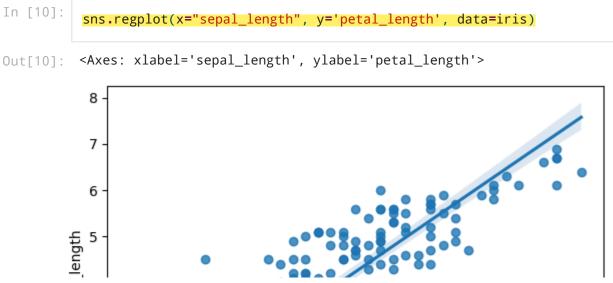
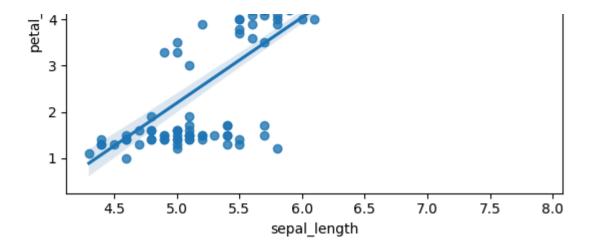


```
In [3]:
           import pandas as pd
           import numpy as np
           import seaborn as sns
           iris = sns.load_dataset("iris")
           iris
                sepal_length sepal_width petal_length petal_width
 Out[3]:
                                                                  species
             0
                        5.1
                                    3.5
                                                 1.4
                                                             0.2
                                                                   setosa
             1
                        4.9
                                    3.0
                                                 1.4
                                                             0.2
                                                                   setosa
             2
                        4.7
                                    3.2
                                                 1.3
                                                             0.2
                                                                   setosa
             3
                        4.6
                                    3.1
                                                 1.5
                                                             0.2
                                                                   setosa
             4
                        5.0
                                    3.6
                                                 1.4
                                                             0.2
                                                                   setosa
                                                  •••
           145
                        6.7
                                    3.0
                                                 5.2
                                                             2.3
                                                                 virginica
           146
                                    2.5
                                                 5.0
                        6.3
                                                             1.9
                                                                 virginica
           147
                        6.5
                                    3.0
                                                 5.2
                                                                 virginica
                                                             2.0
           148
                        6.2
                                    3.4
                                                 5.4
                                                             2.3
                                                                 virginica
           149
                        5.9
                                    3.0
                                                 5.1
                                                             1.8 virginica
          150 rows × 5 columns
 In [4]:
           #!pip install seaborn
 In [7]:
           iris.columns
 Out[7]:
           Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
                   'species'],
                  dtype='object')
In [10]:
           sns.regplot(x="sepal_length", y='petal_length', data=iris)
```





In [16]: sns.distplot(iris.sepal_length,kde=True)

C:\Users\HP\AppData\Local\Temp\ipykernel_17516\4260748696.py:1: UserWarnin
g:

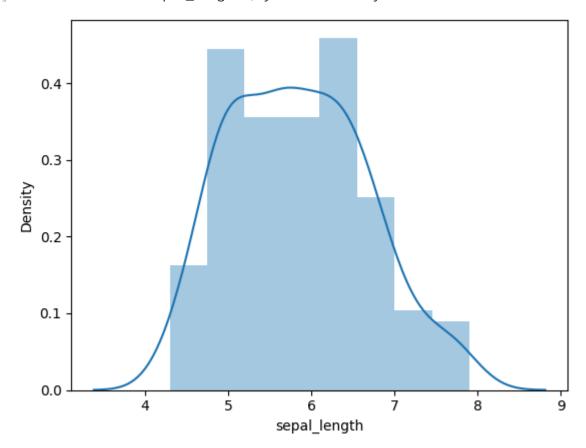
`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751

sns.distplot(iris.sepal_length,kde=True)

Out[16]: <Axes: xlabel='sepal_length', ylabel='Density'>



```
In [23]:
          iris.columns
          Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
Out[23]:
                  'species'],
                dtype='object')
In [27]:
          sns.heatmap(iris[['sepal_length', 'sepal_width', 'petal_length', 'petal_wi
Out[27]:
          <Axes: >
           0
           6
          12
          18
                                                                          - 7
          24
          30 -
          36
42
                                                                          - 6
          48
          54
                                                                           5
          60
          66
          72
          78
          84
          90
                                                                           3
          96
         102
         108
         114
         120
         126
         132
         138
         144
              sepal_length sepal_width petal_length petal_width
In [28]:
          sns.stripplot(x="species", y='petal_length', data=iris)
Out[28]: <Axes: xlabel='species', ylabel='petal_length'>
            7
            6
           5
         petal_length
```

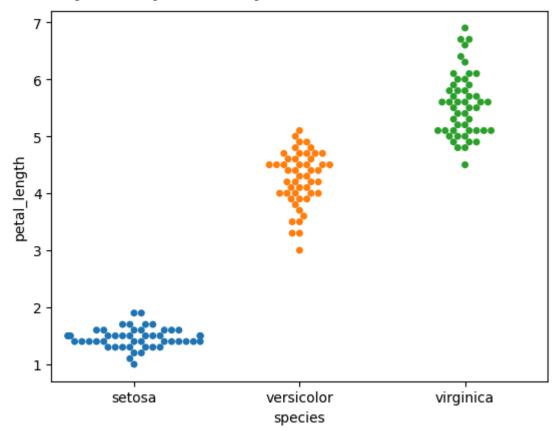
```
2 - 1 - setosa versicolor virginica species
```

In [36]: sns.swarmplot(x="species", y='petal_length', hue='species',data=iris)

Out[36]: <Axes: xlabel='species', ylabel='petal_length'>

C:\Python312\Lib\site-packages\seaborn\categorical.py:3399: UserWarning: 1
2.0% of the points cannot be placed; you may want to decrease the size of t
he markers or use stripplot.

warnings.warn(msg, UserWarning)



In [30]: titanic = sns.load_dataset("titanic")

In [31]: titanic

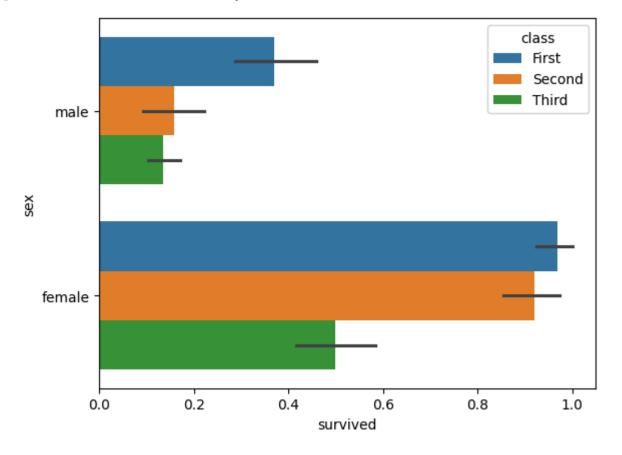
survived pclass fare embarked class who a Out[31]: age sibsp parch sex 0 0 3 male 22.0 7.2500 S Third man 1 1 female 38.0 71.2833 С 1 1 First woman 2 7.9250 S 1 3 female 26.0 0 Third woman

woman	First	S	53.1000	0	1	35.0	female	1	1	3
man	Third	S	8.0500	0	0	35.0	male	3	0	4
•••	•••						•••	•••		•••
man	Second	S	13.0000	0	0	27.0	male	2	0	886
woman	First	S	30.0000	0	0	19.0	female	1	1	887
woman	Third	S	23.4500	2	1	NaN	female	3	0	888
man	First	С	30.0000	0	0	26.0	male	1	1	889
man	Third	Q	7.7500	0	0	32.0	male	3	0	890

891 rows × 15 columns

In [35]: sns.barplot(x="survived", y="sex",hue="class",data=titanic)

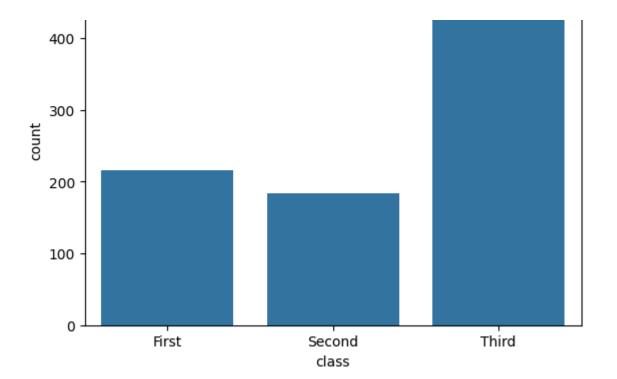
Out[35]: <Axes: xlabel='survived', ylabel='sex'>



In [40]: sns.countplot(x="class",
 data=titanic)

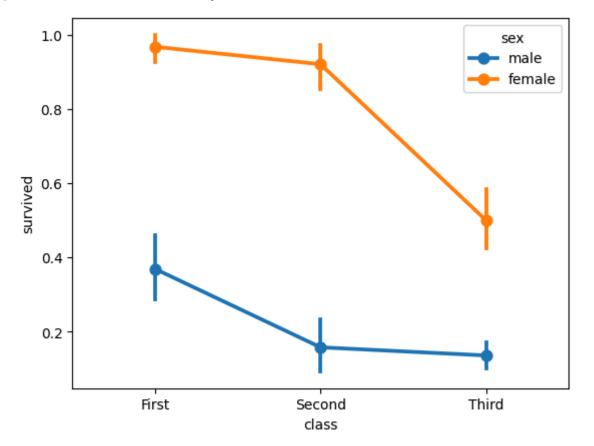
Out[40]: <Axes: xlabel='class', ylabel='count'>

500 -



In [41]:
sns.pointplot(x="class", y="survived", hue="sex", data=titanic)

Out[41]: <Axes: xlabel='class', ylabel='survived'>



```
In [42]: sns.boxplot(data=iris)
```

Out[42]: <Axes: >

