SmartInternz Externships

Applied Data Science

Assignment: 2

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Campus: VIT-AP

In [1]: import numpy as np
 import pandas as pd
 import seaborn as sns

2) Load the dataset

```
In [2]: df = pd.read_csv("titanic.csv")
```

In [3]: df.head()

Out[3]:		survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_tov
	0	0	3	male	22.0	1	0	7.2500	S	Third	man	True	NaN	Southampto
	1	1	1	female	38.0	1	0	71.2833	С	First	woman	False	С	Cherbou
	2	1	3	female	26.0	0	0	7.9250	S	Third	woman	False	NaN	Southampto
	3	1	1	female	35.0	1	0	53.1000	S	First	woman	False	С	Southampto
	4	0	3	male	35.0	0	0	8.0500	S	Third	man	True	NaN	Southampto

In [4]: df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 891 entries, 0 to 890
Data columns (total 15 columns):

#	Column	Non-Null Cou	nt Dtype
0	survived	891 non-null	int64
1	pclass	891 non-null	int64
2	sex	891 non-null	object
3	age	714 non-null	float64
4	sibsp	891 non-null	int64
5	parch	891 non-null	int64
6	fare	891 non-null	float64
7	embarked	889 non-null	object
8	class	891 non-null	object
9	who	891 non-null	object
10	adult_male	891 non-null	bool
11	deck	203 non-null	object
12	embark_town	889 non-null	object
13	alive	891 non-null	object
14	alone	891 non-null	bool

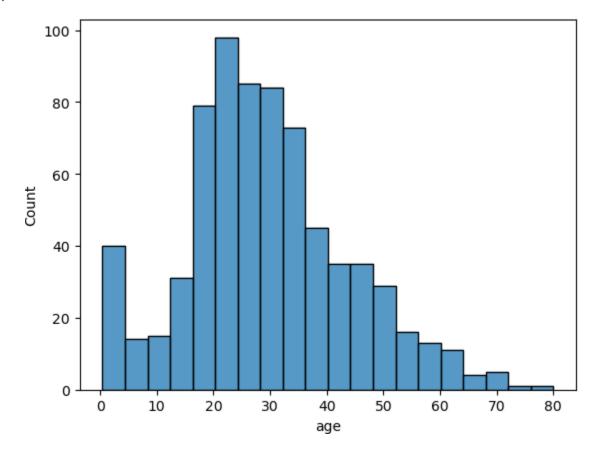
dtypes: bool(2), float64(2), int64(4), object(7)
memory usage: 92.4+ KB

3. Perform Below Visualizations.

- Univariate Analysis
- Bi Variate Analysis
- Multi Variate Analysis

univariate analysis

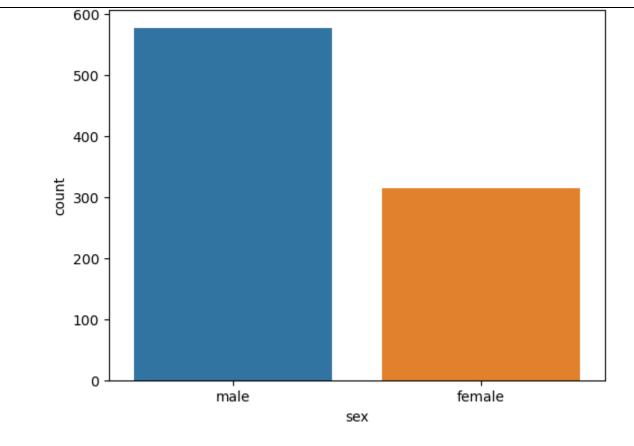
```
In [5]: sns.histplot(df['age'])
Out[5]: <Axes: xlabel='age', ylabel='Count'>
```



```
sns.countplot(x = df['sex'])
```

In [6]: <Axes: xlabel='sex', ylabel='count'>

Out[6]:

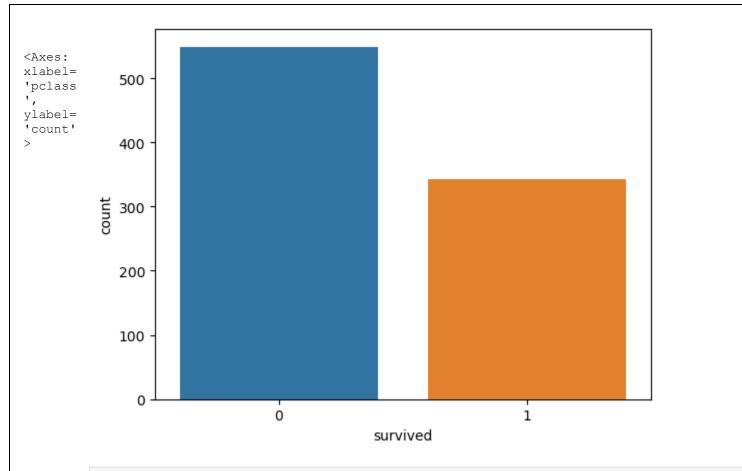


```
In [7]: sns.countplot(x = df['survived'])
```

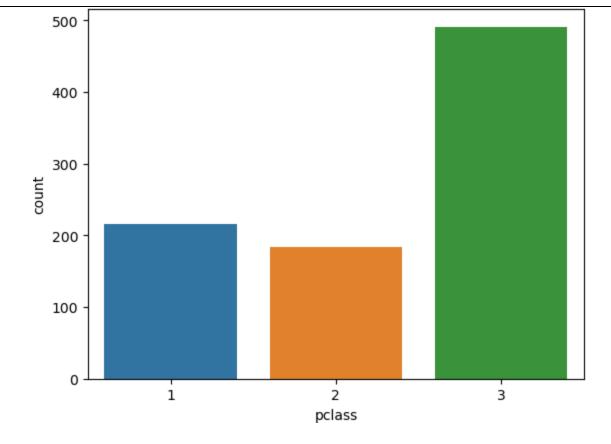
Out[7]: <Axes: xlabel='survived', ylabel='count'>

In [8]:

Out[8]:



sns.countplot(x = df['pclass'])



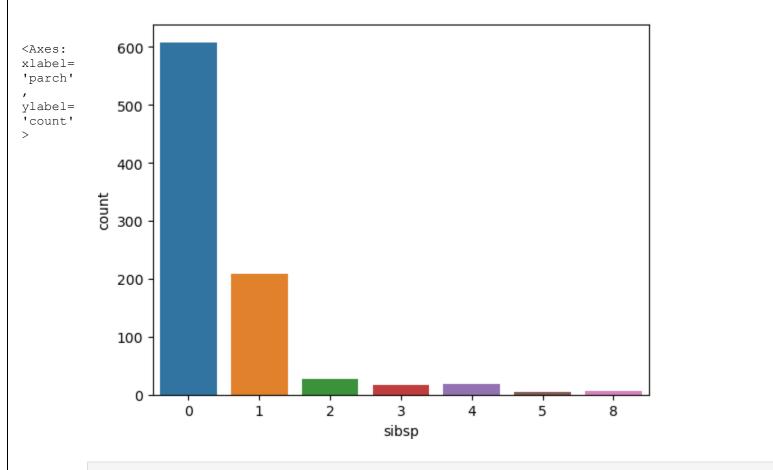
```
In [9]: sns.countplot(x = df['sibsp'])

<Axes: xlabel='sibsp', ylabel='count'>
```

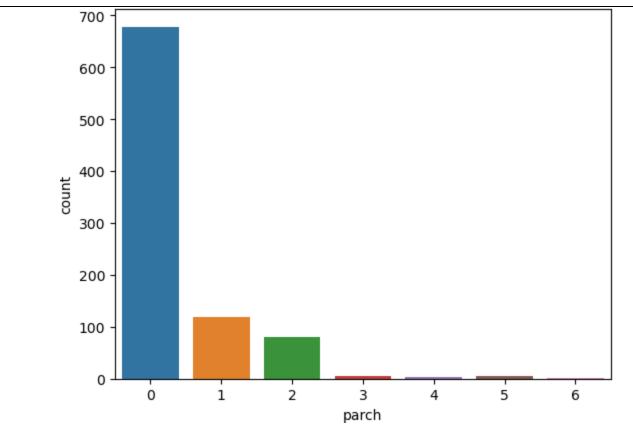
Out[9]: <Axes: xlabel='sibsp', ylabel='count'>

In [10]:

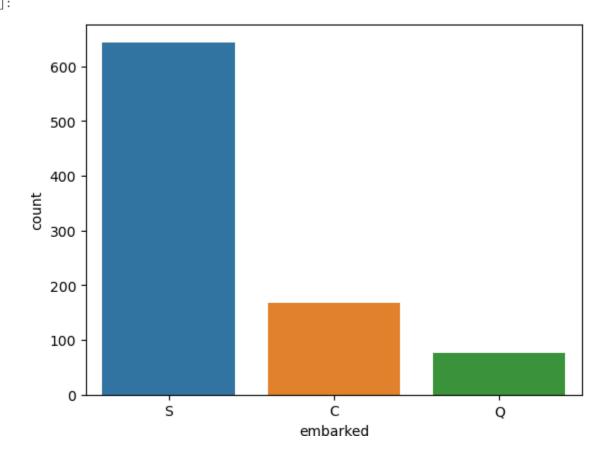
Out[10]:



sns.countplot(x = df['parch'])



```
In [11]: sns.countplot(x = df['embarked'])
Out[11]: <Axes: xlabel='embarked', ylabel='count'>
```

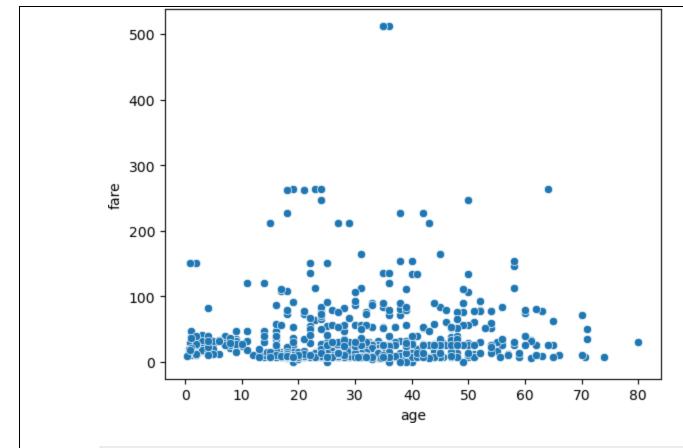


bivariate analysis

In [12]:

Out[12]:

	<pre>sns.scatterplot(data = df, x = 'age', y = 'fare')</pre>											
<axes:< th=""><th></th></axes:<>												
<pre>xlabel= 'age',</pre>												
ylabel= 'fare'>												
'fare'>												

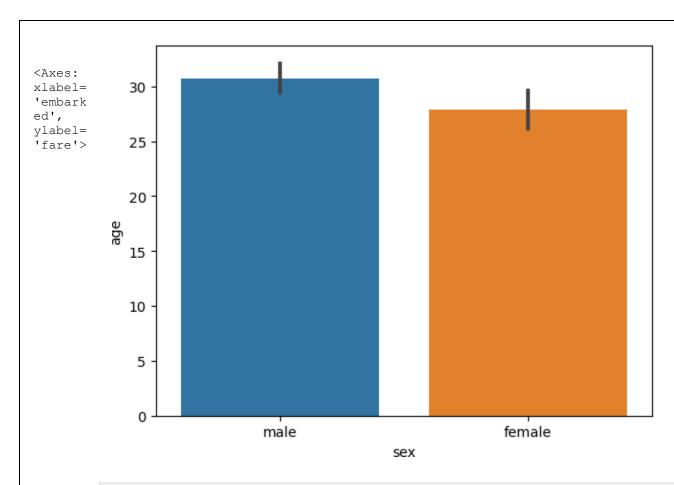


```
In [13]: sns.barplot(data = df, x = 'sex', y = 'age')
```

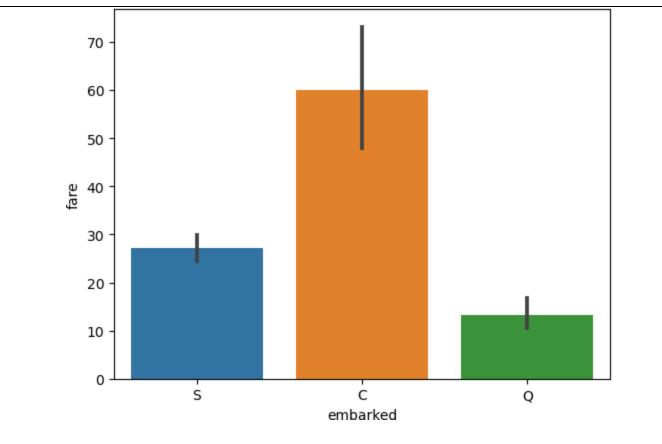
Out[13]: <Axes: xlabel='sex', ylabel='age'>

In [14]:

Out[14]:



```
sns.barplot(data = df, x = 'embarked', y = 'fare')
```

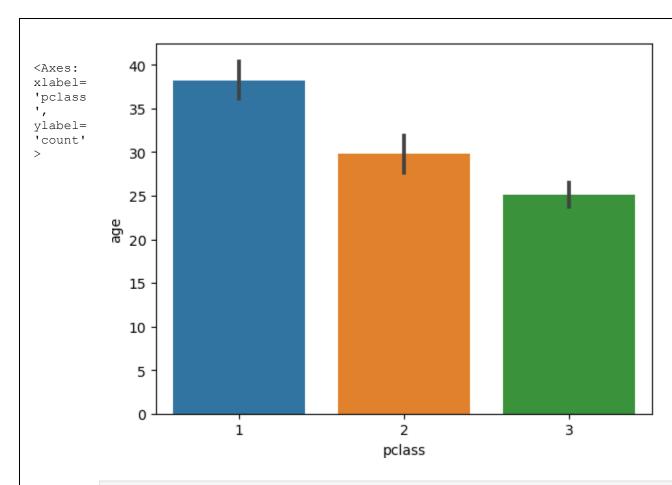


```
In [15]: sns.barplot(data = df, x = 'pclass', y = 'age')
```

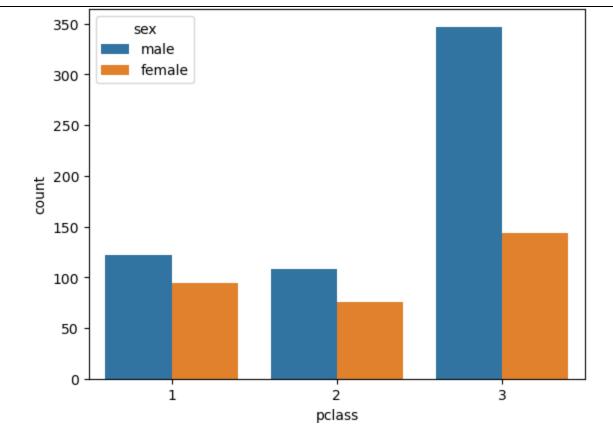
Out[15]: <Axes: xlabel='pclass', ylabel='age'>

In [16]:

Out[16]:



```
sns.countplot(x = df['pclass'], hue = df['sex'])
```

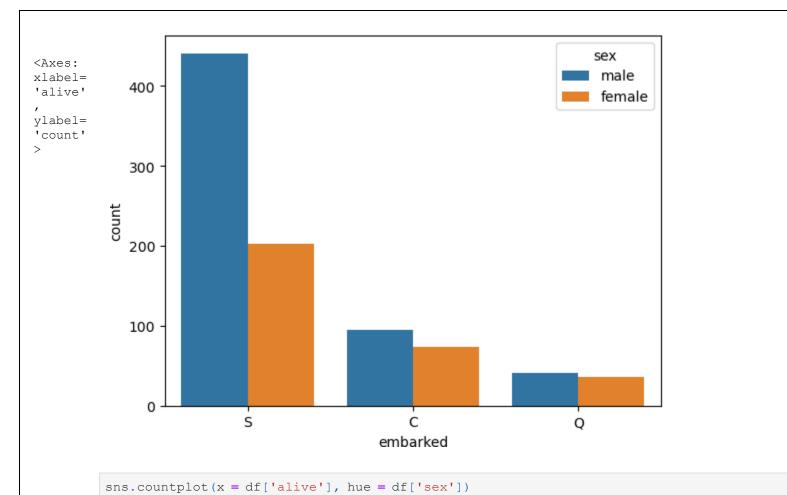


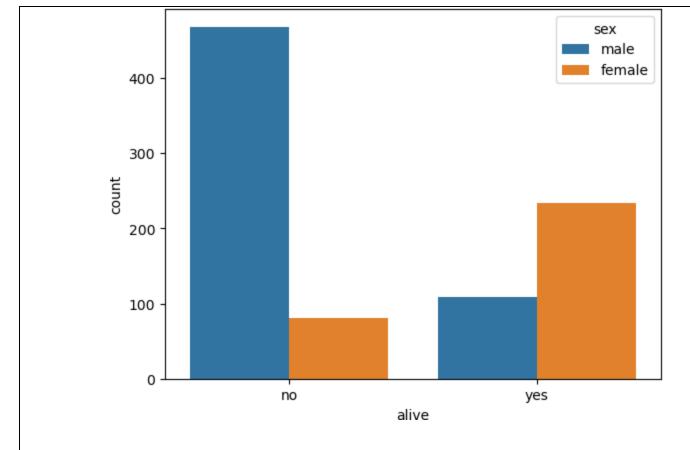
```
sns.countplot(x = df['embarked'], hue = df['sex'])
In [17]:
         <Axes: xlabel='embarked', ylabel='count'>
```

Out[17]:

In [18]:

Out[18]:

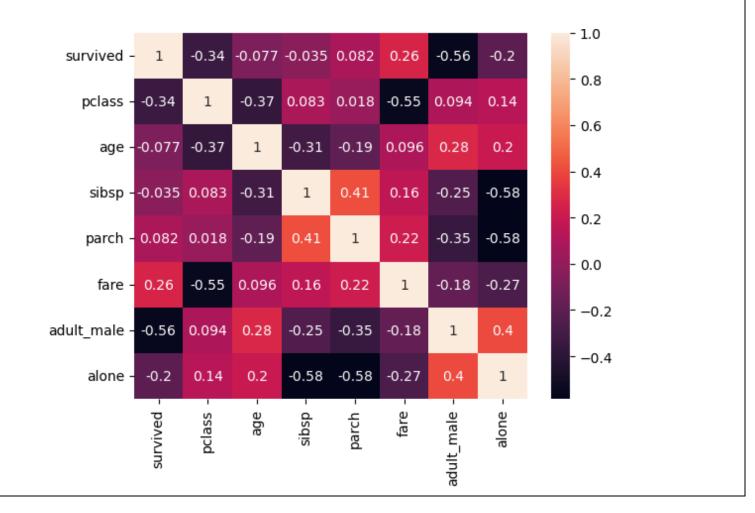




multivariate analysis

In [19]: sns.heatmap(df.corr(numeric_only=True), annot = True)

Out[19]: <Axes: >

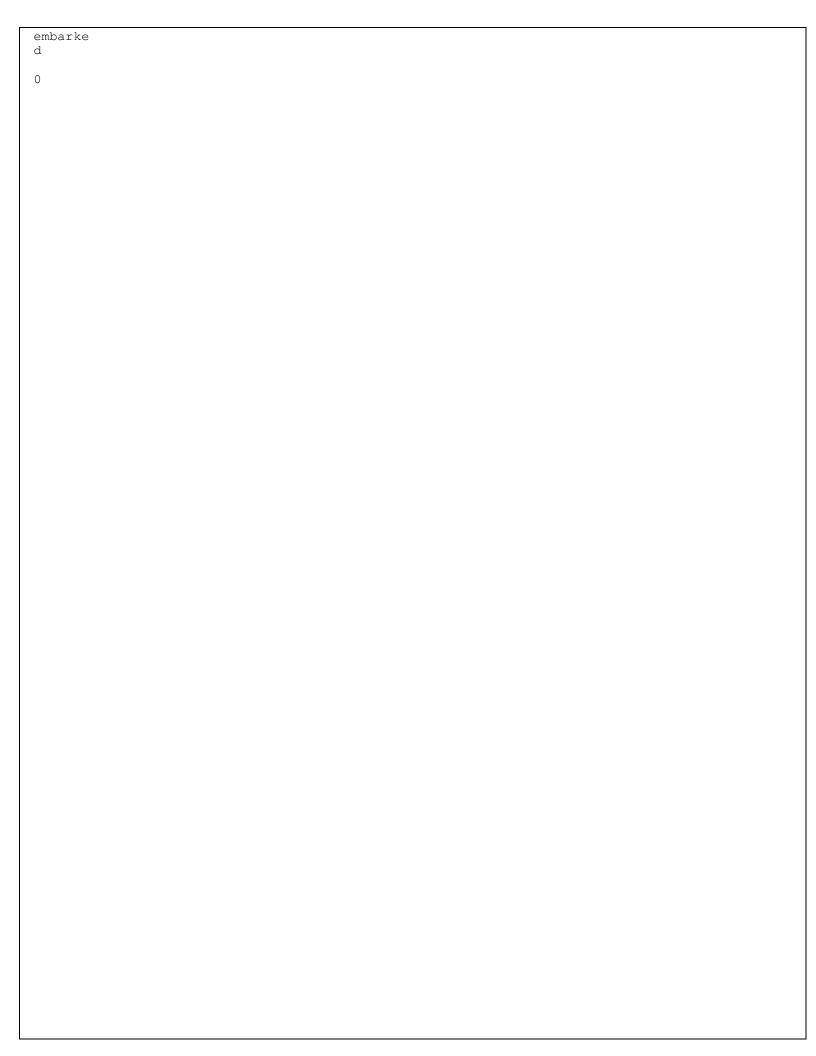


4. Perform descriptive statistics on the dataset.

In [20]:	df.de	scribe()					
Out[20]:		survived	pclass	age	sibsp	parch	fare
	count	891.000000	891.000000	714.000000	891.000000	891.000000	891.000000
	mean	0.383838	2.308642	29.699118	0.523008	0.381594	32.204208
	std	0.486592	0.836071	14.526497	1.102743	0.806057	49.693429
	min	0.000000	1.000000	0.420000	0.000000	0.000000	0.000000
	25%	0.000000	2.000000	20.125000	0.000000	0.000000	7.910400
	50%	0.000000	3.000000	28.000000	0.000000	0.000000	14.454200
	75%	1.000000	3.000000	38.000000	1.000000	0.000000	31.000000
	max	1.000000	3.000000	80.000000	8.000000	6.000000	512.329200

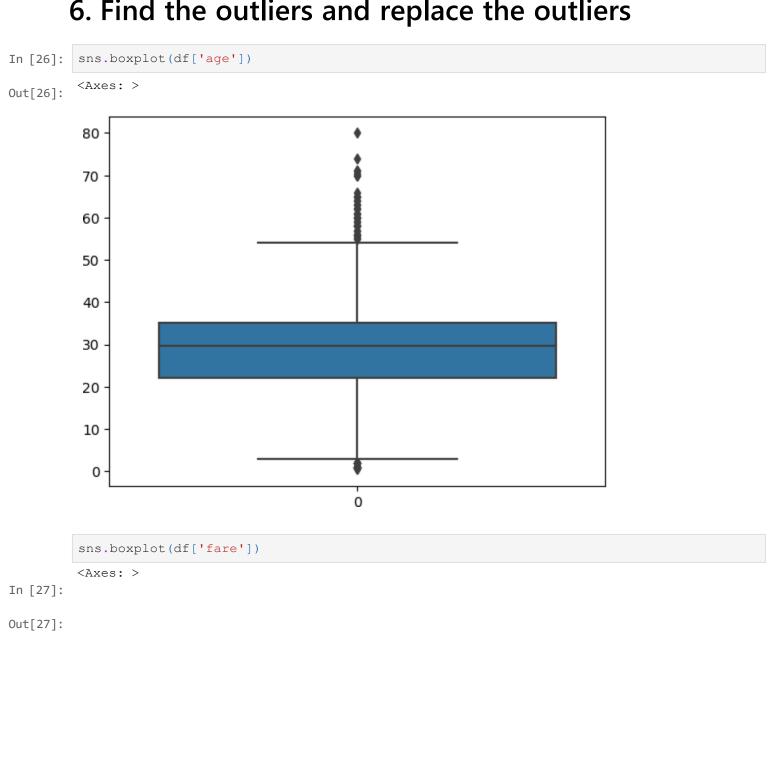
5. Handle the Missing values.

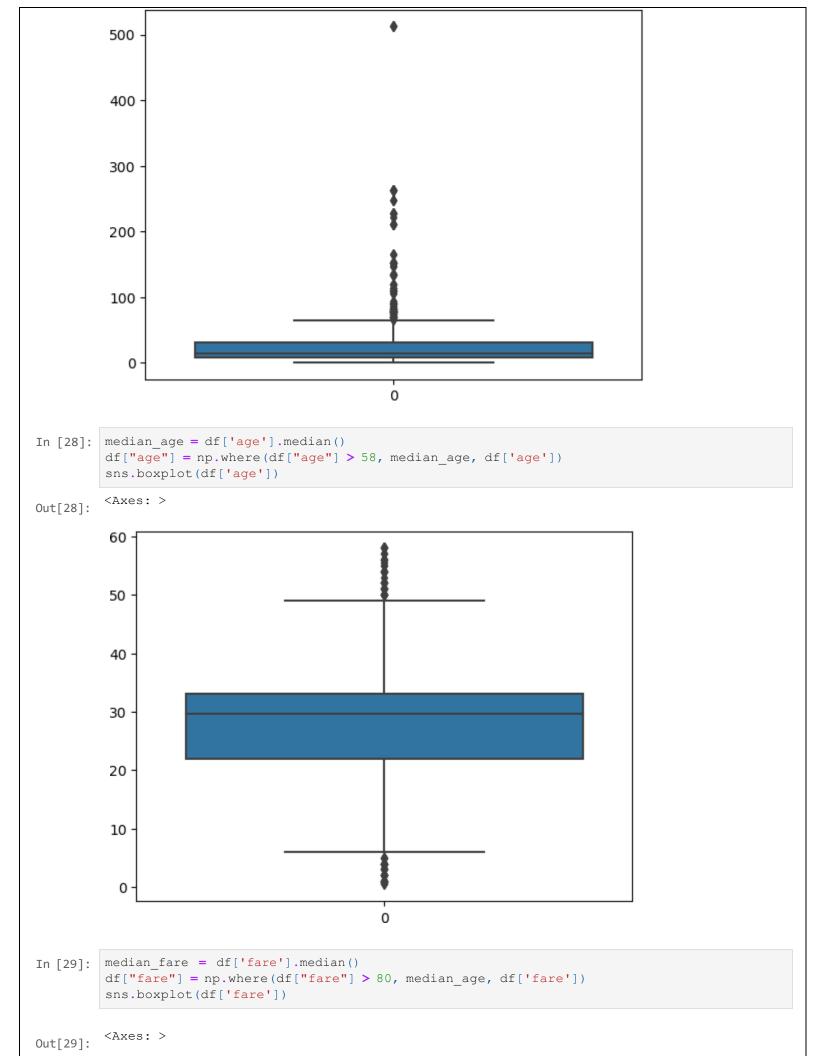
```
In [21]: | df.isnull().sum()
        survived
Out[21]:
        pclass
        sex
                      177
        age
        sibsp
        parch
                        0
        fare
        embarked
        class
        who
                      0
        adult_male
        deck
                      688
                      2
        embark_town
                        0
        alive
        alone
        dtype: int64
In [22]: df.dropna(subset=['embark_town'], how='all', inplace = True)
In [23]: | #for age column we will fill with the average
         df['age'] = df['age'].fillna(df['age'].mean())
In [24]: #only 203 records have valid values for deck column so we will drop that
        df.drop(['deck'], axis = 1,inplace = True)
In [25]: df.isnull().sum()
                       0
        survived
Out[25]:
                       0
        pclass
        sex
                       0
        age
        sibsp
                       0
        parch
        fare
```

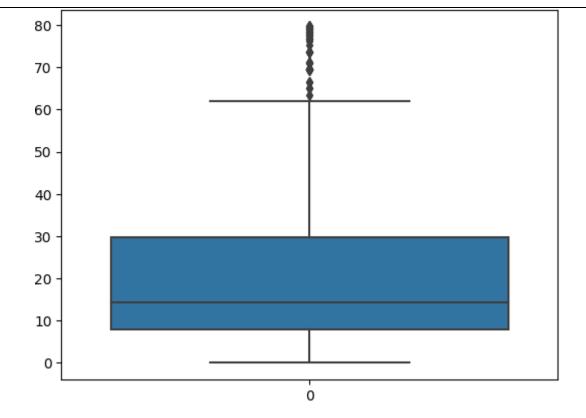


```
class
who
adult male
embark_town
alive
               0
alone
dtype: int64
```

6. Find the outliers and replace the outliers







7. Check for Categorical columns and perform encoding.

In [30]:	<pre>from sklearn.preprocessing import OneHotEncoder</pre>													
In [31]:	<pre>encoding = pd.get_dummies(df, columns = ['sex','embarked','class','who','adult_male', 'a</pre>													', 'a
In [32]:	encoding.head()													
Out[32]:	\$	survived	pclass	age	sibsp	parch	fare	alive	sex_female	sex_male	embarked_C	•••	who_child	who_m
	0	0	3	22.0	1	0	7.2500	no	0	1	0		0	
	1	1	1	38.0	1	0	71.2833	yes	1	0	1		0	
	2	1	3	26.0	0	0	7.9250	yes	1	0	0		0	
	3	1	1	35.0	1	0	53.1000	yes	1	0	0		0	
	4	0	3	35.0	0	0	8.0500	no	0	1	0		0	

5 rows × 25 columns

8. Split the data into dependent and independent variables

```
# independent variables
In [34]:
         X = encoding.drop(['survived', 'alive'], axis = 1)
         X.head()
Out[34]:
            pclass age sibsp parch
                                    fare sex_female sex_male embarked_C embarked_Q embarked_S ... who_chi
               3 22.0
                                   7.2500
                                                0
               1 38.0
                               0 71.2833
         2
                                                                    0
               3 26.0
                                   7.9250
                                                1
                                                         0
                                                                               0
               1 35.0
                               0 53.1000
                         0
                                   8.0500
                                                0
                                                                    0
                                                                               0
               3 35.0
                                                         1
                                                                                          1 ...
        5 rows × 23 columns
In [35]:
         # dependent variables
         y = df[['survived', 'alive']]
         y.head()
Out[35]:
            survived alive
                 0
                     no
                 1
                     yes
                     yes
                     yes
                 0
                     no
        9. Scaling the independent variables
In [36]:
         from sklearn.preprocessing import StandardScaler
         scaler = StandardScaler()
         x_std = scaler.fit_transform(X)
In [37]:
         x std
         array([[ 0.82520863, -0.57985934, 0.43135024, ..., -0.48271079,
Out[37]:
                 -0.30794088, 0.61679395],
                [-1.57221121, 0.83108889, 0.43135024, ..., 2.07163382,
                 -0.30794088, -1.62128697],
```

[0.82520863, -0.22712228, -0.47519908, ..., -0.48271079,

[0.82520863, 0.09405298, 0.43135024, ..., -0.48271079,

[-1.57221121, -0.22712228, -0.47519908, ..., 2.07163382,

[0.82520863, 0.3019833, -0.47519908, ..., -0.48271079,

-0.30794088, 0.61679395],

-0.30794088, 0.61679395],

-0.30794088, -1.62128697],

3.24737656, -1.62128697]])

'alone'],
dtype='object')

10. Split the data into training and testing

```
In [38]: from sklearn.model_selection import train_test_split
    X_train, X_test, y_train, y_test = train_test_split(X, y['survived'], test_size=0.33, ra

In []:
```