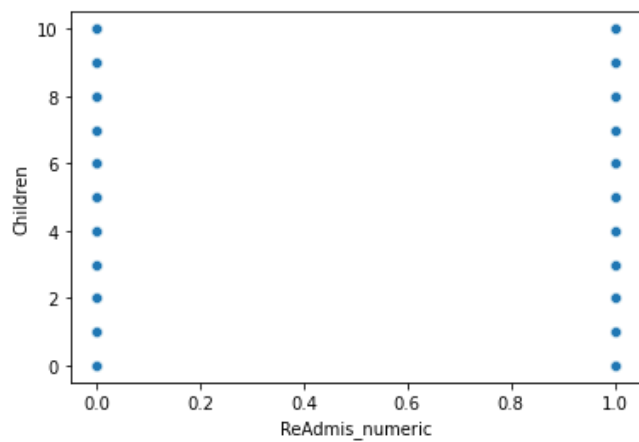
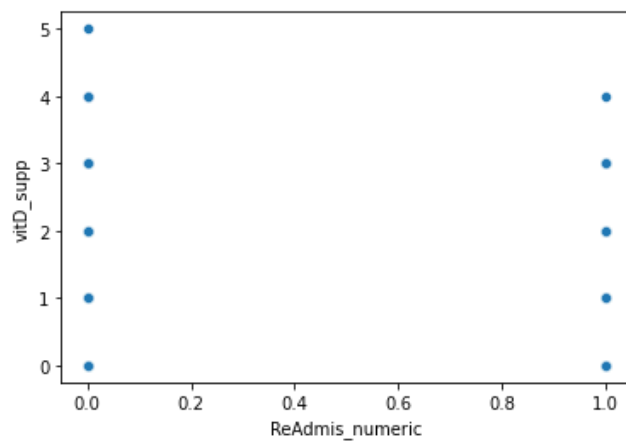
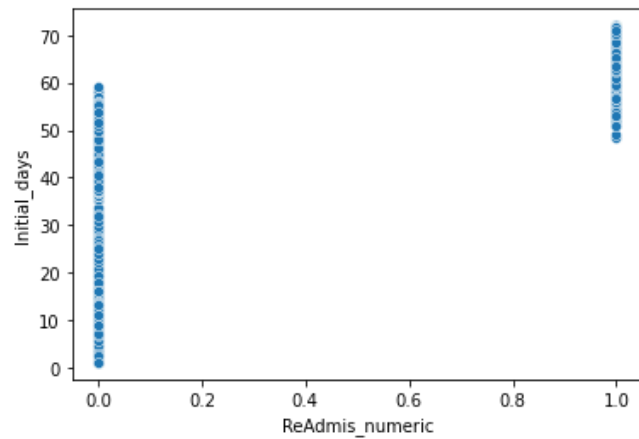
sns.scatterplot(data=df, y="Initial_admin_Elective_Admission", x="ReAdmis_numeric")
plt.show()

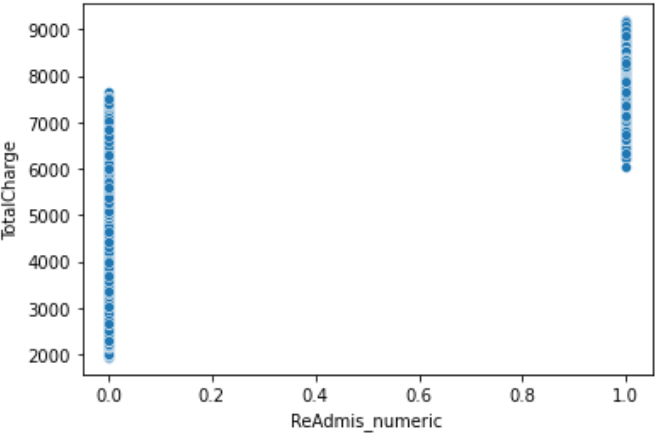
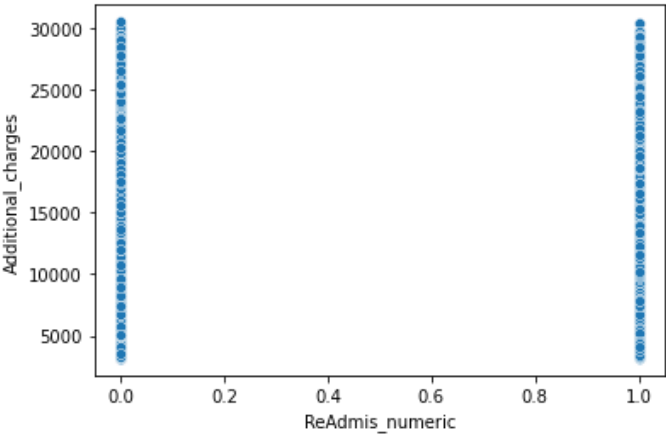
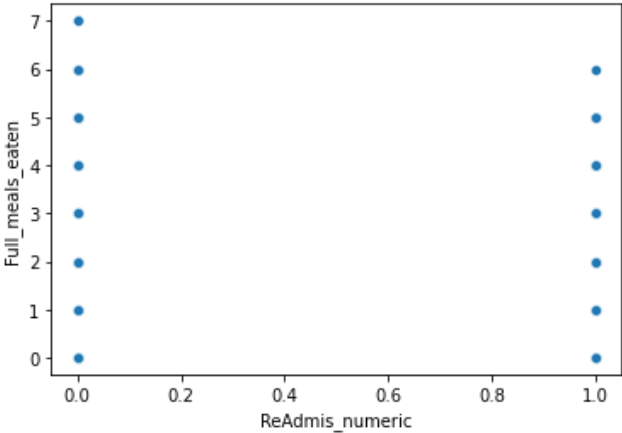
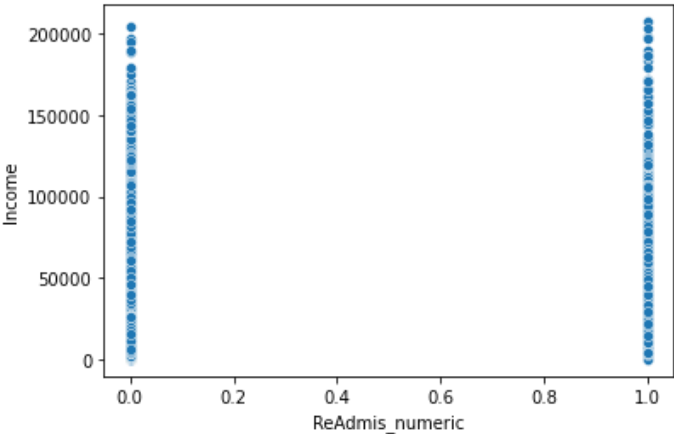
sns.scatterplot(data=df, y="Initial_admin_Emergency_Admission", x="ReAdmis_numeric")
plt.show()

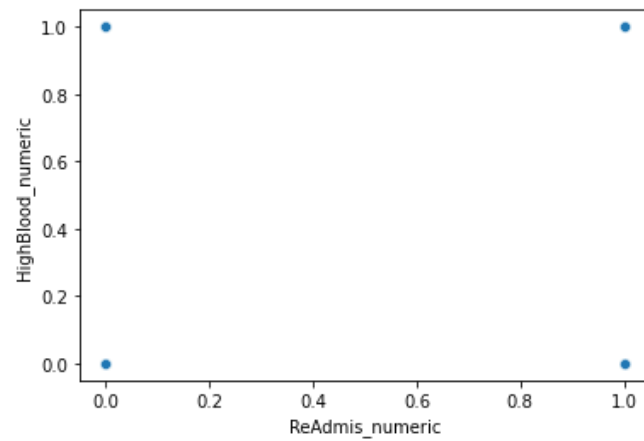
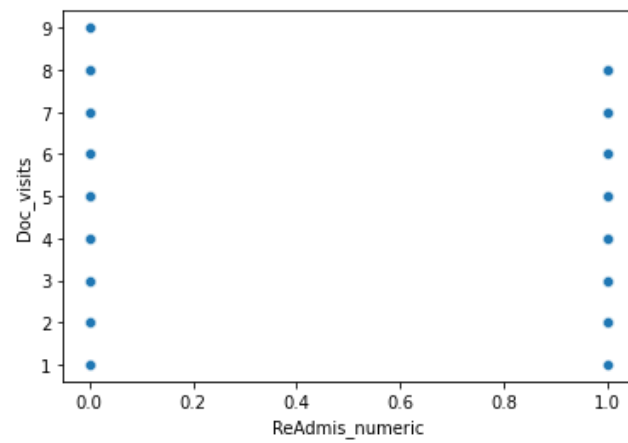
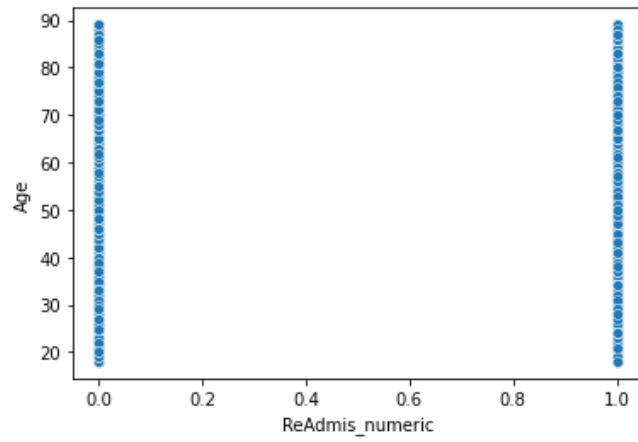
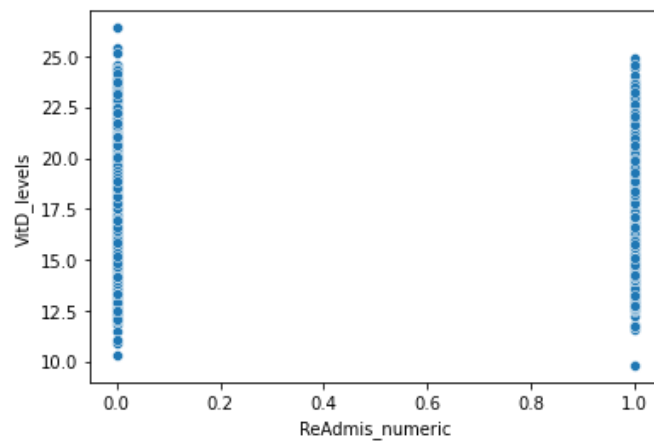
sns.scatterplot(data=df, y="Initial_admin_Observation_Admission", x="ReAdmis_numeric")
plt.show()

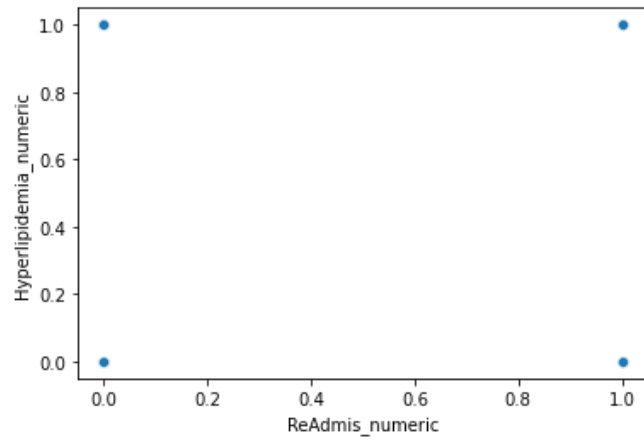
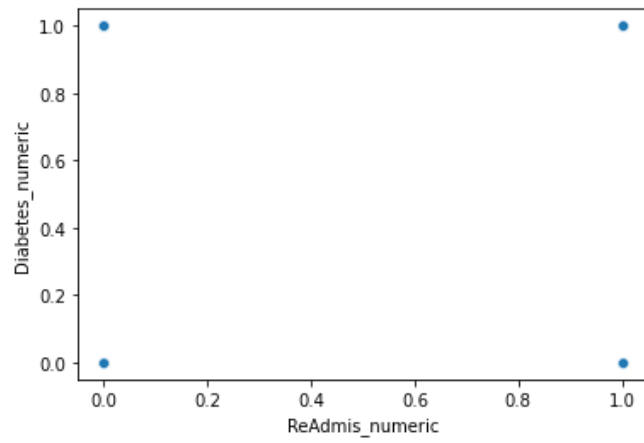
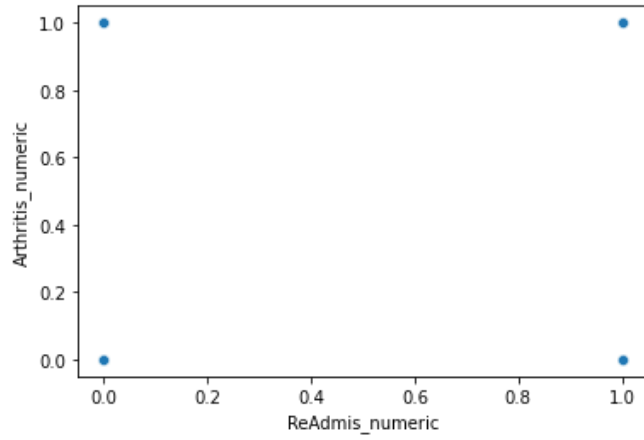
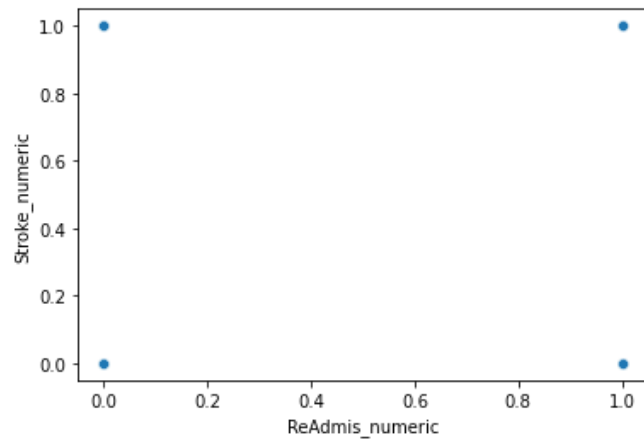
sns.scatterplot(data=df, y="Complication_risk_High", x="ReAdmis_numeric")
plt.show()
```

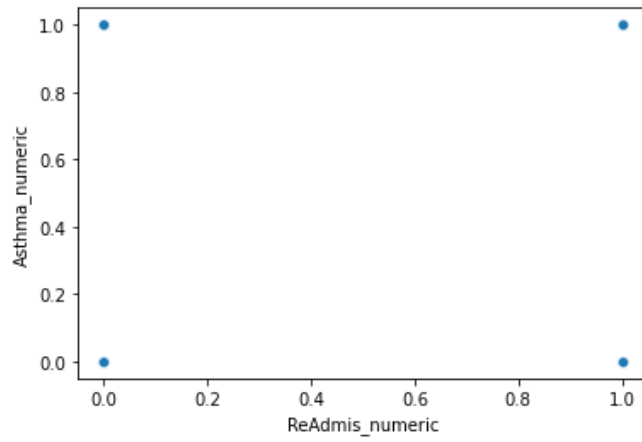
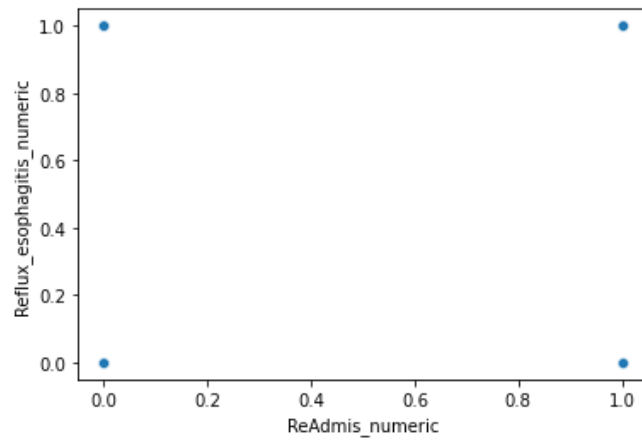
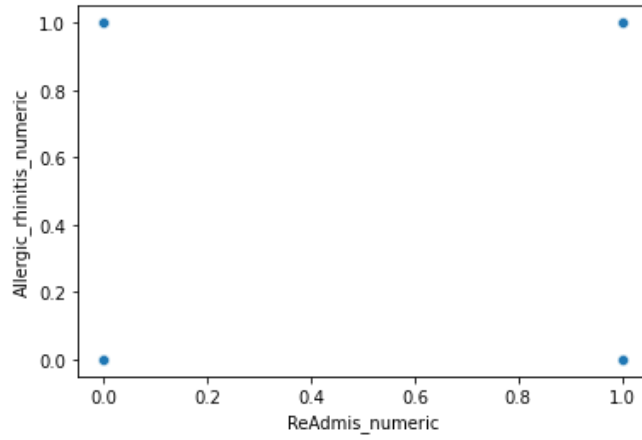
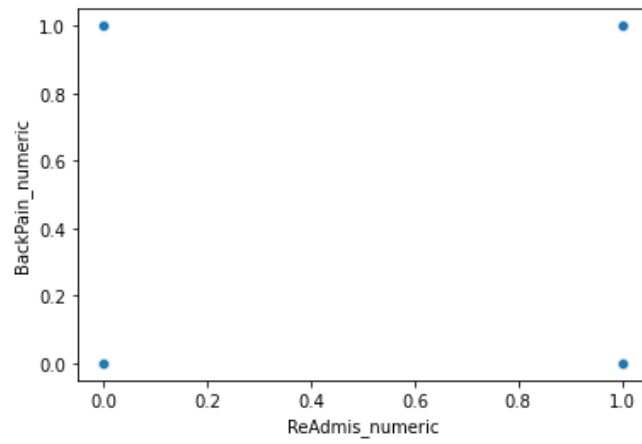
```
sns.scatterplot(data=df, y="Complication_risk_Medium", x="ReAdmis_numeric")  
plt.show()  
  
sns.scatterplot(data=df, y="Complication_risk_Low", x="ReAdmis_numeric")  
plt.show()
```

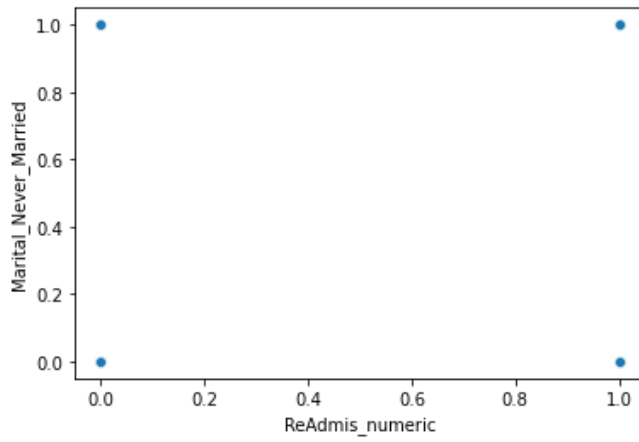
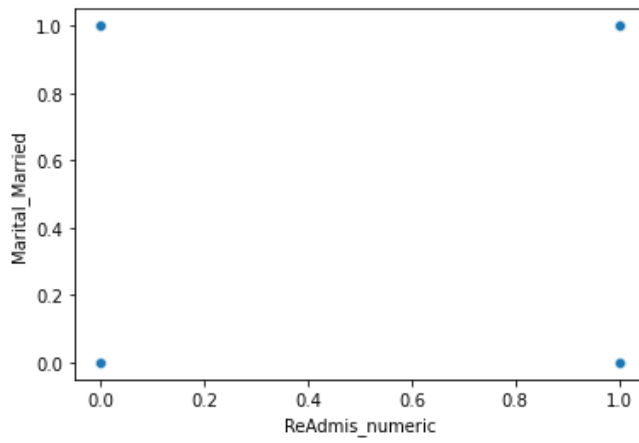
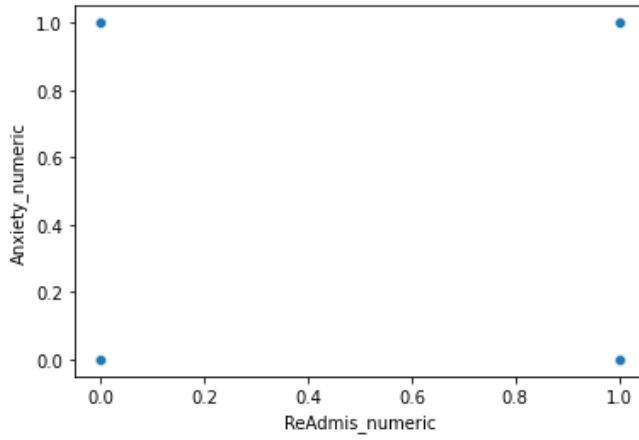
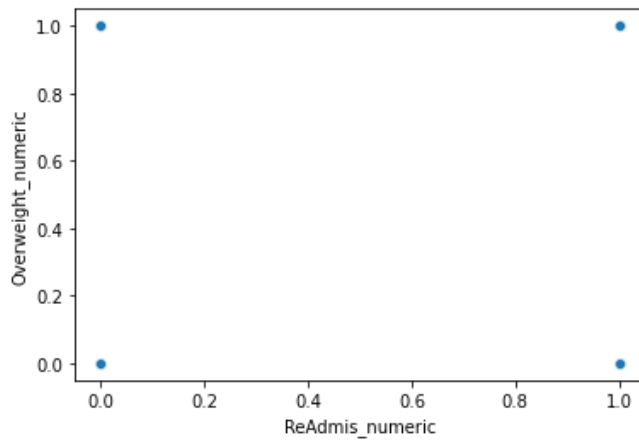


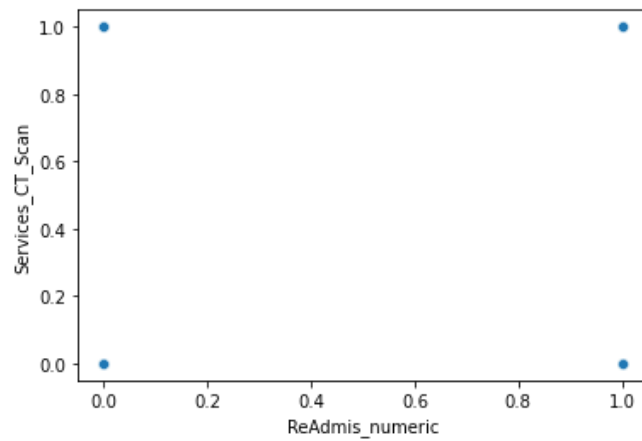
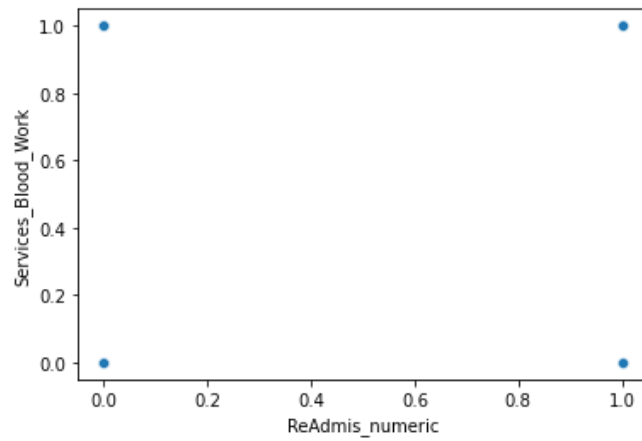
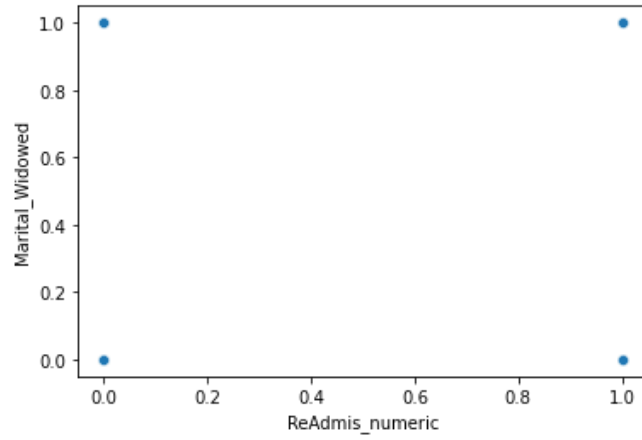
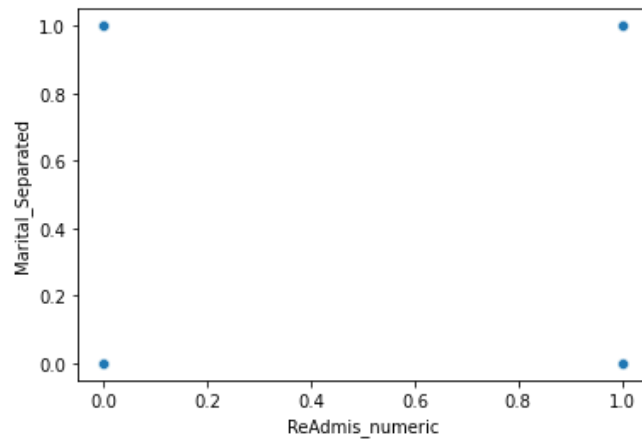




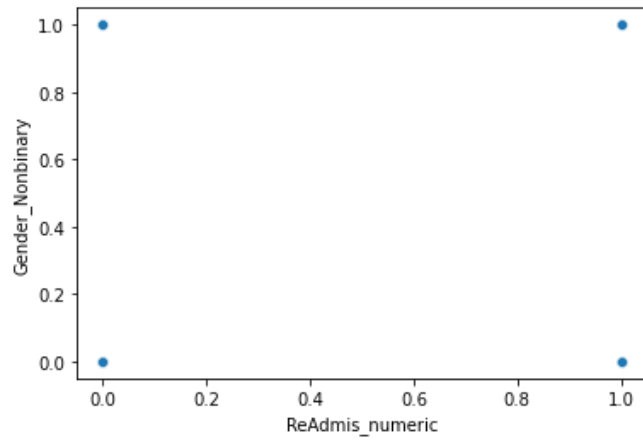
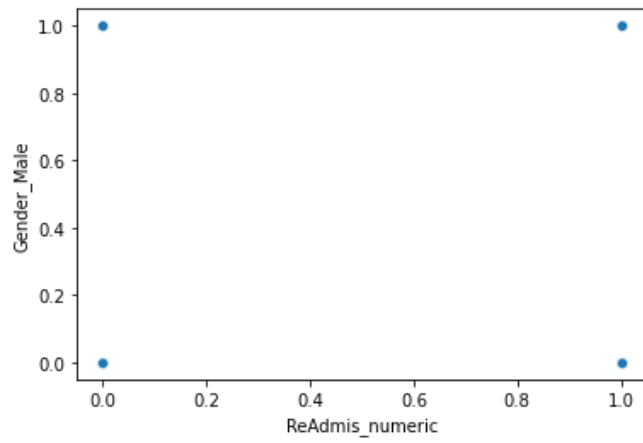
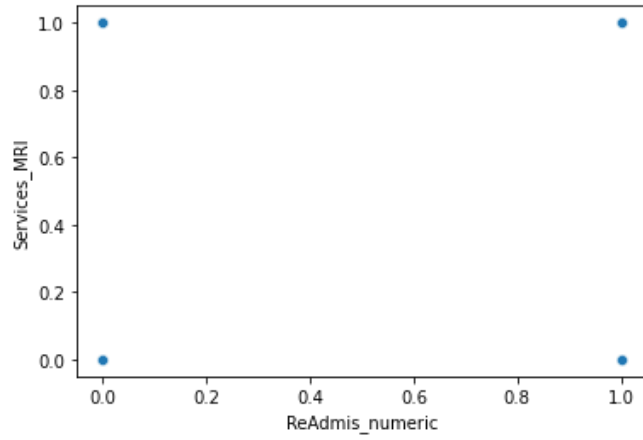
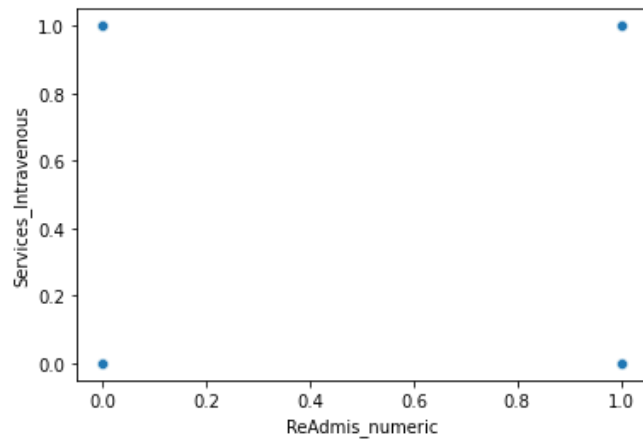


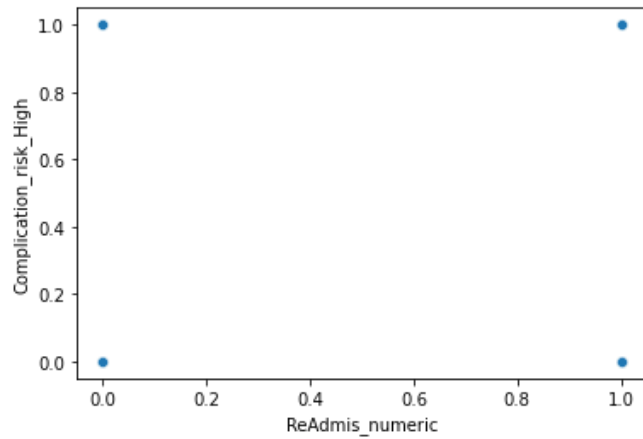
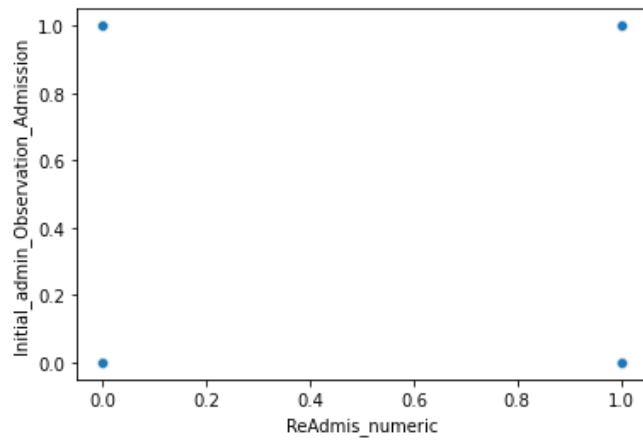
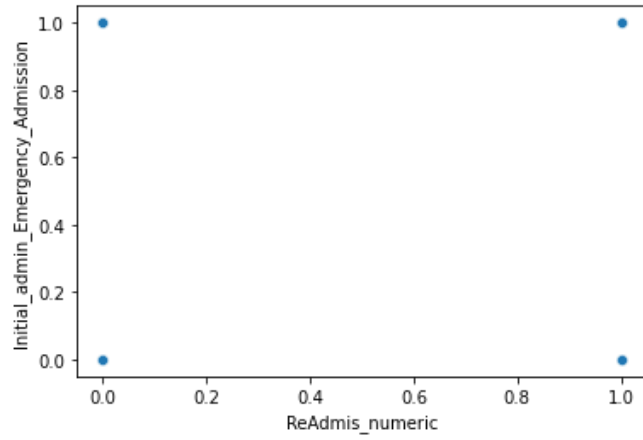
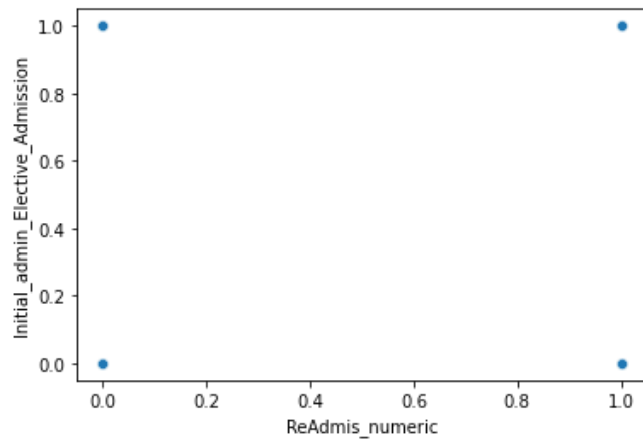


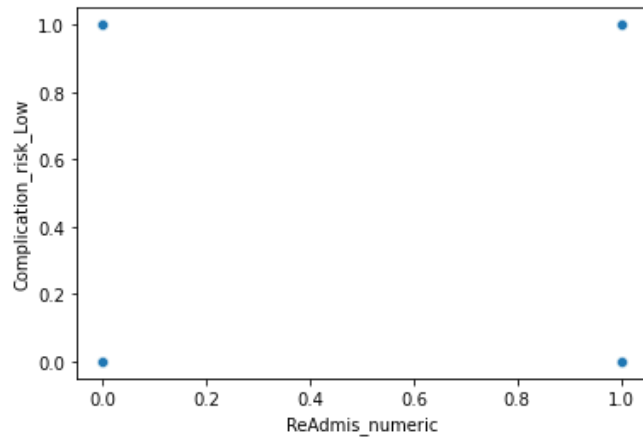
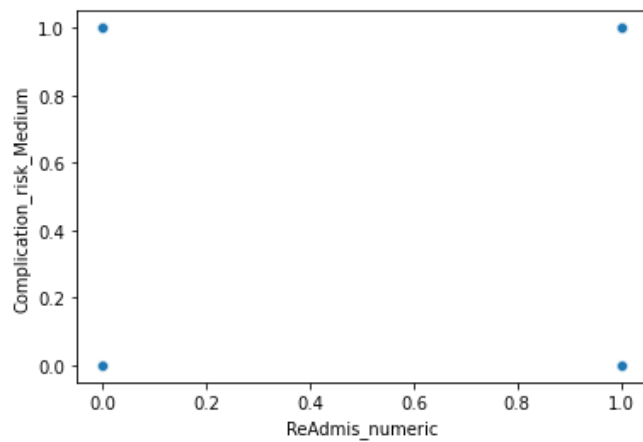












```
In [10]: df.to_csv(r'C:\Users\mmorg\Desktop\D208 Assessment Files\Cleaned208data.csv')
```

```
In [11]: #Initial Logistic Regression Model
mdl_readmis_vs_variables = logit("ReAdmis_numeric ~ Initial_days + vitD_supp + Children + Income + Full_
print(mdl_readmis_vs_variables.params)
mdl_readmis_vs_variables.summary()
```

Warning: Maximum number of iterations has been exceeded.

Current function value: 0.032914

Iterations: 35

Intercept	-78.391168
Initial_days	-1.143612
vitD_supp	-0.105465
Children	0.088933
Income	0.000002
Full_meals_eaten	0.048269
Additional_charges	0.000047
TotalCharge	0.032097
VitD_levels	0.029913
Age	-0.008115
Doc_visits	0.006554
HighBlood_numeric	-3.125420
Stroke_numeric	1.651497
Arthritis_numeric	-3.669106
Diabetes_numeric	-1.908568
Hyperlipidemia_numeric	-2.757763
BackPain_numeric	-2.508272
Allergic_rhinitis_numeric	-2.268662
Reflux_esophagitis_numeric	-2.342286
Asthma_numeric	-1.389130
Overweight_numeric	-0.286489
Anxiety_numeric	-3.817203
Marital_Married	0.268609
Marital_Never_Married	0.356522
Marital_Separated	-0.127193
Marital_Widowed	0.136171
Services_Blood_Work	-20.676842
Services_CT_Scan	-19.061885
Services_Intravenous	-20.683935
Services_MRI	-17.968342
Gender_Male	0.169142
Gender_Nonbinary	0.365800
Initial_admin_Elective_Admission	-21.750667
Initial_admin_Emergency_Admission	-35.680140
Initial_admin_Observation_Admission	-20.960246
Complication_risk_High	-34.251935
Complication_risk_Low	-22.802367
Complication_risk_Medium	-21.336813

dtype: float64

C:\Users\mmorg\anaconda3\lib\site-packages\statsmodels\base\model.py:604: ConvergenceWarning: Maximum Likelihood optimization failed to converge. Check mle\_retvals  
 warnings.warn("Maximum Likelihood optimization failed to "

Out[11]:

## Logit Regression Results

<b>Dep. Variable:</b>	ReAdmis_numeric	<b>No. Observations:</b>	10000
<b>Model:</b>	Logit	<b>Df Residuals:</b>	9965
<b>Method:</b>	MLE	<b>Df Model:</b>	34
<b>Date:</b>	Thu, 24 Nov 2022	<b>Pseudo R-squ.:</b>	0.9499
<b>Time:</b>	22:22:07	<b>Log-Likelihood:</b>	-329.14
<b>converged:</b>	False	<b>LL-Null:</b>	-6572.9
<b>Covariance Type:</b>	nonrobust	<b>LLR p-value:</b>	0.000

	coef	std err	z	P> z	[0.025	0.975]
<b>Intercept</b>	-78.3912	nan	nan	nan	nan	nan
<b>Initial_days</b>	-1.1436	1.5e+04	-7.62e-05	1.000	-2.94e+04	2.94e+04
<b>vitD_supp</b>	-0.1055	0.167	-0.632	0.528	-0.433	0.222
<b>Children</b>	0.0889	0.046	1.913	0.056	-0.002	0.180
<b>Income</b>	1.572e-06	3.62e-06	0.434	0.664	-5.53e-06	8.67e-06
<b>Full_meals_eaten</b>	0.0483	0.103	0.469	0.639	-0.153	0.250
<b>Additional_charges</b>	4.664e-05	6.25e-05	0.746	0.455	-7.58e-05	0.000
<b>TotalCharge</b>	0.0321	183.195	0.000	1.000	-359.024	359.088
<b>VitD_levels</b>	0.0299	0.049	0.613	0.540	-0.066	0.126
<b>Age</b>	-0.0081	0.015	-0.549	0.583	-0.037	0.021
<b>Doc_visits</b>	0.0066	0.098	0.067	0.947	-0.186	0.200
<b>HighBlood_numeric</b>	-3.1254	2.06e+04	-0.000	1.000	-4.03e+04	4.03e+04
<b>Stroke_numeric</b>	1.6515	0.272	6.069	0.000	1.118	2.185
<b>Arthritis_numeric</b>	-3.6691	1.32e+04	-0.000	1.000	-2.58e+04	2.58e+04
<b>Diabetes_numeric</b>	-1.9086	1.38e+04	-0.000	1.000	-2.7e+04	2.7e+04
<b>Hyperlipidemia_numeric</b>	-2.7578	1.72e+04	-0.000	1.000	-3.38e+04	3.37e+04
<b>BackPain_numeric</b>	-2.5083	1.56e+04	-0.000	1.000	-3.06e+04	3.06e+04
<b>Allergic_rhinitis_numeric</b>	-2.2687	1.11e+04	-0.000	1.000	-2.18e+04	2.17e+04
<b>Reflux_esophagitis_numeric</b>	-2.3423	1.09e+04	-0.000	1.000	-2.14e+04	2.14e+04
<b>Asthma_numeric</b>	-1.3891	0.237	-5.852	0.000	-1.854	-0.924
<b>Overweight_numeric</b>	-0.2865	0.229	-1.250	0.211	-0.736	0.163
<b>Anxiety_numeric</b>	-3.8172	1.58e+04	-0.000	1.000	-3.09e+04	3.09e+04
<b>Marital_Married</b>	0.2686	0.331	0.811	0.418	-0.381	0.918
<b>Marital_Never_Married</b>	0.3565	0.338	1.055	0.292	-0.306	1.019
<b>Marital_Separated</b>	-0.1272	0.344	-0.369	0.712	-0.802	0.548
<b>Marital_Widowed</b>	0.1362	0.333	0.409	0.682	-0.516	0.788
<b>Services_Blood_Work</b>	-20.6768	4.07e+06	-5.08e-06	1.000	-7.98e+06	7.98e+06
<b>Services_CT_Scan</b>	-19.0619	4.07e+06	-4.68e-06	1.000	-7.98e+06	7.98e+06
<b>Services_Intravenous</b>	-20.6839	4.07e+06	-5.08e-06	1.000	-7.98e+06	7.98e+06
<b>Services_MRI</b>	-17.9683	4.07e+06	-4.41e-06	1.000	-7.99e+06	7.99e+06
<b>Gender_Male</b>	0.1691	0.210	0.807	0.420	-0.242	0.580
<b>Gender_Nonbinary</b>	0.3658	0.714	0.512	0.608	-1.033	1.765

<b>Initial_admin_Elective_Admission</b>	-21.7507	nan	nan	nan	nan	nan
<b>Initial_admin_Emergency_Admission</b>	-35.6801	nan	nan	nan	nan	nan
<b>Initial_admin_Observation_Admission</b>	-20.9602	nan	nan	nan	nan	nan
<b>Complication_risk_High</b>	-34.2519	2.15e+06	-1.59e-05	1.000	-4.22e+06	4.22e+06
<b>Complication_risk_Low</b>	-22.8024	2.34e+06	-9.75e-06	1.000	-4.59e+06	4.59e+06
<b>Complication_risk_Medium</b>	-21.3368	2.34e+06	-9.12e-06	1.000	-4.59e+06	4.59e+06

Possibly complete quasi-separation: A fraction 0.82 of observations can be perfectly predicted. This might indicate that there is complete quasi-separation. In this case some parameters will not be identified.

```
In [12]: #Confusion Matrix for Initial Model

conf_matrix = mdl_readmis_vs_variables.pred_table()
print(conf_matrix)

from statsmodels.graphics.mosaicplot import mosaic
mosaic(conf_matrix)

#Calculating accuracy: the proportion of correct predictions

TN = conf_matrix[0,0]
TP = conf_matrix[1,1]
FN = conf_matrix[1,0]
FP = conf_matrix[0,1]

acc = (TN + TP) / (TN + TP + FN + FP)
print('Accuracy:', acc)

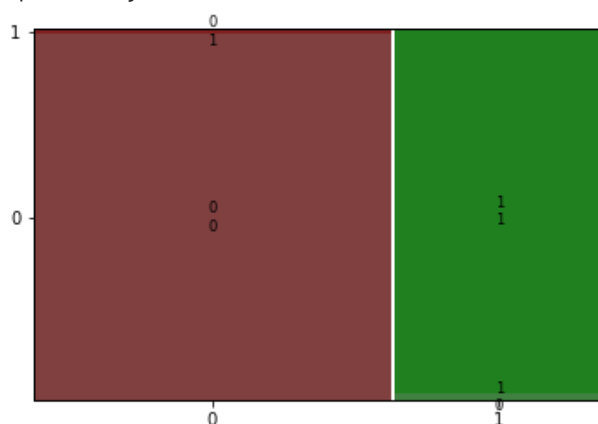
#Sensitivity: proportion of true positives

sens = TP / (FN + TP)
print('Sensitivity:', sens)

#Specificity: proportion of true negatives

spec = TN / (TN + FP)
print('Specificity:', spec)
```

```
[[6261.  70.]
 [ 60. 3609.]]
Accuracy: 0.987
Sensitivity: 0.9836467702371219
Specificity: 0.9889432948981204
```



```
In [13]: #Variable Selection
# Checking for the VIF values of the variables.
from statsmodels.stats.outliers_influence import variance_inflation_factor

X = df[['Initial_days', 'vitD_supp', 'Children', 'Income', 'Full_meals_eaten', 'Additional_charges', 'To

# VIF dataframe
vif_data = pd.DataFrame()
vif_data["feature"] = X.columns

# calculating VIF for each feature
vif_data["VIF"] = [variance_inflation_factor(X.values, i)
                    for i in range(len(X.columns))]

print(vif_data)
```

	feature	VIF
0	Initial_days	2880.163153
1	vitD_supp	1.003676
2	Children	1.003506
3	Income	1.002683
4	Full_meals_eaten	1.004107
5	Additional_charges	16.303881
6	TotalCharge	2944.078834
7	VitD_levels	1.003914
8	Age	9.273563
9	Doc_visits	1.003377
10	HighBlood_numeric	9.711378
11	Stroke_numeric	1.010014
12	Arthritis_numeric	1.760819
13	Diabetes_numeric	1.696512
14	Hyperlipidemia_numeric	2.198017
15	BackPain_numeric	2.112077
16	Allergic_rhinitis_numeric	1.551215
17	Reflux_esophagitis_numeric	1.527188
18	Asthma_numeric	1.003104
19	Marital_Married	1.627325
20	Marital_Never_Married	1.618488
21	Marital_Separated	1.617166
22	Marital_Widowed	1.630512
23	Services_Blood_Work	inf
24	Services_CT_Scan	inf
25	Services_Intravenous	inf
26	Services_MRI	inf
27	Gender_Male	1.026146
28	Gender_Nonbinary	1.023726
29	Initial_admin_Elective_Admission	inf
30	Initial_admin_Emergency_Admission	inf
31	Initial_admin_Observation_Admission	inf
32	Complication_risk_High	inf
33	Complication_risk_Low	inf
34	Complication_risk_Medium	inf

C:\Users\mmorg\anaconda3\lib\site-packages\statsmodels\stats\outliers\_influence.py:195: RuntimeWarning: divide by zero encountered in double\_scalars  
 vif = 1. / (1. - r\_squared\_i)

```
In [14]: #Reduced Model removing complication risk, initial admin, services, demographics, and charges columns du

mdl_readmis_vs_variables1 = logit("ReAdmis_numeric ~ Initial_days + Children + Stroke_numeric + Asthma_n

print(mdl_readmis_vs_variables1.params)

mdl_readmis_vs_variables1.summary()
```

Optimization terminated successfully.  
 Current function value: 0.045704  
 Iterations 13  
 Intercept -57.974445  
 Initial\_days 1.065053  
 Children 0.069841  
 Stroke\_numeric 1.273473  
 Asthma\_numeric -0.944019  
 dtype: float64

Out[14]:

Logit Regression Results

<b>Dep. Variable:</b>	ReAdmis_numeric	<b>No. Observations:</b>	10000
<b>Model:</b>	Logit	<b>Df Residuals:</b>	9995
<b>Method:</b>	MLE	<b>Df Model:</b>	4
<b>Date:</b>	Thu, 24 Nov 2022	<b>Pseudo R-squ.:</b>	0.9305
<b>Time:</b>	22:23:54	<b>Log-Likelihood:</b>	-457.04
<b>converged:</b>	True	<b>LL-Null:</b>	-6572.9
<b>Covariance Type:</b>	nonrobust	<b>LLR p-value:</b>	0.000

	coef	std err	z	P> z	[0.025	0.975]
<b>Intercept</b>	-57.9744	2.749	-21.087	0.000	-63.363	-52.586
<b>Initial_days</b>	1.0651	0.050	21.125	0.000	0.966	1.164
<b>Children</b>	0.0698	0.038	1.839	0.066	-0.005	0.144
<b>Stroke_numeric</b>	1.2735	0.223	5.706	0.000	0.836	1.711
<b>Asthma_numeric</b>	-0.9440	0.190	-4.975	0.000	-1.316	-0.572

Possibly complete quasi-separation: A fraction 0.75 of observations can be perfectly predicted. This might indicate that there is complete quasi-separation. In this case some parameters will not be identified.

In [15]:

```
#Confusion Matrix for Reduced Model

conf_matrix = mdl_readmis_vs_variables1.pred_table()

print(conf_matrix)

from statsmodels.graphics.mosaicplot import mosaic

mosaic(conf_matrix)

#Calculating accuracy: the proportion of correct predictions

TN = conf_matrix[0,0]
TP = conf_matrix[1,1]
FN = conf_matrix[1,0]
FP = conf_matrix[0,1]

acc = (TN + TP) / (TN + TP + FN + FP)
print('Accuracy:', acc)

#Sensitivity: proportion of true positives

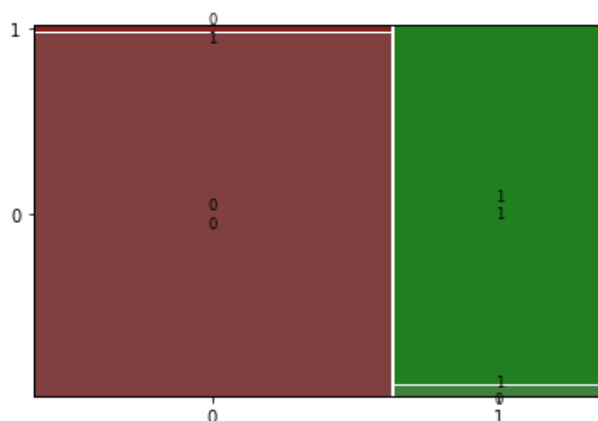
sens = TP / (FN + TP)
print('Sensitivity:', sens)

#Specificity: proportion of true negatives

spec = TN / (TN + FP)
print('Specificity:', spec)
```



```
[[6228. 103.]
 [ 102. 3567.]]
Accuracy: 0.9795
Sensitivity: 0.9721995094031071
Specificity: 0.9837308482072342
```



```
In [16]: #Calculating AIC of Initial Model
from sklearn.linear_model import LinearRegression
import statsmodels.api as sm

#define response variable
y = df['ReAdmis_numeric']

#define predictor variables
x = df[['Initial_days', 'vitD_supp', 'Children', 'Income', 'Full_meals_eaten', 'Additional_charges', 'To

#add constant to predictor variables
x = sm.add_constant(x)

#fit regression model
model = sm.OLS(y, x).fit()

#view AIC of model
print(model.aic)
```

```
897.5899676679401
```

```
In [17]: #Calculating AIC of Reduced Model #1

#define response variable
y = df['ReAdmis_numeric']

#define predictor variables
x = df[['Initial_days', 'Children', 'Stroke_numeric', 'Asthma_numeric', 'Overweight_numeric']]

#add constant to predictor variables
x = sm.add_constant(x)

#fit regression model
model = sm.OLS(y, x).fit()

#view AIC of model
print(model.aic)
```

```
918.392404996368
```

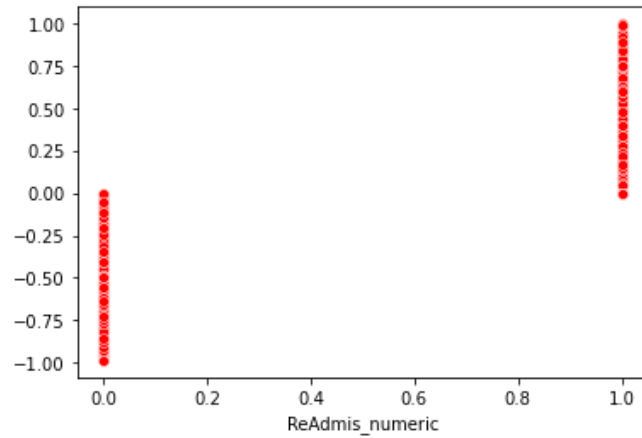
```
In [18]: #Residual Standard Error for reduced model
np.sqrt mdl_readmis_vs_variables1.scale)
```

```
Out[18]: 1.0
```

```
In [19]: #Residual plot for reduced model
df['intercept'] = 1
```

```
residuals = df['ReAdmis_numeric'] - mdl_readmis_vs_variables1.predict(df[['Initial_days', 'Children', 'S  
sns.scatterplot(x=df['ReAdmis_numeric'], y=residuals, color='red')
```

Out[19]: <AxesSubplot:xlabel='ReAdmis\_numeric'>



In [ ]: