

# Seaborn Cheat Sheet

This cheat sheet covers the essential and advanced features of Seaborn, a powerful library for creating informative and attractive statistical graphics in Python.

## 1. Introduction to Seaborn

### 1.1 What is Seaborn?

Seaborn is a Python data visualization library based on Matplotlib. It provides a high-level interface for drawing attractive and informative statistical graphics.

### 1.2 Installing Seaborn

You can install Seaborn using pip:

```
pip install seaborn
```

Or, if you are using Anaconda:

```
conda install seaborn
```

### 1.3 Importing Seaborn

To use Seaborn, you need to import it in your script:

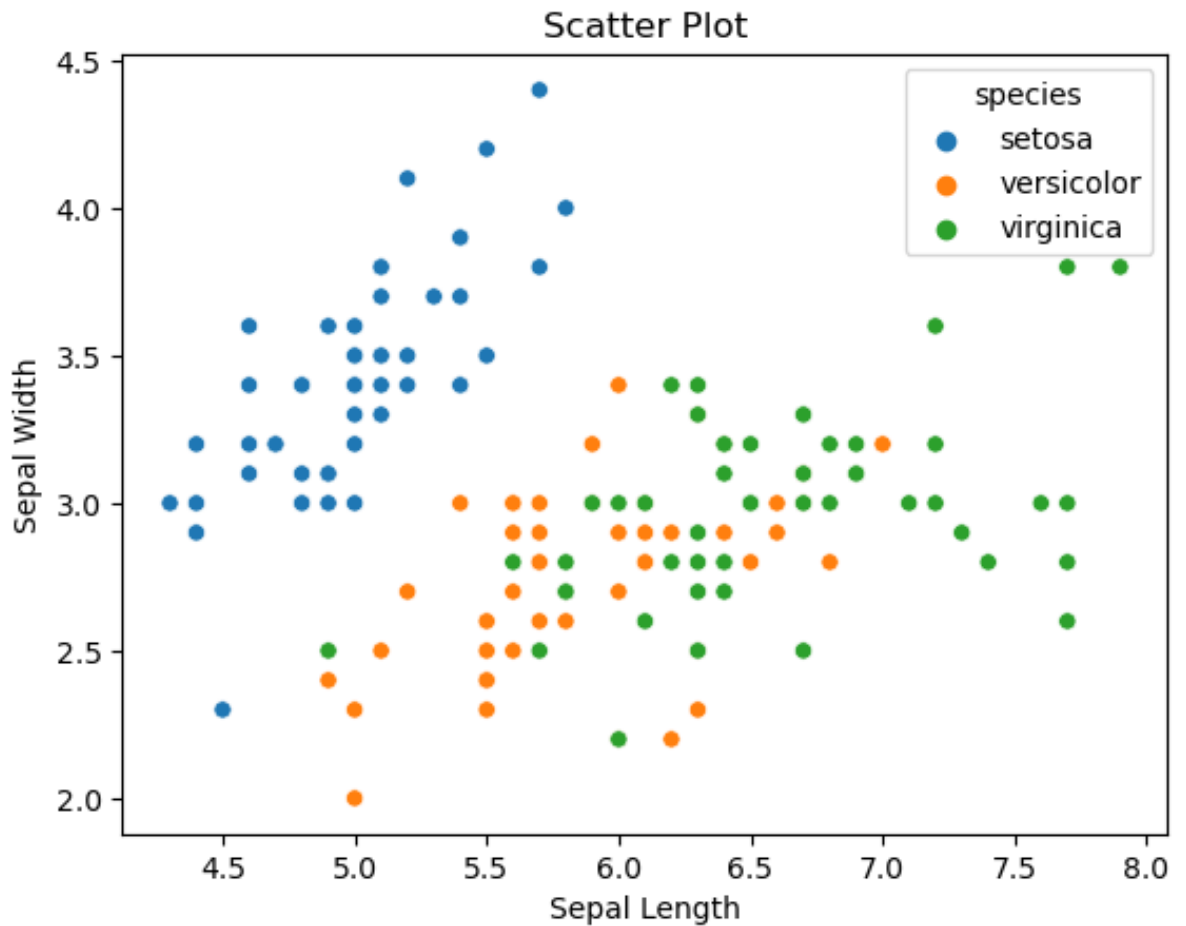
```
In [1]: import seaborn as sns
import matplotlib.pyplot as plt
# Now you can use Seaborn functions with the sns prefix
```

## 2. Basic Plots

### 2.1 Scatter Plot

Scatter plots are used to represent individual data points.

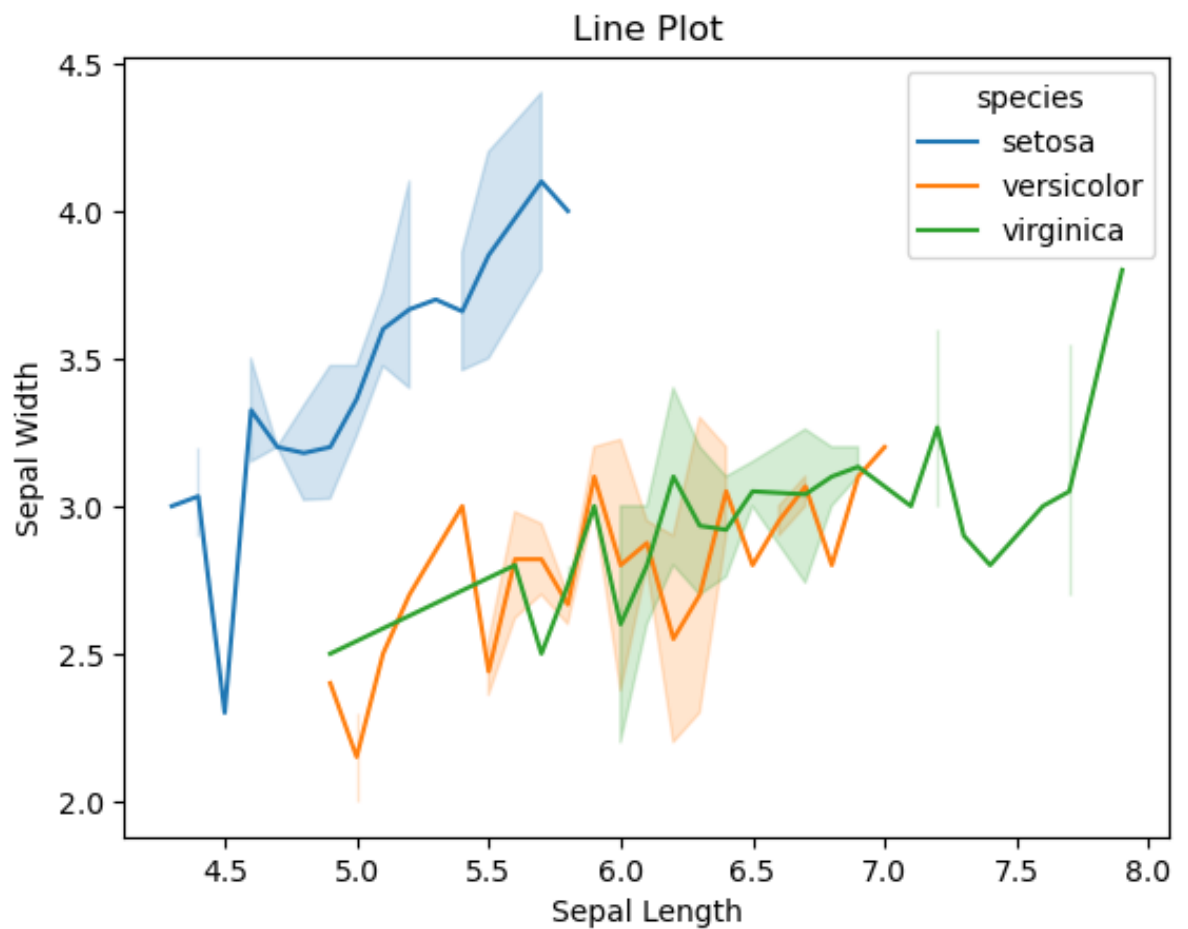
```
In [2]: data = sns.load_dataset('iris')
sns.scatterplot(x='sepal_length', y='sepal_width', hue='species', data=data)
plt.title('Scatter Plot')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.show()
```



## 2.2 Line Plot

Line plots are used to represent data points connected by straight lines.

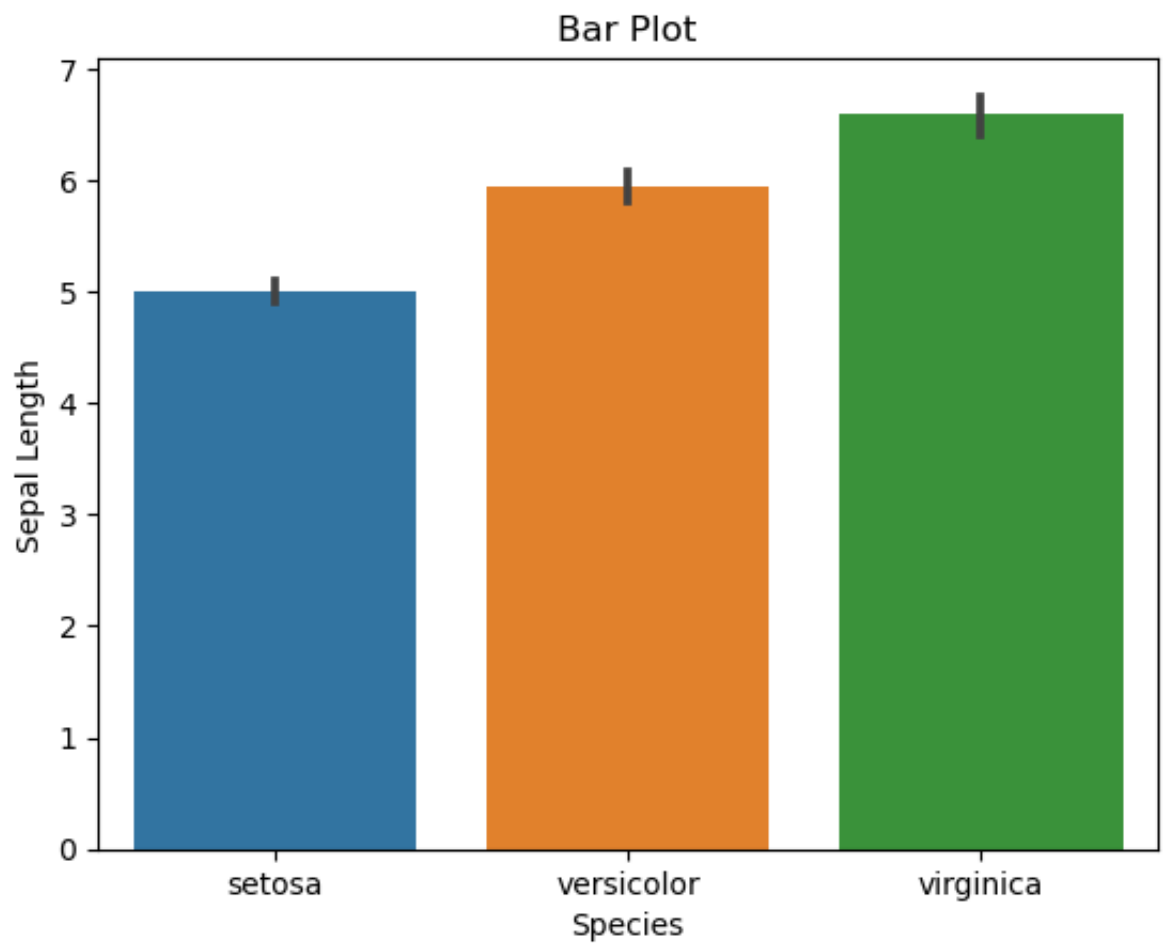
```
In [3]: sns.lineplot(x='sepal_length', y='sepal_width', hue='species', data=
plt.title('Line Plot')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.show()
```



## 2.3 Bar Plot

Bar plots are used to represent data with rectangular bars.

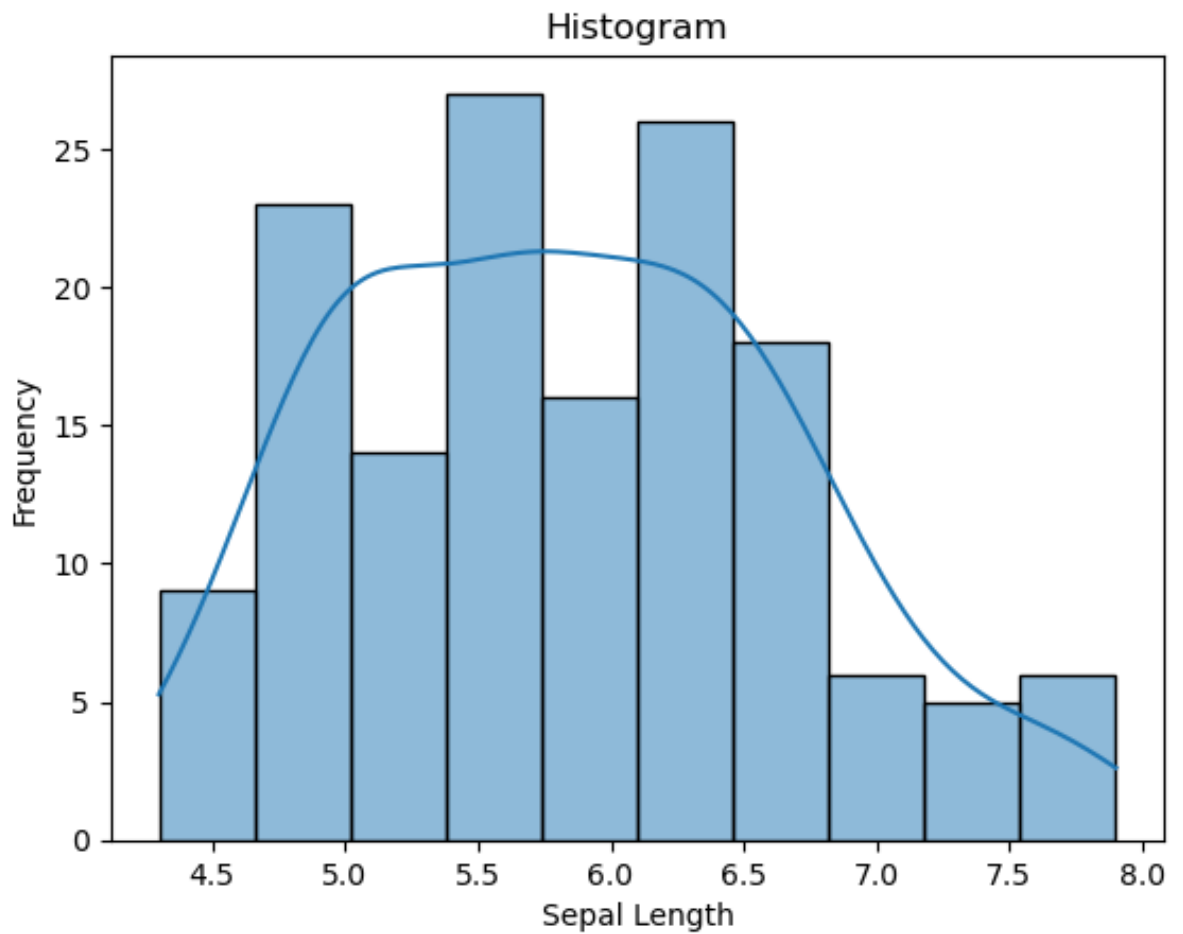
```
In [4]: sns.barplot(x='species', y='sepal_length', data=data)
plt.title('Bar Plot')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```



## 2.4 Histogram

Histograms are used to represent the distribution of a dataset.

```
In [5]: sns.histplot(data['sepal_length'], bins=10, kde=True)
plt.title('Histogram')
plt.xlabel('Sepal Length')
plt.ylabel('Frequency')
plt.show()
```

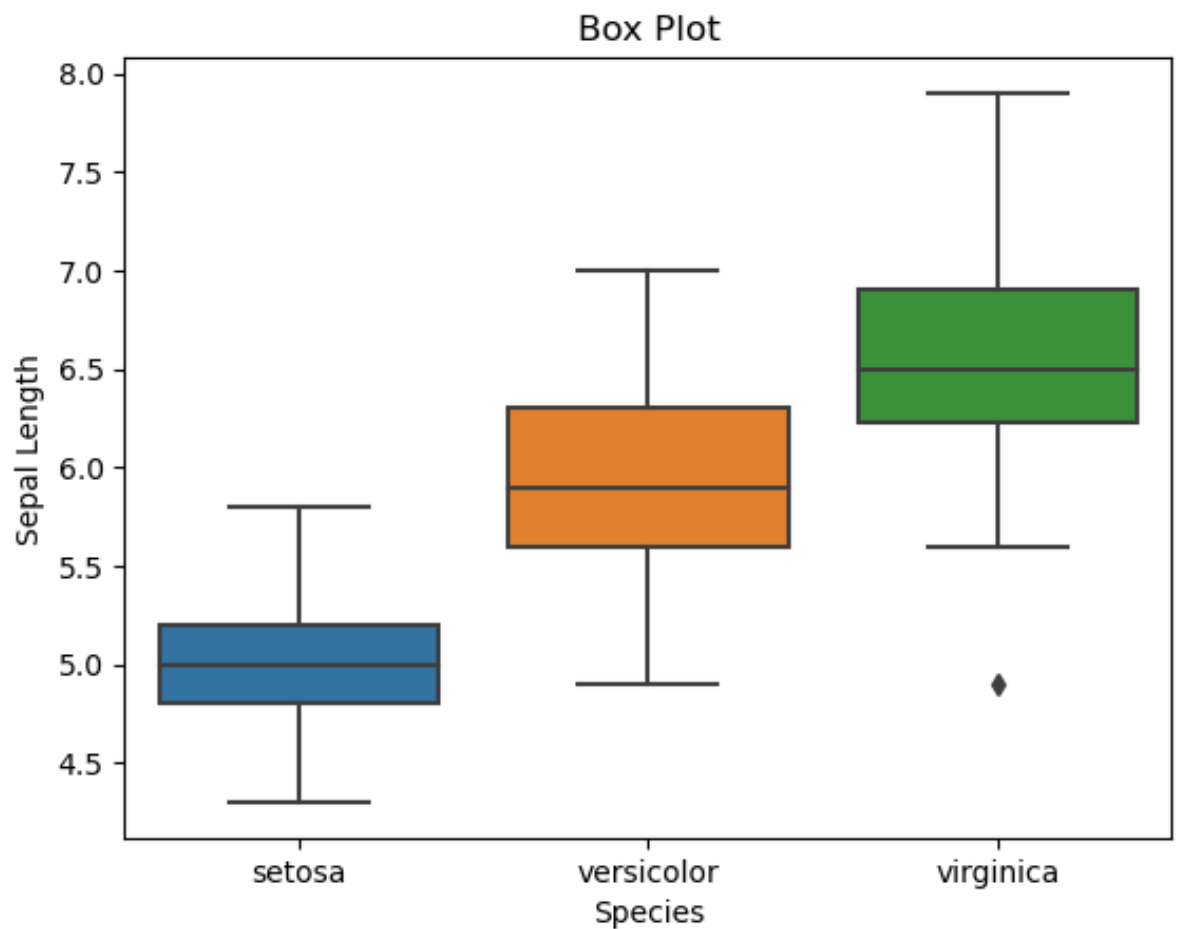


## 3. Statistical Plots

### 3.1 Box Plot

Box plots are used to visualize the distribution of data based on a five-number summary.

```
In [6]: sns.boxplot(x='species', y='sepal_length', data=data)
plt.title('Box Plot')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```



### 3.2 Violin Plot

Violin plots are similar to box plots but also show the density of the data at different values.

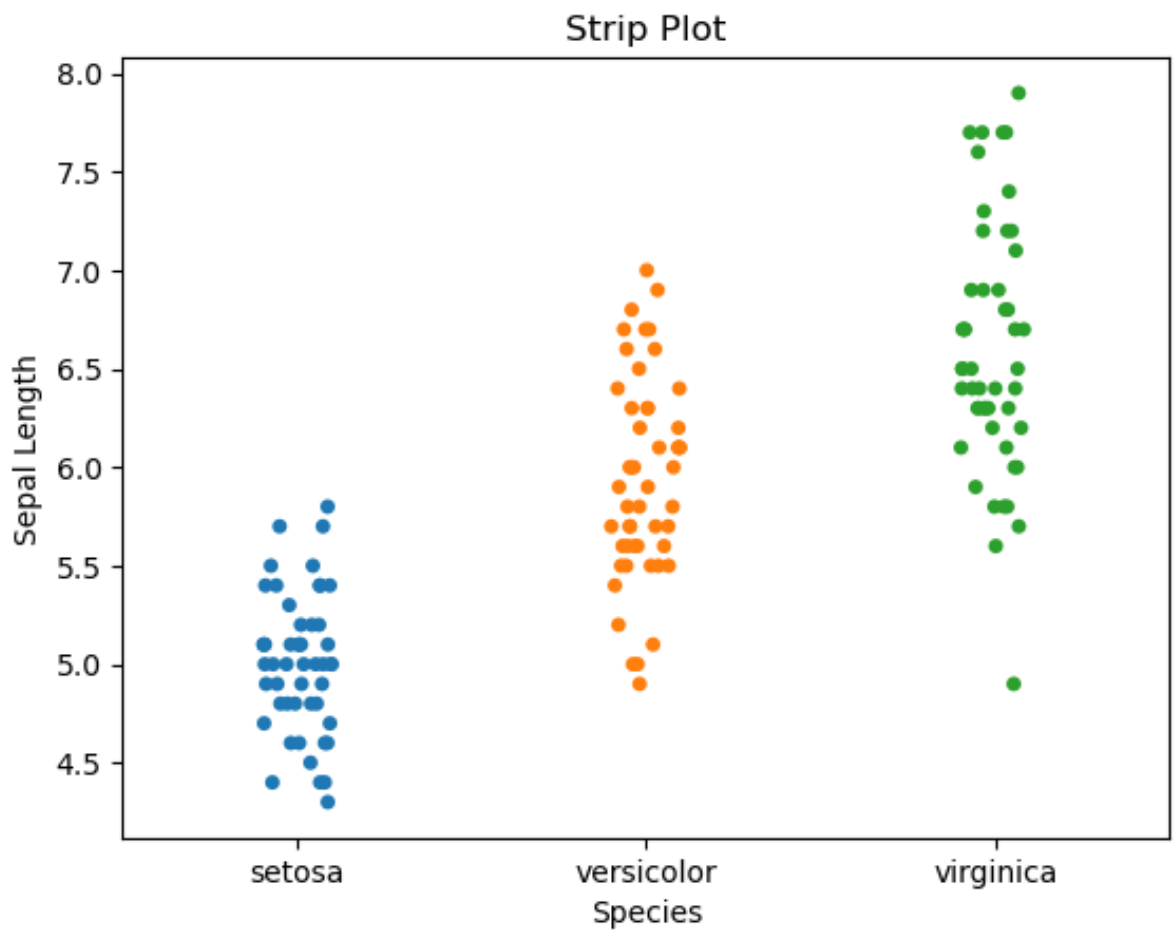
```
In [7]: sns.violinplot(x='species', y='sepal_length', data=data)
plt.title('Violin Plot')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```



### 3.3 Strip Plot

Strip plots are used to represent individual data points with a jitter to avoid overlap.

```
In [8]: sns.stripplot(x='species', y='sepal_length', data=data)
plt.title('Strip Plot')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```

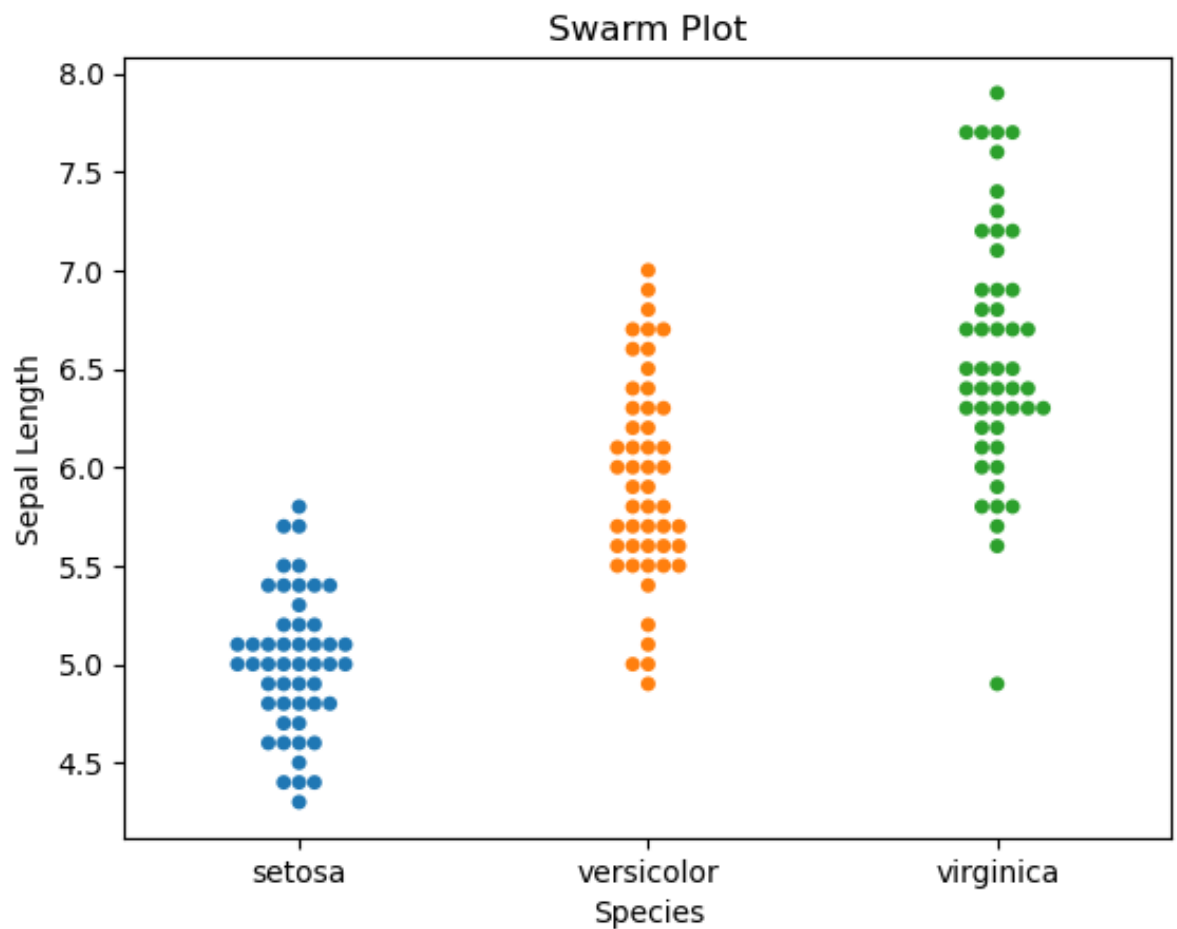


### 3.4 Swarm Plot

Swarm plots are similar to strip plots but avoid overlapping data points.



```
In [9]: sns.swarmplot(x='species', y='sepal_length', data=data)
plt.title('Swarm Plot')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```

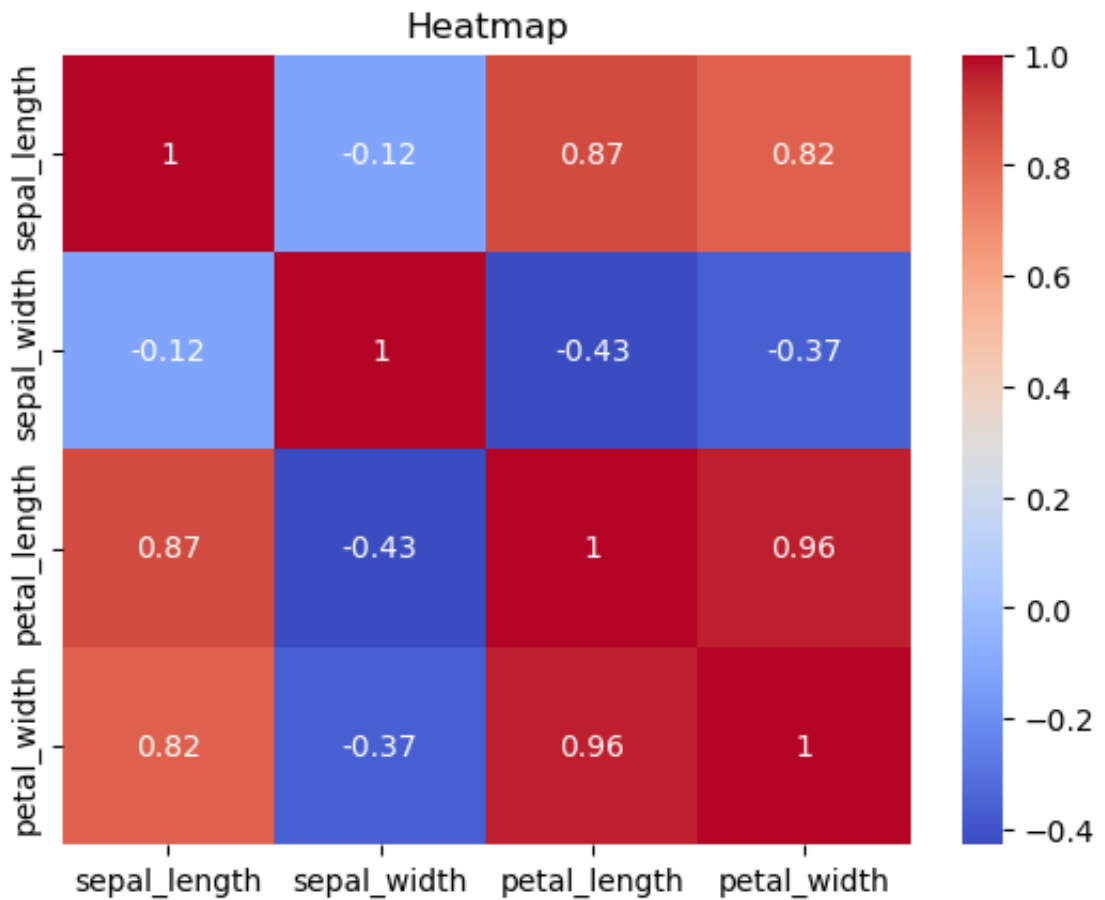


## 4. Matrix Plots

### 4.1 Heatmap

Heatmaps are used to represent data values as colors in a matrix.

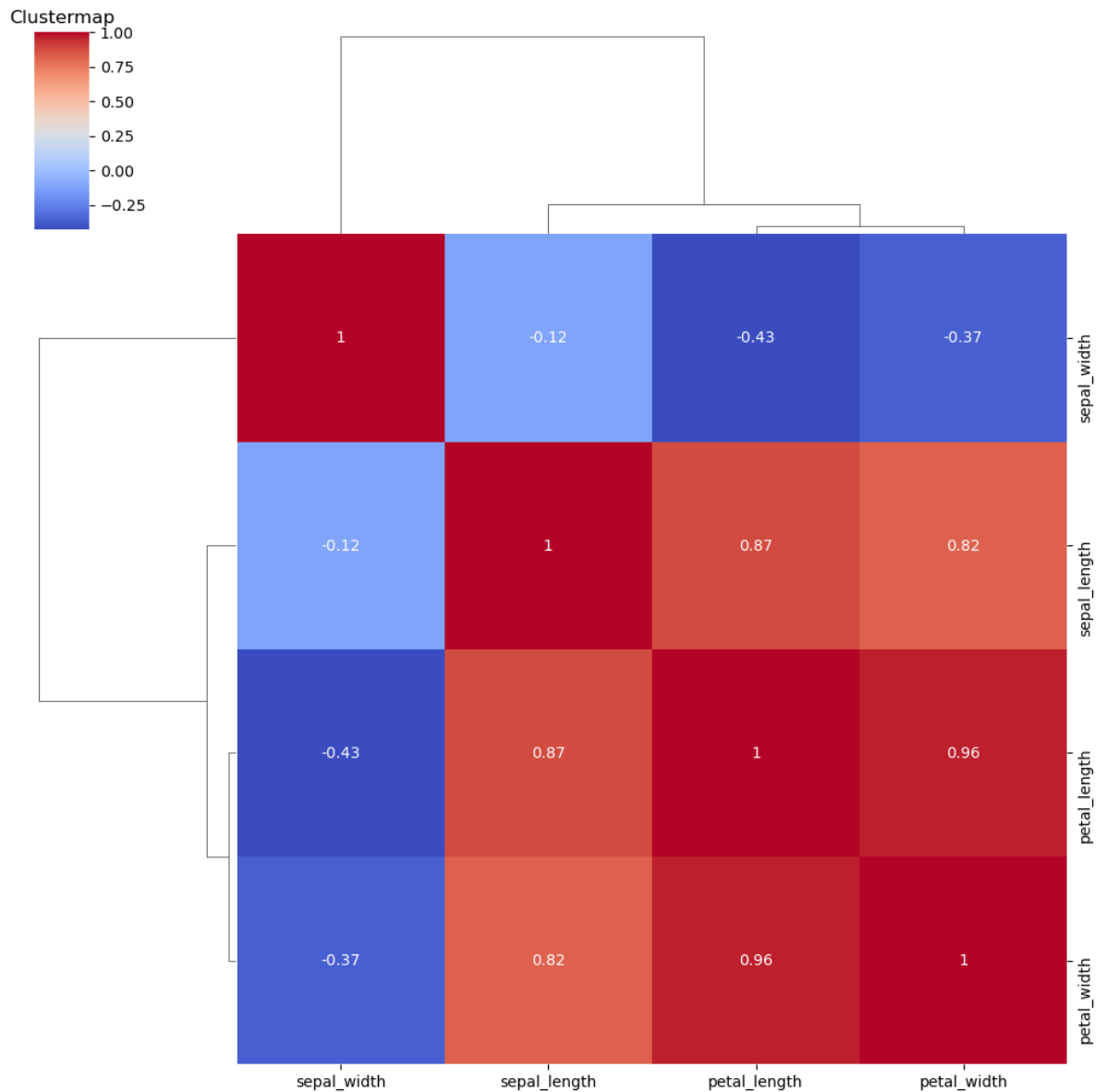
```
In [10]: corr = data.corr()
sns.heatmap(corr, annot=True, cmap='coolwarm')
plt.title('Heatmap')
plt.show()
```



## 4.2 Clustermap

Clustermaps are used to represent data values as colors in a matrix with hierarchical clustering.

```
In [11]: sns.clustermap(corr, annot=True, cmap='coolwarm')  
plt.title('Clustermap')  
plt.show()
```

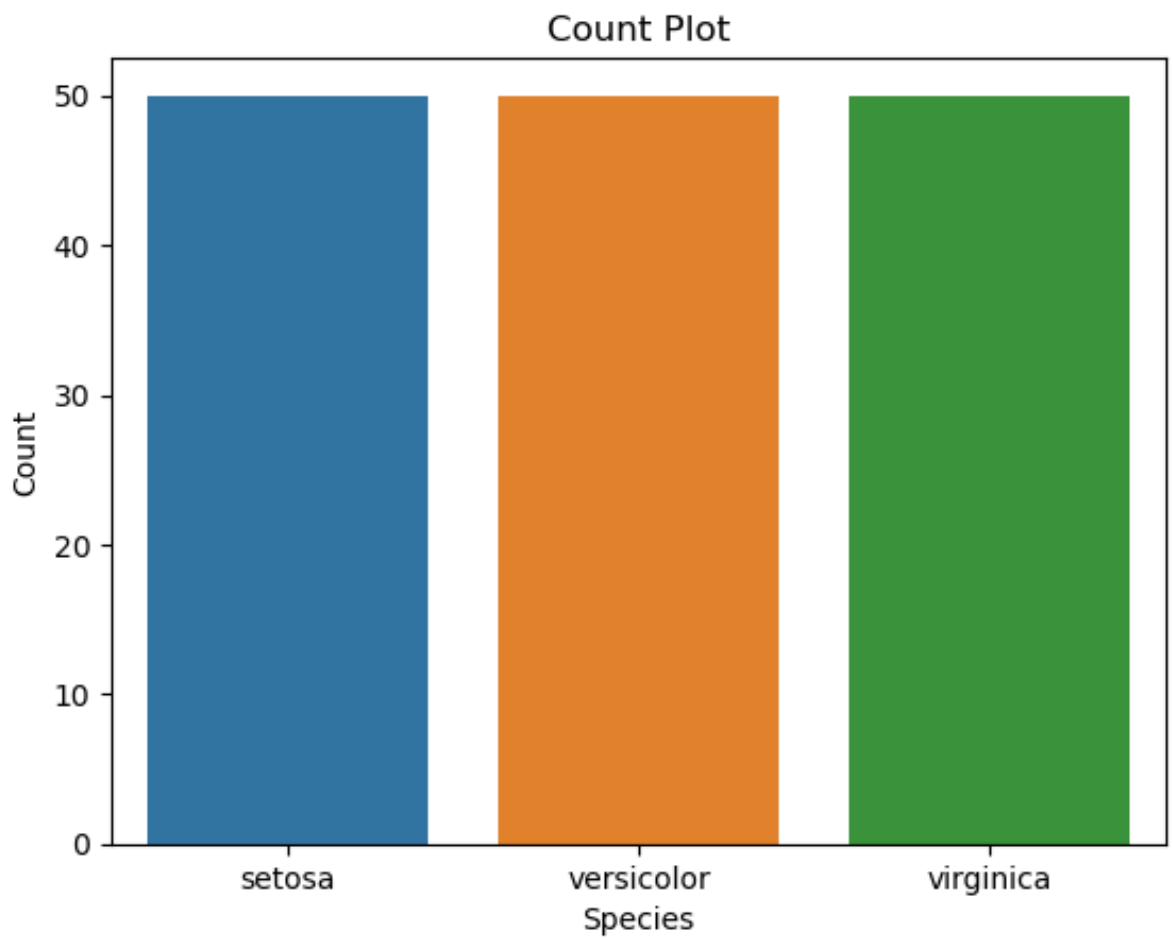


## 5. Categorical Plots

### 5.1 Count Plot

Count plots are used to show the counts of observations in each categorical bin using bars.

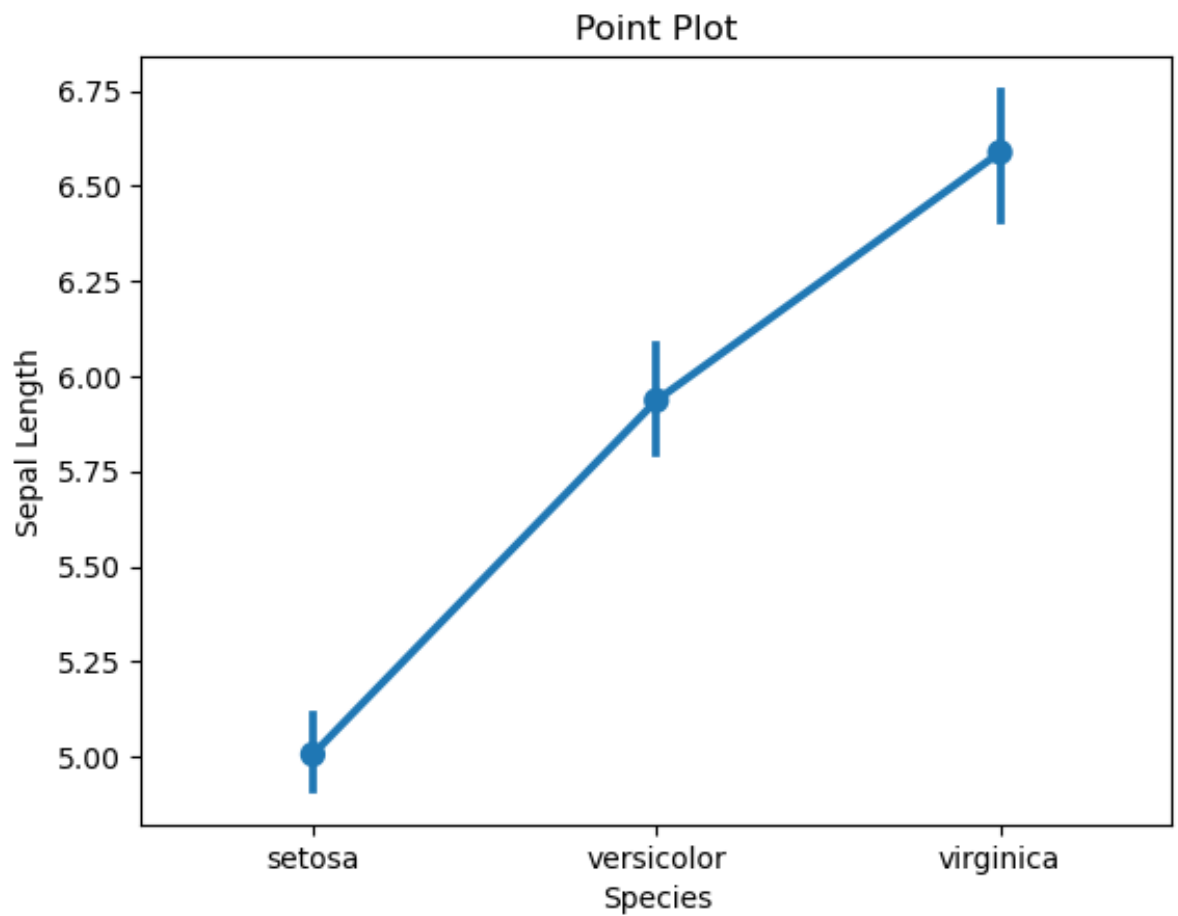
```
In [12]: sns.countplot(x='species', data=data)
plt.title('Count Plot')
plt.xlabel('Species')
plt.ylabel('Count')
plt.show()
```



## 5.2 Point Plot

Point plots are used to show point estimates and confidence intervals using scatter points.

```
In [13]: sns.pointplot(x='species', y='sepal_length', data=data)
plt.title('Point Plot')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```



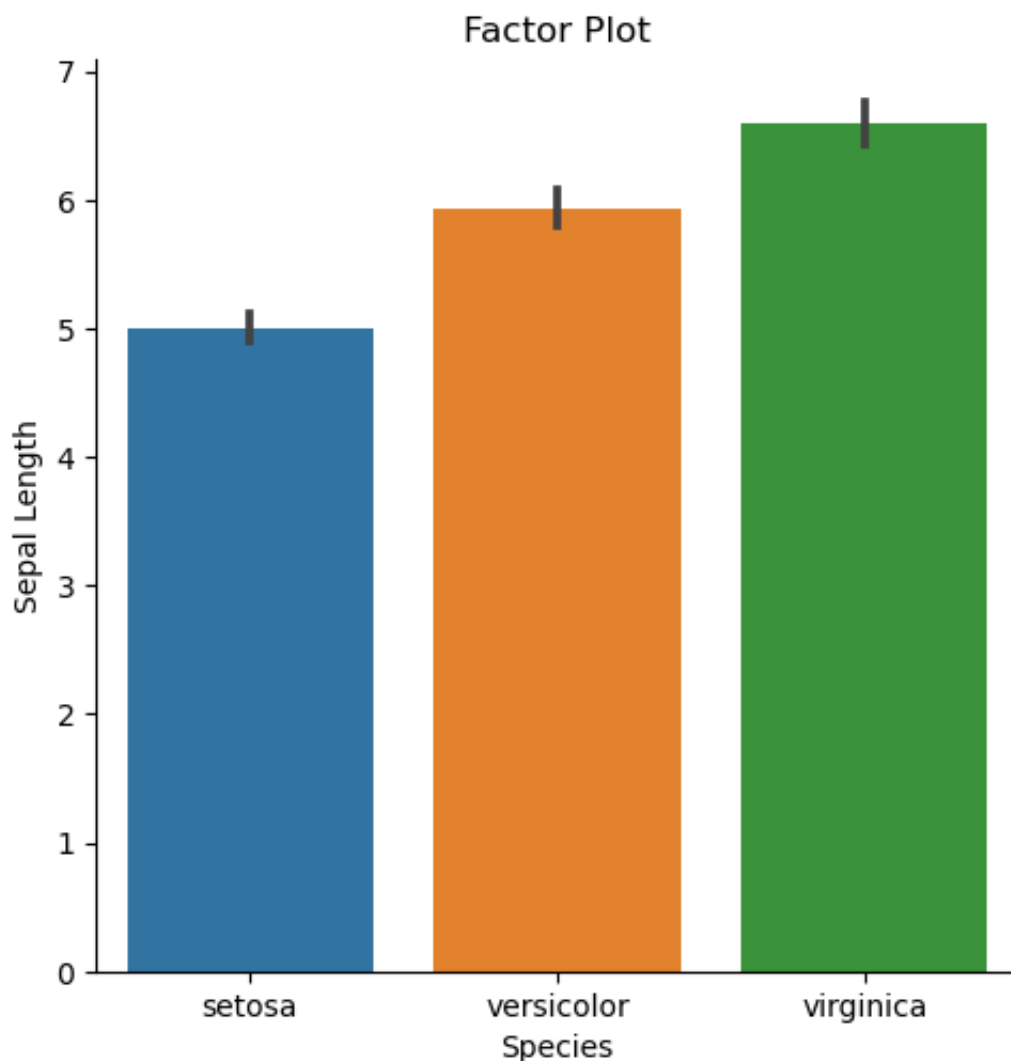
### 5.3 Factor Plot

Factor plots are used to draw multi-faceted categorical plots.

```
In [14]: sns.factorplot(x='species', y='sepal_length', data=data, kind='bar')
plt.title('Factor Plot')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```

/Users/ashishzangra/opt/anaconda3/lib/python3.9/site-packages/seaborn/categorical.py:3717: UserWarning: The `factorplot` function has been renamed to `catplot`. The original name will be removed in a future release. Please update your code. Note that the default `kind` in `factorplot` (`'point'`) has changed to `strip` in `catplot`.

warnings.warn(msg)

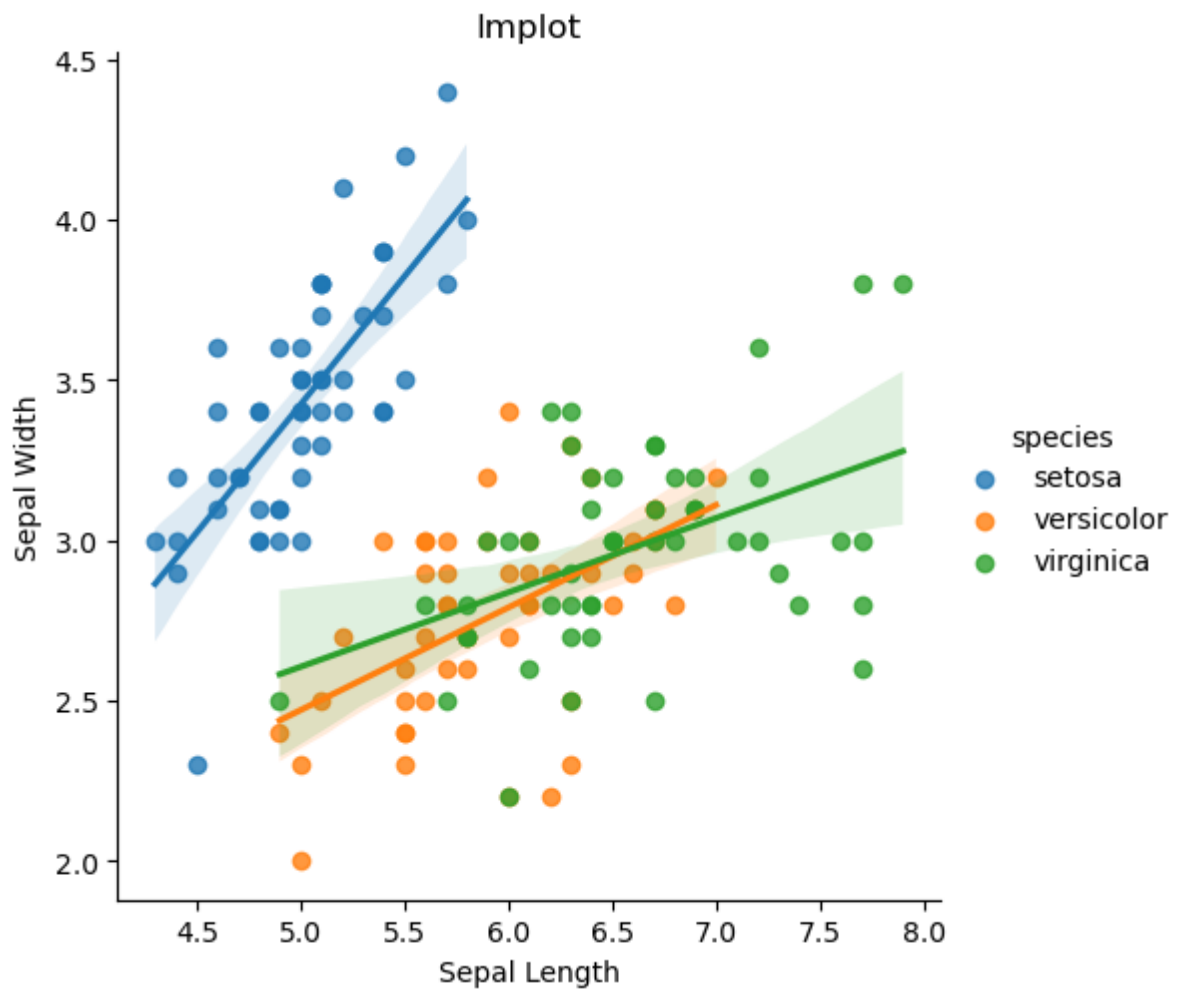


## 6. Regression Plots

### 6.1 Implot

Implot is used to draw a scatter plot with a linear regression model fit.

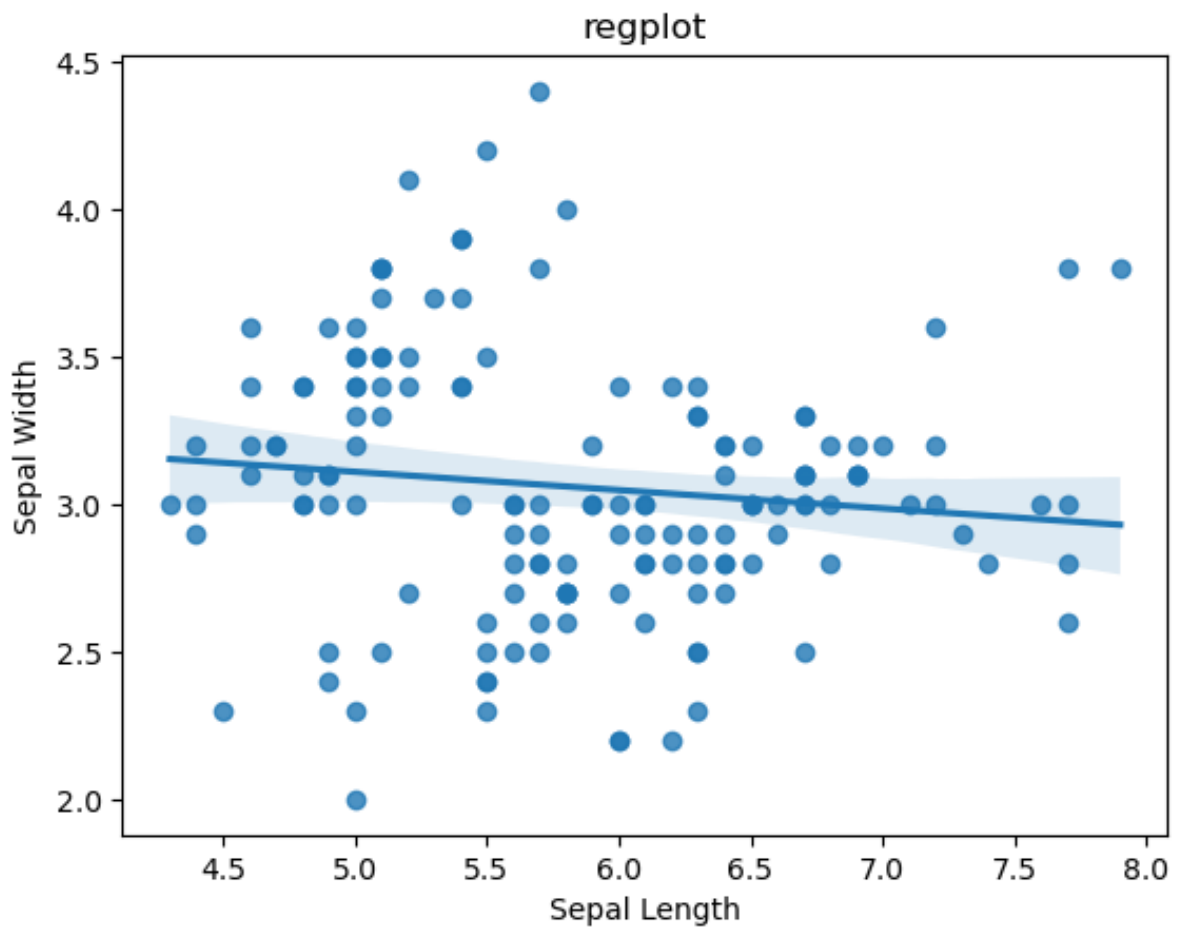
```
In [15]: sns.lmplot(x='sepal_length', y='sepal_width', hue='species', data=d)
plt.title('lmplot')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.show()
```



## 6.2 regplot

regplot is used to draw a scatter plot with a linear regression model fit.

```
In [16]: sns.regplot(x='sepal_length', y='sepal_width', data=data)
plt.title('regplot')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.show()
```

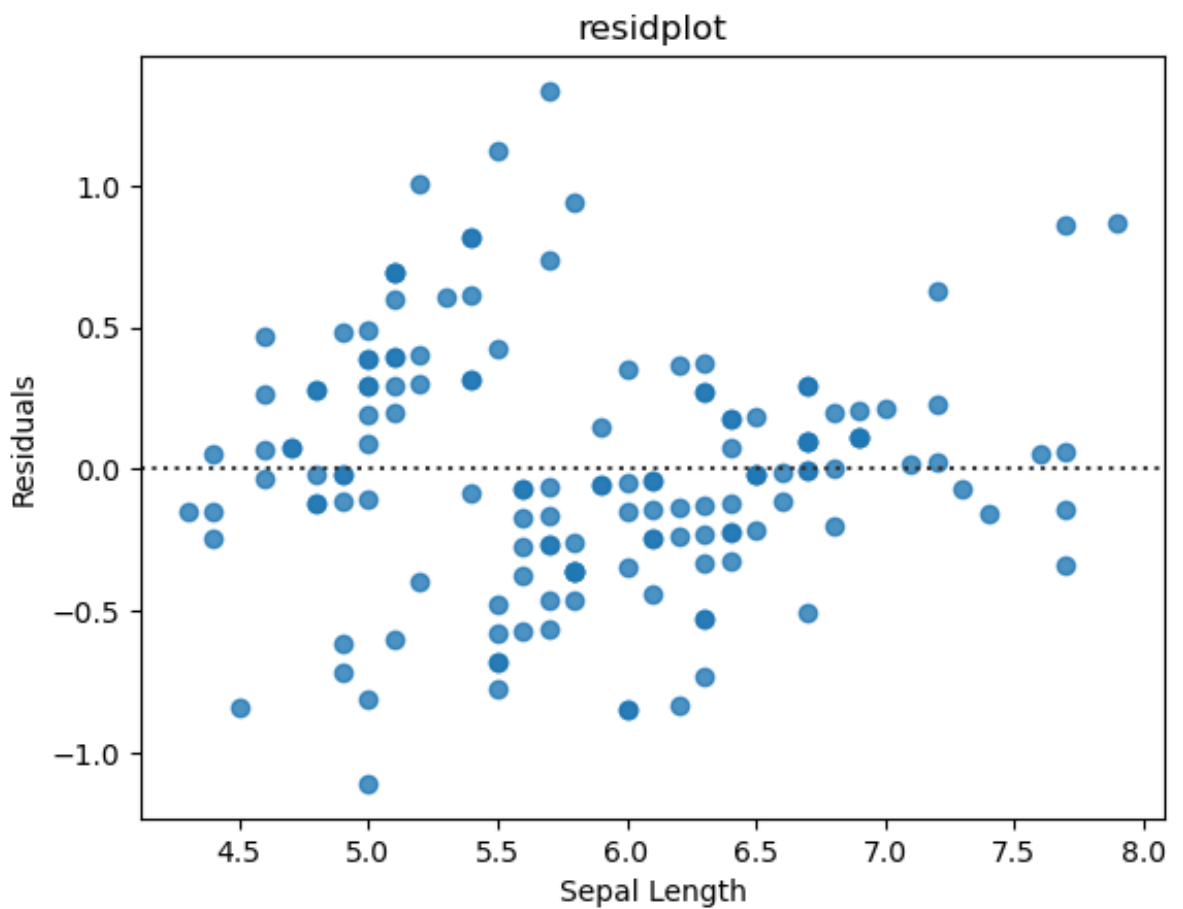


### 6.3 residplot

residplot is used to draw a scatter plot of the residuals of a linear regression.



```
In [17]: sns.residplot(x='sepal_length', y='sepal_width', data=data)
plt.title('residplot')
plt.xlabel('Sepal Length')
plt.ylabel('Residuals')
plt.show()
```



## 7. Distribution Plots

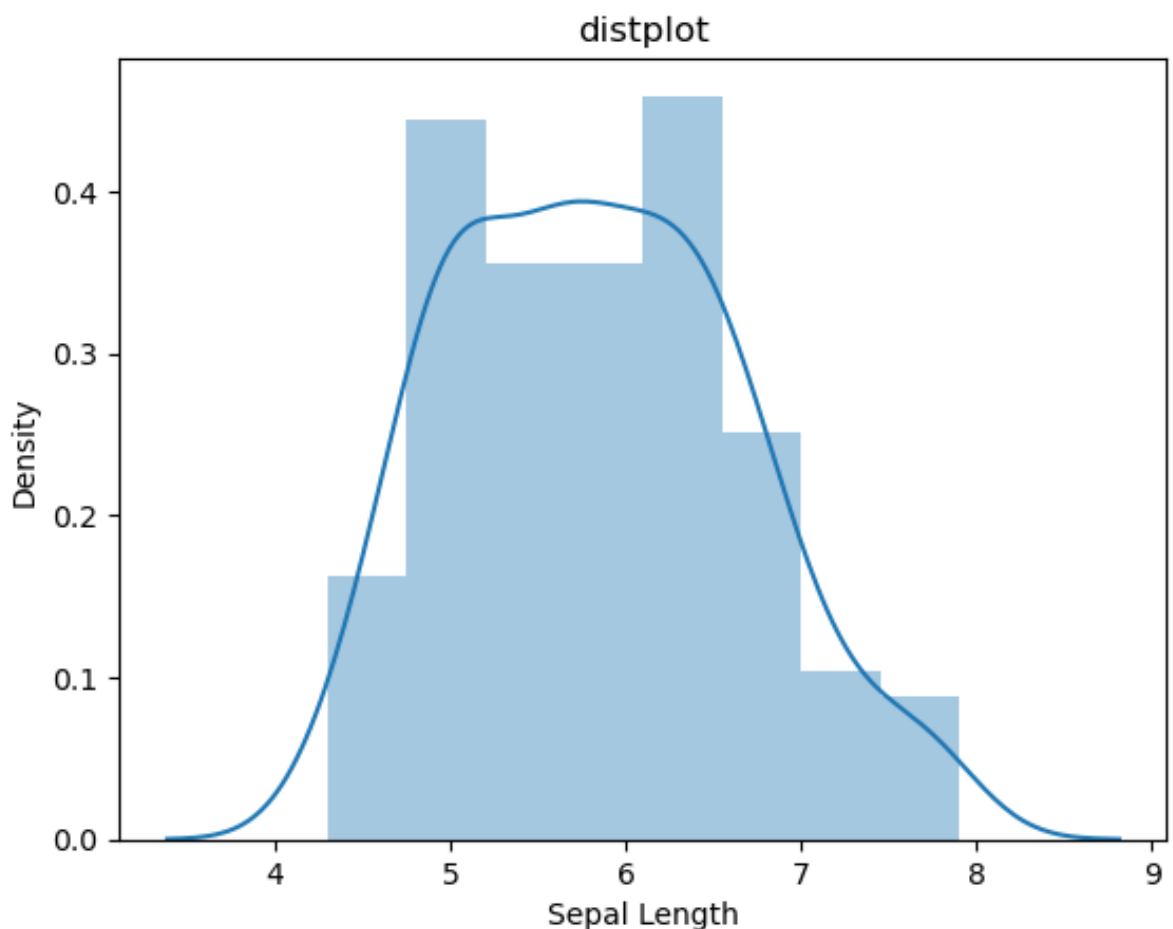
### 7.1 distplot

distplot is used to plot a univariate distribution of observations.

```
In [18]: sns.distplot(data['sepal_length'])  
plt.title('distplot')  
plt.xlabel('Sepal Length')  
plt.ylabel('Density')  
plt.show()
```

/Users/ashishzangra/opt/anaconda3/lib/python3.9/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).

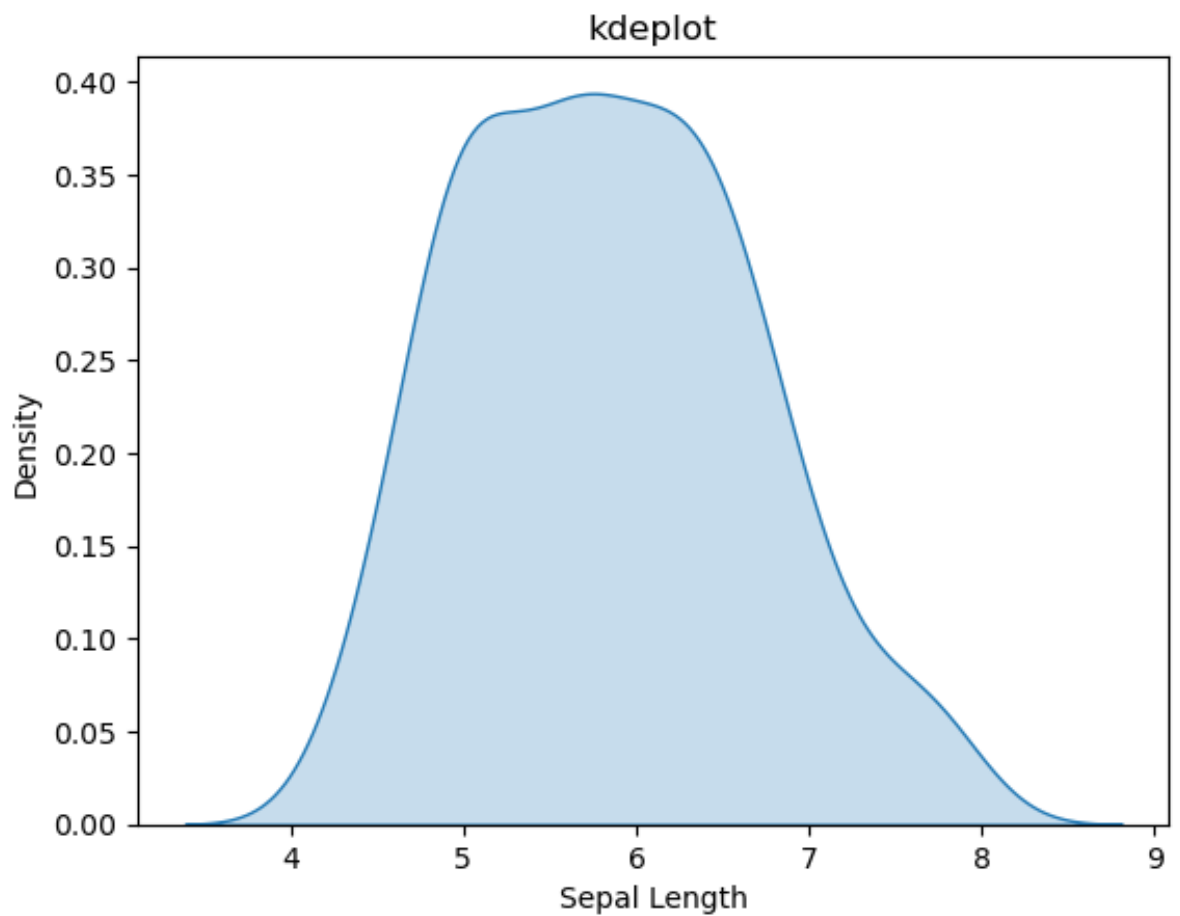
warnings.warn(msg, FutureWarning)



## 7.2 kdeplot

kdeplot is used to plot a univariate or bivariate kernel density estimate.

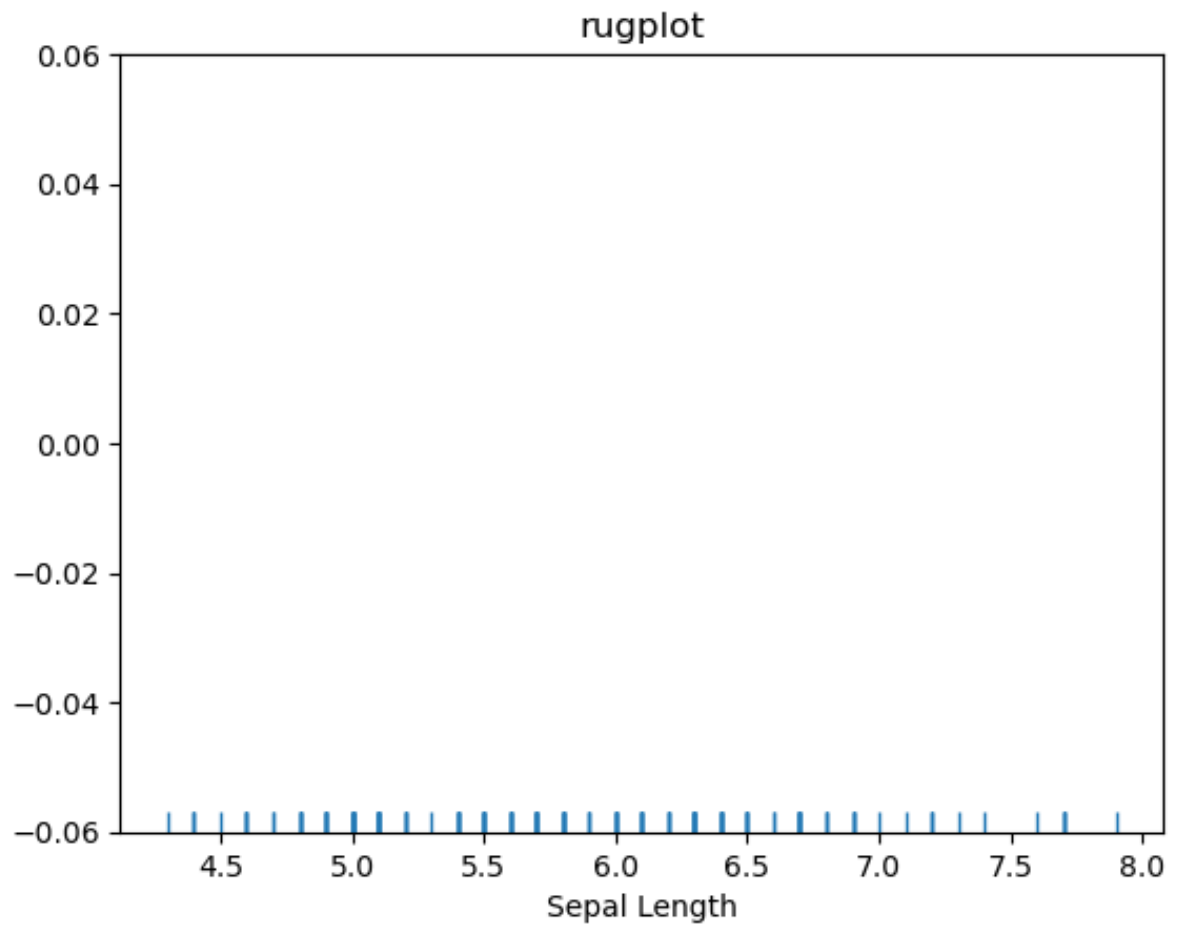
```
In [19]: sns.kdeplot(data['sepal_length'], shade=True)
plt.title('kdeplot')
plt.xlabel('Sepal Length')
plt.ylabel('Density')
plt.show()
```



## 7.3 rugplot

rugplot is used to plot datapoints in an array as sticks on an axis.

```
In [20]: sns.rugplot(data['sepal_length'])  
plt.title('rugplot')  
plt.xlabel('Sepal Length')  
plt.show()
```

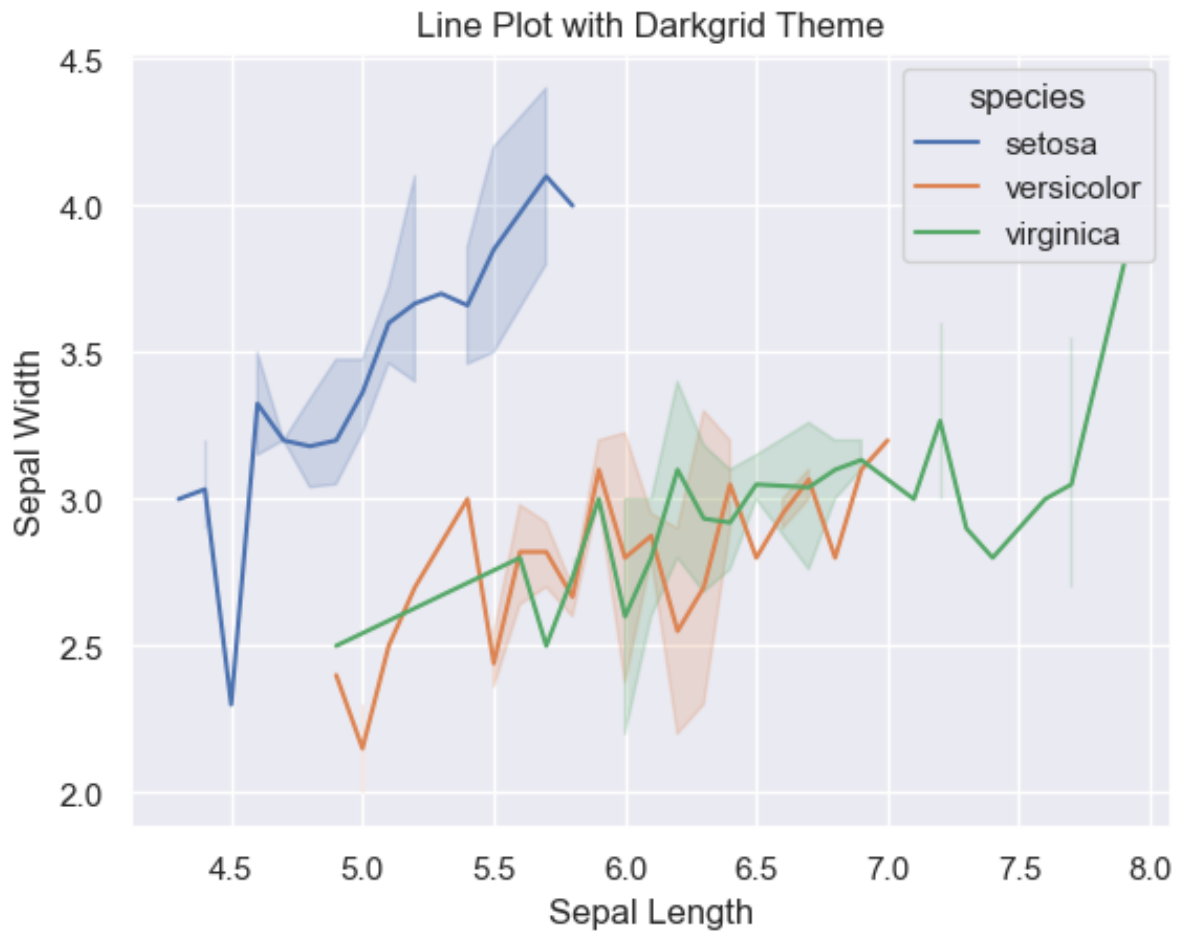


## 8. Customizing Plots

### 8.1 Themes and Styles

Seaborn allows you to set different themes and styles for your plots.

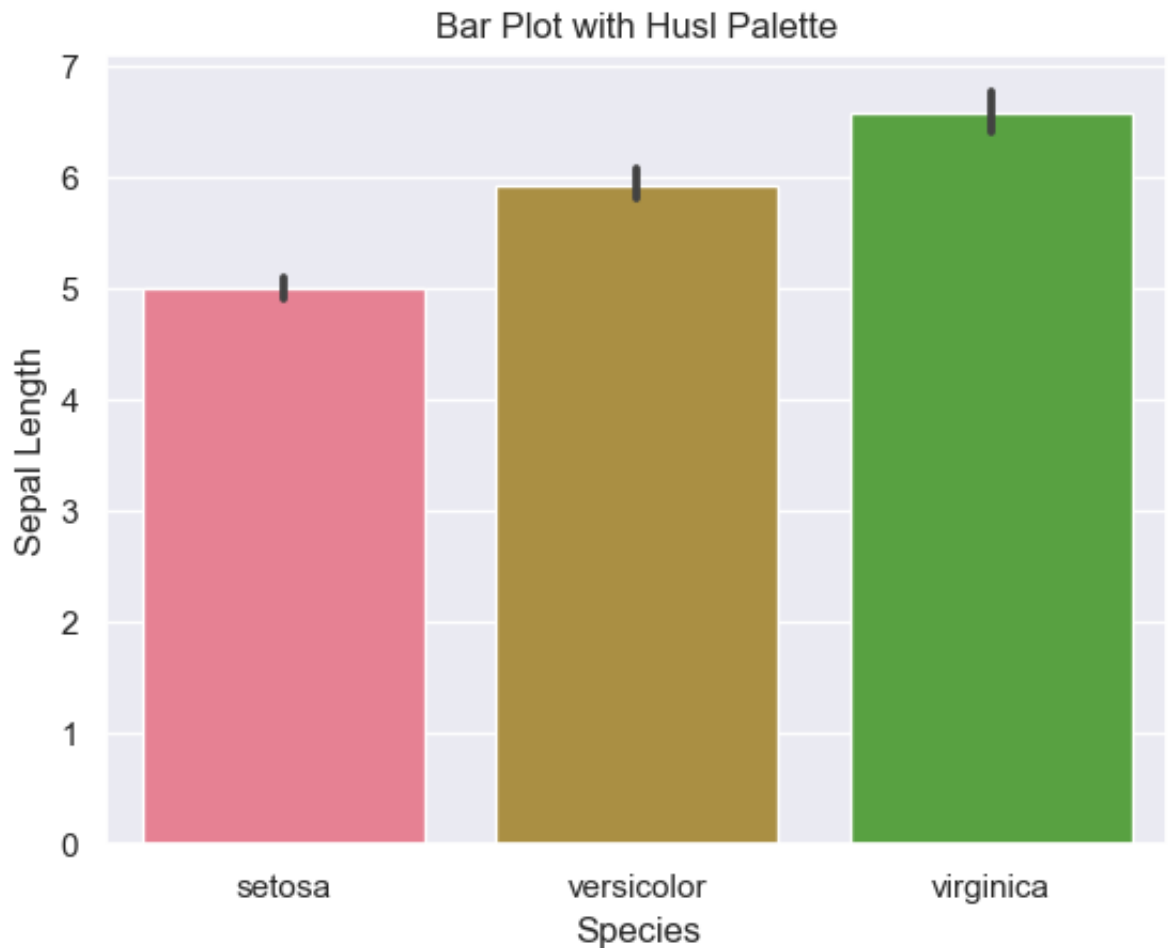
```
In [21]: sns.set_theme(style='darkgrid')
sns.lineplot(x='sepal_length', y='sepal_width', hue='species', data=
plt.title('Line Plot with Darkgrid Theme')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.show()
```



## 8.2 Color Palettes

Seaborn provides various color palettes to enhance the visual appeal of your plots.

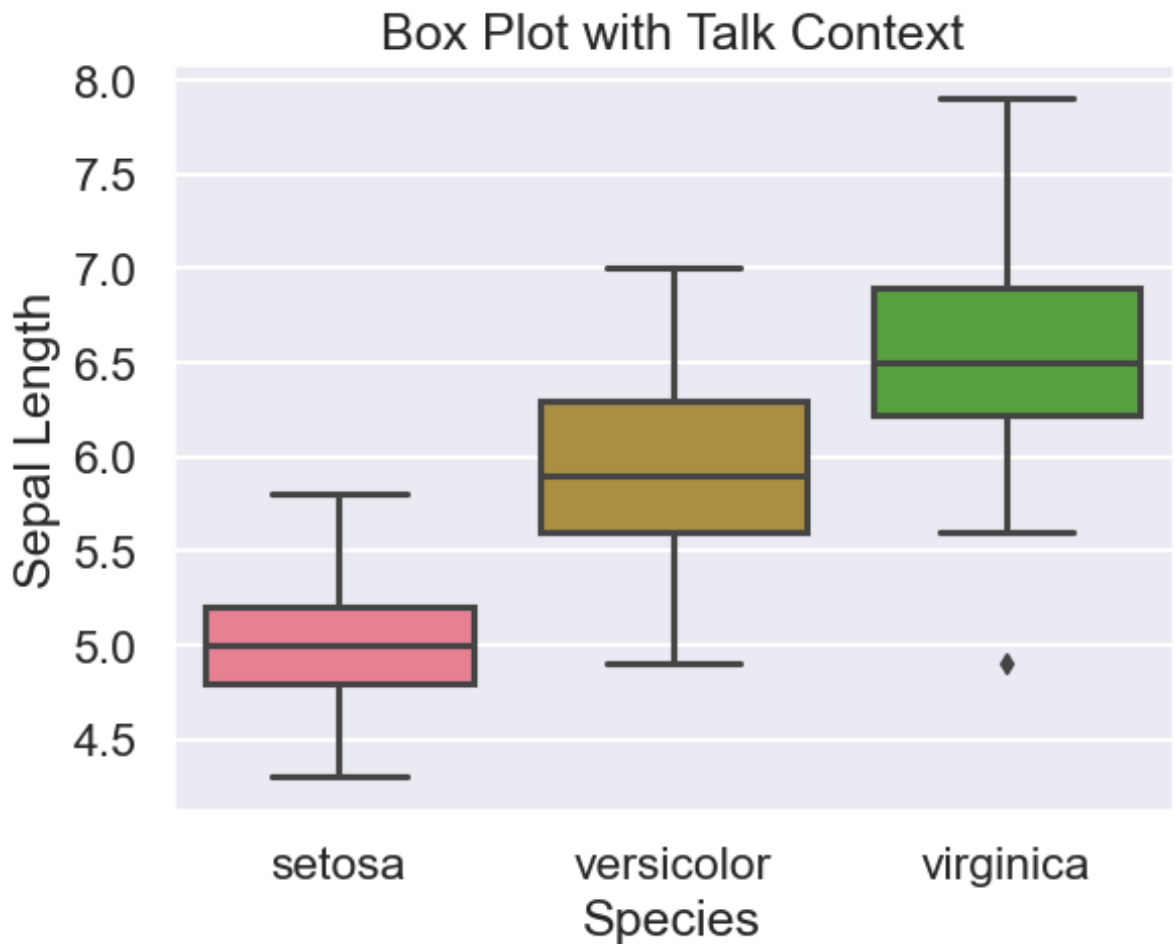
```
In [22]: sns.set_palette('husl')
sns.barplot(x='species', y='sepal_length', data=data)
plt.title('Bar Plot with Husl Palette')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```



### 8.3 Contexts

Seaborn contexts allow you to customize the scaling of plot elements to improve readability.

```
In [23]: sns.set_context('talk')
sns.boxplot(x='species', y='sepal_length', data=data)
plt.title('Box Plot with Talk Context')
plt.xlabel('Species')
plt.ylabel('Sepal Length')
plt.show()
```



## 9. Working with Inbuilt Datasets

Seaborn provides several built-in datasets for practice and demonstration purposes.

### 9.1 Tips Dataset

The Tips dataset contains information about tips received based on various factors.

```
In [24]: tips = sns.load_dataset('tips')
print(tips.head())
```

	total_bill	tip	sex	smoker	day	time	size
0	16.99	1.01	Female	No	Sun	Dinner	2
1	10.34	1.66	Male	No	Sun	Dinner	3
2	21.01	3.50	Male	No	Sun	Dinner	3
3	23.68	3.31	Male	No	Sun	Dinner	2
4	24.59	3.61	Female	No	Sun	Dinner	4

## 9.2 Flights Dataset

The Flights dataset contains information about the number of passengers on flights over time.

```
In [25]: flights = sns.load_dataset('flights')
print(flights.head())
```

	year	month	passengers
0	1949	Jan	112
1	1949	Feb	118
2	1949	Mar	132
3	1949	Apr	129
4	1949	May	121

## 9.3 Iris Dataset

The Iris dataset contains information about the measurements of iris flowers.

```
In [26]: iris = sns.load_dataset('iris')
print(iris.head())
```

	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

# 10. Advanced Customizations

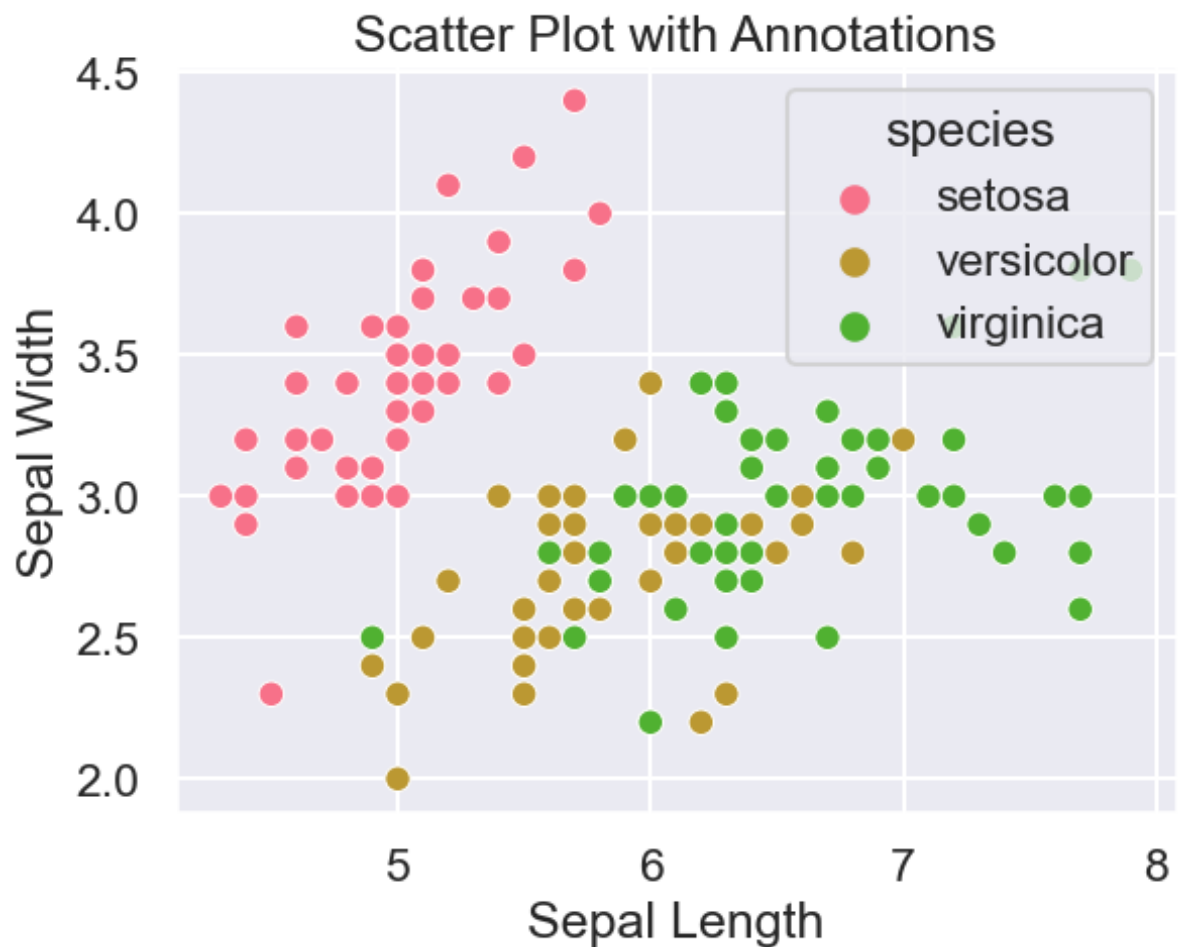
Learn how to customize your plots to make them more informative and visually appealing.



## 10.1 Adding Annotations

You can add annotations to your plots to highlight important data points.

```
In [29]: sns.scatterplot(x='sepal_length', y='sepal_width', hue='species', data=iris,
plt.title('Scatter Plot with Annotations')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
# for i in range(len(iris)):
#     plt.text(iris['sepal_length'][i], iris['sepal_width'][i], iris['species'][i])
plt.show()
```



## 10.2 Custom Legends

Customize the legends of your plots to improve clarity.

```
In [28]: sns.lineplot(x='sepal_length', y='sepal_width', hue='species', data=
plt.title('Line Plot with Custom Legends')
plt.xlabel('Sepal Length')
plt.ylabel('Sepal Width')
plt.legend(title='Species', loc='upper left', labels=['Setosa', 'Ve
plt.show()
```

