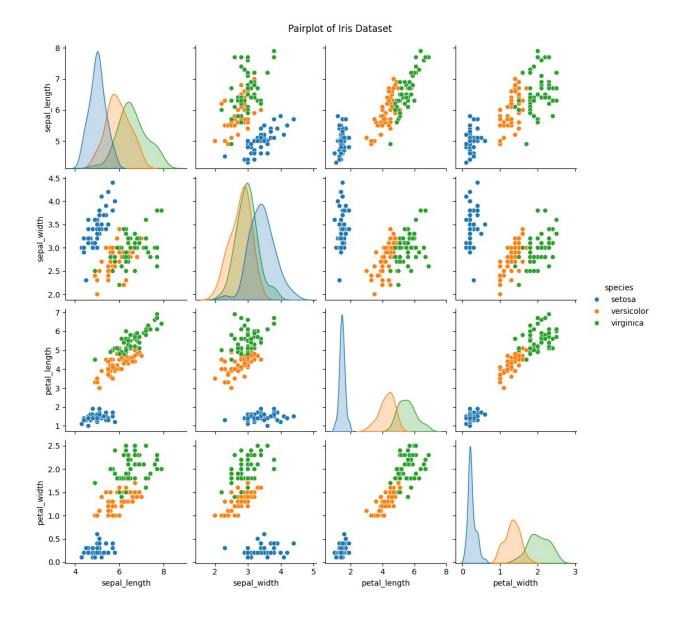
General Statistics Plot (Matplotlib or Seaborn):-

```
import seaborn as sns
import matplotlib.pyplot as plt
import pandas as pd
# Load Iris dataset
iris = sns.load dataset('iris')
# Using pandas describe to summarize statistics
summary stats = iris.describe()
print(summary_stats)
# Using Seaborn pairplot to visualize relationships
sns.pairplot(iris, hue='species')
plt.suptitle("Pairplot of Iris Dataset", y=1.02)
plt.show()
       sepal_length
                     sepal_width
                                  petal_length
                                                 petal width
         150.000000
                      150.000000
                                     150.000000
                                                  150.000000
count
           5.843333
                        3.057333
                                       3.758000
                                                    1.199333
mean
std
           0.828066
                        0.435866
                                       1.765298
                                                    0.762238
           4.300000
                        2.000000
                                       1.000000
                                                    0.100000
min
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
           7.900000
                        4.400000
                                       6.900000
                                                    2.500000
max
```

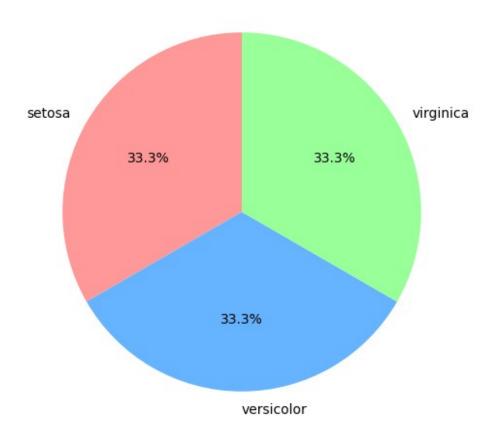


*** Pie Plot for Species Frequency:-***

```
# Create a pie chart of species frequency
species_counts = iris['species'].value_counts()

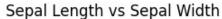
# Plot Pie Chart
plt.figure(figsize=(6, 6))
plt.pie(species_counts, labels=species_counts.index, autopct='%1.1f%%', startangle=90, colors=['#ff9999','#66b3ff','#99ff99'])
plt.title('Frequency of Iris Species')
plt.show()
```

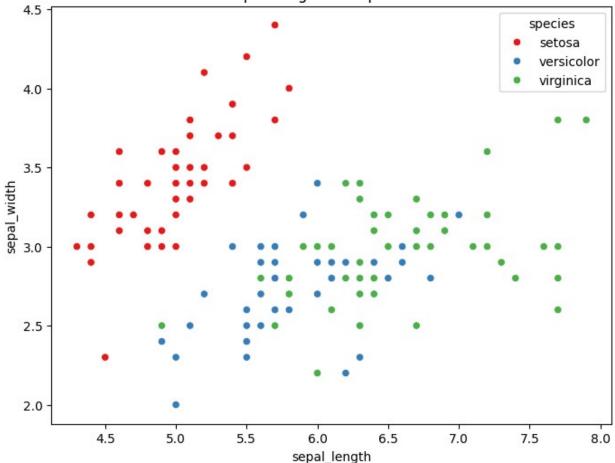
Frequency of Iris Species



Scatter Plot: Relationship Between Sepal Length and Width:-

```
# Scatter plot for sepal length vs sepal width
plt.figure(figsize=(8, 6))
sns.scatterplot(x='sepal_length', y='sepal_width', hue='species',
data=iris, palette='Set1')
plt.title('Sepal Length vs Sepal Width')
plt.show()
```



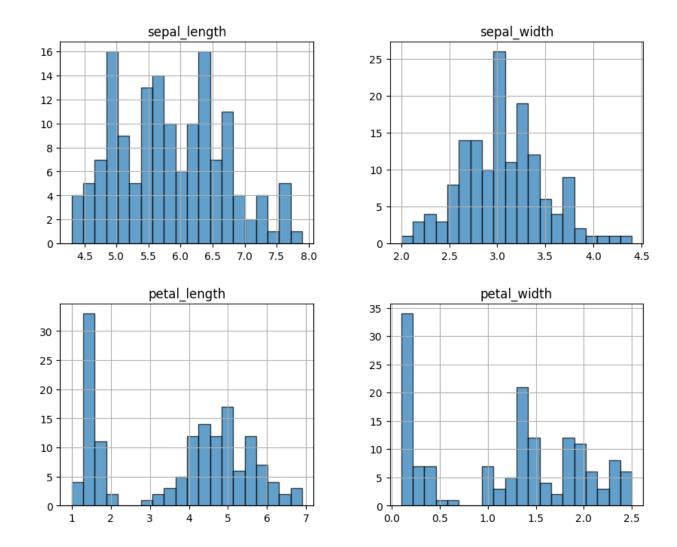


Distribution of Sepal and Petal Features:-

```
# Distribution of Sepal and Petal features
plt.figure(figsize=(10, 8))
iris[['sepal_length', 'sepal_width', 'petal_length',
    'petal_width']].hist(bins=20, edgecolor='black', alpha=0.7,
    figsize=(10, 8))
plt.suptitle('Distribution of Sepal and Petal Features', y=1.02)
plt.show()

    # Distribution of Sepal length',
    'petal_length',
    'petal_length',
    'petal_length',
    'petal_length',
    'petal_length',
    'petal_length',
    'petal_length',
    'sepal_width', 'petal_length',
    'petal_width', 'petal_length',
    'petal_width', 'petal_length',
    'sepal_width', 'petal_length',
    'petal_width', 'petal_length',
    'sepal_width', 'petal_length',
    'petal_width', 'petal_length',
    'petal_width', 'petal_length',
    'sepal_width', 'petal_length',
    'petal_width', 'petal_length',
    'sepal_width', 'petal_length',
    'sepal_width',
    'sepal_w
```

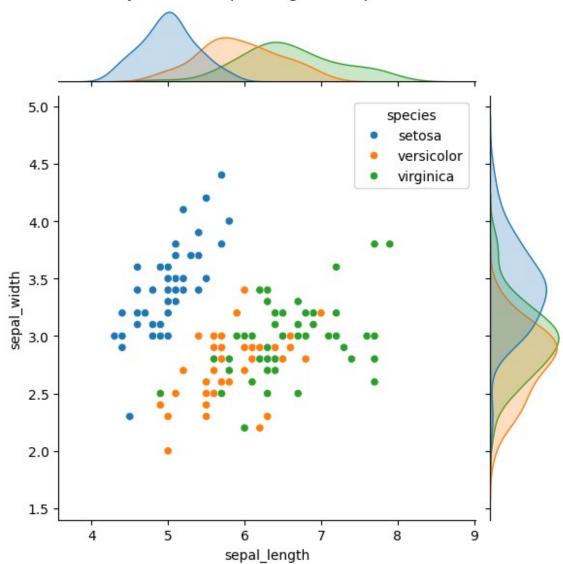
Distribution of Sepal and Petal Features



Jointplot of Sepal Length vs Sepal Width:-

```
# Jointplot for sepal length vs sepal width
sns.jointplot(x='sepal_length', y='sepal_width', data=iris,
kind='scatter', hue='species')
plt.suptitle("Joint Plot: Sepal Length vs Sepal Width", y=1.02)
plt.show()
```



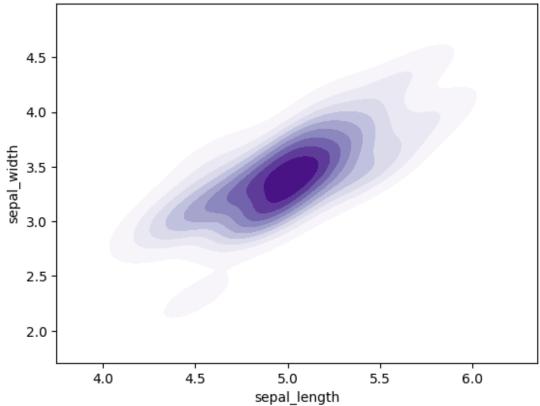


KDE Plot for Setosa Species (Sepal Length vs Sepal Width):-

```
# Filter dataset for Setosa species
setosa = iris[iris['species'] == 'setosa']

# KDE plot for Sepal Length vs Sepal Width
sns.kdeplot(x='sepal_length', y='sepal_width', data=setosa,
cmap='Purples', fill=True)
plt.title("KDE Plot: Setosa (Sepal Length vs Sepal Width)")
plt.show()
```





KDE Plot for Setosa Species (Petal Length vs Petal Width):-

```
# KDE plot for Petal Length vs Petal Width
sns.kdeplot(x='petal_length', y='petal_width', data=setosa,
cmap='Purples', fill=True)
plt.title("KDE Plot: Setosa (Petal Length vs Petal Width)")
plt.show()
```

KDE Plot: Setosa (Petal Length vs Petal Width)

