



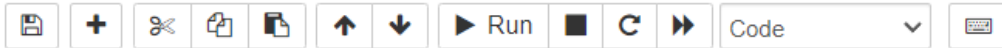
```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import statistics
```

```
In [2]: train=pd.read_csv('C:/Users/Ranjan kumar/Downloads/healthcare/train_data.csv')
```

```
In [3]: train
```

Out[3]:

	case_id	Hospital_code	Hospital_type_code	City_Code_Hospital	Hospital_region_code	Available Extra Rooms in Hospital	Department	Ward_Type	Ward_Facility_Code	Bed Grade
0	1	8	c	3	Z	3	radiotherapy	R	F	2.0
1	2	2	c	5	Z	2	radiotherapy	S	F	2.0
2	3	10	e	1	X	2	anesthesia	S	E	2.0
3	4	26	b	2	Y	2	radiotherapy	R	D	2.0
4	5	26	b	2	Y	2	radiotherapy	S	D	2.0
...	...	...	...	...	...	...	...	...	...	...
318433	318434	6	a	6	X	3	radiotherapy	Q	F	4.0
318434	318435	24	c	1	X	2	anesthesia	Q	F	4.0



```
In [10]: train.isnull().sum()
```

```
Out[10]: case_id                0
         Hospital_code          0
         Hospital_type_code      0
         City_Code_Hospital      0
         Hospital_region_code    0
         Available Extra Rooms in Hospital  0
         Department              0
         Ward_Type               0
         Ward_Facility_Code      0
         Bed Grade               0
         patientid              0
         City_Code_Patient       0
         Type of Admission       0
         Severity of Illness     0
         Visitors with Patient   0
         Age                    0
         Admission_Deposit       0
         Stay                   0
         dtype: int64
```

```
In [11]: train.drop(['case_id', 'patientid'], axis=1, inplace=True)
```

```
In [12]: category=[]
         numer=[]

         for col in train.columns:
             if train[col].dtypes=='object':
                 category.append(col)

         for col in train.columns:
             if train[col].dtypes!='object':
                 numer.append(col)

         print(category)
```

```
In [12]: category=[]
        number=[]

        for col in train.columns:
            if train[col].dtypes=='object':
                category.append(col)

        for col in train.columns:
            if train[col].dtypes!='object':
                number.append(col)

        print(category)
        print(number)

['Hospital_type_code', 'Hospital_region_code', 'Department', 'Ward_Type', 'Ward_Facility_Code', 'Type of Admission', 'Severity of Illness', 'Age', 'Stay']
['Hospital_code', 'City_Code_Hospital', 'Available Extra Rooms in Hospital', 'Bed Grade', 'City_Code_Patient', 'Visitors with Patient', 'Admission_Deposit']
```

```
In [13]: i=1
        plt.figure(figsize=(16,18))
        for col in category:
            plt.subplot(6,2,i)
            sns.countplot(train[col])
            i=i+1
        plt.show()
```

C:\ProgramData\Anaconda3\lib\site-packages\seaborn\\_decorators.py:36: FutureWarning: Pass the following variable as a keyword argument: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

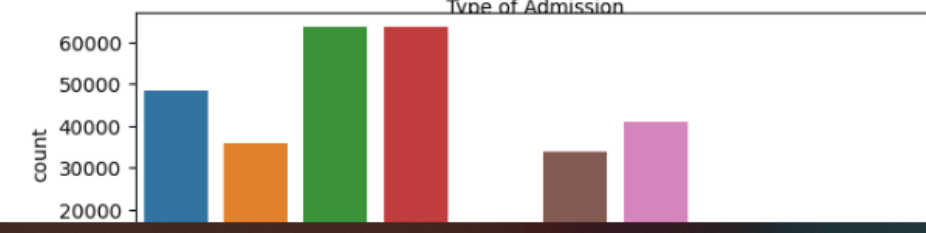
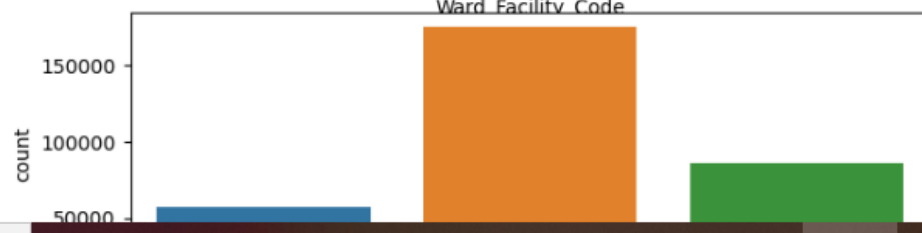
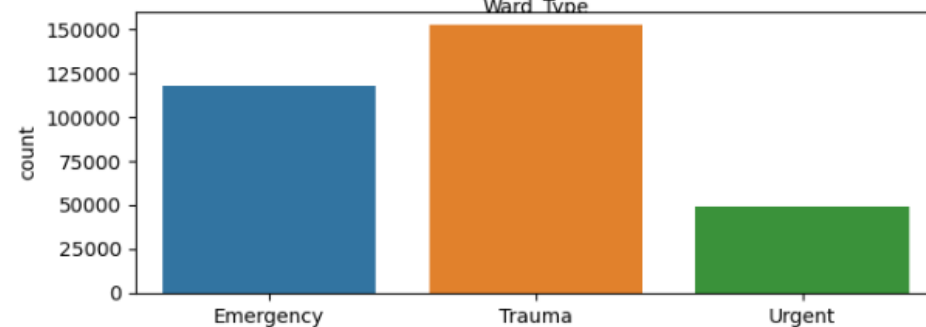
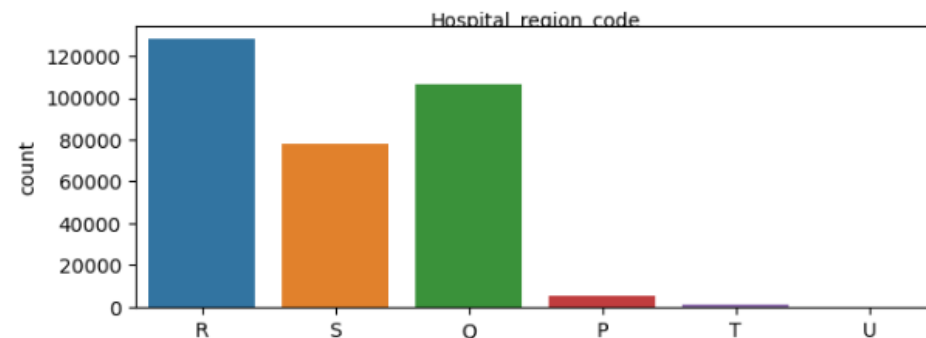
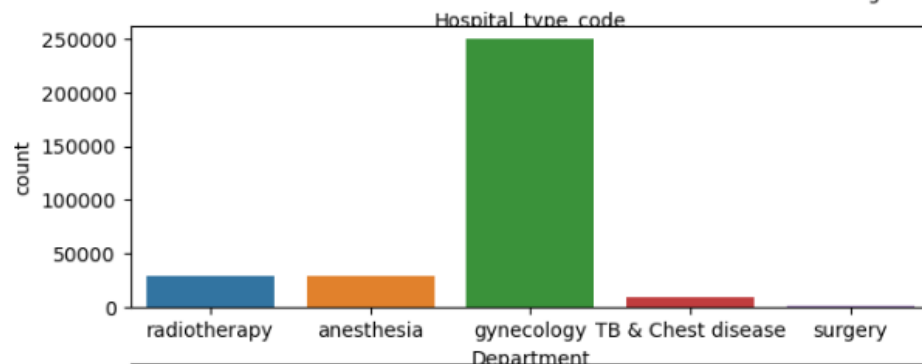
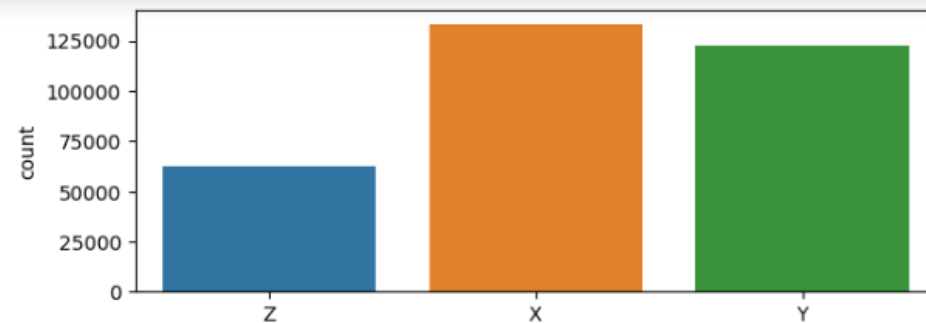
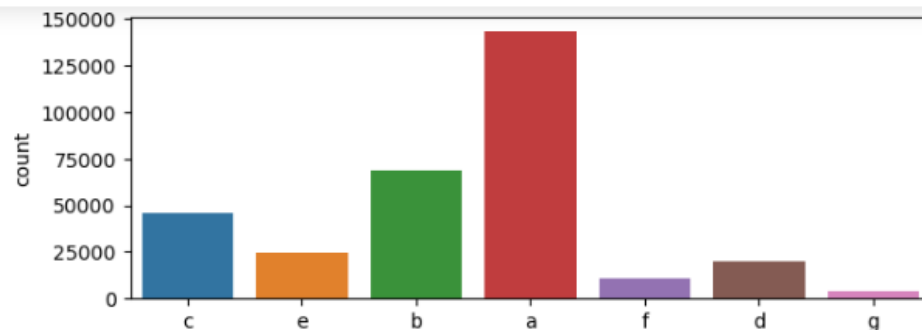
C:\ProgramData\Anaconda3\lib\site-packages\seaborn\\_decorators.py:36: FutureWarning: Pass the following variable as a keyword argument: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

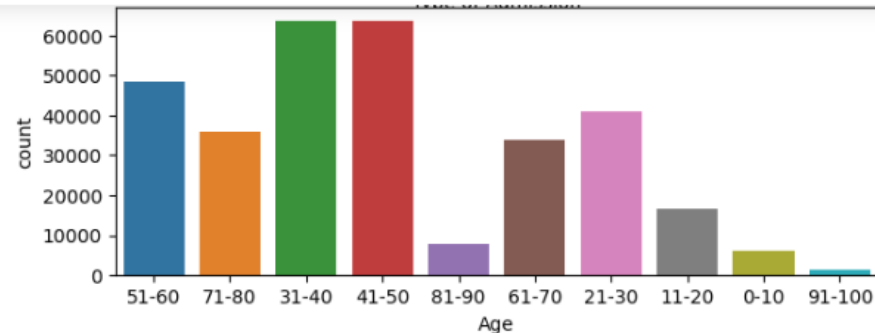
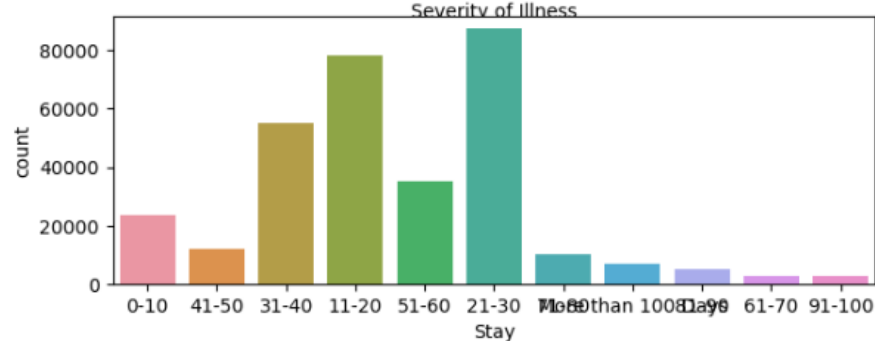
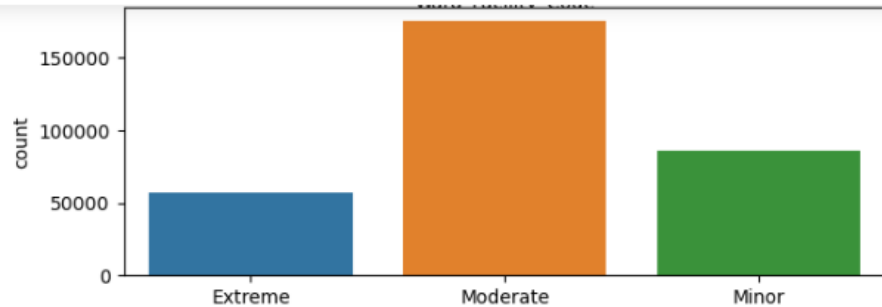
warnings.warn(

C:\ProgramData\Anaconda3\lib\site-packages\seaborn\\_decorators.py:36: FutureWarning: Pass the following variable as a keyword argument: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(







```
In [14]: train['Stay'].value_counts()
```

```
Out[14]: 21-30          87491
11-20          78139
31-40          55159
51-60          35018
0-10           23604
41-50          11743
71-80          10254
More than 100 Days  6683
81-90           4838
91-100          2765
61-70           2744
Name: Stay, dtype: int64
```



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Python 3 (ipykernel)

Code

```
In [15]: train['Stay'].replace('More than 100 Days', '>100', inplace=True)
```

```
In [16]: train['Stay'].value_counts()
```

```
Out[16]: 21-30      87491
         11-20      78139
         31-40      55159
         51-60      35018
         0-10       23604
         41-50      11743
         71-80      10254
         >100       6683
         81-90       4838
         91-100      2765
         61-70       2744
         Name: Stay, dtype: int64
```

```
In [17]: i=1
plt.figure(figsize=(15,20))
for col in number:
    plt.subplot(4,2,i)
    sns.distplot(train[col])
    i=i+1

plt.show()
```

C:\ProgramData\Anaconda3\lib\site-packages\seaborn\distributions.py:2619: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)

C:\ProgramData\Anaconda3\lib\site-packages\seaborn\distributions.py:2619: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)

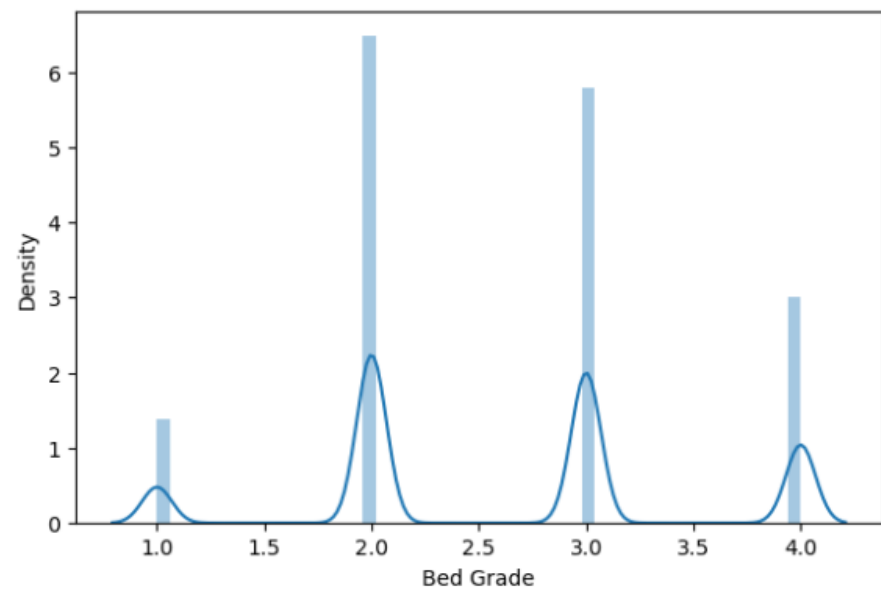
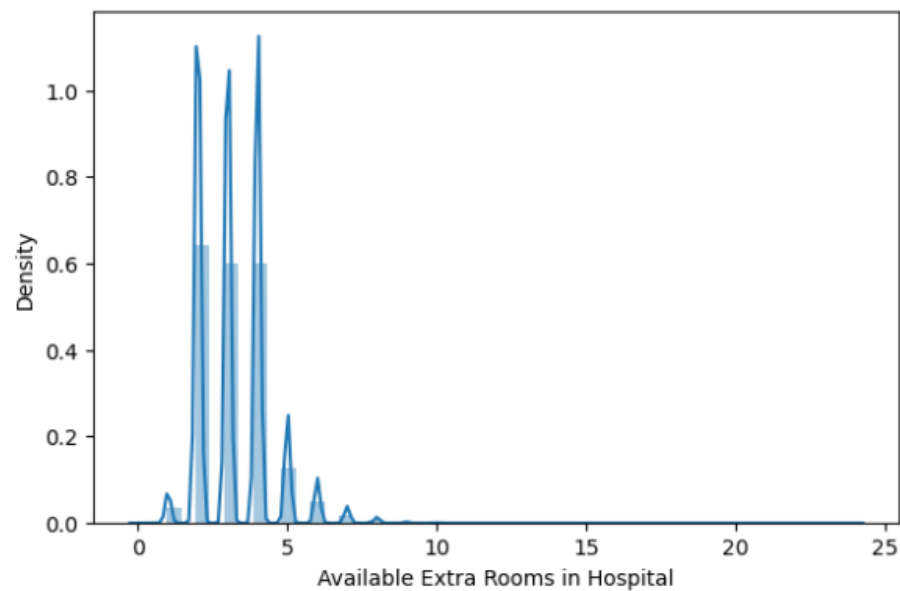
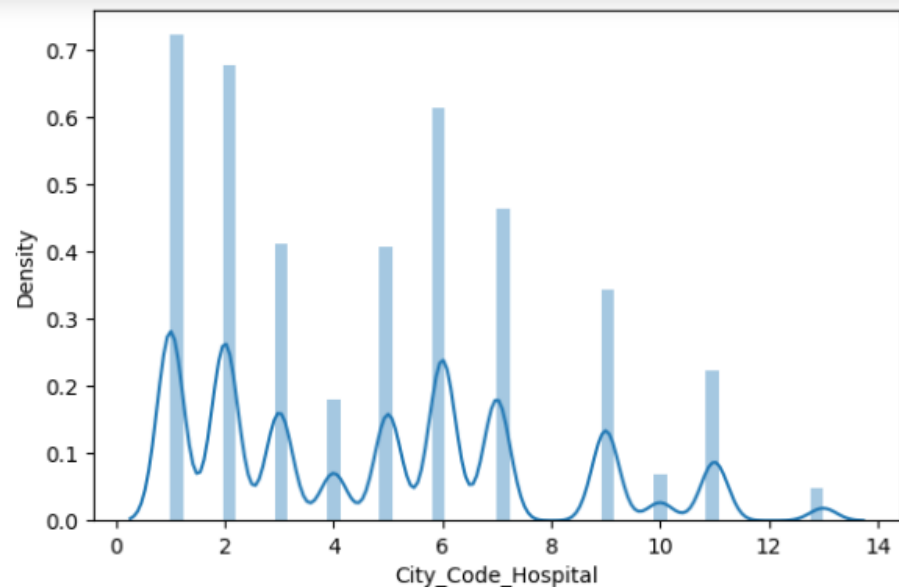
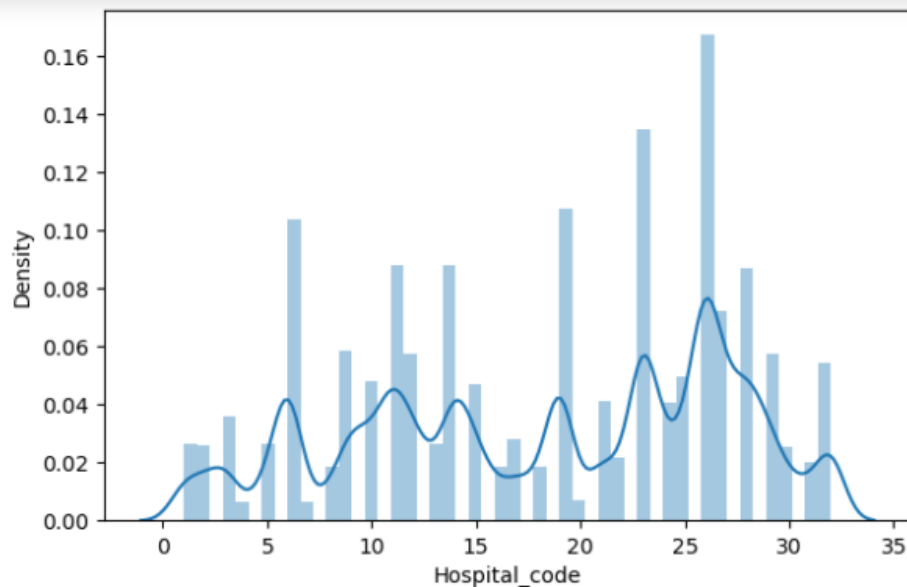
C:\ProgramData\Anaconda3\lib\site-packages\seaborn\distributions.py:2619: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

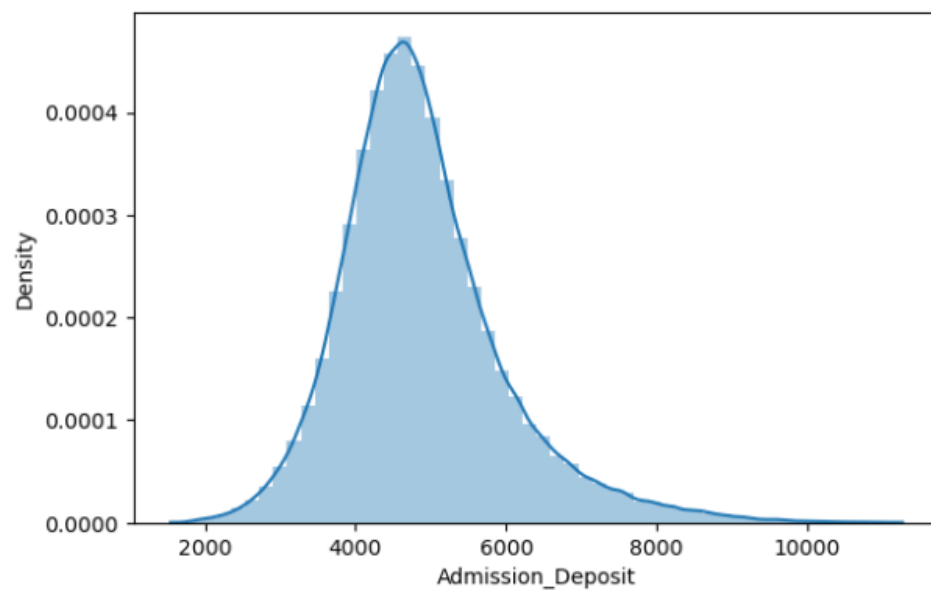
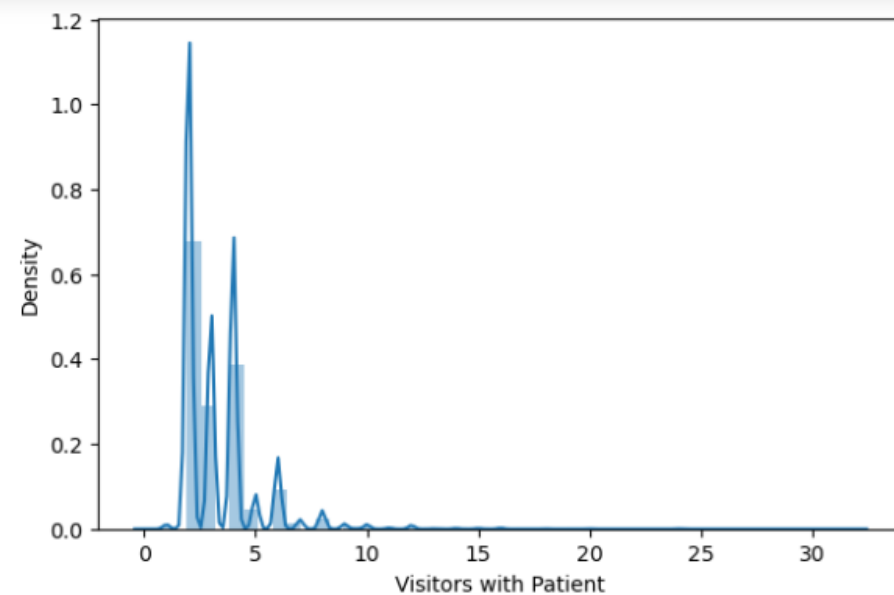
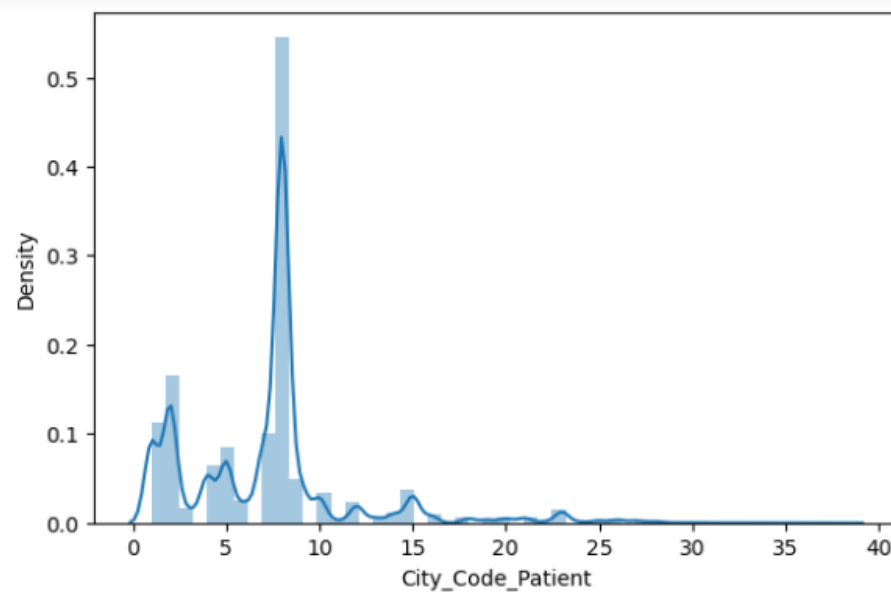
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Python 3 (ipykernel)

Run Code







```
In [19]: df_beds=train[['Hospital_code','Available Extra Rooms in Hospital','Hospital_region_code']]
df_beds['Hospital_region_code']=df_beds['Hospital_region_code'].map({'X':1,'Y':2,'Z':3})
df_beds_grouped=df_beds.groupby('Hospital_code')[['Available Extra Rooms in Hospital','Hospital_region_code']].median().reset_index()
```

C:\Users\Ranjan kumar\AppData\Local\Temp\ipykernel\_9944\3549231326.py:2: SettingWithCopyWarning:  
A value is trying to be set on a copy of a slice from a DataFrame.  
Try using .loc[row\_indexer,col\_indexer] = value instead

See the caveats in the documentation: [https://pandas.pydata.org/pandas-docs/stable/user\\_guide/indexing.html#returning-a-view-versus-a-copy](https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy)

```
df_beds['Hospital_region_code']=df_beds['Hospital_region_code'].map({'X':1,'Y':2,'Z':3})
```

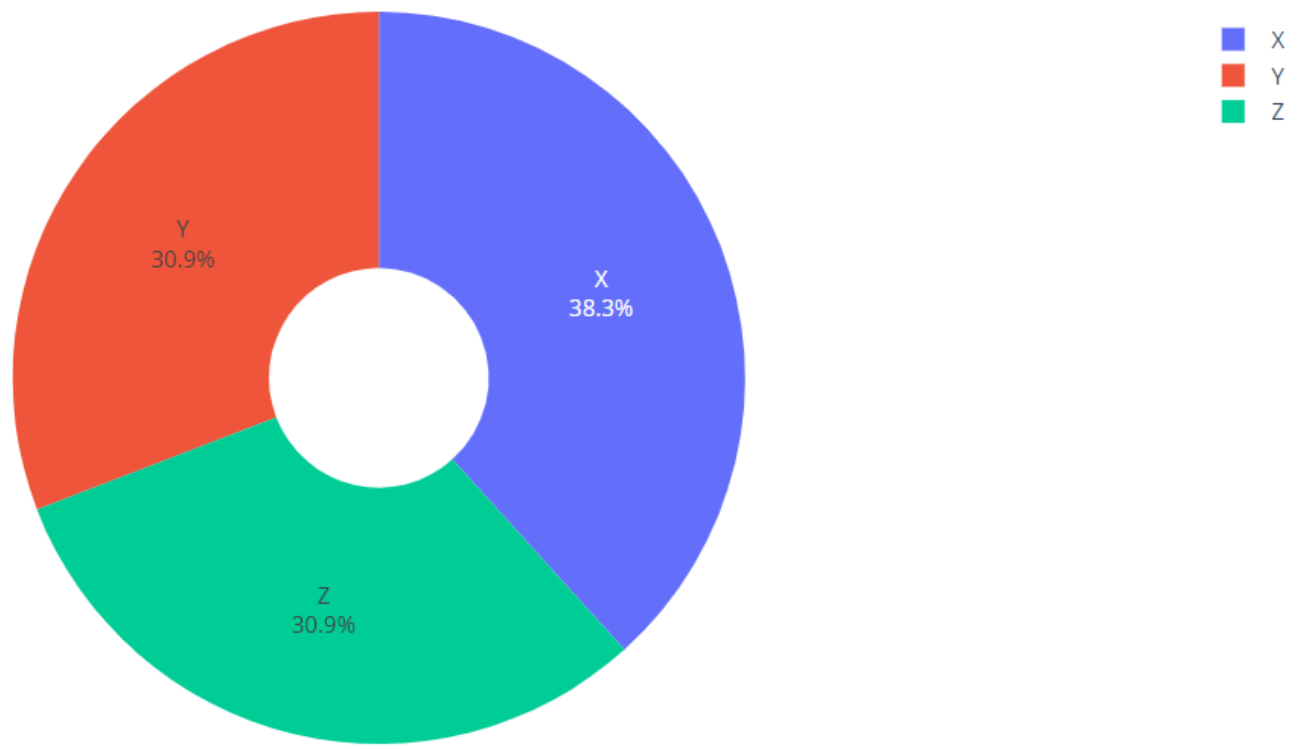
```
In [20]: df_beds_grouped['Hospital_region_code'].map({1:'X',2:'Y',3:'Z'})
```

```
Out[20]: 0      Y
1      Z
2      Z
3      X
4      X
5      X
6      X
7      Z
8      Z
9      X
10     Y
11     Y
12     Z
13     X
14     Z
15     Z
16     X
17     Y
18     Y
19     Y
20     Z
```

In [22]: `import plotly.express as px`

In [23]: `df_beds_1=df_beds_grouped.groupby('Hospital_region_code')['Available Extra Rooms in Hospital'].sum().reset_index()  
fig=px.pie(df_beds_1,values='Available Extra Rooms in Hospital',names='Hospital_region_code',hole=0.3)  
fig.update_layout(title='Number of extra rooms in each region code',title_x=0.5)  
fig.update_traces(textinfo='percent+label')`

Number of extra rooms in each region code

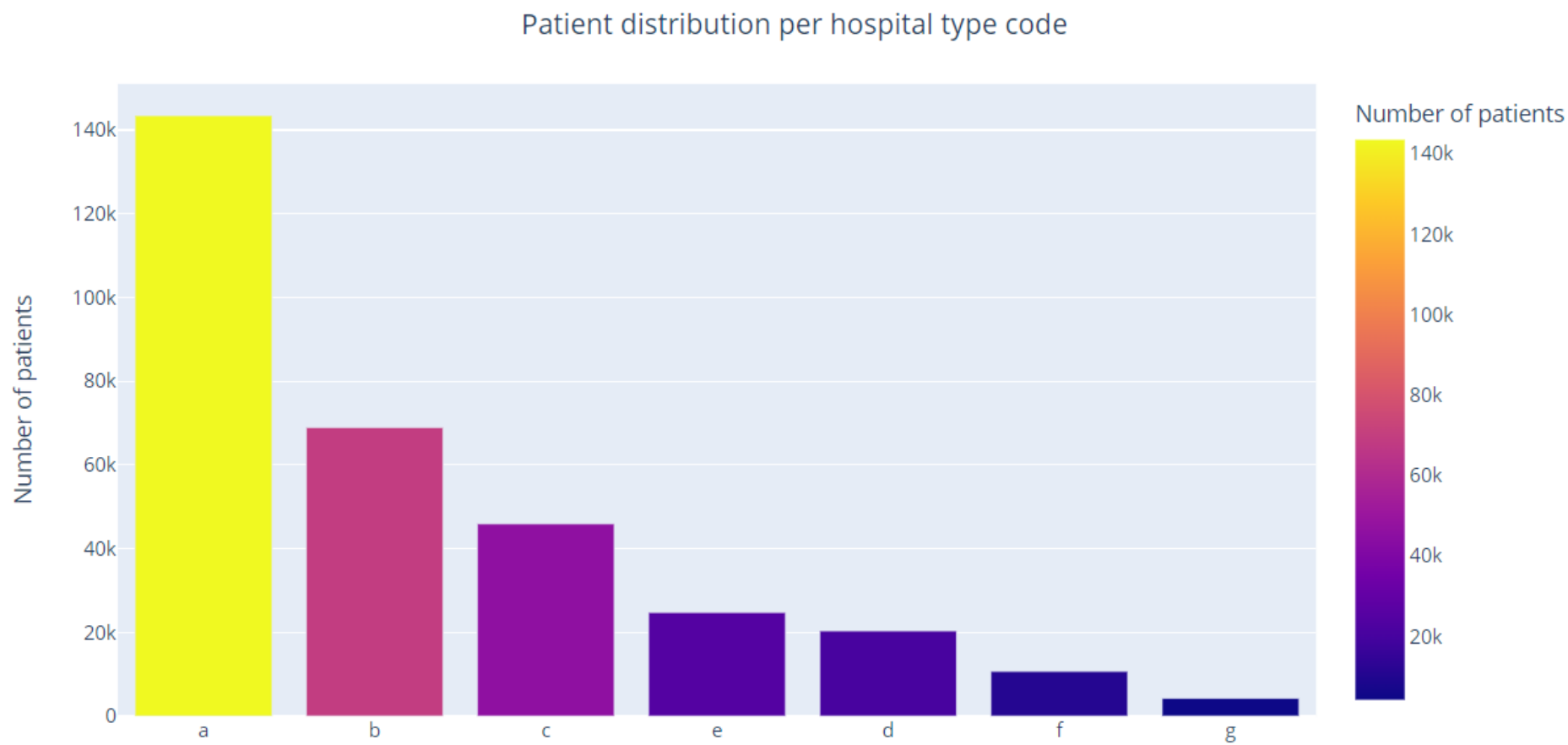




```
In [24]: train['train_flag']=1
```

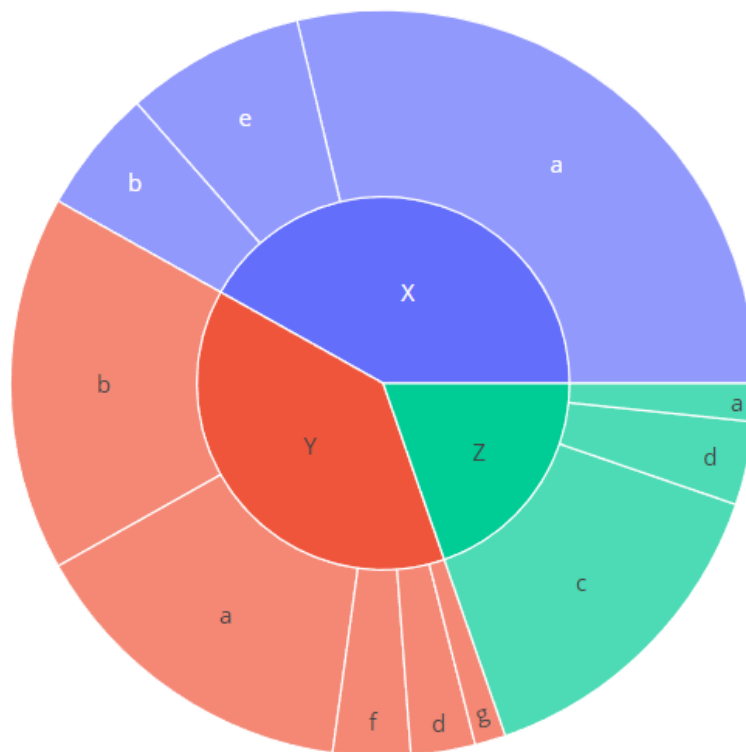
```
In [25]: train_index=train.groupby('Hospital_type_code')['train_flag'].sum().reset_index().sort_values(by='train_flag',ascending=False)
```

```
In [26]: plt.figure(figsize=(10,5)).bar(train_index,x='Hospital_type_code',y='train_flag',color='train_flag',labels={'Hospital_type_code':'Hospital Type Code','train_flag':'Number of patients'},title='Patient distribution per hospital type code',title_x=0.5)
```



```
In [27]: fig3=px.sunburst(train,path=['Hospital_region_code','Hospital_type_code'])
fig3.update_layout(title='Hospital region case load distribution',title_x=0.5)
fig3.show()
```

Hospital region case load distribution



```
91-100      2765
61-70      2744
Name: Stay, dtype: int64
```

```
In [29]: from sklearn.preprocessing import LabelEncoder
encoder = LabelEncoder()
for col in category:
    train[col] = encoder.fit_transform(train[col])
```

```
In [30]: train[category]
```

Out[30]:

	Hospital_type_code	Hospital_region_code	Department	Ward_Type	Ward_Facility_Code	Type of Admission	Severity of Illness	Age	Stay
0	2	2	3	2	5	0	0	5	0
1	2	2	3	3	5	1	0	5	4
2	4	0	1	3	4	1	0	5	3
3	1	1	3	2	3	1	0	5	4
4	1	1	3	3	3	1	0	5	4
...	...	...	...	...	...	...	...	...	...
318433	0	0	3	1	5	0	2	4	1
318434	0	0	1	1	4	2	2	8	3
318435	0	0	2	2	5	0	1	7	1
318436	1	1	1	1	3	1	1	1	1
318437	0	1	2	1	2	0	1	1	0

318438 rows × 9 columns

```
In [31]: train['City_Code_Hospital'].value_counts()
```

```
Out[31]: 1    55351
         2    51809
```

```
In [32]: from sklearn.preprocessing import StandardScaler
        scaler= StandardScaler()
        train[numer]= scaler.fit_transform(train[numer].values)
```

In [33]: train

Out[33]:

	Hospital_code	Hospital_type_code	City_Code_Hospital	Hospital_region_code	Available Extra Rooms in Hospital	Department	Ward_Type	Ward_Facility_Code	Bed Grade	City_C
0	-1.195176	2	-0.571055	2	-0.169177	3	2	5	-0.716535	
1	-1.890124	2	0.073580	2	-1.025217	3	3	5	-0.716535	
2	-0.963527	4	-1.215691	0	-1.025217	1	3	4	-0.716535	
3	0.889668	1	-0.893373	1	-1.025217	3	2	3	-0.716535	
4	0.889668	1	-0.893373	1	-1.025217	3	3	3	-0.716535	
...	...	...	...	...	...	...	...	...	...	...
318433	-1.426825	0	0.395897	0	-0.169177	3	1	5	1.574234	
318434	0.658018	0	-1.215691	0	-1.025217	1	1	4	1.574234	
318435	-1.311001	0	-0.248738	0	-0.169177	2	2	5	1.574234	
318436	-0.847702	1	-0.893373	1	-0.169177	1	1	3	0.428849	
318437	0.078895	0	0.718215	1	1.542903	2	1	2	-0.716535	

318438 rows × 17 columns

```
In [34]: from sklearn.model_selection import train_test_split
        y= train['Stay']
```



```
In [34]: from sklearn.model_selection import train_test_split
y= train['Stay']
X= train.drop('Stay', axis=1)
```

In [35]: X

Out[35]:

	Hospital_code	Hospital_type_code	City_Code_Hospital	Hospital_region_code	Available Extra Rooms in Hospital	Department	Ward_Type	Ward_Facility_Code	Bed Grade	City_Code_P
0	-1.195176	2	-0.571055	2	-0.169177	3	2	5	-0.716535	-0.0
1	-1.890124	2	0.073580	2	-1.025217	3	3	5	-0.716535	-0.0
2	-0.963527	4	-1.215691	0	-1.025217	1	3	4	-0.716535	-0.0
3	0.889668	1	-0.893373	1	-1.025217	3	2	3	-0.716535	-0.0
4	0.889668	1	-0.893373	1	-1.025217	3	3	3	-0.716535	-0.0
...	...	...	...	...	...	...	...	...	...	...
133	-1.426825	0	0.395897	0	-0.169177	3	1	5	1.574234	3.3
134	0.658018	0	-1.215691	0	-1.025217	1	1	4	1.574234	0.1
135	-1.311001	0	-0.248738	0	-0.169177	2	2	5	1.574234	0.5
136	-0.847702	1	-0.893373	1	-0.169177	1	1	3	0.428849	0.1
137	0.078895	0	0.718215	1	1.542903	2	1	2	-0.716535	0.1

38 rows × 16 columns

```
In [ ]: X_train, X_test, y_train,y_test= train_test_split(X,y,test_size= 0.2, stratify=y, random_state=42)
```



Code

134	0.658018	0	-1.215691	0	-1.025217	1	1	4	1.574234	0.1
135	-1.311001	0	-0.248738	0	-0.169177	2	2	5	1.574234	0.5
136	-0.847702	1	-0.893373	1	-0.169177	1	1	3	0.428849	0.1
137	0.078895	0	0.718215	1	1.542903	2	1	2	-0.716535	0.1

38 rows × 16 columns



```
In [ ]: X_train, X_test, y_train,y_test= train_test_split(X,y,test_size= 0.2, stratify=y, random_state=42)
```

```
In [ ]: from sklearn.linear_model import LogisticRegression
from sklearn.metrics import classification_report
from sklearn.metrics import accuracy_score
```

```
In [40]: reg = LogisticRegression()
reg.fit(X_train,y_train)

y_pred=reg.predict(X_test)
accuracy_score(y_test,y_pred)
```

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\linear\_model\\_logistic.py:814: ConvergenceWarning:

lbfgs failed to converge (status=1):  
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max\_iter) or scale the data as shown in:

<https://scikit-learn.org/stable/modules/preprocessing.html>

Please also refer to the documentation for alternative solver options:

[https://scikit-learn.org/stable/modules/linear\\_model.html#logistic-regression](https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression)

Out[40]: 0.3780932043713101

```
In [42]: print (classification_report(y_test,y_pred))
```





STOP: TOTAL NO. OF ITERATIONS REACHED LIMIT.

Increase the number of iterations (max\_iter) or scale the data as shown in:  
<https://scikit-learn.org/stable/modules/preprocessing.html>  
 Please also refer to the documentation for alternative solver options:  
[https://scikit-learn.org/stable/modules/linear\\_model.html#logistic-regression](https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression)

Out[40]: 0.3780932043713101

In [42]: `print (classification_report(y_test,y_pred))`

	precision	recall	f1-score	support
0	0.44	0.03	0.06	4721
1	0.37	0.40	0.38	15628
2	0.40	0.68	0.50	17498
3	0.33	0.18	0.23	11032
4	0.00	0.00	0.00	2349
5	0.35	0.49	0.41	7004
6	0.00	0.00	0.00	549
7	0.00	0.00	0.00	2051
8	0.00	0.00	0.00	967
9	0.00	0.00	0.00	553
10	0.54	0.29	0.38	1336
accuracy				0.38 63688
macro avg				0.22 0.19 0.18 63688
weighted avg				0.34 0.38 0.33 63688

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\metrics\\_classification.py:1318: UndefinedMetricWarning:

Precision and F-score are ill-defined and being set to 0.0 in labels with no predicted samples. Use `zero\_division` parameter to control this behavior.

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\metrics\\_classification.py:1318: UndefinedMetricWarning:



```
In [43]: from sklearn.ensemble import AdaBoostClassifier
from sklearn.metrics import confusion_matrix, classification_report, make_scorer, accuracy_score
from sklearn.model_selection import cross_val_score, train_test_split
```

```
In [44]: ada_classifier = AdaBoostClassifier(n_estimators = 19, random_state = 10)
ada_classifier.fit(X_train, y_train)
pred_ada = ada_classifier.predict(X_test)
accuracy_score(y_test, pred_ada)
```

Out[44]: 0.3595182765984173

```
In [45]: print (classification_report(y_test, pred_ada))
```

	precision	recall	f1-score	support
0	0.23	0.04	0.07	4721
1	0.33	0.32	0.33	15628
2	0.39	0.68	0.49	17498
3	0.32	0.19	0.24	11032
4	0.00	0.00	0.00	2349
5	0.35	0.47	0.40	7004
6	0.00	0.00	0.00	549
7	0.00	0.00	0.00	2051
8	0.16	0.02	0.04	967
9	0.00	0.00	0.00	553
10	0.47	0.30	0.37	1336
accuracy			0.36	63688
macro avg	0.20	0.18	0.18	63688
weighted avg	0.31	0.36	0.31	63688

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\metrics\\_classification.py:1318: UndefinedMetricWarning:

Precision and F-score are ill-defined and being set to 0.0 in labels with no predicted samples. Use `zero\_division` parameter to control this behavior.



```
In [46]: from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn import tree
from sklearn.model_selection import GridSearchCV
```

```
In [47]: deci= DecisionTreeClassifier(criterion = 'gini', max_depth=10, random_state = 42)

decision_tree = deci.fit(X_train, y_train)
```

```
In [48]: y_preddd = decision_tree.predict(X_test)
accuracy_score(y_test,y_preddd)
```

Out[48]: 0.4109879412134154

```
In [49]: print(accuracy_score(y_test,y_preddd))

0.4109879412134154
```

```
In [50]: print (classification_report(y_test,y_preddd ))
```

	precision	recall	f1-score	support
0	0.35	0.13	0.19	4721
1	0.42	0.46	0.44	15628
2	0.41	0.67	0.51	17498
3	0.41	0.25	0.31	11032
4	0.16	0.00	0.01	2349
5	0.41	0.45	0.43	7004
6	0.00	0.00	0.00	549
7	0.31	0.02	0.04	2051
8	0.35	0.20	0.25	967
9	0.18	0.01	0.02	553
10	0.54	0.39	0.45	1336
accuracy				0.41 63688
macro avg				0.32 0.23 0.24 63688
weighted avg				0.39 0.41 0.38 63688



```
In [54]: Random = RandomForestClassifier(criterion = 'entropy', n_estimators = 61, random_state = 142)

random_model =Random.fit(X_train, y_train)
```

```
In [55]: y_predddd = random_model.predict(X_test)
accuracy_score(y_test,y_predddd)
```

```
Out[55]: 0.38200288908428587
```

```
In [56]: print (classification_report(y_test,y_predddd ))
```

	precision	recall	f1-score	support
0	0.28	0.19	0.22	4721
1	0.39	0.44	0.41	15628
2	0.41	0.54	0.47	17498
3	0.34	0.27	0.30	11032
4	0.08	0.02	0.03	2349
5	0.39	0.44	0.41	7004
6	0.09	0.02	0.03	549
7	0.27	0.10	0.15	2051
8	0.37	0.22	0.27	967
9	0.27	0.05	0.09	553
10	0.54	0.46	0.49	1336
accuracy				0.38 63688
macro avg				0.31 0.25 0.26 63688
weighted avg				0.36 0.38 0.37 63688

```
In [57]: import xgboost as xgb
```

```
In [58]: xg=xgb.XGBClassifier(max_depth=11, objective='multi:softmax', n_estimators=59,random_state=42)
xg.fit(X_train,y_train)
xgbpred=xgbcl.predict(X_test)
```



```
In [54]: Random = RandomForestClassifier(criterion = 'entropy', n_estimators = 61, random_state = 142)
        random_model =Random.fit(X_train, y_train)
```

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10	0.54	0.46	0.49	1336
accuracy			0.38	63688
macro avg	0.31	0.25	0.26	63688
weighted avg	0.36	0.38	0.37	63688

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        xg.fit(X_train,y_train)
        xgbpred=xgbcl.predict(X_test)
```

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import statistics
```

```
In [2]: train=pd.read_csv('C:/Users/Ranjan kumar/Downloads/healthcare/train_data.csv')
```

```
In [3]: train.head(5)
```

Out[3]:

	case_id	Hospital_code	Hospital_type_code	City_Code_Hospital	Hospital_region_code	Available Extra Rooms in Hospital	Department	Ward_Type	Ward_Facility_Code	Bed Grade	patient
0	1	8	c	3	Z	3	radiotherapy	R	F	2.0	313
1	2	2	c	5	Z	2	radiotherapy	S	F	2.0	313
2	3	10	e	1	X	2	anesthesia	S	E	2.0	313
3	4	26	b	2	Y	2	radiotherapy	R	D	2.0	313
4	5	26	b	2	Y	2	radiotherapy	S	D	2.0	313

```
In [4]: test=pd.read_csv('C:/Users/Ranjan kumar/Downloads/healthcare/test_data.csv')
```

```
In [5]: train.describe()
```

Out[5]:

e	1	X	2	anesthesia	S	E	2.0	31397	7.0	Trauma	Extreme
b	2	Y	2	radiotherapy	R	D	2.0	31397	7.0	Trauma	Extreme
b	2	Y	2	radiotherapy	S	D	2.0	31397	7.0	Trauma	Extreme

```
In [4]: test=pd.read_csv('C:/Users/Ranjan kumar/Downloads/healthcare/test_data.csv')
```

```
In [5]: test.head(5)
```

Out[5]:

	case_id	Hospital_code	Hospital_type_code	City_Code_Hospital	Hospital_region_code	Available Extra Rooms in Hospital	Department	Ward_Type	Ward_Facility_Code	Bed Grade	patient
0	318439	21	c	3	Z	3	gynecology	S	A	2.0	1700
1	318440	29	a	4	X	2	gynecology	S	F	2.0	1700
2	318441	26	b	2	Y	3	gynecology	Q	D	4.0	1700
3	318442	6	a	6	X	3	gynecology	Q	F	2.0	1700
4	318443	28	b	11	X	2	gynecology	R	F	2.0	1700



weighted avg 0.39 0.41 0.38 63688

```
In [*]: Random = RandomForestClassifier(criterion = 'entropy', n_estimators = 61, random_state = 142)

random_model =Random.fit(X_train, y_train)
```

```
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```
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In [*]: xg=xgb.XGBClassifier(max_depth=11, objective='multi:softmax', n_estimators=59,random_state=42)
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xgbpred=xgbcl.predict(X_test)
```

```
In [*]: accuracy_score(y_test,xgbpred)
```

```
In [*]: print (classification_report(y_test,xgbpred ))
```

```
In [ ]:
```

```
In [ ]:
```