In [4]:
```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
```

In [5]:
```
col_names  =  ['Sepal_Length','Sepal_Width','Petal_Length','Petal_Width','Species']
```

In [6]:
```
iris  =  pd.read_csv('C:/Users/Lenovo/Downloads/iris.csv',names=col_names)
```

In [7]:
```
iris
```

Out[7]:

	Sepal_Length	Sepal_Width	Petal_Length	Petal_Width	Species
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa
...	...	...	...	...	...
145	6.7	3.0	5.2	2.3	Iris-virginica
146	6.3	2.5	5.0	1.9	Iris-virginica
147	6.5	3.0	5.2	2.0	Iris-virginica
148	6.2	3.4	5.4	2.3	Iris-virginica
149	5.9	3.0	5.1	1.8	Iris-virginica

150 rows × 5 columns

In [8]:
```
len(list(iris))
```

Out[8]:
```
5
```

In [9]:
```
iris.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
 #   Column        Non-Null Count  Dtype  
---  ------------------  -----  
 0   Sepal_Length  150 non-null    float64
 1   Sepal_Width   150 non-null    float64
 2   Petal_Length  150 non-null    float64
 3   Petal_Width   150 non-null    float64
 4   Species       150 non-null    object 
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

In [10]:
```
np.unique(iris["Species"])
```

Out[10]:
```
array(['Iris-setosa', 'Iris-versicolor', 'Iris-virginica'], dtype=object)
```

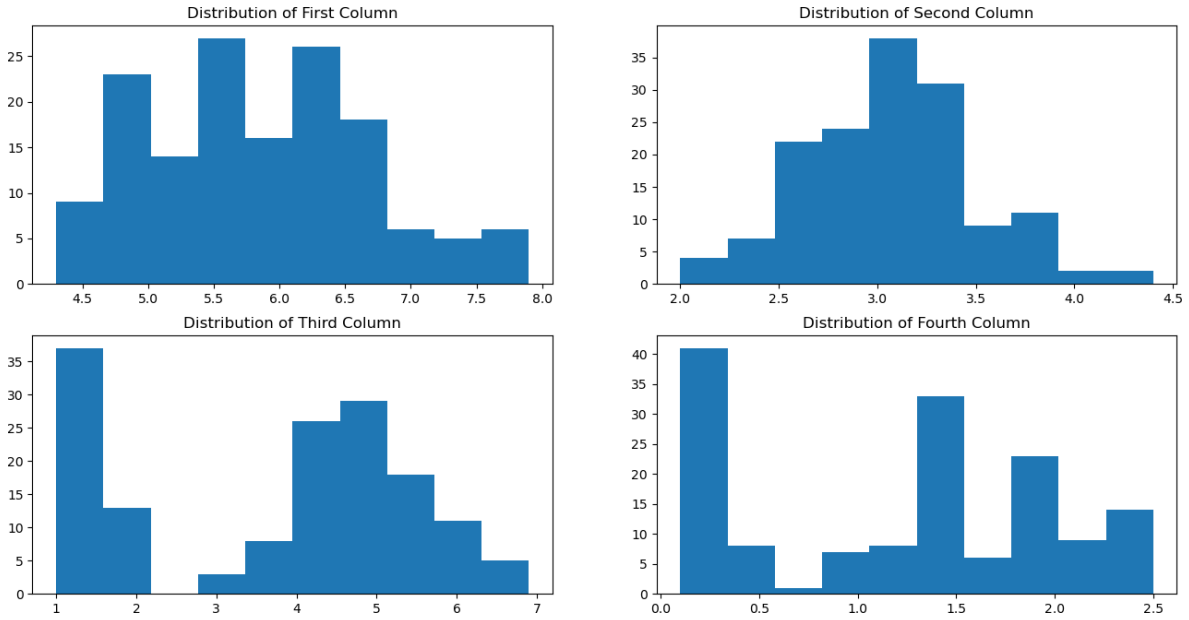
In [11]:
```
iris.describe()
```

Out[11]:

	Sepal_Length	Sepal_Width	Petal_Length	Petal_Width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.054000	3.758667	1.198667
std	0.828066	0.433594	1.764420	0.763161
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.350000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

In [12]:
```
import matplotlib.pyplot as plt
%matplotlib inline
```

In [13]:
```
fig, axes = plt.subplots(2, 2, figsize=(16, 8))
axes[0,0].set_title("Distribution of First Column")
axes[0,0].hist(iris["Sepal_Length"]);
axes[0,1].set_title("Distribution of Second Column")
axes[0,1].hist(iris["Sepal_Width"]);
axes[1,0].set_title("Distribution of Third Column")
axes[1,0].hist(iris["Petal_Length"]);
axes[1,1].set_title("Distribution of Fourth Column")
axes[1,1].hist(iris["Petal_Width"]);
```

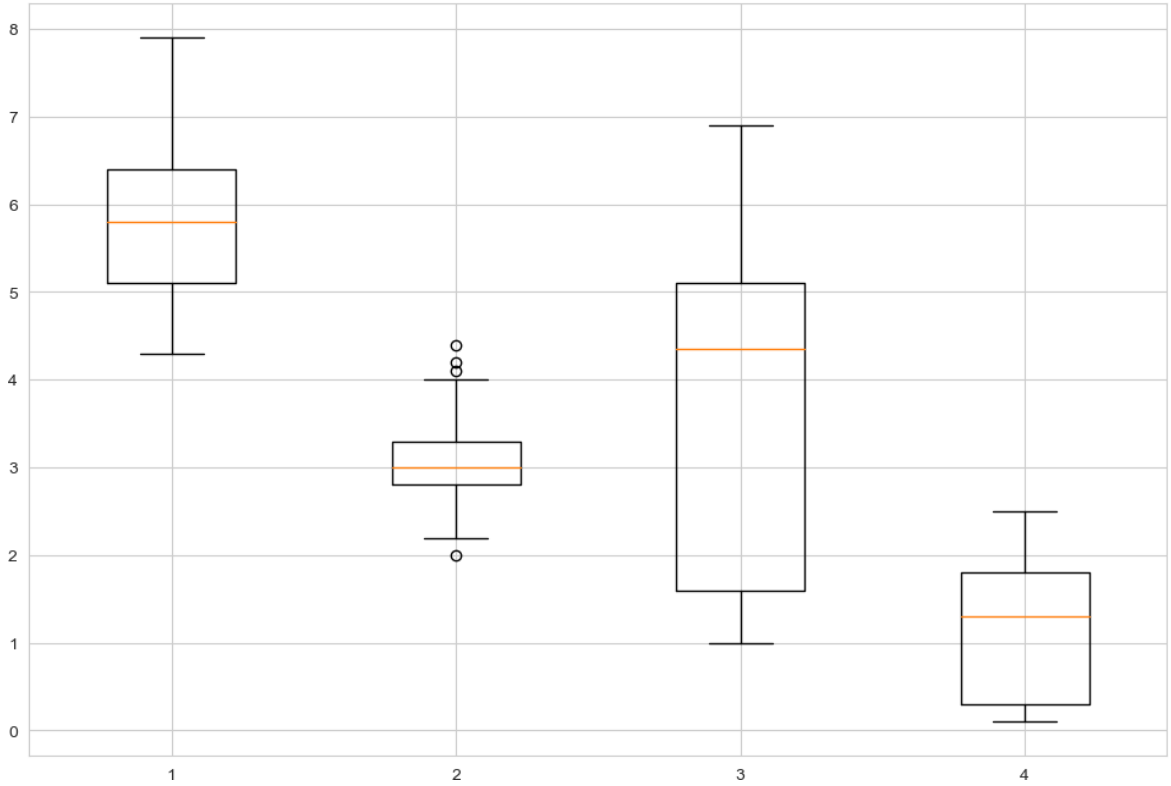


In [14]:
```
data_to_plot = [iris["Sepal_Length"],iris["Sepal_Width"],iris["Petal_Length"],iris
sns.set_style("whitegrid")

fig = plt.figure(1,

figsize=(12,8)) ax =

fig.add_subplot(111)
```



In [15]:
```
from scipy import stats
```

In [16]:
```
z = np.abs(stats.zscore(iris['Sepal_Length']))
```

In [17]:
```
print(z)
```

```
0      0.900681
1      1.143017
2      1.385353
3      1.506521
4      1.021849
         ...
145    1.038005
146    0.553333
147    0.795669
148    0.432165
149    0.068662
Name: Sepal_Length, Length: 150, dtype:
      float64
```

In [18]:
```
threshold = 0.5
```

In [19]:
```
sample_outliers = np.where(z <threshold)
sample_outliers
```

Out[19]:
```
(array([ 14,  15,  18,  33,  36,  53,  55,  61,  62,  63,  64,  66,  67,
         68,  69,  70,  71,  73,  78,  79,  80,  81,  82,  83,  85,  88,
         89,  90,  91,  92,  94,  95,  96,  97,  99, 101, 113, 114, 119,
        121, 126, 127, 134, 138, 142, 148, 149],
                        dtype=int64),)
```

In [20]:
```
sorted_rscore=  sorted(iris['Sepal_Length'])
```

In [21]:
```
sorted_rscore
```

Out[21]:
```
[4.3,
 4.4,
 4.4,
 4.4,
 4.5,
 4.6,
 4.6,
 4.6,
 4.6,
 4.7,
 4.7,
 4.8,
 4.8,
 4.8,
 4.8,
 4.8,
 4.9,
 4.9,
 4.9,
 4.9,
 4.9,
 4.9,
 5.5,
 5.5,
 5.6,
 5.6,
 5.6,
 5.6,
 5.6,
```

In [22]:
```
q1 = np.percentile(sorted_rscore, 25)
q3 = np.percentile(sorted_rscore, 75)
print(q1,q3)
```

```
5.1 6.4
```

In [23]:
```
IQR = q3-q1
```

In [24]:
```
IQR
```

Out[24]:
```
1.3000000000000007
```

In [25]:
```
lwr_bound = q1-
(1.5*IQR) upr_bound =
q3+(1.5*IQR)
```

```
3.1499999999999986  8.350000000000001
```

In [26]:
```
r_outliers = []
for i in sorted_rscore:
        if (i<lwr_bound or
                i>upr_bound):
                r_outliers.append(i)
```

In [27]:
```
print(r_outliers)
```

```
[]
```

In [28]:
```
z = np.abs(stats.zscore(iris['Sepal_Width']))
```

In [29]:
```
print(z)
```

```
0      1.032057
1      0.124958
2      0.337848
3      0.106445
4      1.263460
         ...
145    0.124958
146    1.281972
147    0.124958
148    0.800654
149    0.124958
Name: Sepal_Width, Length: 150, dtype:
      float64
```

In [30]:
```
threshold = 0.5
```

In [31]:
```
sample_outliers = np.where(z <threshold)
sample_outliers
```

Out[31]:
```
(array([  1,   2,   3,   8,   9,  12,  13,  25,  29,  30,  34,  35,  37,
         38,  42,  45,  47,  50,  51,  52,  58,  61,  63,  64,  65,  66,
         70,  74,  75,  77,  78,  84,  86,  88,  91,  95,  96,  97, 102,
        103, 104, 105, 107, 110, 112, 115, 116, 120, 125, 127, 129, 135,
        137, 138, 139, 140, 141, 143, 145, 147, 149],
       dtype=int64),)
```

In [32]:
```
sorted_rscore=  sorted(iris['Sepal_Width'])
```

In [33]:
```
sorted_rscore
```

Out[33]:
```
[2.0,
 2.2,
 2.2,
 2.2,
 2.3,
 2.3,
 2.3,
 2.3,
 2.4,
 2.4,
 2.4,
 2.5,
 2.5,
 2.5,
 2.5,
 2.5,
 2.5,
 2.5,
 2.5,
 2.6,
 2.6,
 2.6,
 2.6,
 2.6,
 2.7,
 2.7,
 2.7,
 2.7,
 2.7,
 2.7,
 2.7,
 2.7,
 2.7,
 2.8,
 2.8,
```

In [34]:
```
q1 = np.percentile(sorted_rscore, 25)
q3 = np.percentile(sorted_rscore, 75)
print(q1,q3)
```

```
2.8 3.3
```

In [35]:
```
IQR = q3-q1
```

In [36]:
```
IQR
```

Out[36]:
```
0.5
```

In [37]:
```
lwr_bound = q1-
(1.5*IQR) upr_bound =
q3+(1.5*IQR)
```

```
2.05 4.05
```

In [38]:
```
r_outliers = []
for i in sorted_rscore:
        if (i<lwr_bound or
                i>upr_bound):
                r_outliers.append(i)
print(r_outliers)
```

```
[2.0, 4.1, 4.2, 4.4]
```

In [39]:
```
from scipy import stats
```

In [40]:
```
z = np.abs(stats.zscore(iris['Petal_Length']))
```

In [41]:
```
print(z)
```

```
0      1.341272
1      1.341272
2      1.398138
3      1.284407
4      1.341272
         ...
145    0.819624
146    0.705893
147    0.819624
148    0.933356
149    0.762759
Name: Petal_Length, Length: 150, dtype:
float64
```

In [42]:
```
threshold = 0.5
```

In [43]:
```
sample_outliers = np.where(z <threshold)
sample_outliers
```

Out[43]:
```
(array([ 51,  53,  54,  55,  57,  58,  59,  60,  61,  62,  64,  65,  66,
         67,  68,  69,  71,  74,  75,  78,  79,  80,  81,  82,  84,  85,
         87,  88,  89,  90,  91,  92,  93,  94,  95,  96,  97,  98,  99,
        106], dtype=int64),)
```

In [44]:
```
sorted_rscore= sorted(iris['Petal_Length'])
```

In [45]:
```
q1 = np.percentile(sorted_rscore, 25)
q3 = np.percentile(sorted_rscore, 75)
print(q1,q3)
```

```
1.6 5.1
```

In [46]:
```
IQR = q3-q1
```

In [47]:
```
lwr_bound = q1-(1.5*IQR)
upr_bound = q3+(1.5*IQR)
print(lwr_bound,  upr_bound)
```

```
-3.649999999999999  10.349999999999998
```

In [60]:
```
r_outliers = []
for i in sorted_rscore:
        if (i<lwr_bound or i>upr_bound):
                r_outliers.append(i)
print(r_outliers)
```

```
[]
```

In [61]:
```
print(r_outliers)
```

```
[]
```

In [62]:
```
z  =  np.abs(stats.zscore(iris['Petal_Width']))
```

In [63]:
```
print(z)
```

```
0      1.312977
1      1.312977
2      1.312977
3      1.312977
4      1.312977
         ...
145    1.447956
146    0.922064
147    1.053537
148    1.447956
149    0.790591
Name: Petal_Width, Length: 150, dtype:
float64
```

In [64]:
```
threshold = 0.5
```

In [65]:
```
sample_outliers = np.where(z <threshold)
sample_outliers
```

Out[65]:
```
(array([ 50,  51,  52,  53,  54,  55,  57,  58,  59,  60,  61,  62,  63,
         64,  65,  66,  67,  68,  69,  71,  72,  73,  74,  75,  76,  78,
         79,  80,  81,  82,  84,  86,  87,  88,  89,  90,  91,  92,  93,
         94,  95,  96,  97,  98,  99, 119, 133, 134], dtype=int64),)
```

In [66]:
```
sorted_rscore=  sorted(iris['Petal_Width'])
```

In [67]:
```
sorted_rscore
```

Out[67]:
```
[0.1,
 0.1,
 0.1,
 0.1,
 0.1,
 0.1,
 0.2,
 0.2,
 1.1,
 1.2,
 1.2,
 1.2,
 1.2,
 1.8,
 1.8,
 1.8,
 1.8,
 1.8,
 1.8,
 1.9,
 1.9,
 1.9,
 2.0,
 2.0,
 2.0,
 2.0,
 2.0,
 2.0,
 2.1,
 2.3,
 2.3,
 2.3,
 2.3,
 2.3,
 2.3,
 2.3,
 2.3,
 2.4,
 2.4,
 2.4,
 2.5,
 2.5,
 2.5]
```

In [68]:
```
q1 = np.percentile(sorted_rscore, 25)
q3 = np.percentile(sorted_rscore, 75)
print(q1,q3)
```

```
0.3 1.8
```

In [69]:
```
IQR = q3-q1
```

In [70]:
```
lwr_bound = q1-
(1.5*IQR) upr_bound =
q3+(1.5*IQR)
print(lwr_bound, upr_bound)
```

```
-1.95 4.05
```

In [71]:
```
r_outliers = []
for i in sorted_rscore:
        if (i<lwr_bound or
                i>upr_bound):
                r_outliers.append(i)
print(r_outliers)
```

```
[]
```

In [3]:
```
import pandas as pd
import numpy as np

import matplotlib.pyplot as plt
import seaborn as sns

dataset = sns.load_dataset('titanic')

dataset.head()
```

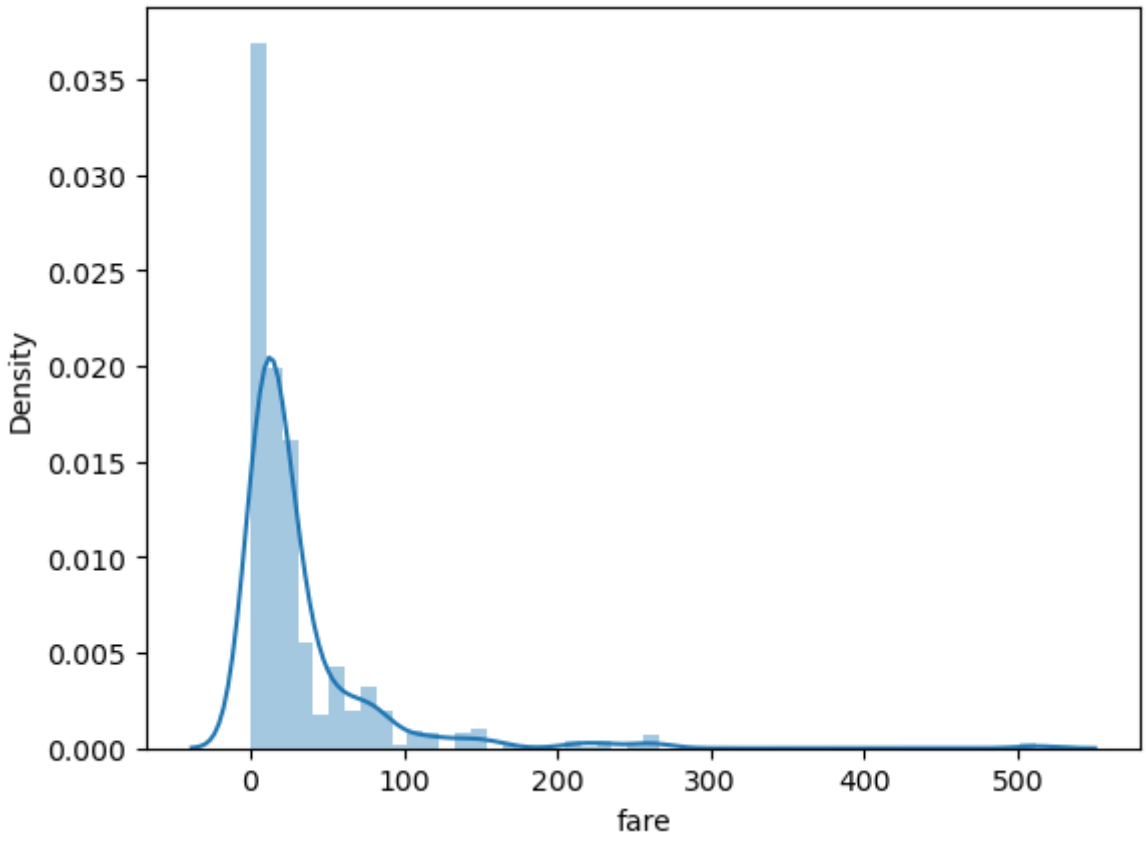
Out[3]:

	survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	de
0	0	3	male	22.0	1	0	7.2500	S	Third	man	True	Na
1	1	1	female	38.0	1	0	71.2833	C	First	woman	False	
2	1	3	female	26.0	0	0	7.9250	S	Third	woman	False	Na
3	1	1	female	35.0	1	0	53.1000	S	First	woman	False	
4	0	3	male	35.0	0	0	8.0500	S	Third	man	True	Na

In [4]:
```
sns.distplot(dataset['fare'])
```

```
C:\Users\Lenovo\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)
```

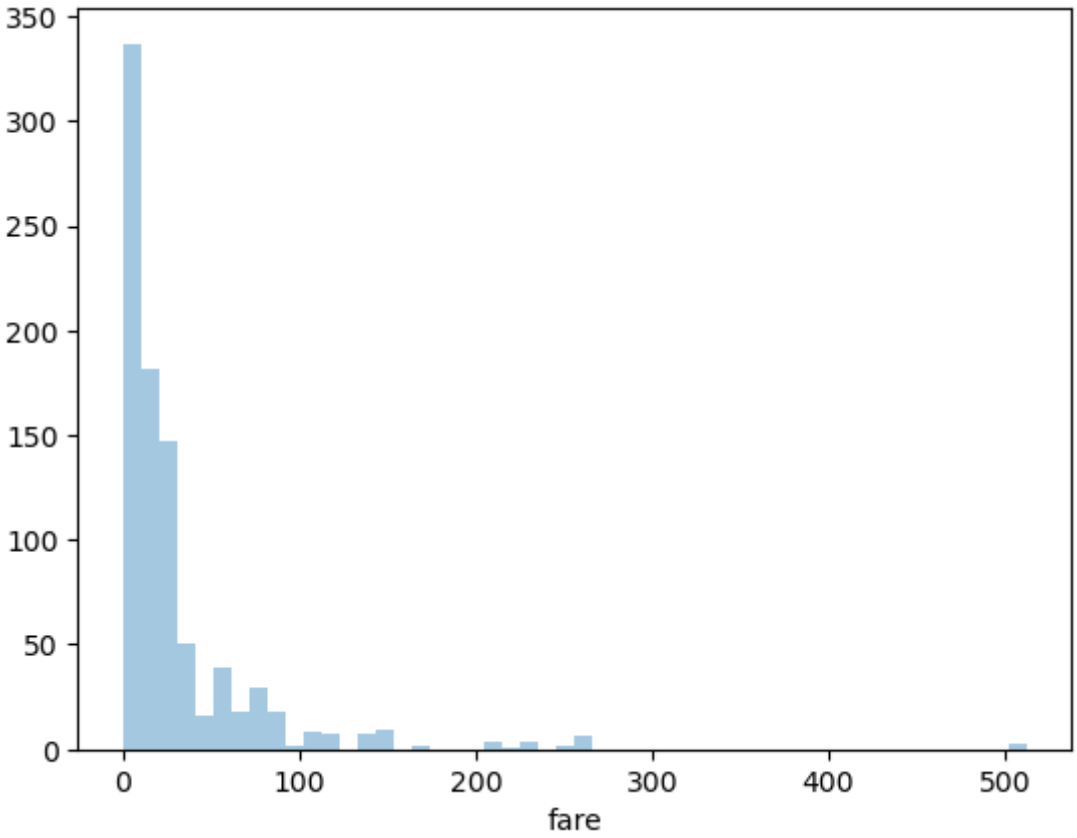
Out[4]:
```
<AxesSubplot:xlabel='fare', ylabel='Density'>
```



In [5]:
```
sns.distplot(dataset['fare'], kde=False)
```

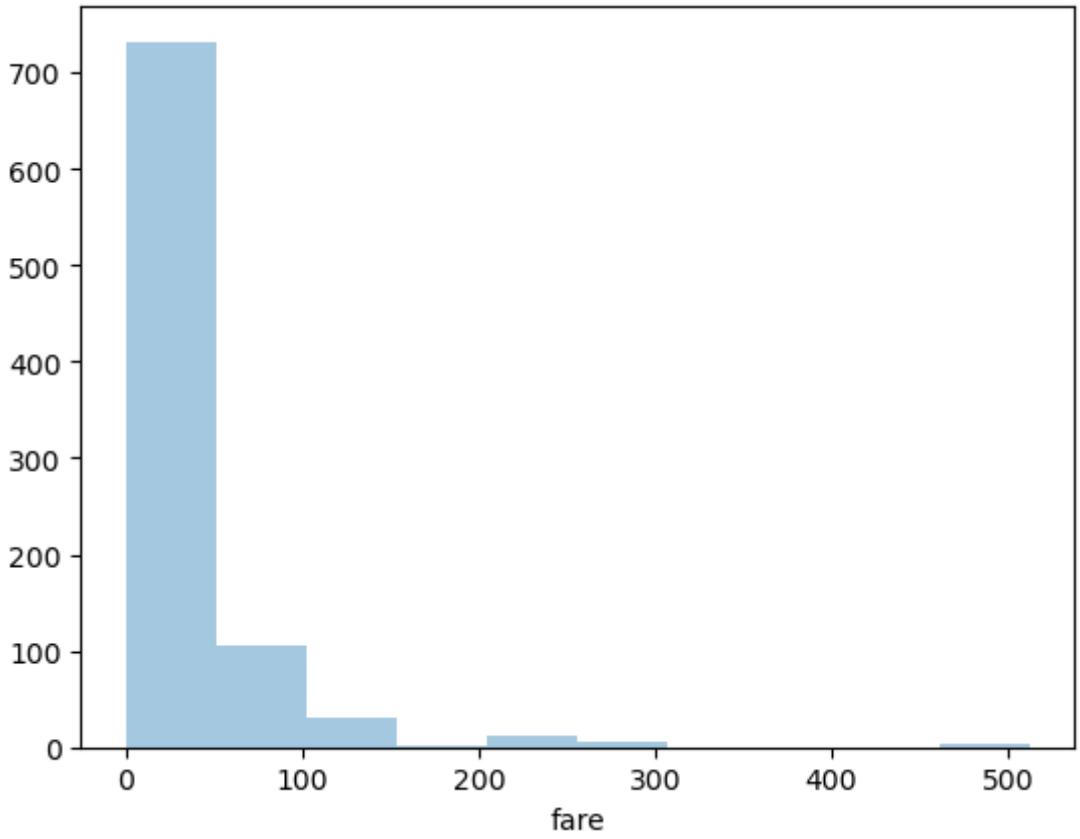
```
C:\Users\Lenovo\anaconda3\lib\site-packages\seaborn\distributions.py:2619: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).
  warnings.warn(msg, FutureWarning)
```

Out[5]: <AxesSubplot:xlabel='fare'>



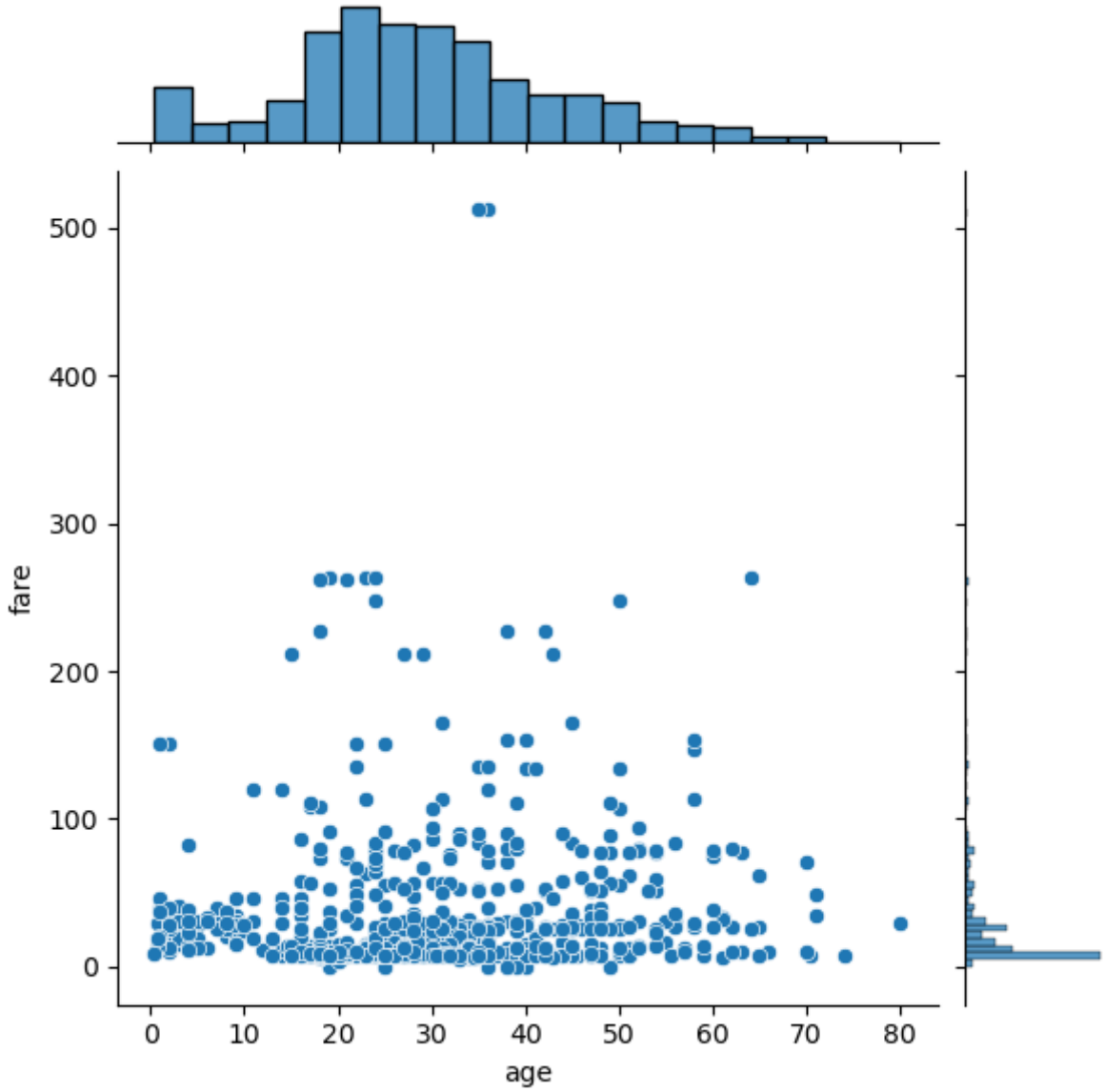
In [6]:
```
sns.distplot(dataset['fare'], kde=False, bins=10)
```

Out[6]: <AxesSubplot:xlabel='fare'>



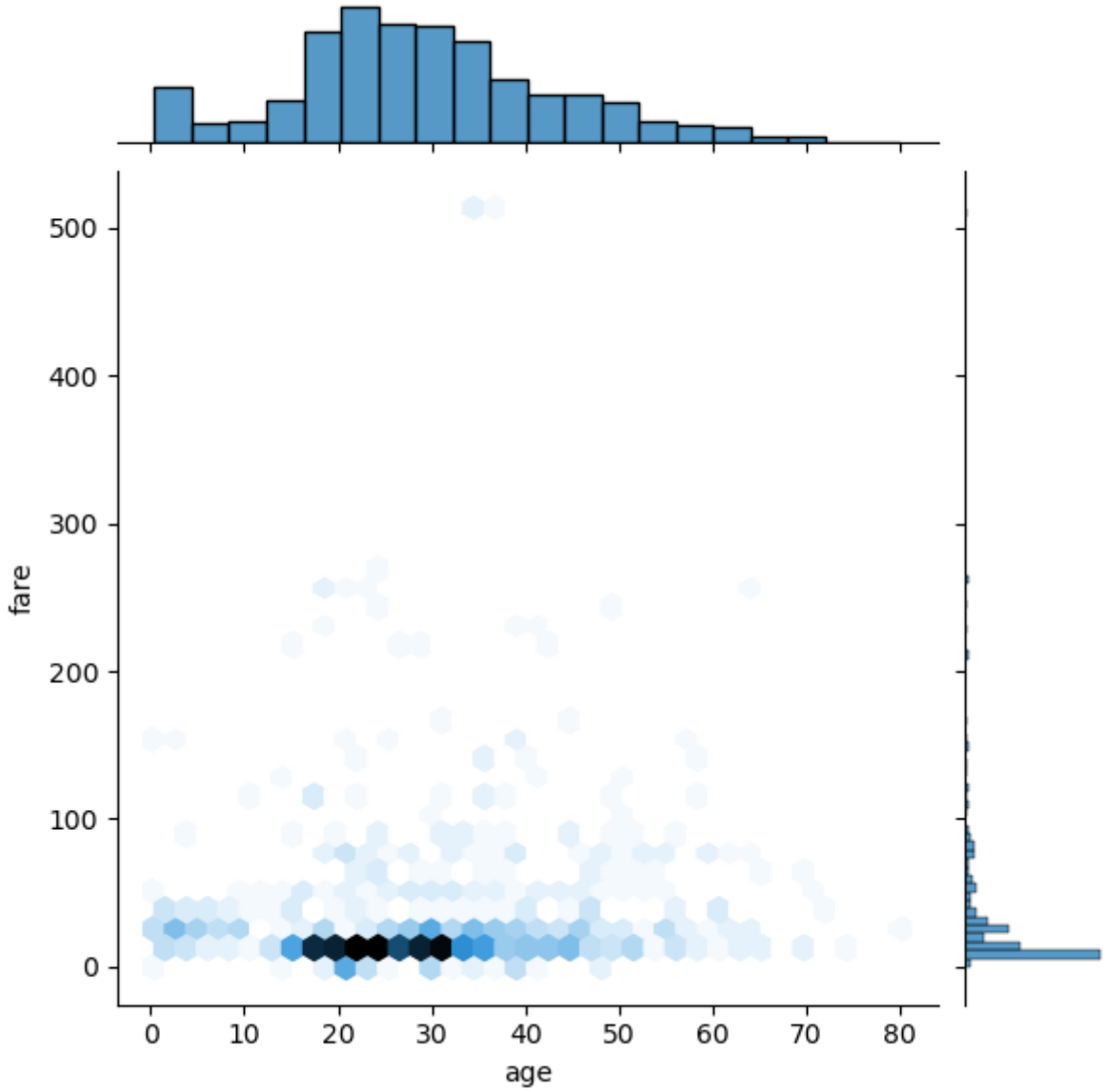
In [7]:
```
sns.jointplot(x='age', y='fare', data=dataset)
```

Out[7]:
```
<seaborn.axisgrid.JointGrid at 0x27a9bc08ac0>
```



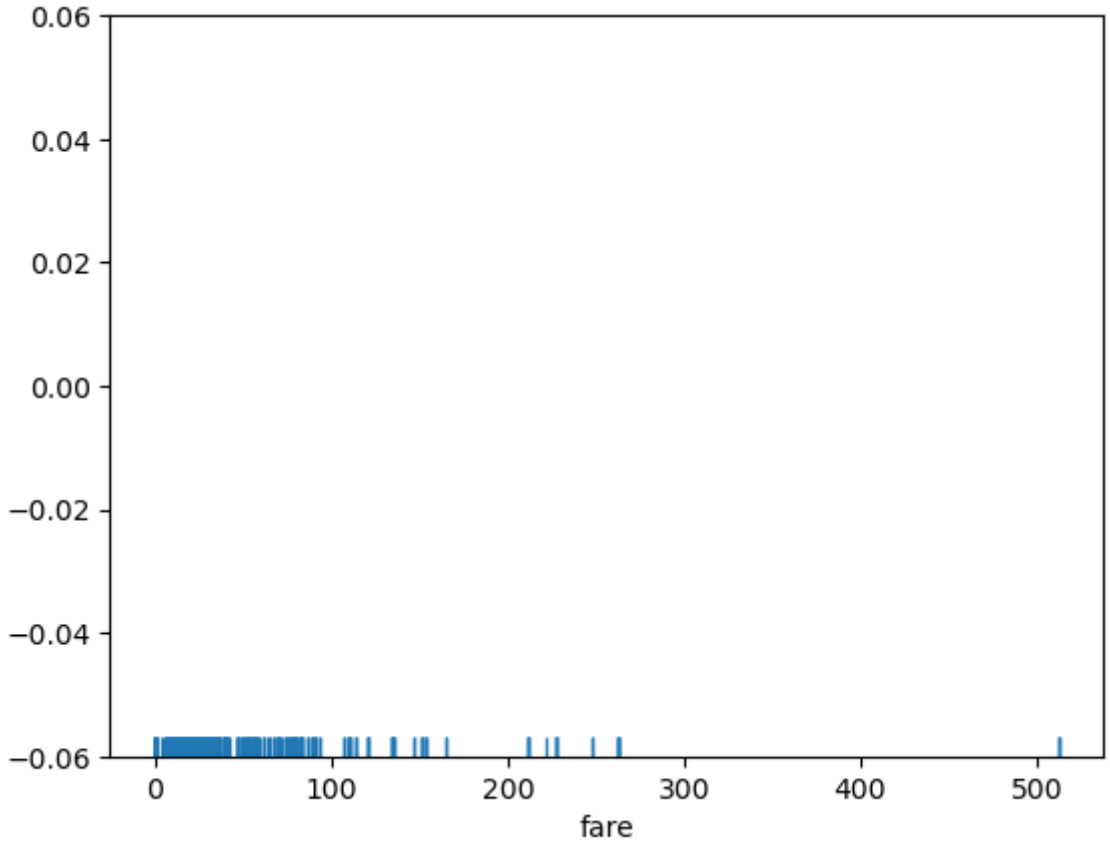
In [8]:
```
sns.jointplot(x='age', y='fare', data=dataset, kind='hex')
```

Out[8]:
```
<seaborn.axisgrid.JointGrid at 0x27a9c0adeb0>
```



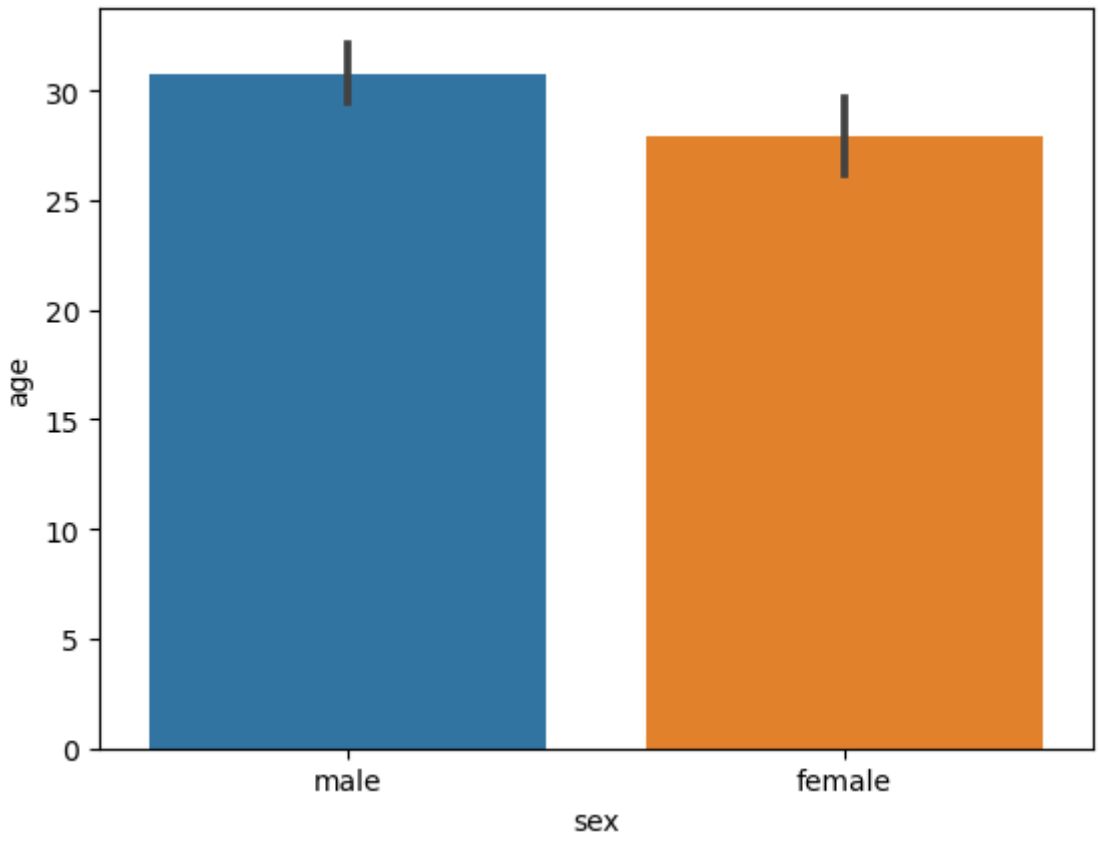
In [9]:
```
sns.rugplot(dataset['fare'])
```

Out[9]:
```
<AxesSubplot:xlabel='fare'>
```



In [10]:
```
sns.barplot(x='sex', y='age', data=dataset)
```

Out[10]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```

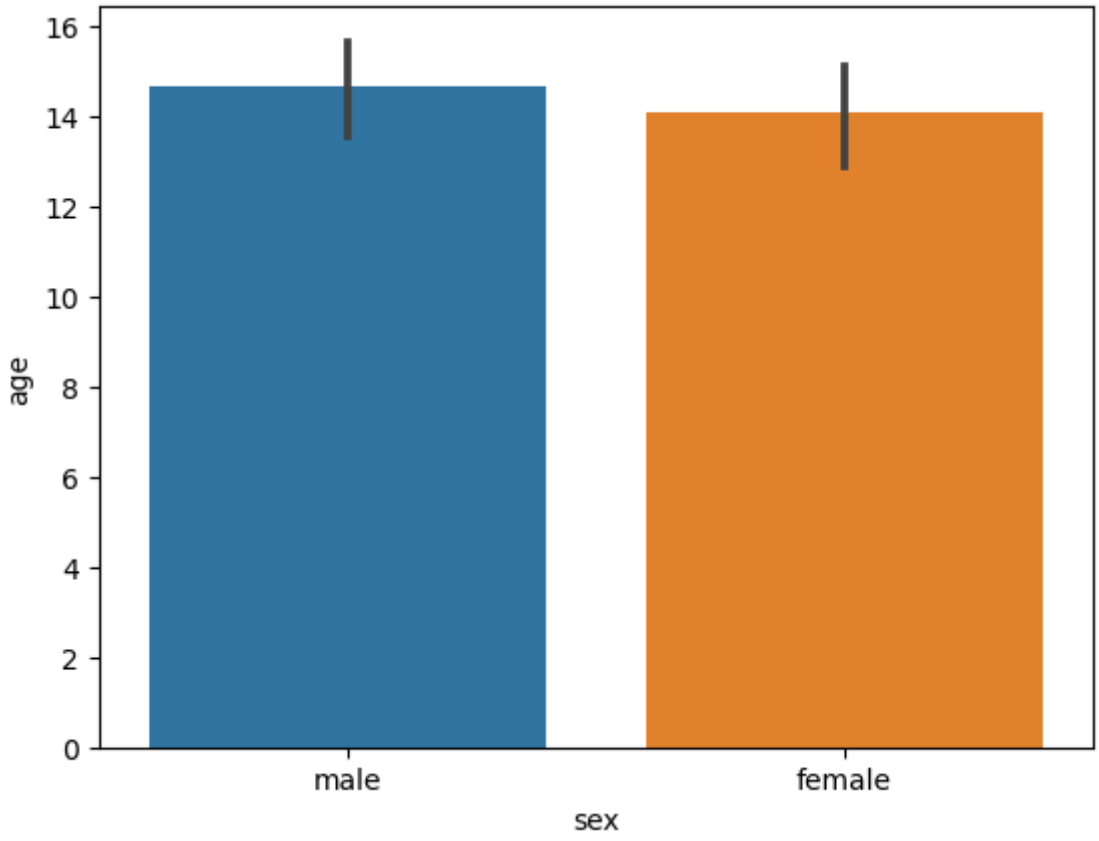


In [11]:
```
import numpy as np

import matplotlib.pyplot as plt
```

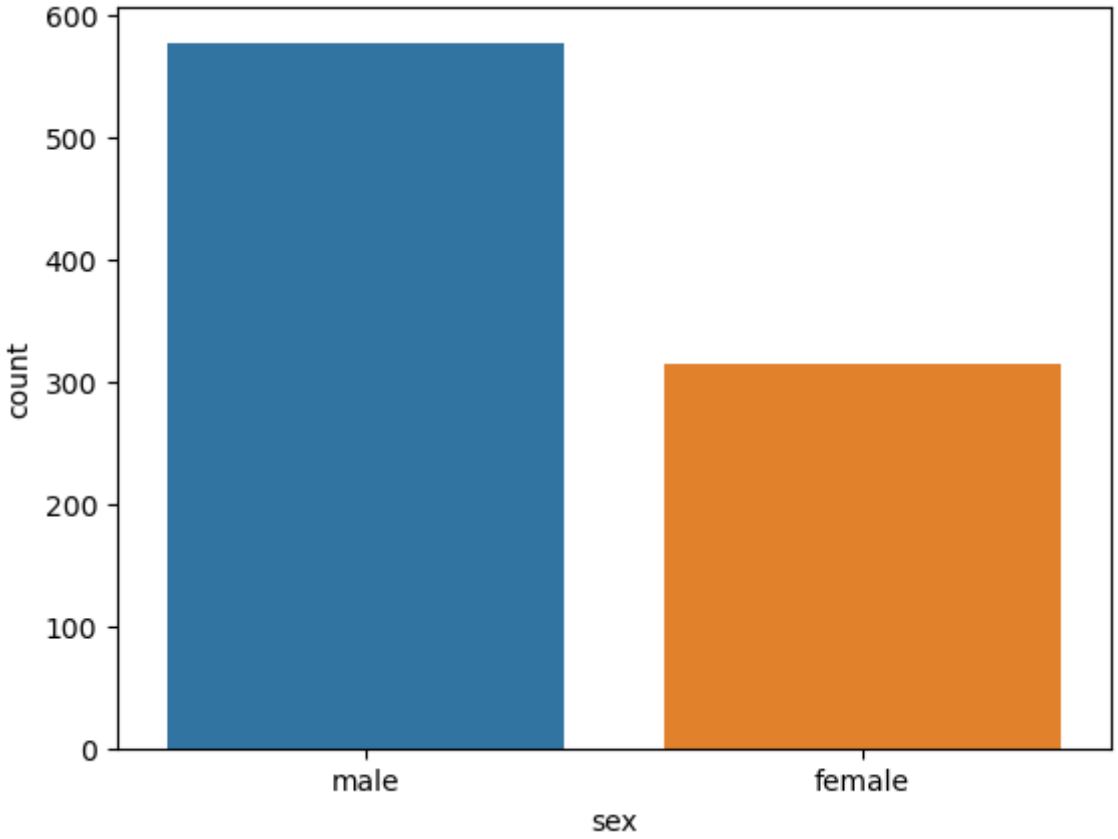
```
sns.barplot(x='sex', y='age', data=dataset, estimator=np.std)
```

Out[11]: <AxesSubplot:xlabel='sex', ylabel='age'>



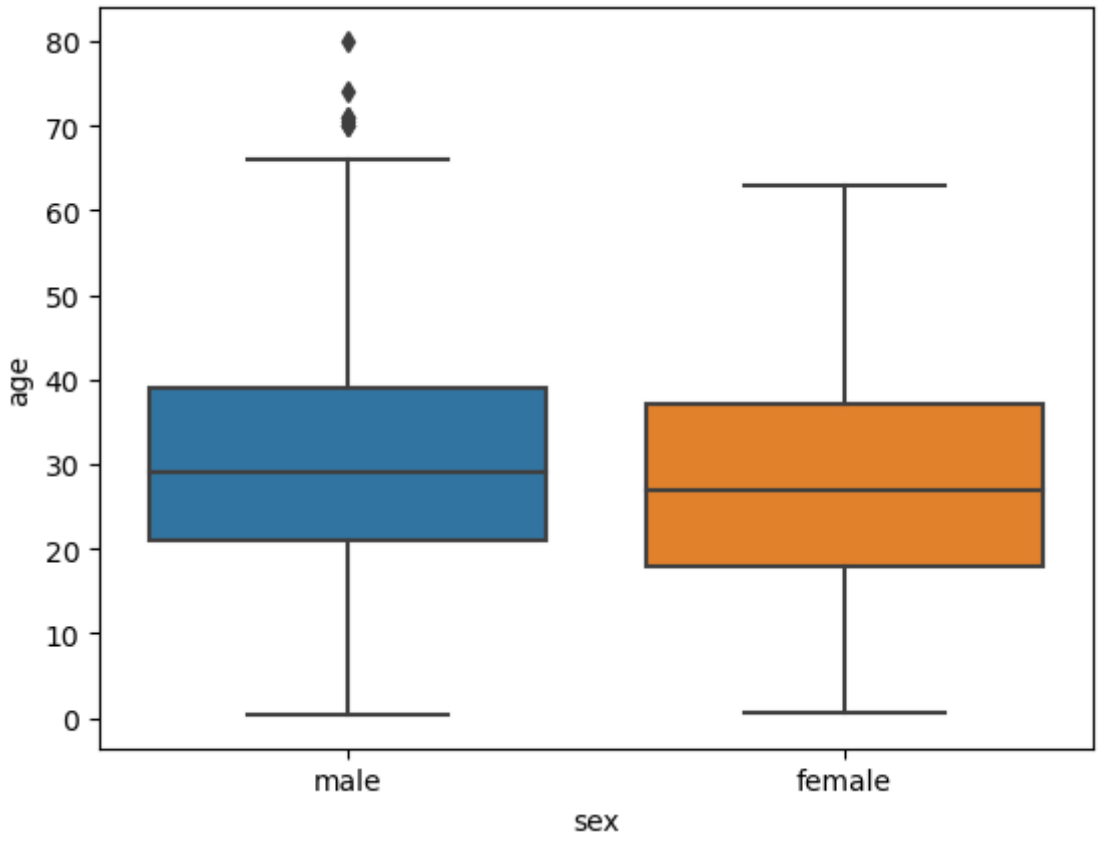
In [12]:
```
sns.countplot(x='sex', data=dataset)
```

Out[12]: <AxesSubplot:xlabel='sex', ylabel='count'>



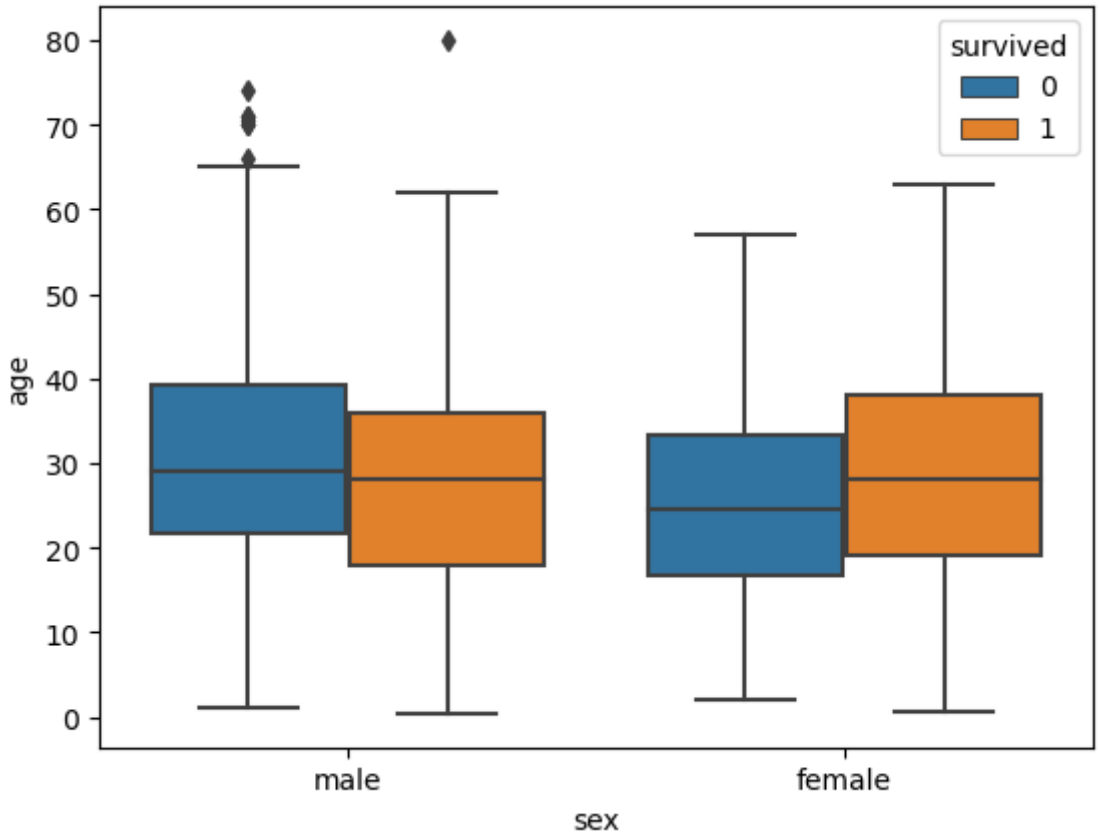
In [13]:
```
sns.boxplot(x='sex', y='age', data=dataset)
```

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



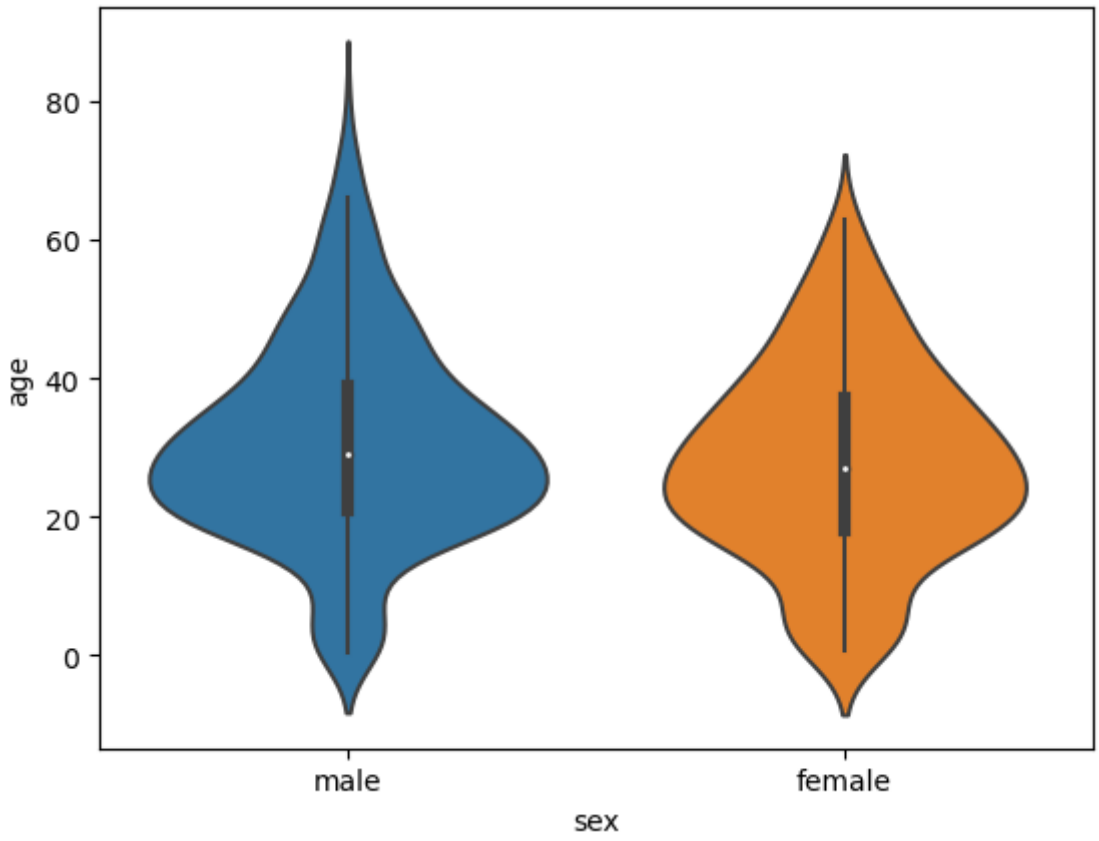
In [14]:
```
sns.boxplot(x='sex', y='age', data=dataset, hue="survived")
```

Out[14]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



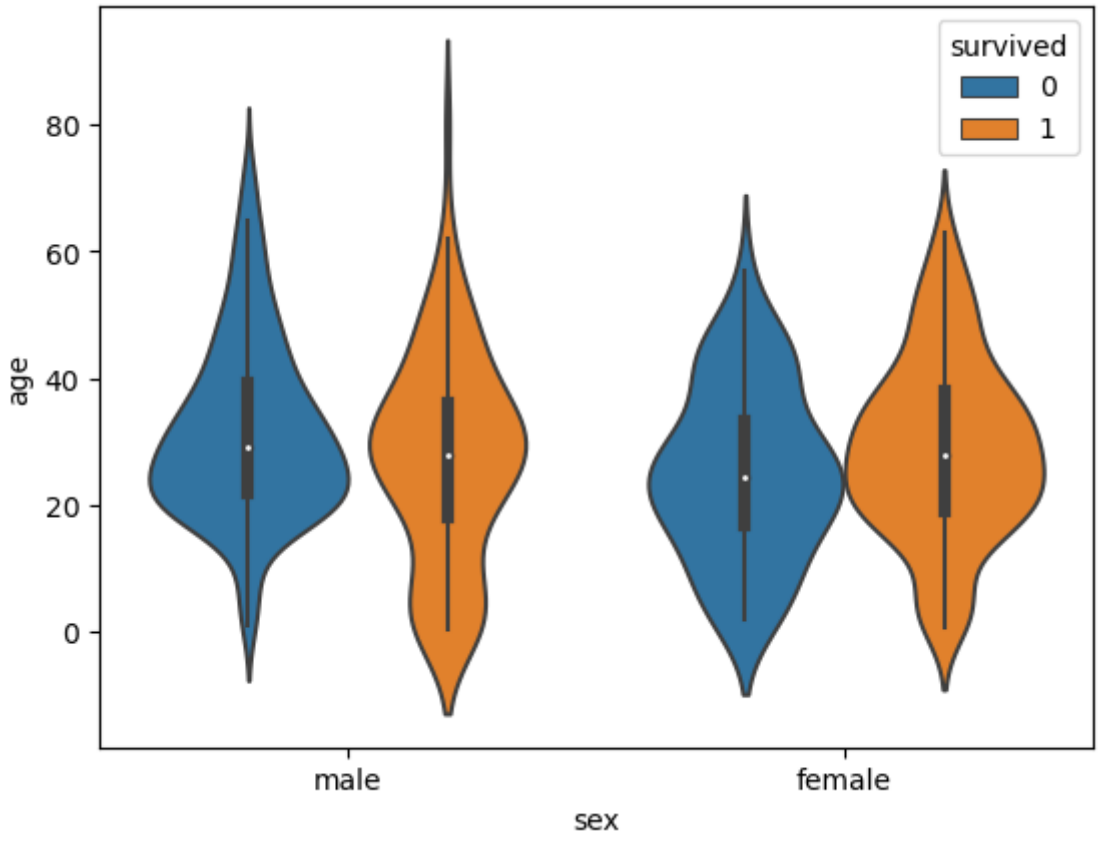
In [15]:
```
sns.violinplot(x='sex', y='age', data=dataset)
```

Out[15]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



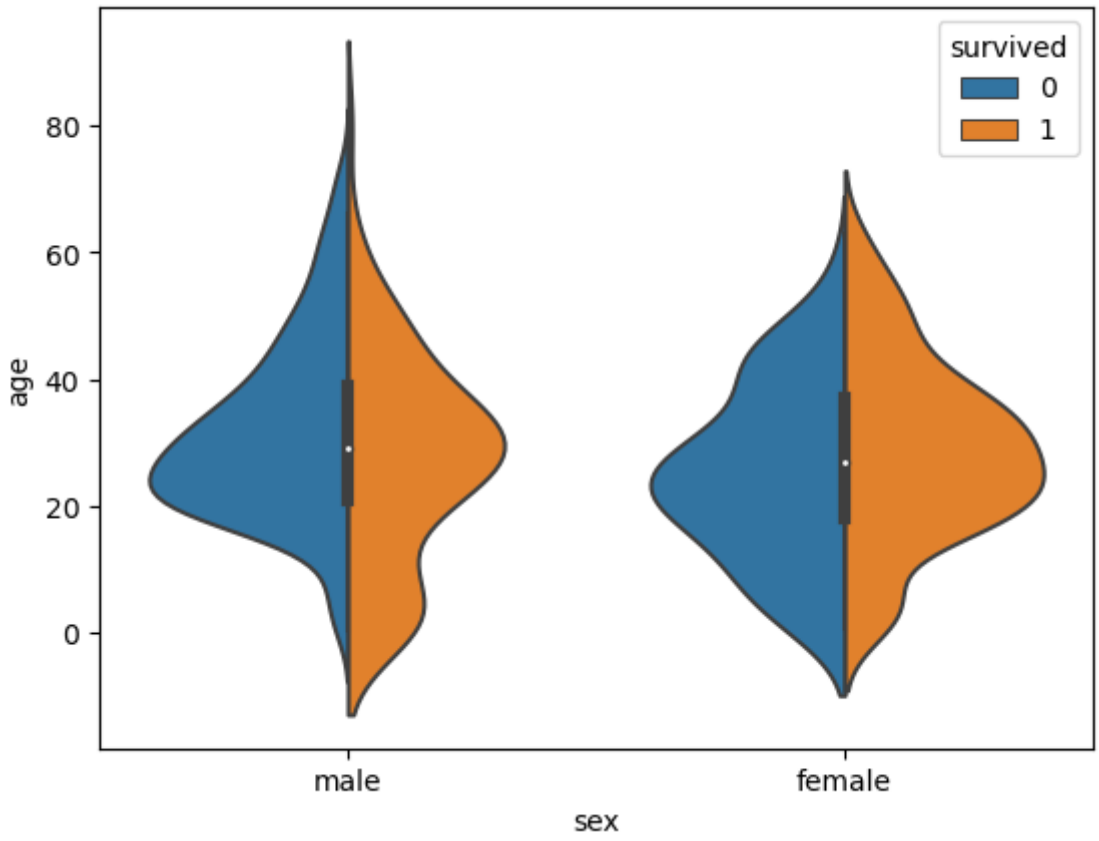
In [16]:
```
sns.violinplot(x='sex', y='age', data=dataset, hue='survived')
```

Out[16]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



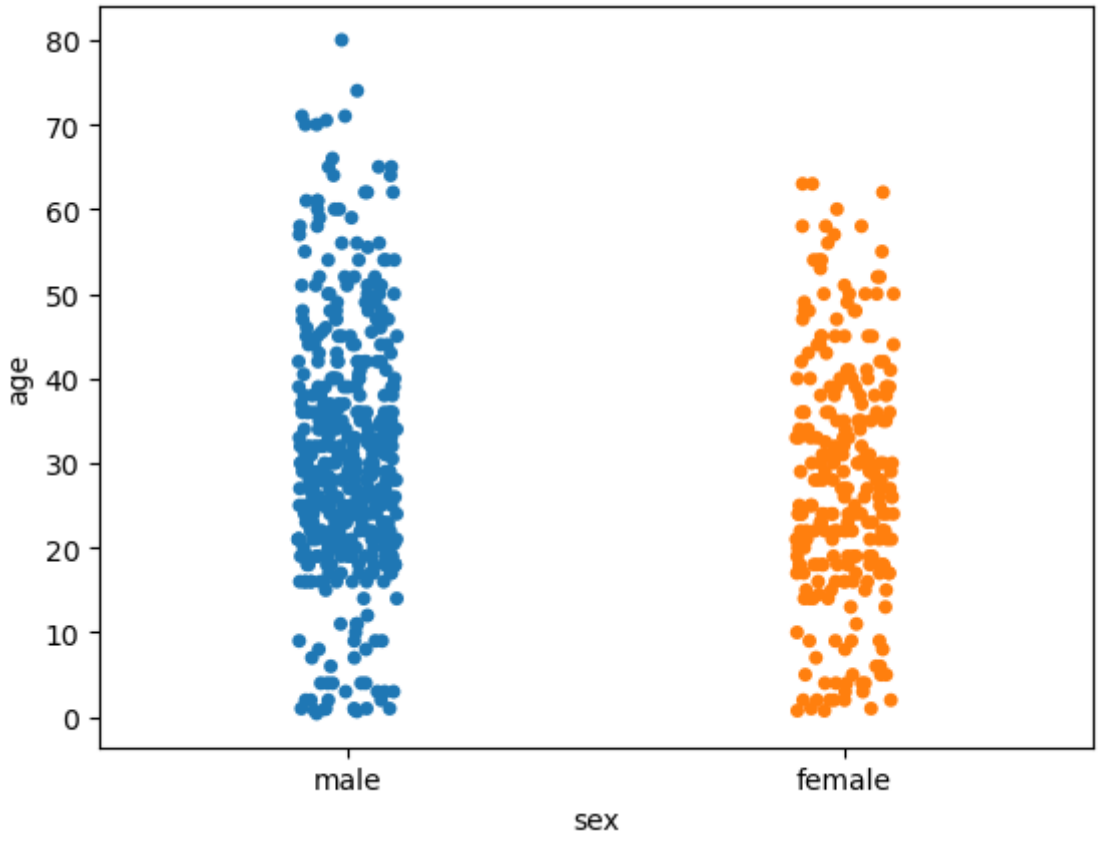
In [17]:
```
sns.violinplot(x='sex', y='age', data=dataset, hue='survived', split=True)
```

Out[17]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



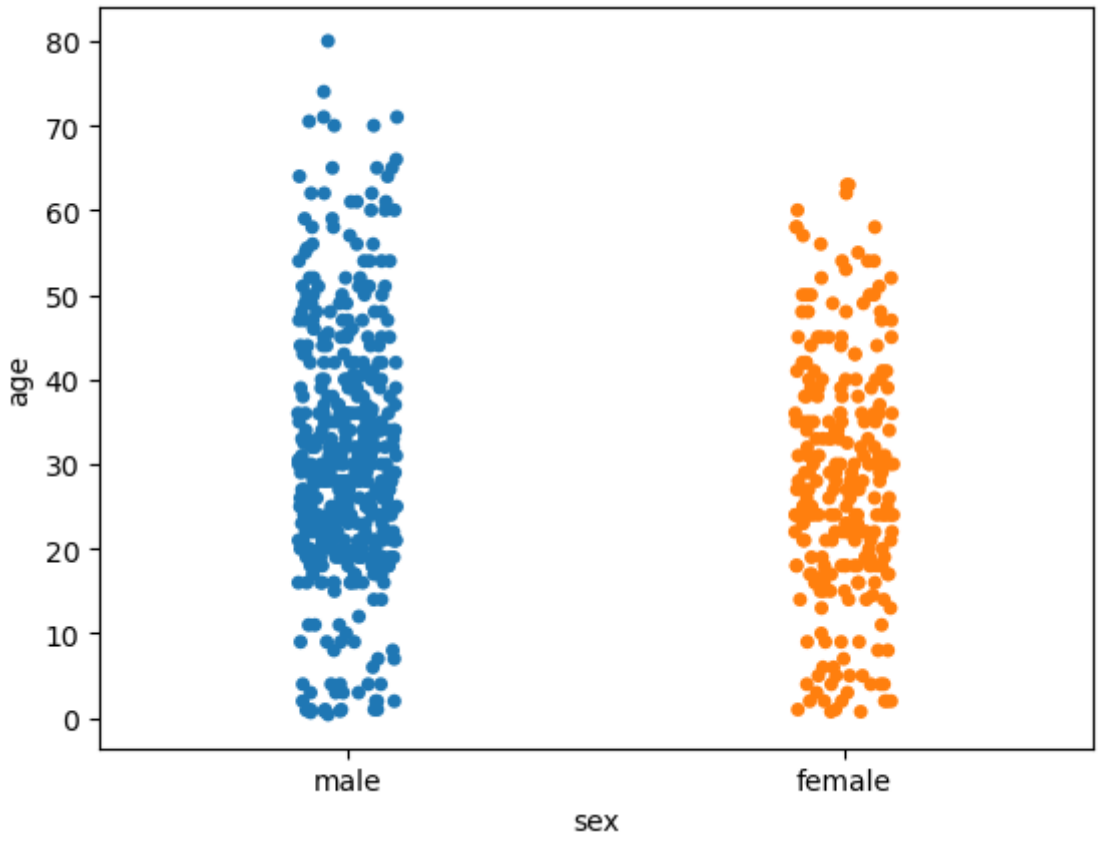
In [18]:
```
sns.stripplot(x='sex', y='age', data=dataset)
```

Out[18]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



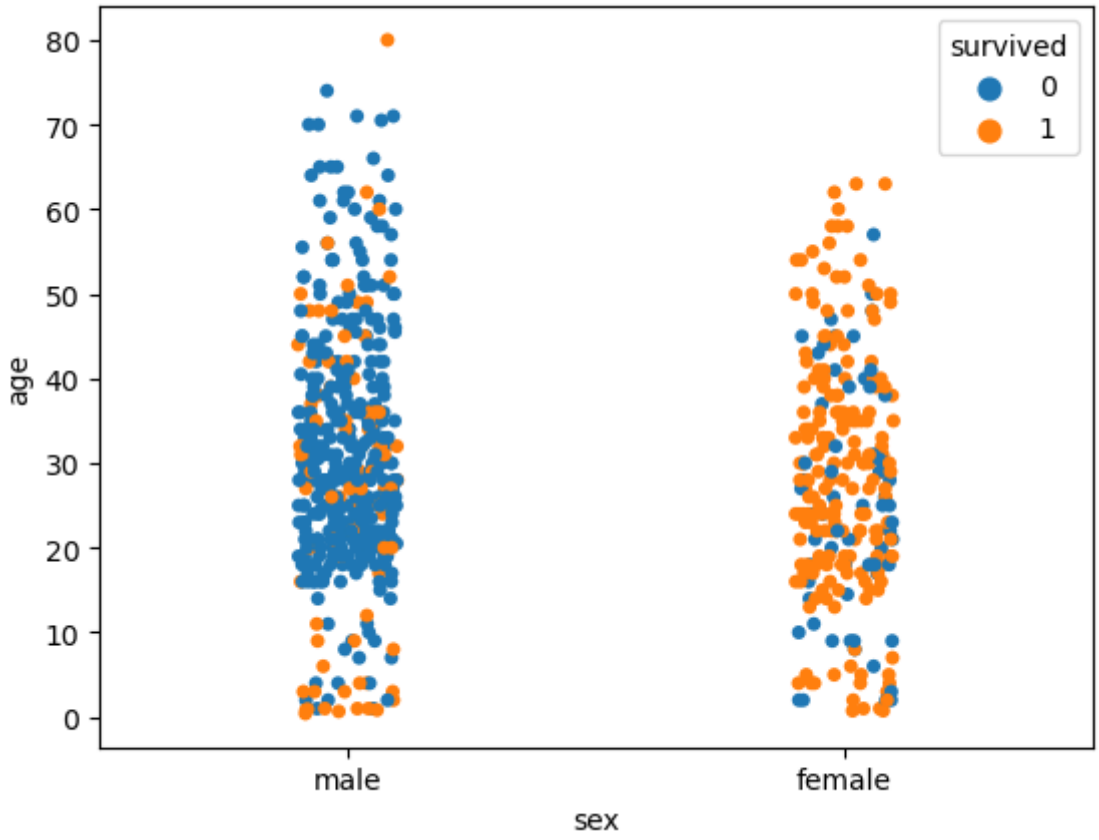
In [19]:
```
sns.stripplot(x='sex', y='age', data=dataset, jitter=True)
```

Out[19]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



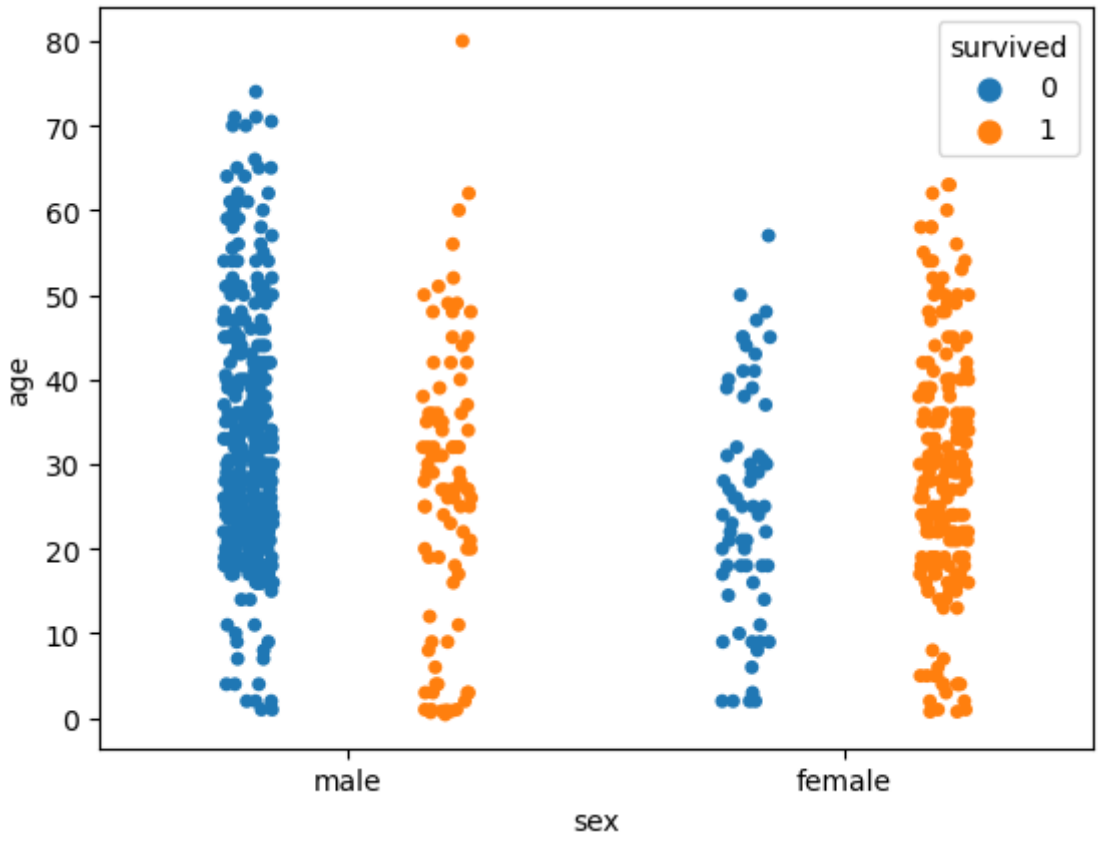
In [20]:
```
sns.stripplot(x='sex', y='age', data=dataset, jitter=True, hue='survived')
```

Out[20]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



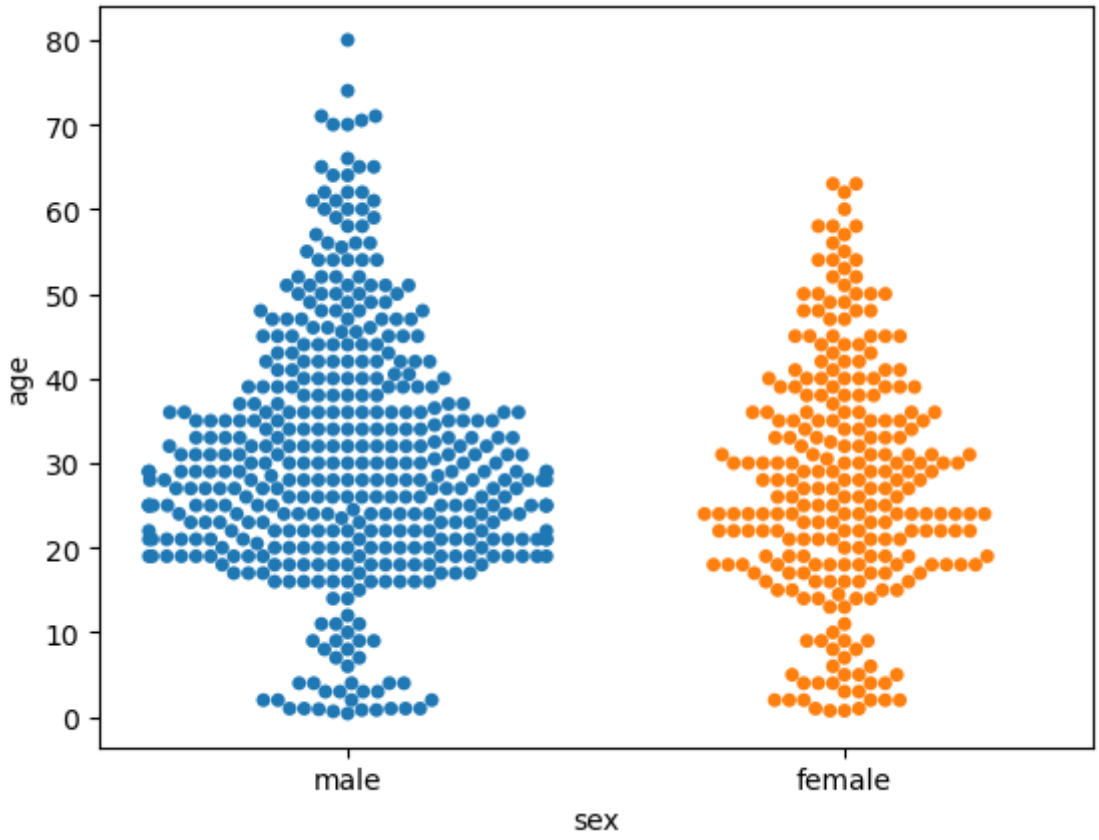
In [21]:
```
sns.stripplot(x='sex', y='age', data=dataset, jitter=True, hue='survived', dodge=T
```

Out[21]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



In [25]:
```
sns.swarmplot(x='sex', y='age', data=dataset)
```

Out[25]:
```
<AxesSubplot:xlabel='sex', ylabel='age'>
```



In [26]:
```
sns.matrixplot(x='sex', y='age', data=dataset)
```

```
---------------------------------------------------------------------------
AttributeError                            Traceback (most recent call last)
~\AppData\Local\Temp\ipykernel_14272\2259532084.py in <module>
----> 1 sns.matrixplot(x='sex', y='age', data=dataset)

AttributeError: module 'seaborn' has no attribute 'matrixplot'
```

In [27]:
```
dataset.corr()
```

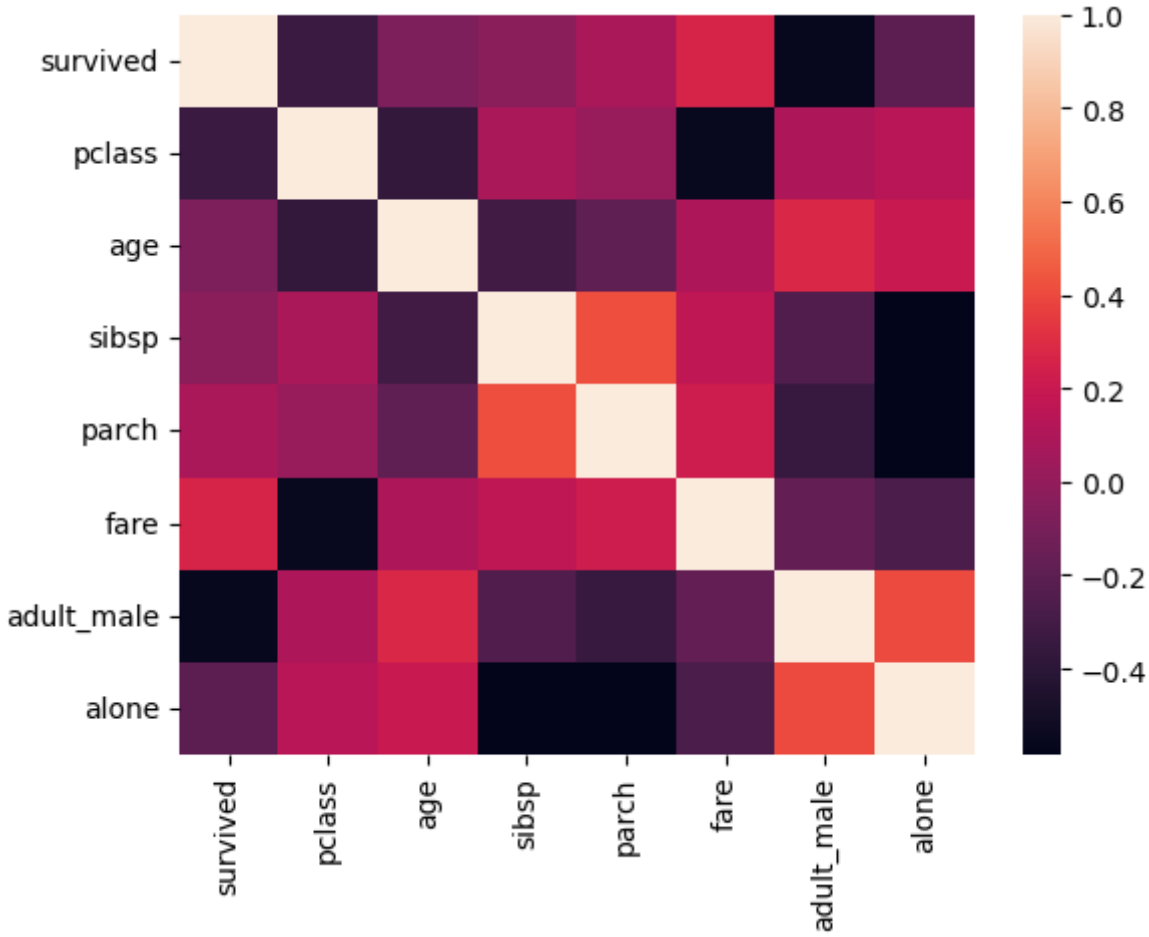
Out[27]:

	survived	pclass	age	sibsp	parch	fare	adult_male	alone
survived	1.000000	-0.338481	-0.077221	-0.035322	0.081629	0.257307	-0.557080	-0.203367
pclass	-0.338481	1.000000	-0.369226	0.083081	0.018443	-0.549500	0.094035	0.135207
age	-0.077221	-0.369226	1.000000	-0.308247	-0.189119	0.096067	0.280328	0.198270
sibsp	-0.035322	0.083081	-0.308247	1.000000	0.414838	0.159651	-0.253586	-0.584471
parch	0.081629	0.018443	-0.189119	0.414838	1.000000	0.216225	-0.349943	-0.583398
fare	0.257307	-0.549500	0.096067	0.159651	0.216225	1.000000	-0.182024	-0.271832
adult_male	-0.557080	0.094035	0.280328	-0.253586	-0.349943	-0.182024	1.000000	0.404744
alone	-0.203367	0.135207	0.198270	-0.584471	-0.583398	-0.271832	0.404744	1.000000

In [28]:
```
corr=dataset.corr()
```

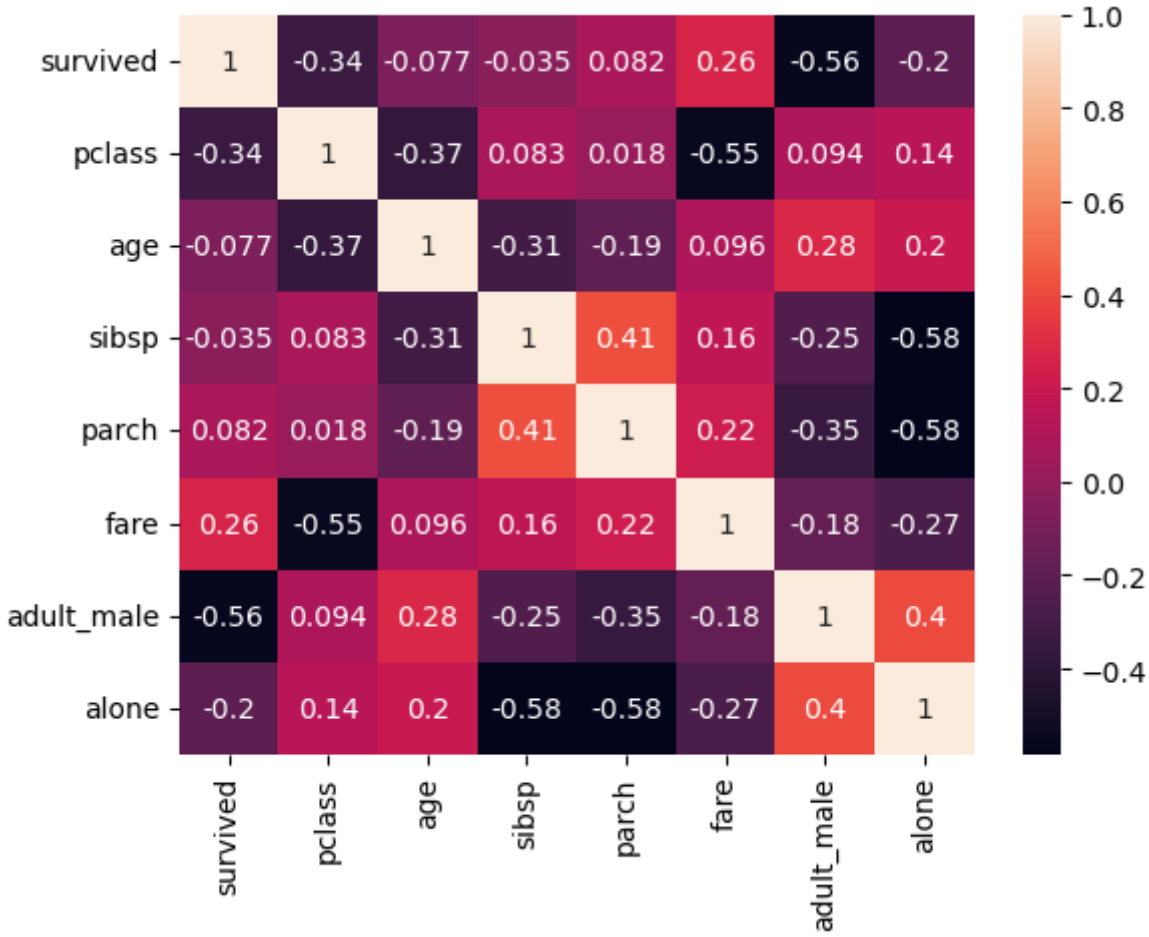
In [31]:
```
sns.heatmap(corr)
```

Out[31]:
```
<AxesSubplot:>
```



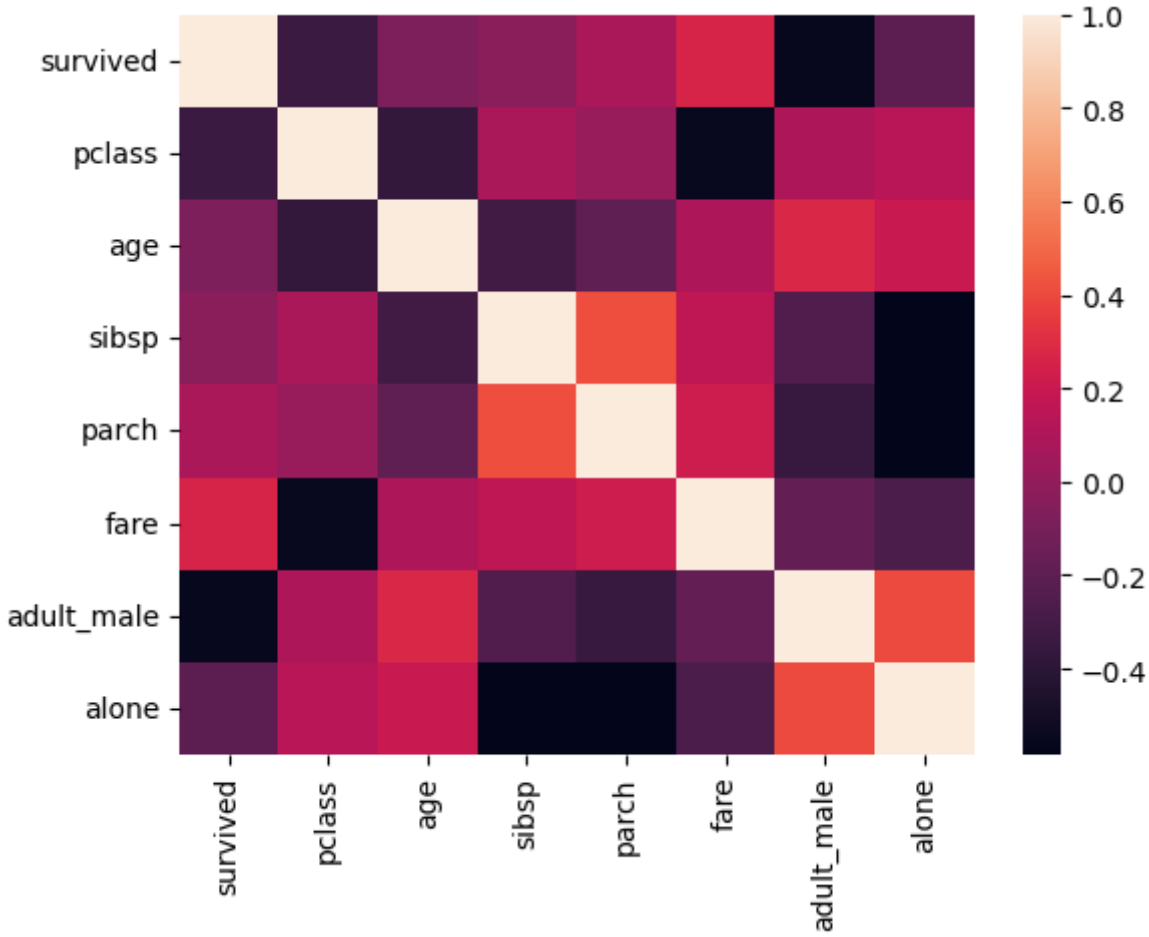
In [33]:
```
sns.heatmap(corr, annot=True)
```

Out[33]:
```
<AxesSubplot:>
```



In [34]:
```
sns.heatmap(corr)
```

Out[34]:
```
<AxesSubplot:>
```



In [ ]: