

D207 - Exploratory Data Analysis

By Krista Moik

```
In [1]: #import packages
import pandas as pd
import numpy as np
from scipy.stats import chi2_contingency
import matplotlib.pyplot as plt
import seaborn as sns

%matplotlib inline
```

```
In [2]: #Load medical_clean CSV
df = pd.read_csv('C:/Users/Kmoik WGU/Desktop/D207/medical_clean.csv')
```

```
In [3]: #View data set
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 10000 entries, 0 to 9999
Data columns (total 50 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  Marital                10000 non-null  object
18  Gender                 10000 non-null  object
19  ReAdmis                10000 non-null  object
20  VitD_levels            10000 non-null  float64
21  Doc_visits             10000 non-null  int64
22  Full_meals_eaten       10000 non-null  int64
23  vitD_supp              10000 non-null  int64
24  Soft_drink             10000 non-null  object
25  Initial_admin          10000 non-null  object
26  HighBlood              10000 non-null  object
27  Stroke                 10000 non-null  object
28  Complication_risk      10000 non-null  object
29  Overweight             10000 non-null  object
30  Arthritis              10000 non-null  object
31  Diabetes               10000 non-null  object
32  Hyperlipidemia         10000 non-null  object
33  BackPain               10000 non-null  object
34  Anxiety                10000 non-null  object
35  Allergic_rhinitis      10000 non-null  object
36  Reflux_esophagitis     10000 non-null  object
37  Asthma                 10000 non-null  object
38  Services               10000 non-null  object
39  Initial_days           10000 non-null  float64
40  TotalCharge             10000 non-null  float64
41  Additional_charges     10000 non-null  float64
42  Item1                  10000 non-null  int64
43  Item2                  10000 non-null  int64
44  Item3                  10000 non-null  int64
45  Item4                  10000 non-null  int64
46  Item5                  10000 non-null  int64
47  Item6                  10000 non-null  int64
48  Item7                  10000 non-null  int64
49  Item8                  10000 non-null  int64
dtypes: float64(7), int64(16), object(27)
memory usage: 3.8+ MB
```

In [4]: *#View variable ReAdmis*

```
df.ReAdmis.describe()
```

```
Out[4]: count      10000
        unique        2
        top          No
        freq       6331
        Name: ReAdmis, dtype: object
```

```
In [5]: #Count responses in ReAdmis
        df['ReAdmis'].value_counts(normalize=True, sort=True, dropna=True)
```

```
Out[5]: ReAdmis
        No      0.6331
        Yes     0.3669
        Name: proportion, dtype: float64
```

```
In [6]: #View variable Complication_risk
        df.Complication_risk.describe()
```

```
Out[6]: count      10000
        unique        3
        top      Medium
        freq       4517
        Name: Complication_risk, dtype: object
```

```
In [7]: #Count responses in Complication_risk
        df['Complication_risk'].value_counts(normalize=True, sort=True, dropna=True)
```

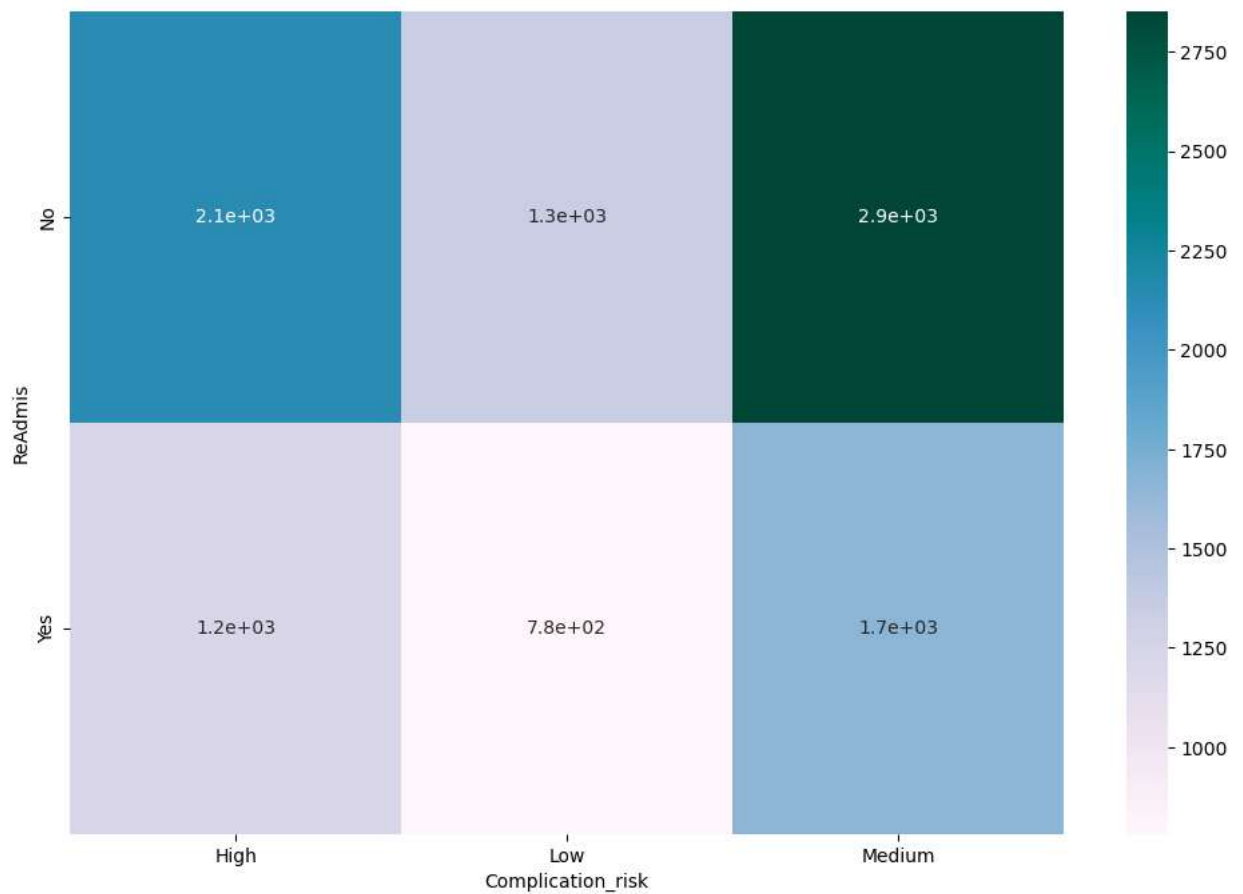
```
Out[7]: Complication_risk
        Medium  0.4517
        High    0.3358
        Low     0.2125
        Name: proportion, dtype: float64
```

```
In [8]: #Create Contingency Table for Chi-Square using code from WGU course materials
        table=pd.crosstab(df.ReAdmis, df.Complication_risk, margins=True)
        print(table)
```

Complication_risk	High	Low	Medium	All
ReAdmis				
No	2135	1343	2853	6331
Yes	1223	782	1664	3669
All	3358	2125	4517	10000

```
In [9]: #Obtain P-Value using code from WGU course materials
        df.head()
        contingency=pd.crosstab(df['ReAdmis'], df['Complication_risk'])
        contingency
        contingency_pct=pd.crosstab(df['ReAdmis'], df['Complication_risk'], normalize='index')
        contingency_pct
        plt.figure(figsize=(12,8))
        sns.heatmap(contingency, annot=True, cmap="PuBuGn")
```

```
Out[9]: <Axes: xlabel='Complication_risk', ylabel='ReAdmis'>
```



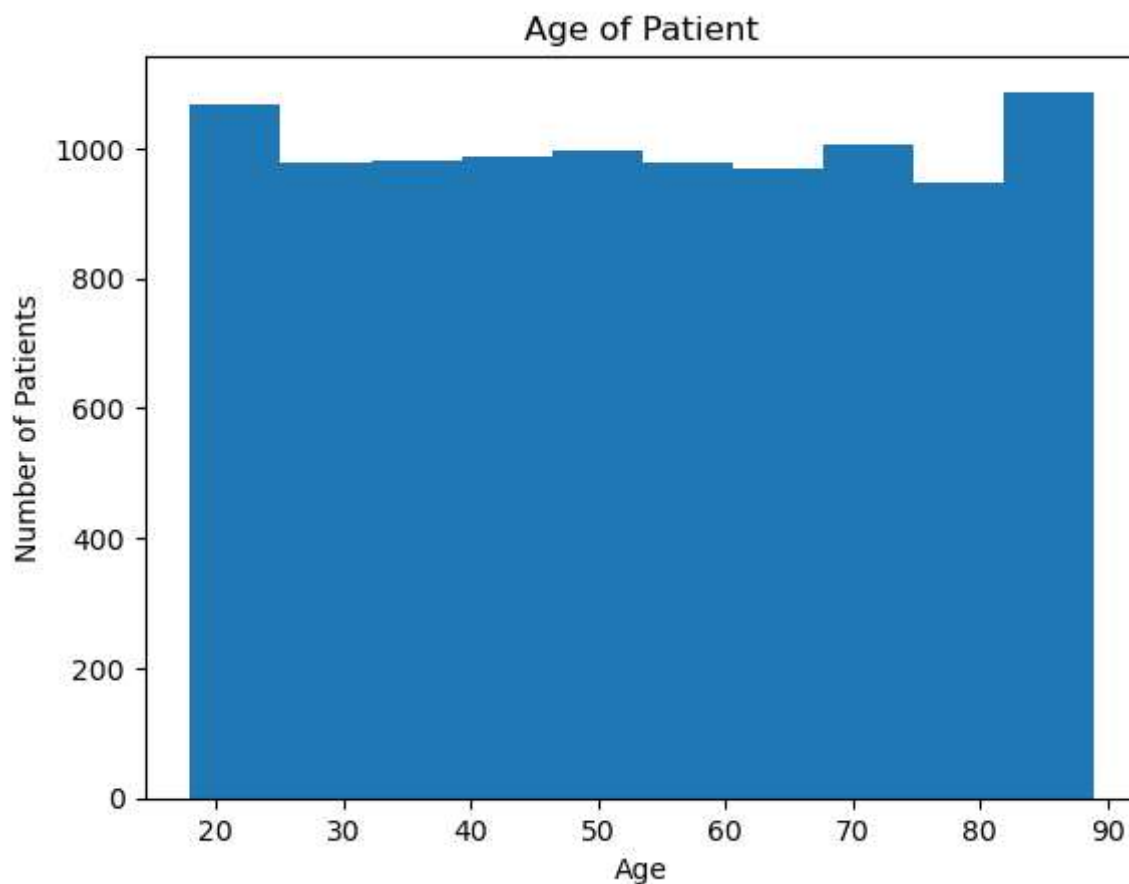
```
In [10]: #Chi-Square test of independence using code from WGU course materials
c, p, dof, expected=chi2_contingency(contingency)
```

```
In [11]: #Print P-value
print(p)
```

0.923567890607327

Univariate Statistics

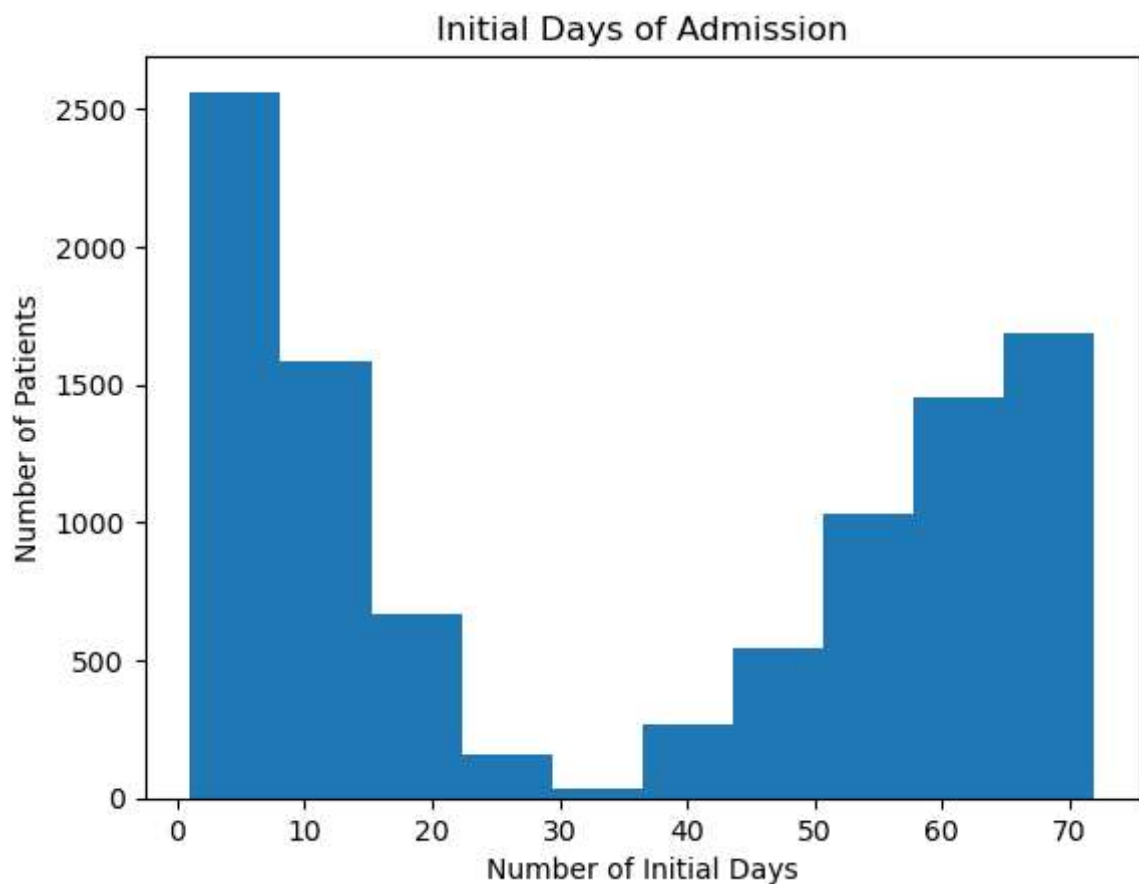
```
In [12]: #Histogram of Age column - continuous
plt.hist(df['Age'])
plt.title("Age of Patient")
plt.xlabel("Age")
plt.ylabel("Number of Patients")
plt.show()
```



```
In [13]: df.Age.describe()
```

```
Out[13]: count    10000.000000
mean       53.511700
std        20.638538
min        18.000000
25%        36.000000
50%        53.000000
75%        71.000000
max        89.000000
Name: Age, dtype: float64
```

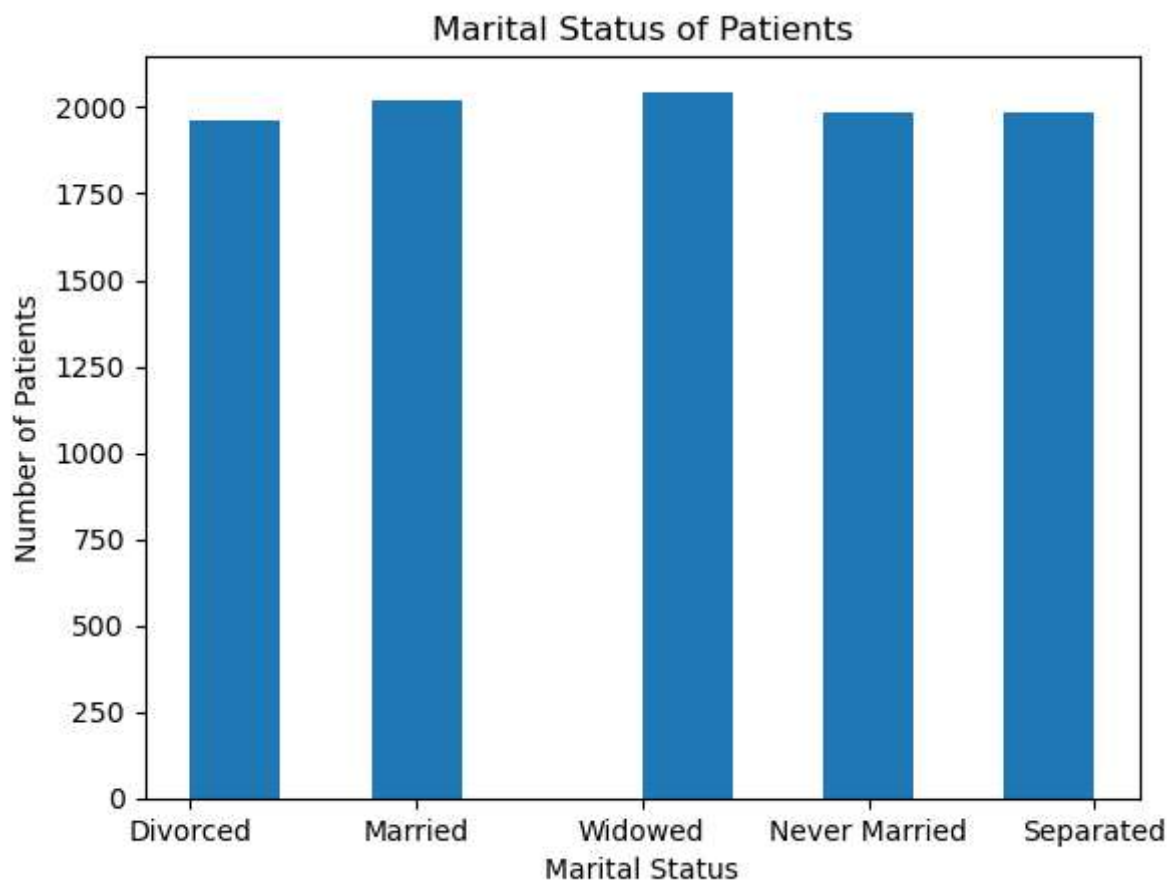
```
In [14]: #Histogram of Initial_days column - continuous
plt.hist(df['Initial_days'])
plt.title("Initial Days of Admission")
plt.xlabel("Number of Initial Days")
plt.ylabel("Number of Patients")
plt.show()
```



```
In [15]: df.Initial_days.describe()
```

```
Out[15]: count    10000.000000
mean       34.455299
std        26.309341
min         1.001981
25%         7.896215
50%        35.836244
75%        61.161020
max        71.981490
Name: Initial_days, dtype: float64
```

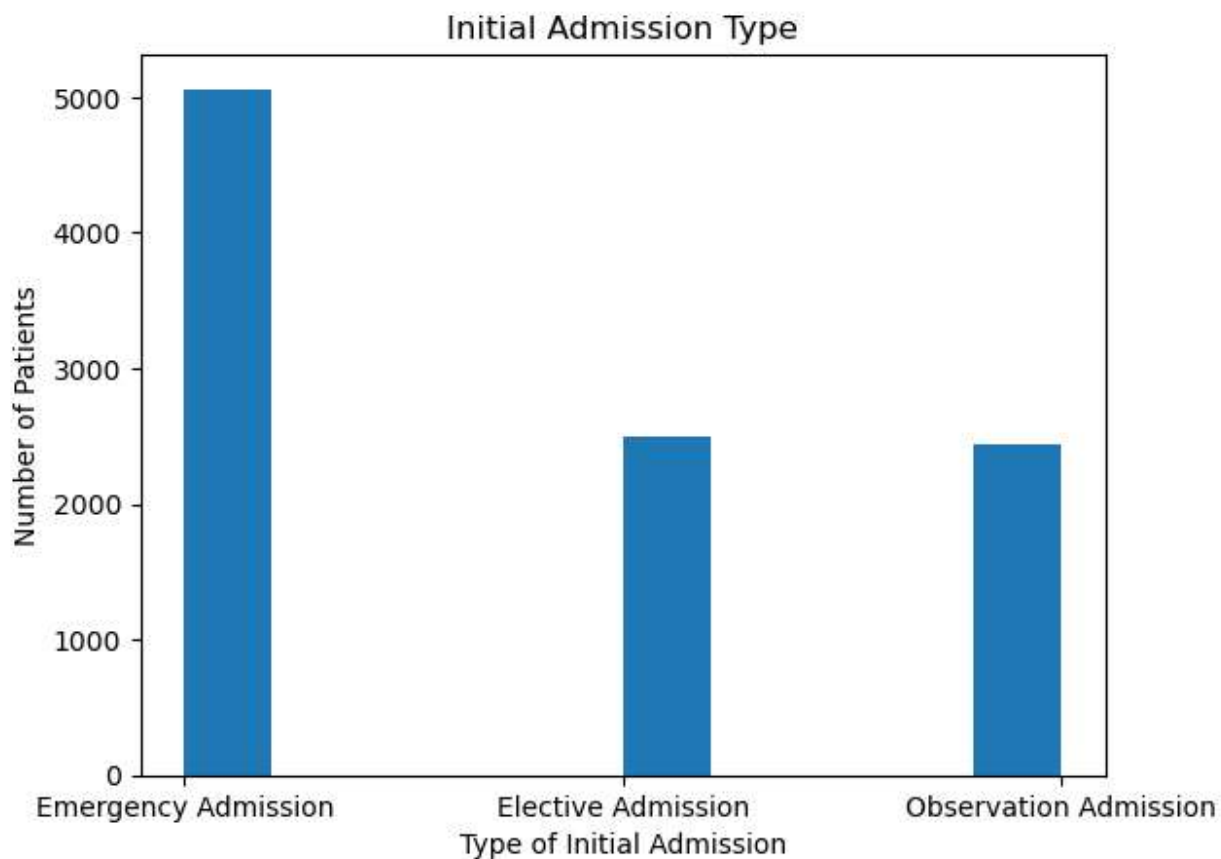
```
In [16]: #Histogram of Marital - categorical
plt.hist(df['Marital'])
plt.title("Marital Status of Patients")
plt.xlabel("Marital Status")
plt.ylabel("Number of Patients")
plt.show()
```



```
In [17]: df.Marital.describe()
```

```
Out[17]: count      10000  
unique         5  
top      Widowed  
freq         2045  
Name: Marital, dtype: object
```

```
In [18]: #Histogram of Initial_admin - categorical  
plt.hist(df['Initial_admin'])  
plt.title("Initial Admission Type")  
plt.xlabel("Type of Initial Admission")  
plt.ylabel("Number of Patients")  
plt.show()
```

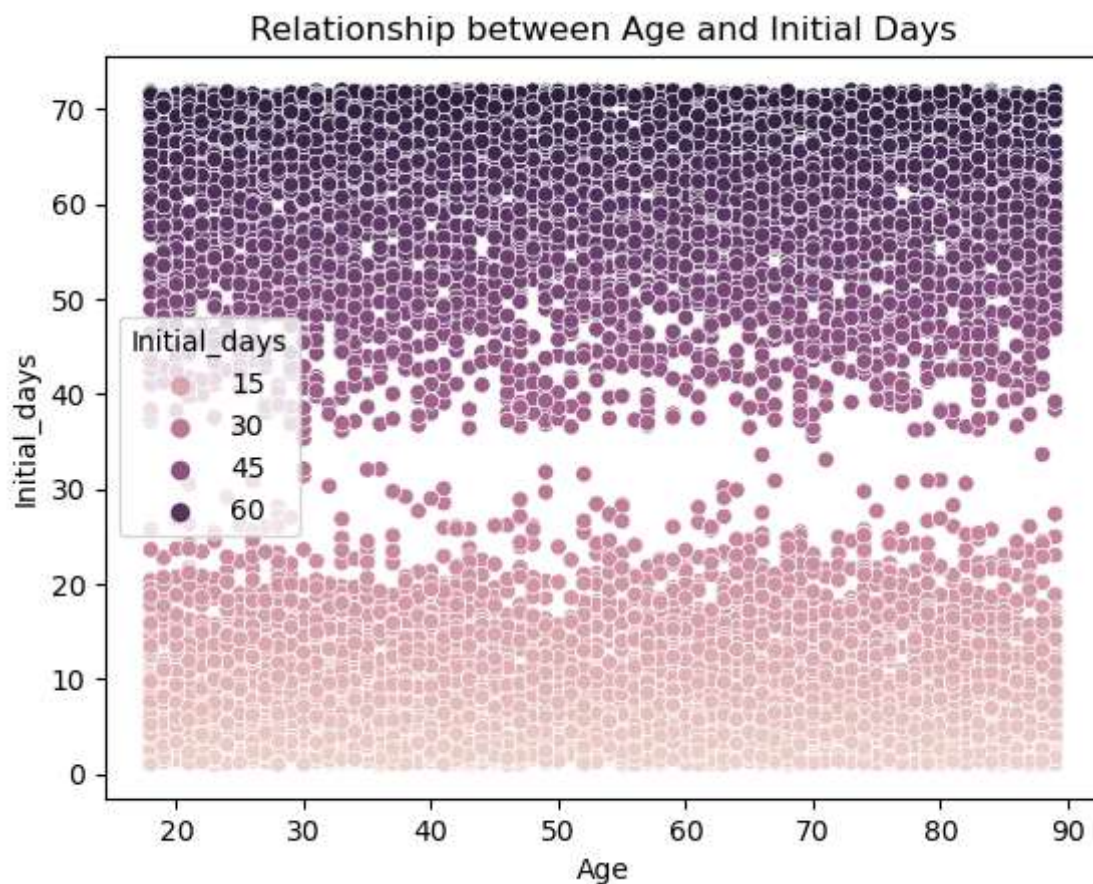


```
In [19]: df.Initial_admin.describe()
```

```
Out[19]: count          10000  
unique           3  
top      Emergency Admission  
freq           5060  
Name: Initial_admin, dtype: object
```

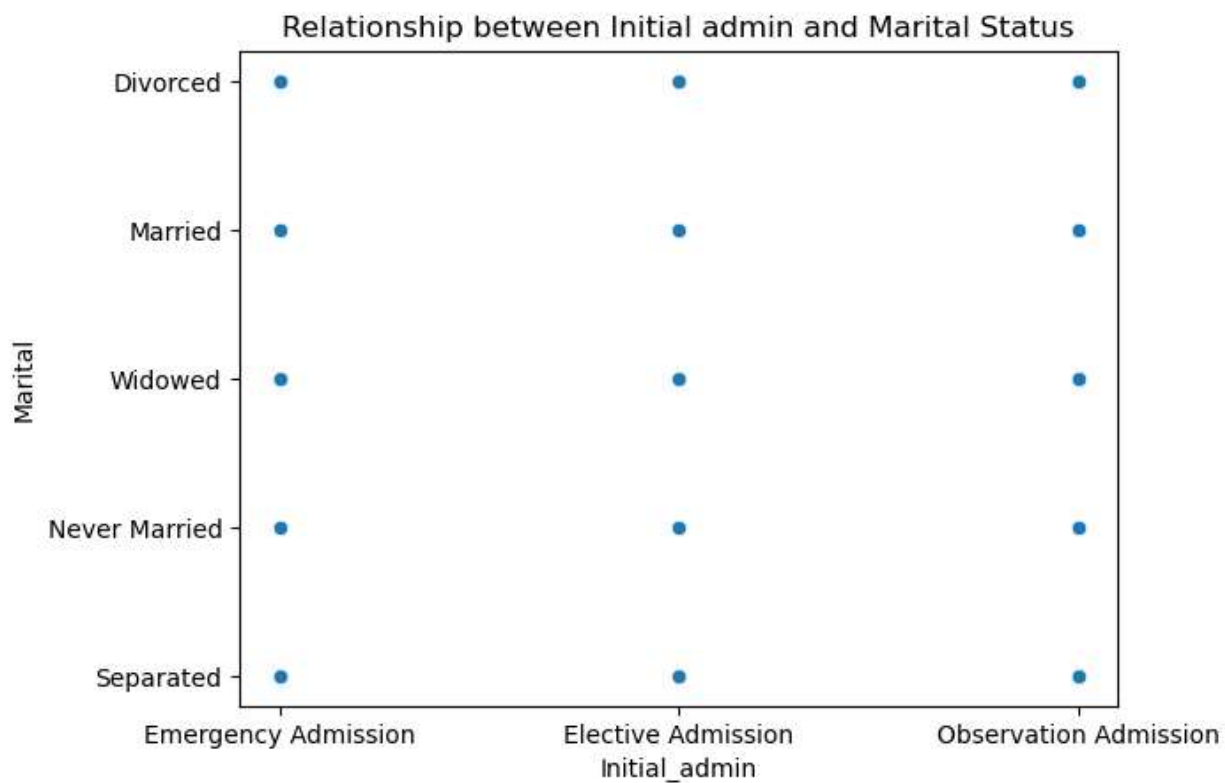
Bivariate Statistics

```
In [20]: #Scatterplot of Age and Initial_days - continuous  
sns.scatterplot(x='Age', y='Initial_days', hue='Initial_days', data=df).set(title='Rel  
plt.show()
```

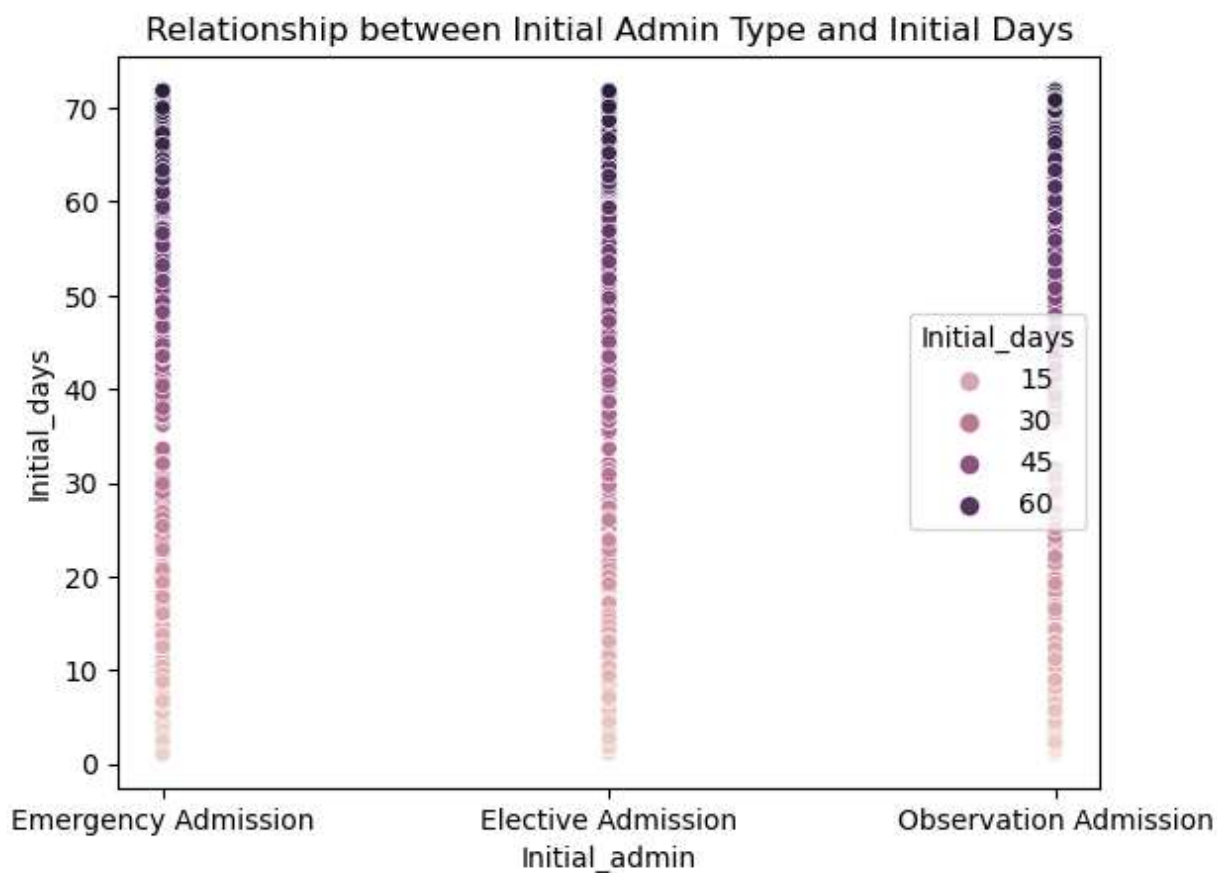



In [21]: `#Scatter plot of Initial_admin and Marital - categorical`
`sns.scatterplot(x='Initial_admin', y='Marital', data=df).set(title='Relationship between`

Out[21]: `[Text(0.5, 1.0, 'Relationship between Initial admin and Marital Status')]`

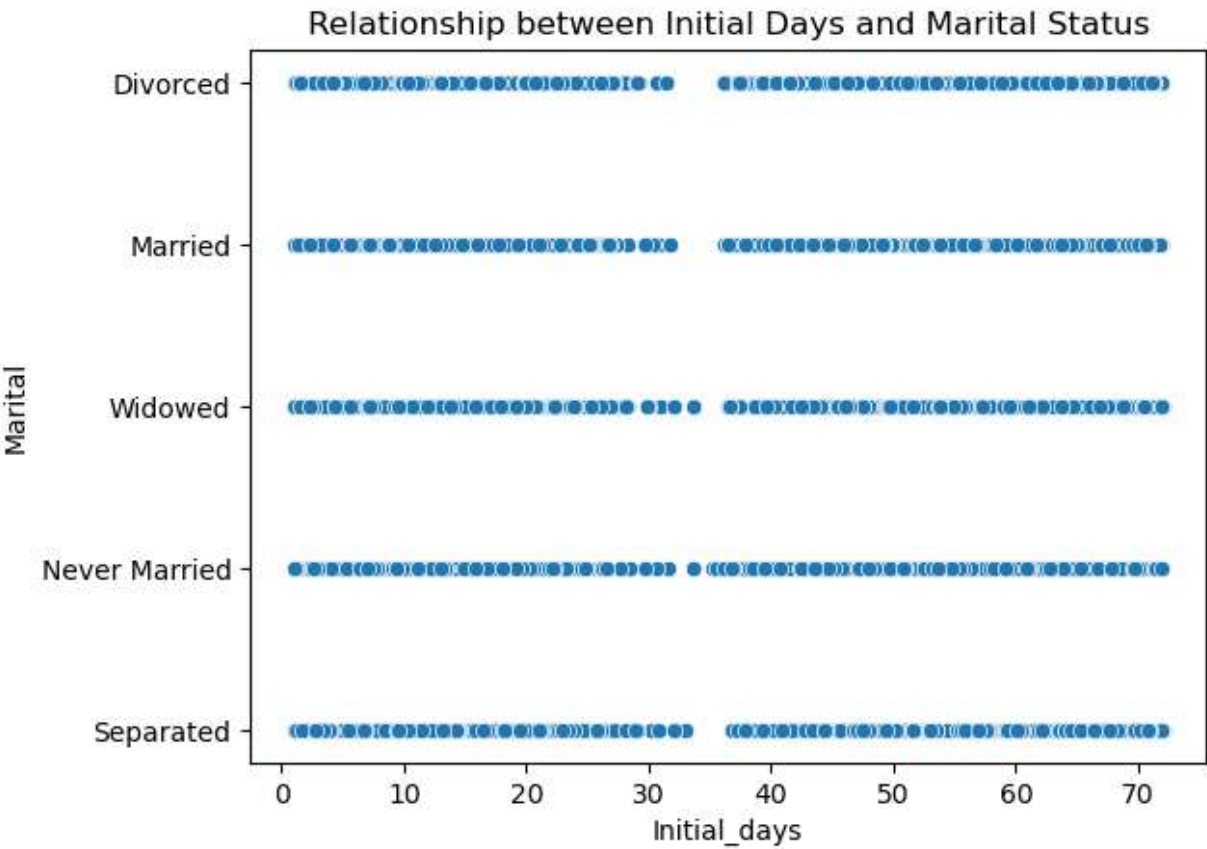


```
In [22]: #Scatterplot of Initial_admin and Initial_days - categorical and continuous
sns.scatterplot(x='Initial_admin', y='Initial_days', hue='Initial_days', data=df).set(
plt.show()
```



```
In [23]: #Scatter plot of Initial_admin and Marital - categorical
sns.scatterplot(x='Initial_days', y='Marital', data=df).set(title='Relationship between
```

```
Out[23]: [Text(0.5, 1.0, 'Relationship between Initial Days and Marital Status')]
```



```
In [ ]:
```