

# Assignment 3 - Matplotlib and Seaborn

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## Importing Libraries

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

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In [ ]:
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## Getting the data

```
In [2]: iris = sns.load_dataset('iris')
```

```
In [6]: iris
```

```
Out[6]:
```

	sepal_length	sepal_width	petal_length	petal_width	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	6.3	2.5	5.0	1.9	virginica
147	6.5	3.0	5.2	2.0	virginica
148	6.2	3.4	5.4	2.3	virginica
149	5.9	3.0	5.1	1.8	virginica

150 rows × 5 columns

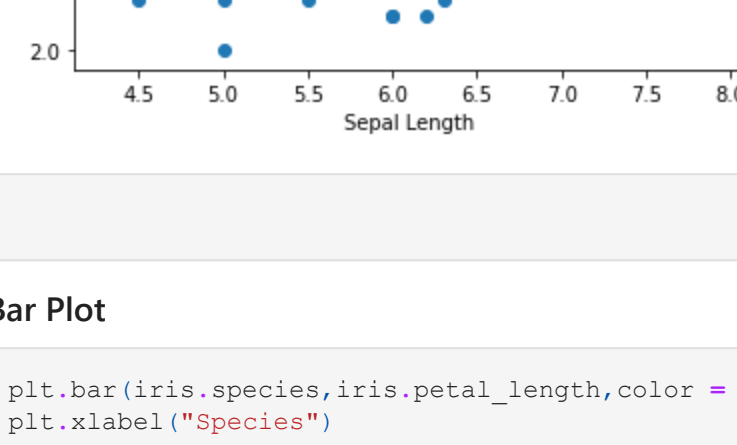
```
In [ ]:
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## Matplotlib Visualizations

### Line Plot

```
In [33]: plt.plot(np.linspace(10,20),
np.linspace(15,25),
color = "blue",
label = "x v/s y")

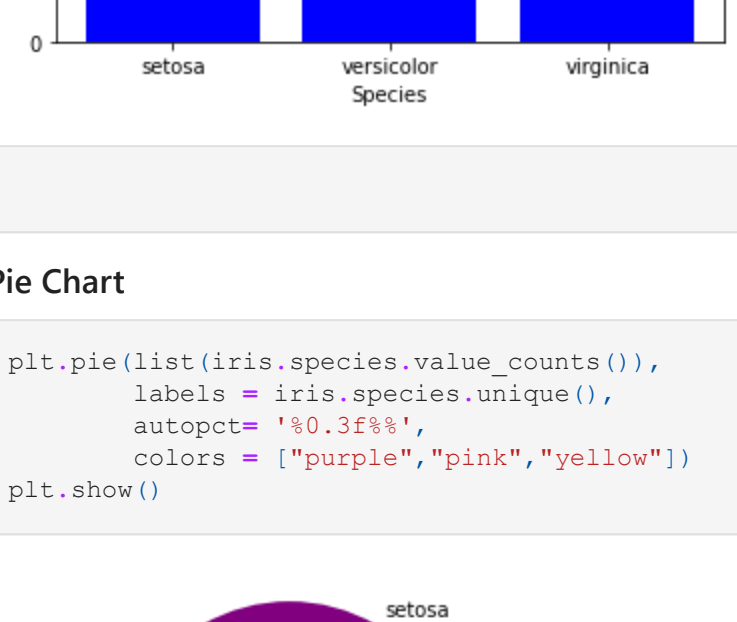
plt.xlabel("x")
plt.ylabel("y")
plt.legend()
plt.show()
```



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### Scatter Plot

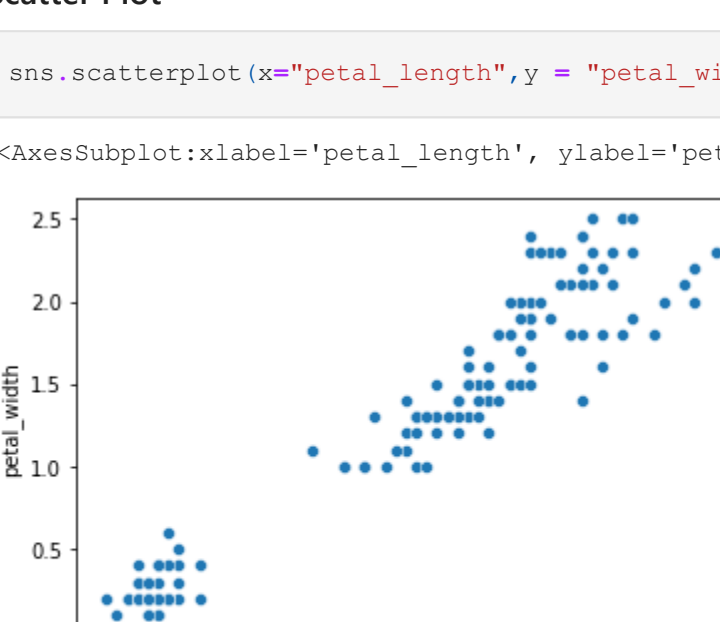
```
In [34]: plt.scatter(iris.sepal_length,iris.sepal_width)
plt.xlabel("Sepal Length")
plt.ylabel("Sepal Width")
plt.title("Sepal Width v/s Length")
plt.show()
```



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### Bar Plot

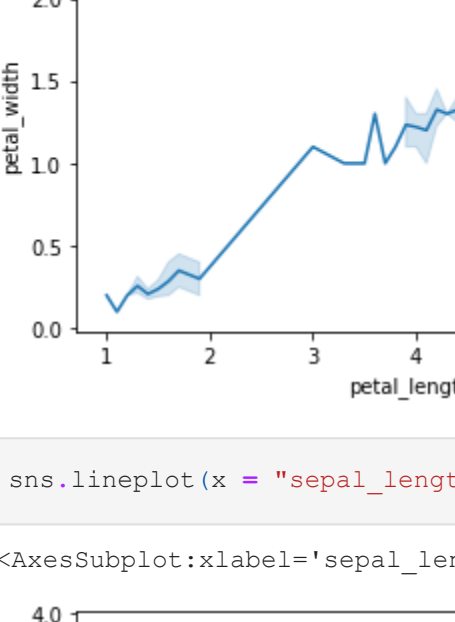
```
In [35]: plt.bar(iris.species,iris.petal_length,color = "blue")
plt.xlabel("Species")
plt.ylabel("Petal Length")
plt.title("Petal Length Distribution of different Iris Species")
plt.show()
```



```
In [ ]:
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### Pie Chart

```
In [38]: plt.pie(list(iris.species.value_counts()),
labels = iris.species.unique(),
autopct = '%0.3f%%',
colors = ["purple","pink","yellow"])
plt.show()
```



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In [ ]:
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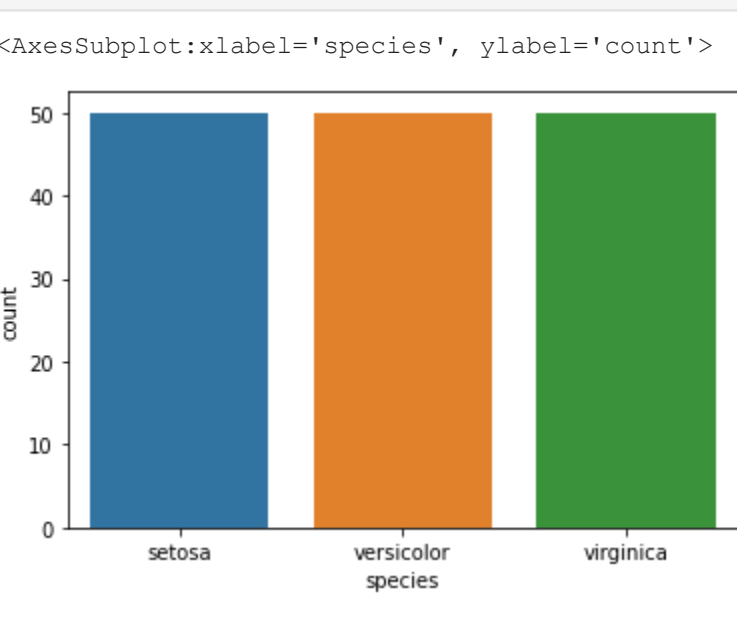
## Seaborn Visualizations

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### Scatter Plot

```
In [37]: sns.scatterplot(x="petal_length",y = "petal_width",data = iris)
```

```
Out[37]: <AxesSubplot:xlabel='petal_length', ylabel='petal_width'>
```



```
In [ ]:
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### Line Plot

```
In [5]: sns.lineplot(x = "petal_length",y = "petal_width",data = iris)
```

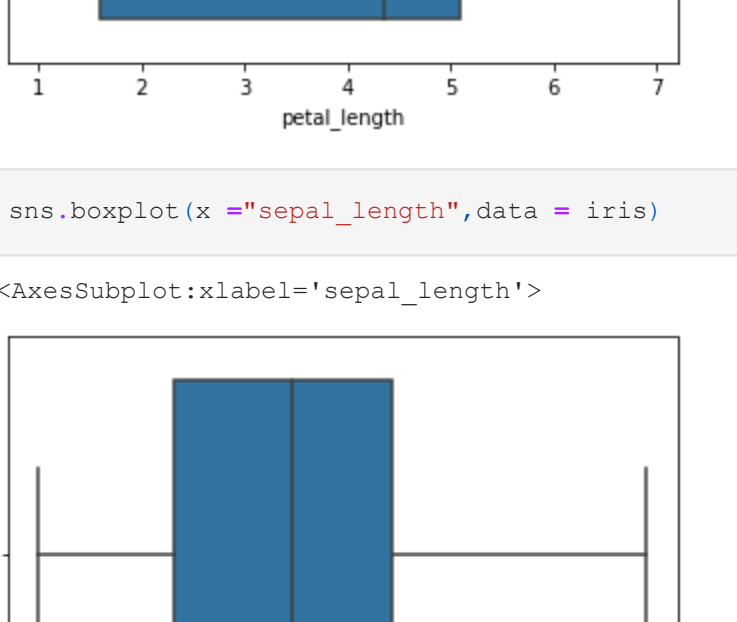
```
Out[5]: <AxesSubplot:xlabel='petal_length', ylabel='petal_width'>
```



```
In [8]:
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```
sns.lineplot(x = "sepal_length",y = "sepal_width",data = iris)
```

```
Out[8]: <AxesSubplot:xlabel='sepal_length', ylabel='sepal_width'>
```

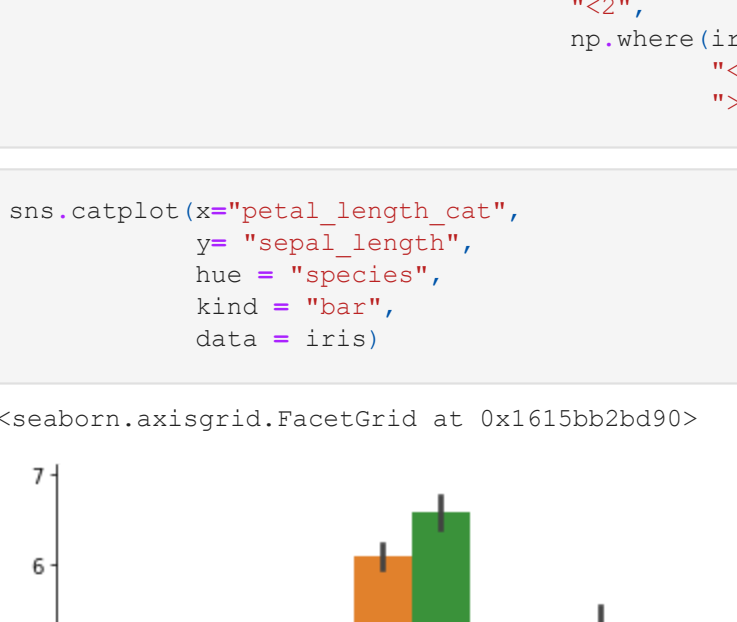


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### Count Plot

```
In [9]: sns.countplot(x = "species",data = iris)
```

```
Out[9]: <AxesSubplot:xlabel='species', ylabel='count'>
```

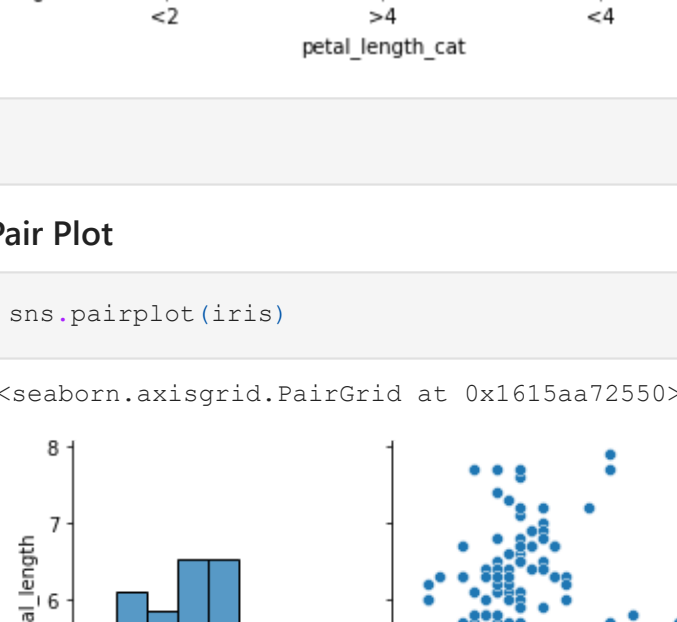


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### Box Plot

```
In [12]: sns.boxplot(x = "petal_width",data = iris)
```

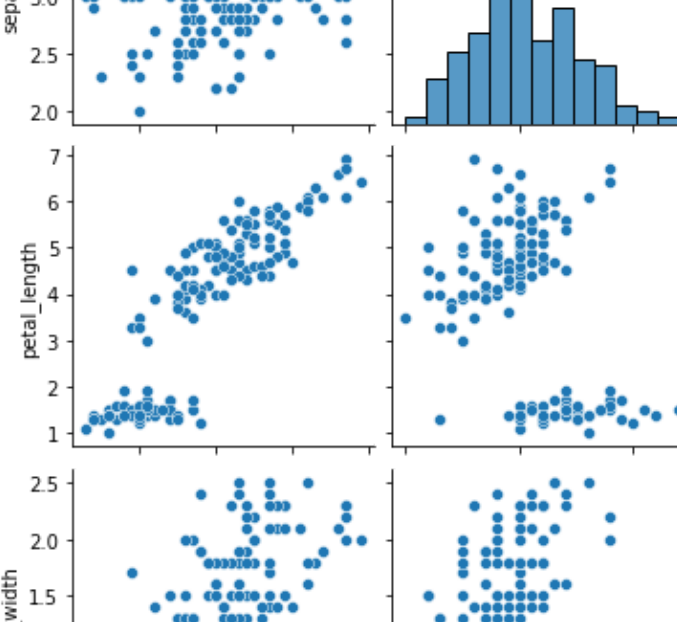
```
Out[12]: <AxesSubplot:xlabel='petal_width'>
```



```
In [13]:
```

```
sns.boxplot(x = "petal_length",data = iris)
```

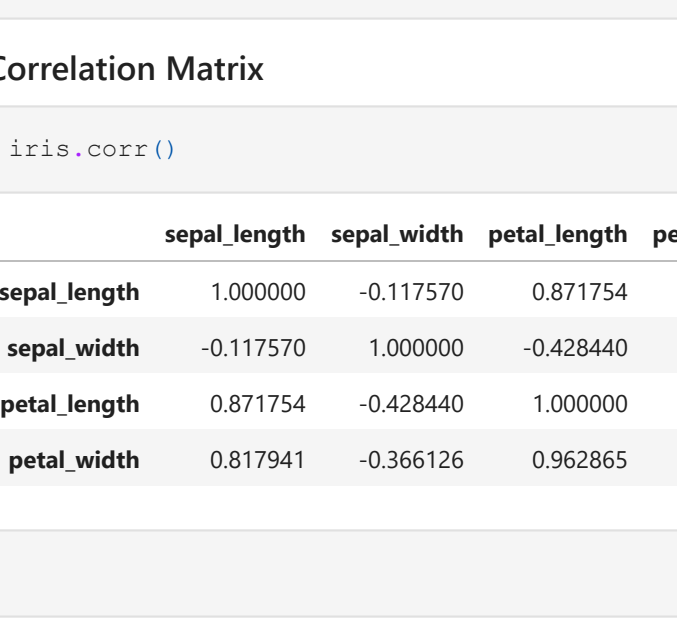
```
Out[13]: <AxesSubplot:xlabel='petal_length'>
```



```
In [14]:
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```
sns.boxplot(x = "sepal_length",data = iris)
```

```
Out[14]: <AxesSubplot:xlabel='sepal_length'>
```



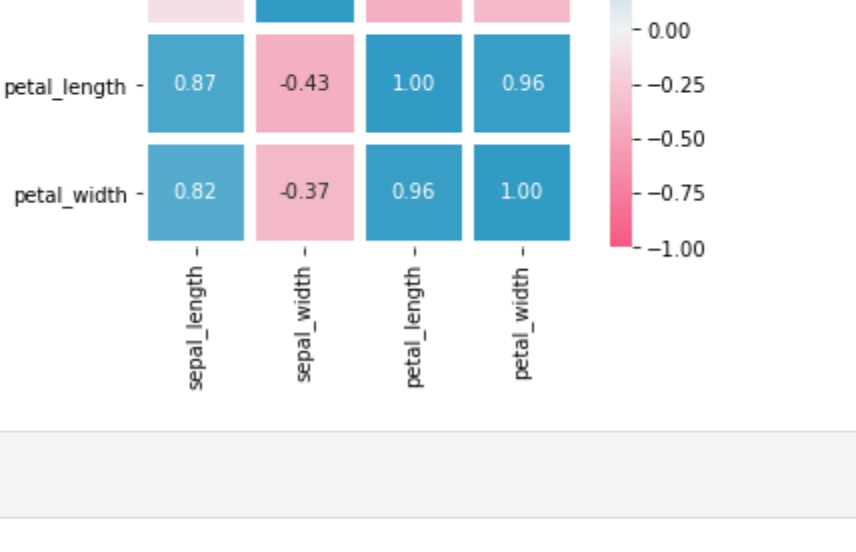
```
In [ ]:
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## Categorical Plot

```
In [27]: iris["petal_length_cat"] = np.where(iris["petal_length"]<2,
"setosa",
np.where(iris["petal_length"]<4,
"versicolor",
">4"))
```

```
In [28]: sns.catplot(x="petal_length_cat",
y = "sepal_length",
hue = "species",
kind = "bar",
data = iris)
```

```
Out[28]: <seaborn.axisgrid.FacetGrid at 0x1615bb2bd90>
```



```
In [ ]:
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## Pair Plot

```
In [29]: sns.pairplot(iris)
```

```
Out[29]: <seaborn.axisgrid.PairGrid at 0x1615aa72550>
```



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## Correlation Matrix

```
In [30]: iris.corr()
```

```
Out[30]:
```

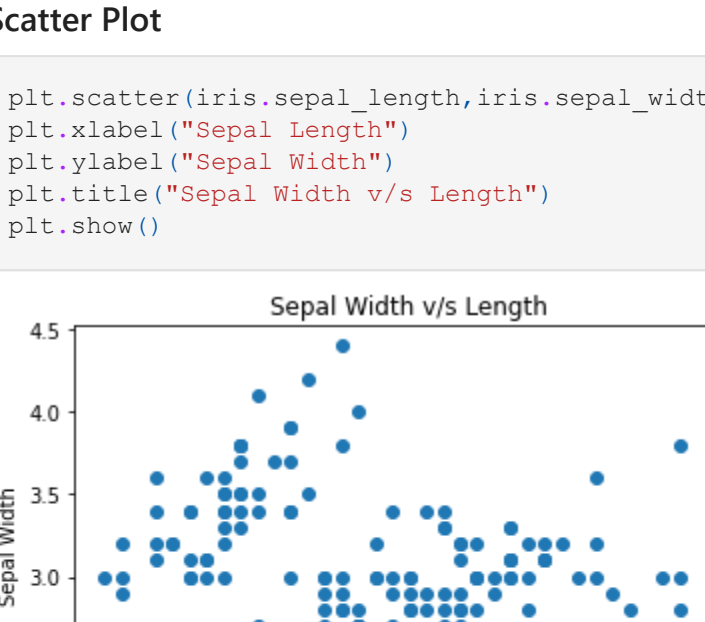
	sepal_length	sepal_width	petal_length	petal_width
sepal_length	1.000000	-0.117570	0.871754	0.817941
sepal_width	-0.117570	1.000000	-0.428440	-0.366126
petal_length	0.871754	-0.428440	1.000000	0.962865
petal_width	0.817941	-0.366126	0.962865	1.000000

```
In [ ]:
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## Heat Map

```
In [32]: sns.heatmap(iris.corr(),
annot=True,
fmt=".2f",
linewidths=5,
cmap=sns.diverging_palette(0,
230,
90,
60,
as_cmap=True),
vmin=-1,
vmax=1,
square=True)
```

```
Out[32]: <AxesSubplot:~>
```



```
In [ ]:
```