

iris-dataset

September 12, 2023

Import Libraries

```
[1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')
%matplotlib inline
```

```
[2]: iris =pd.read_csv(r'D:\DatSets\iris.csv')
iris
```

```
[2]:      Id  SepalLengthCm  SepalWidthCm  PetalLengthCm  PetalWidthCm  \
0      1           5.1           3.5           1.4           0.2
1      2           4.9           3.0           1.4           0.2
2      3           4.7           3.2           1.3           0.2
3      4           4.6           3.1           1.5           0.2
4      5           5.0           3.6           1.4           0.2
..  ...           ...           ...           ...           ...
145  146           6.7           3.0           5.2           2.3
146  147           6.3           2.5           5.0           1.9
147  148           6.5           3.0           5.2           2.0
148  149           6.2           3.4           5.4           2.3
149  150           5.9           3.0           5.1           1.8
```

```
      Species
0      Iris-setosa
1      Iris-setosa
2      Iris-setosa
3      Iris-setosa
4      Iris-setosa
..  ...
145  Iris-virginica
146  Iris-virginica
147  Iris-virginica
148  Iris-virginica
149  Iris-virginica
```

[150 rows x 6 columns]

```
[3]: #We dont require the ID Columns
iris.drop('Id',axis=1, inplace =True)
```

```
[4]: iris.head()
```

```
[4]:   SepalLengthCm  SepalWidthCm  PetalLengthCm  PetalWidthCm   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
```

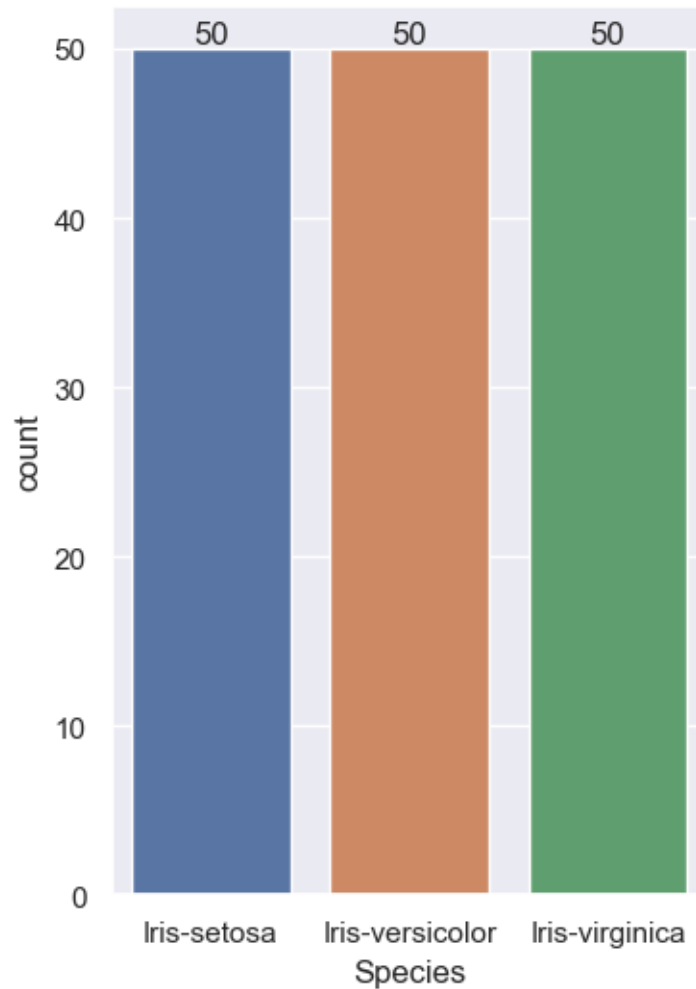
```
[5]: iris.info()
```

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

```
[6]: iris['Species'].value_counts()
```

```
[6]: Iris-setosa      50
     Iris-versicolor  50
     Iris-virginica   50
     Name: Species, dtype: int64
```

```
[14]: aa =plt.subplots(figsize=(4,6))
      aa=sns.countplot(x='Species', data =iris)
      for bars in aa.containers :
          aa.bar_label(bars)
```



We can see that there are 50 samples each of all the Iris Species in the data set.

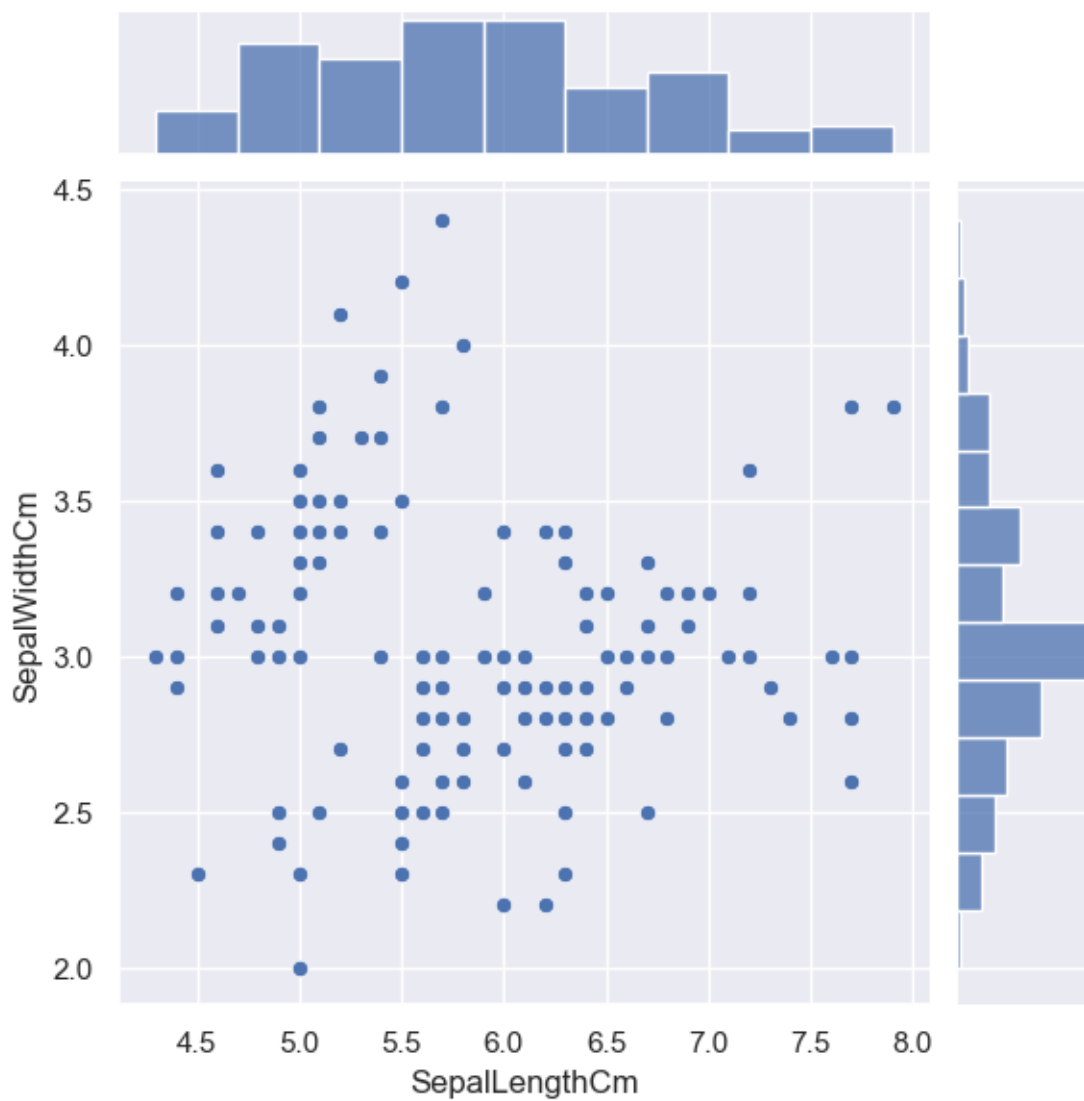
0.0.1 JointPlot

```
[15]: iris.columns
```

```
[15]: Index(['SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm',  
         'Species'],  
         dtype='object')
```

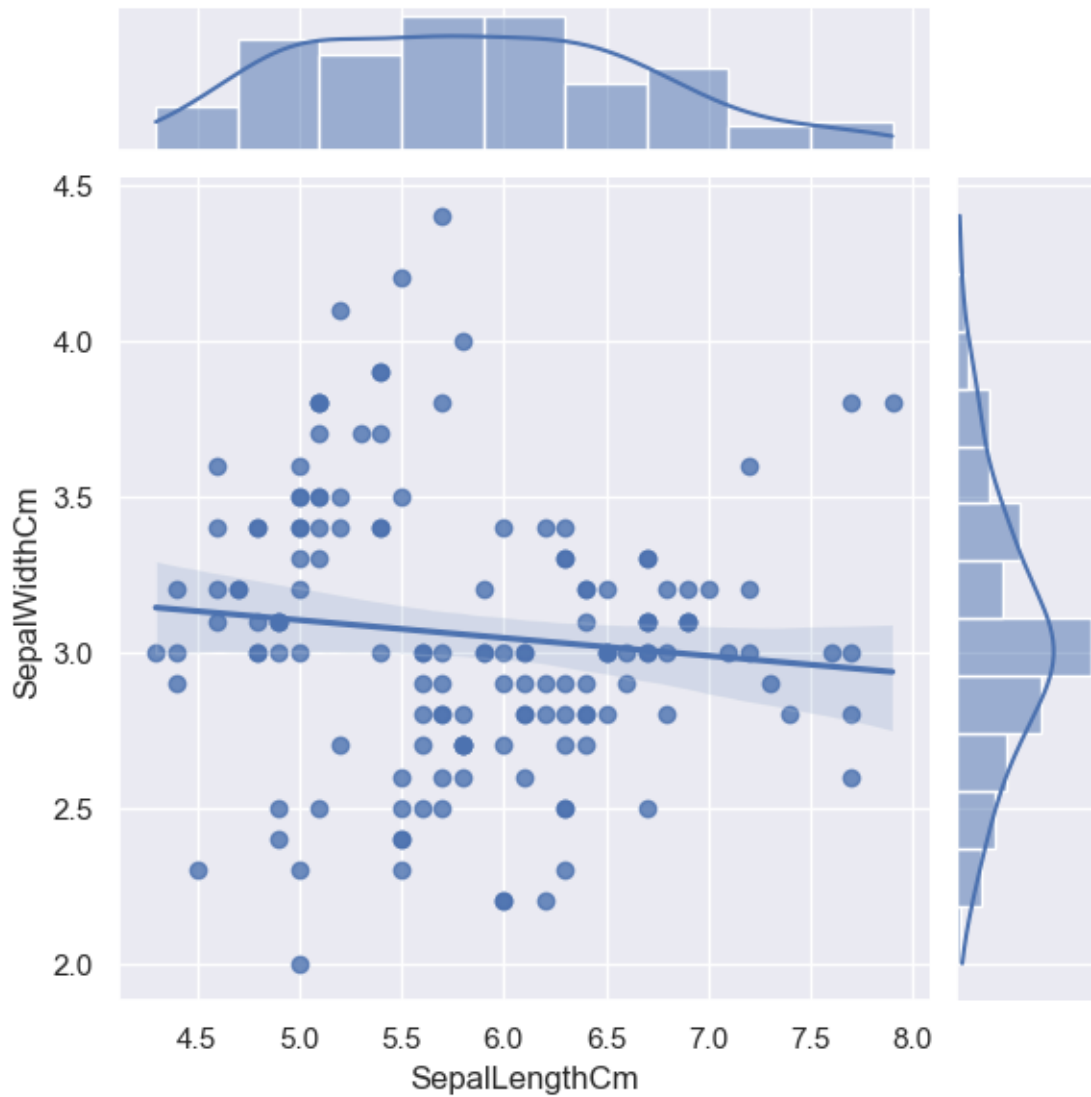
```
[20]: sns.jointplot(x='SepalLengthCm',y='SepalWidthCm',data=iris)
```

```
[20]: <seaborn.axisgrid.JointGrid at 0x29e12525c00>
```



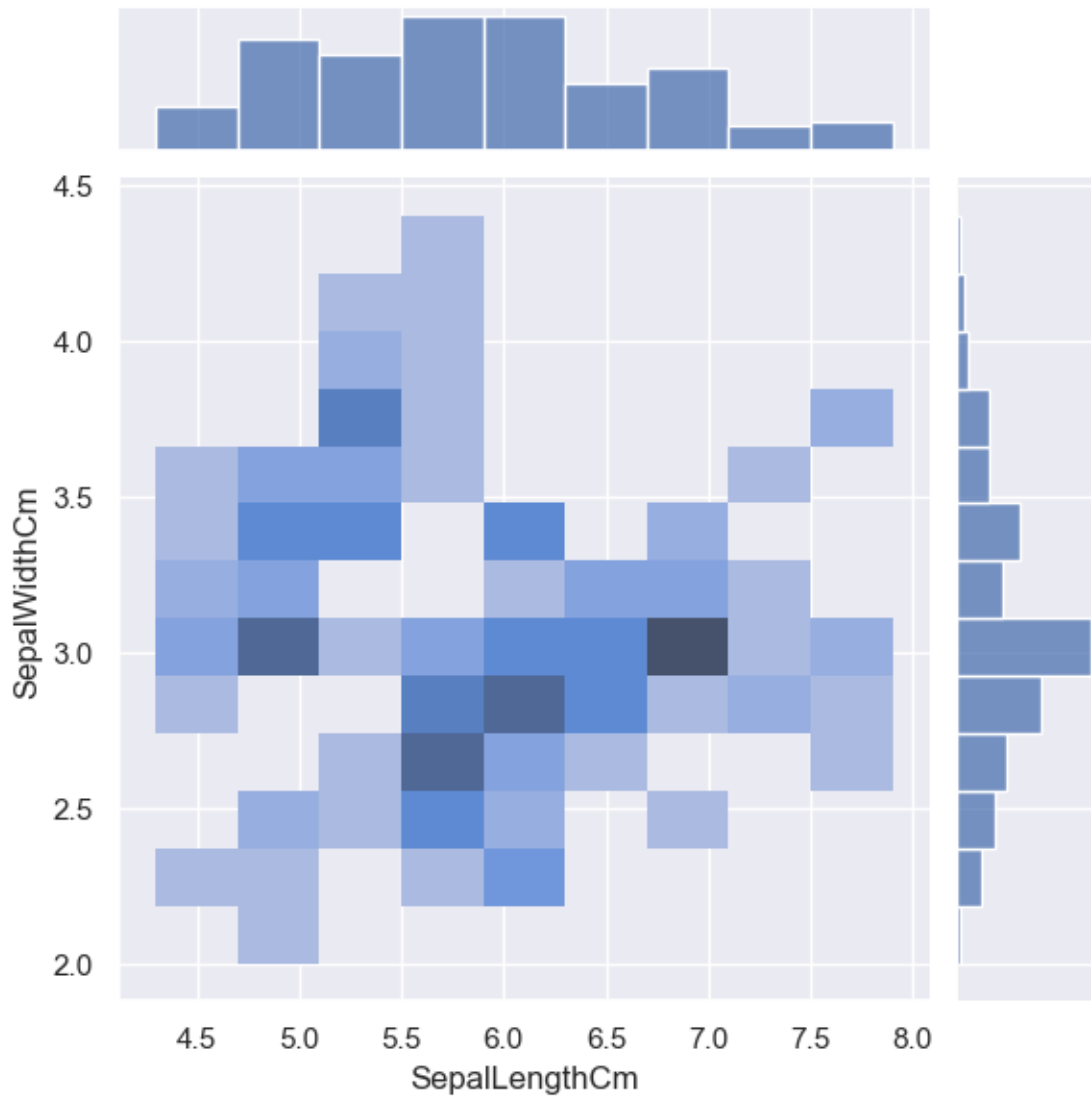
```
[21]: sns.jointplot(x='SepalLengthCm',y='SepalWidthCm',data=iris, kind='reg')  
      #kind : { "scatter" / "kde" / "hist" / "hex" / "reg" / "resid" }
```

```
[21]: <seaborn.axisgrid.JointGrid at 0x29e1312de40>
```



```
[22]: sns.jointplot(x='SepalLengthCm',y='SepalWidthCm',data=iris, kind='hist')
```

```
[22]: <seaborn.axisgrid.JointGrid at 0x29e13b357e0>
```



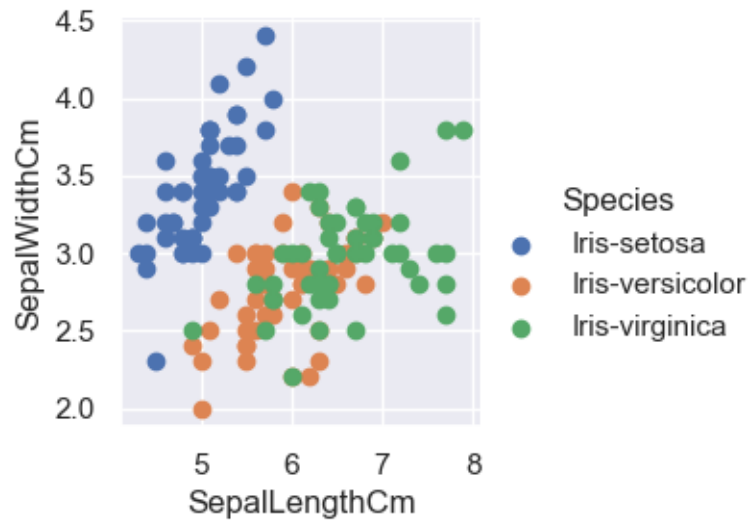
0.0.2 Facegrid Plot

```
[25]: iris.columns
```

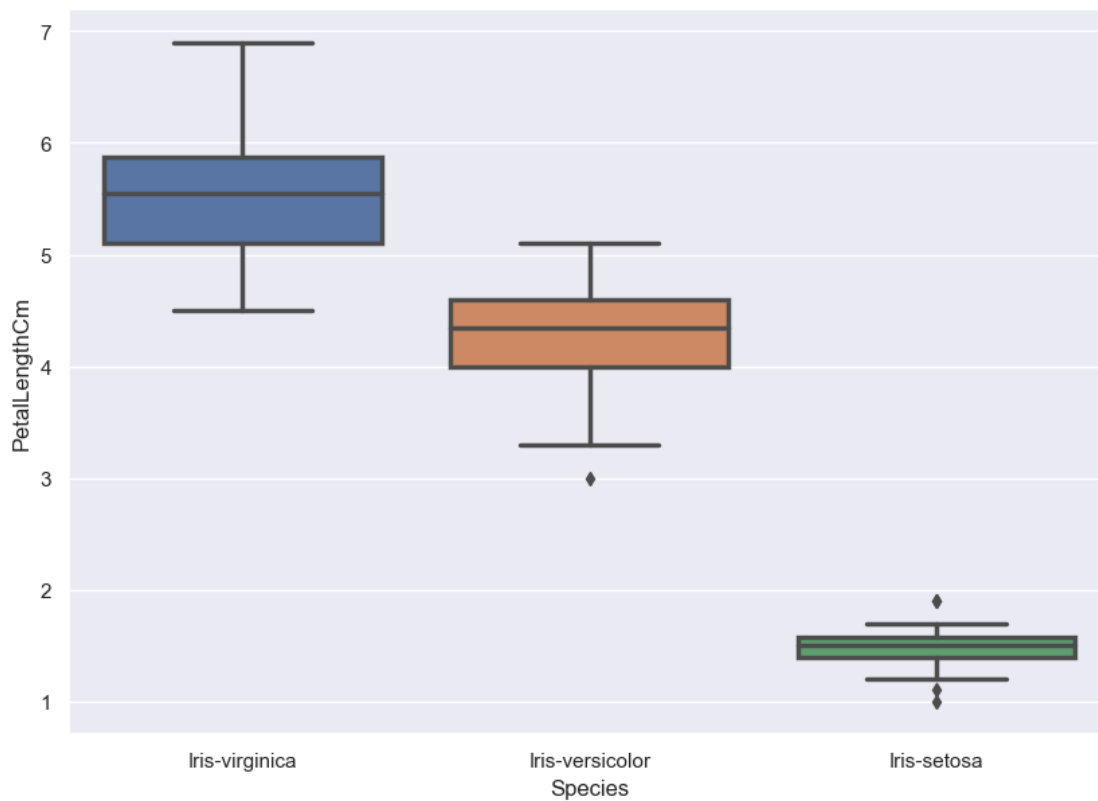
```
[25]: Index(['SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm',
            'Species'],
            dtype='object')
```

```
[26]: sns.FacetGrid(iris, hue= 'Species').map(plt.
        ↳scatter, 'SepalLengthCm', 'SepalWidthCm').add_legend()
```

```
[26]: <seaborn.axisgrid.FacetGrid at 0x29e1305bd00>
```

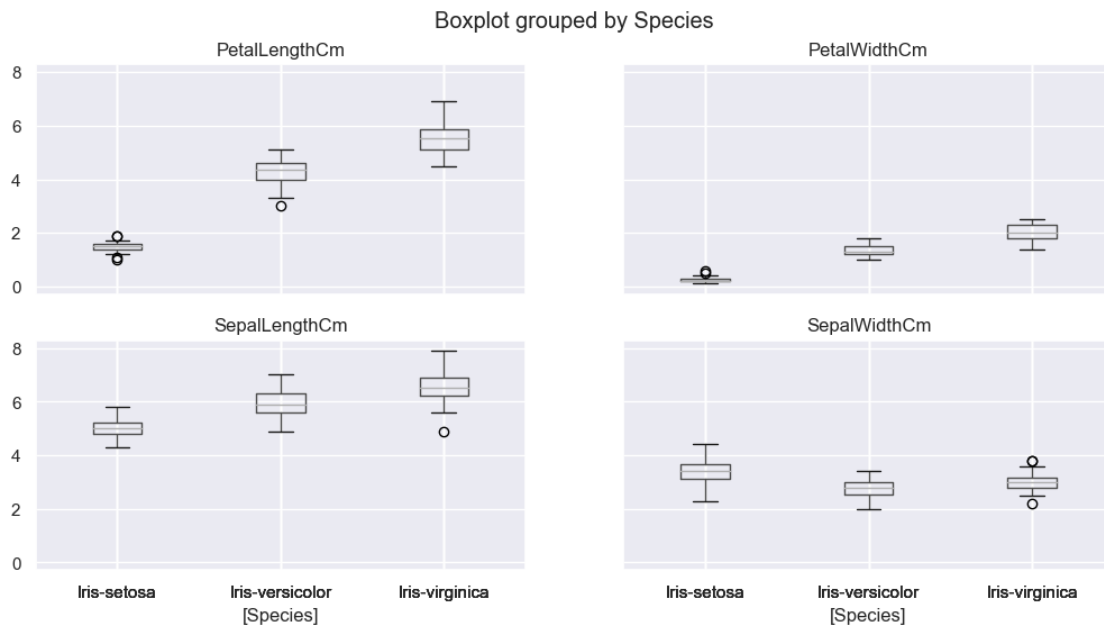


```
[27]: fig=plt.gcf()
fig.set_size_inches(10,7)
fig=sns.
    ↳boxplot(x='Species',y='PetalLengthCm',data=iris,order=['Iris-virginica','Iris-versicolor','Iris-setosa'],
    ↳5,orient='v',dodge=False)
```



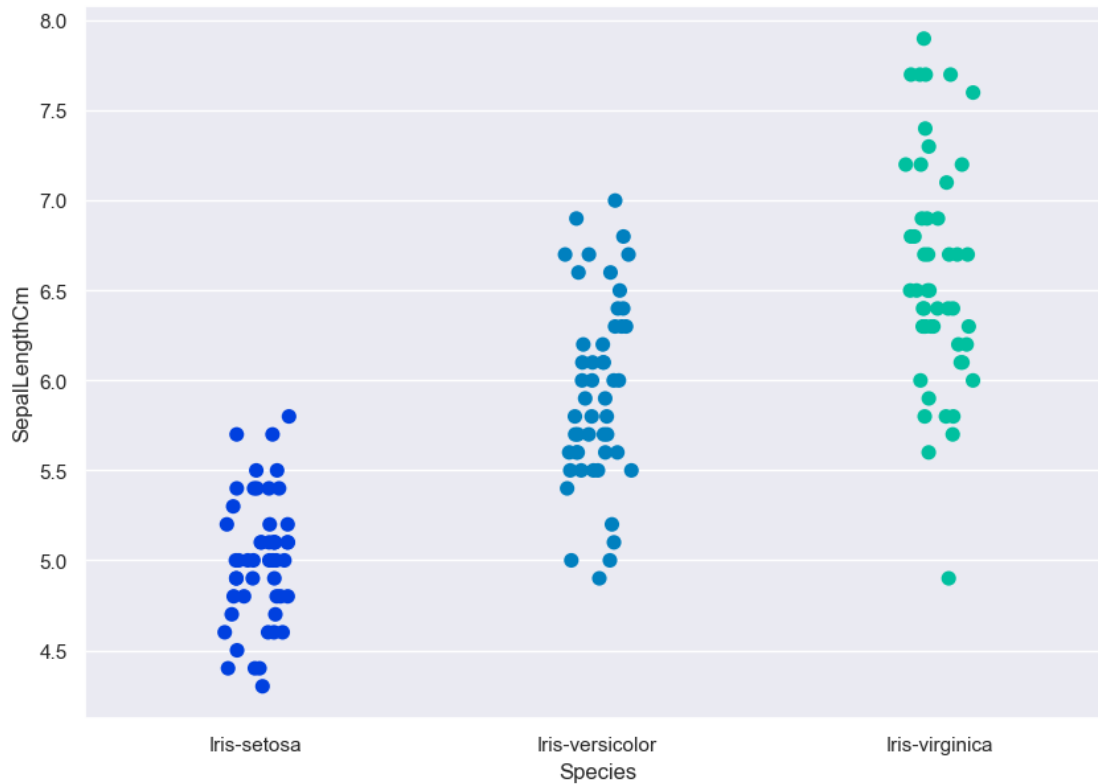
```
[28]: iris.boxplot(by="Species", figsize=(12, 6))
```

```
[28]: array([[<Axes: title={'center': 'PetalLengthCm'}, xlabel='[Species] '>,
  <Axes: title={'center': 'PetalWidthCm'}, xlabel='[Species] '>],
  [<Axes: title={'center': 'SepalLengthCm'}, xlabel='[Species] '>,
  <Axes: title={'center': 'SepalWidthCm'}, xlabel='[Species] '>]],
  dtype=object)
```



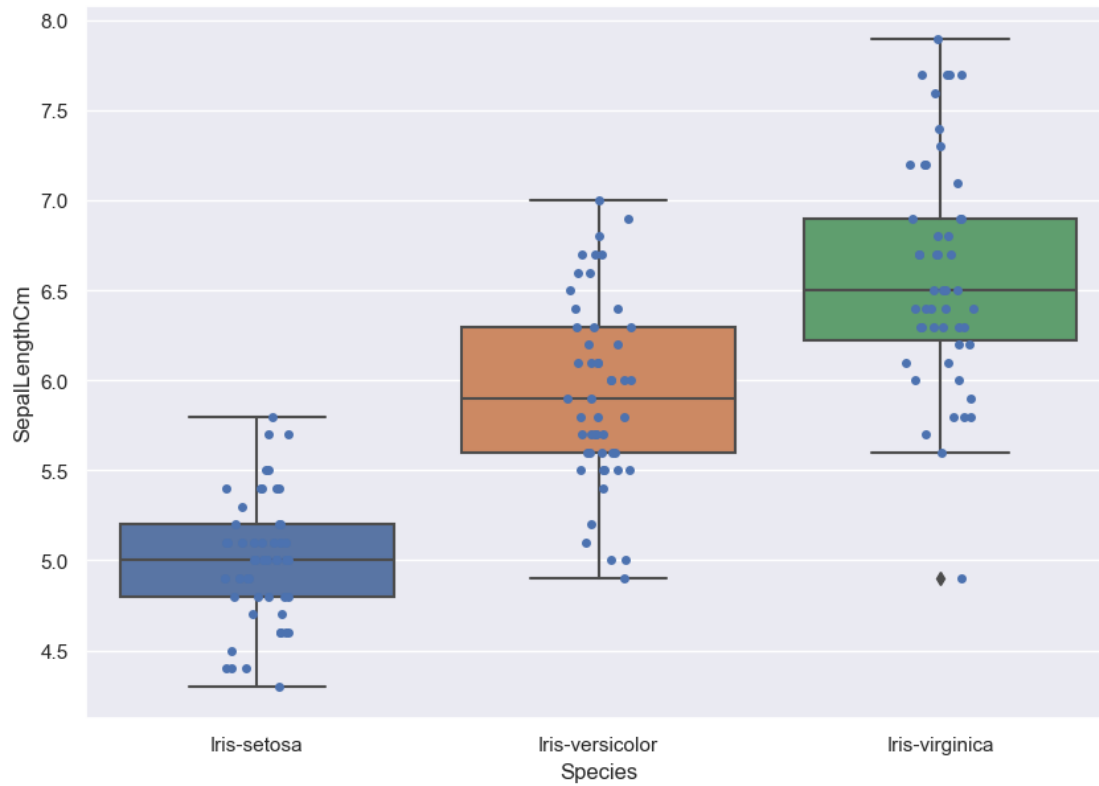
0.1 Strip Plot

```
[29]: fig=plt.gcf()
fig.set_size_inches(10,7)
fig=sns.
↪stripplot(x='Species',y='SepalLengthCm',data=iris,jitter=True,edgecolor='gray',size=8,palet
```

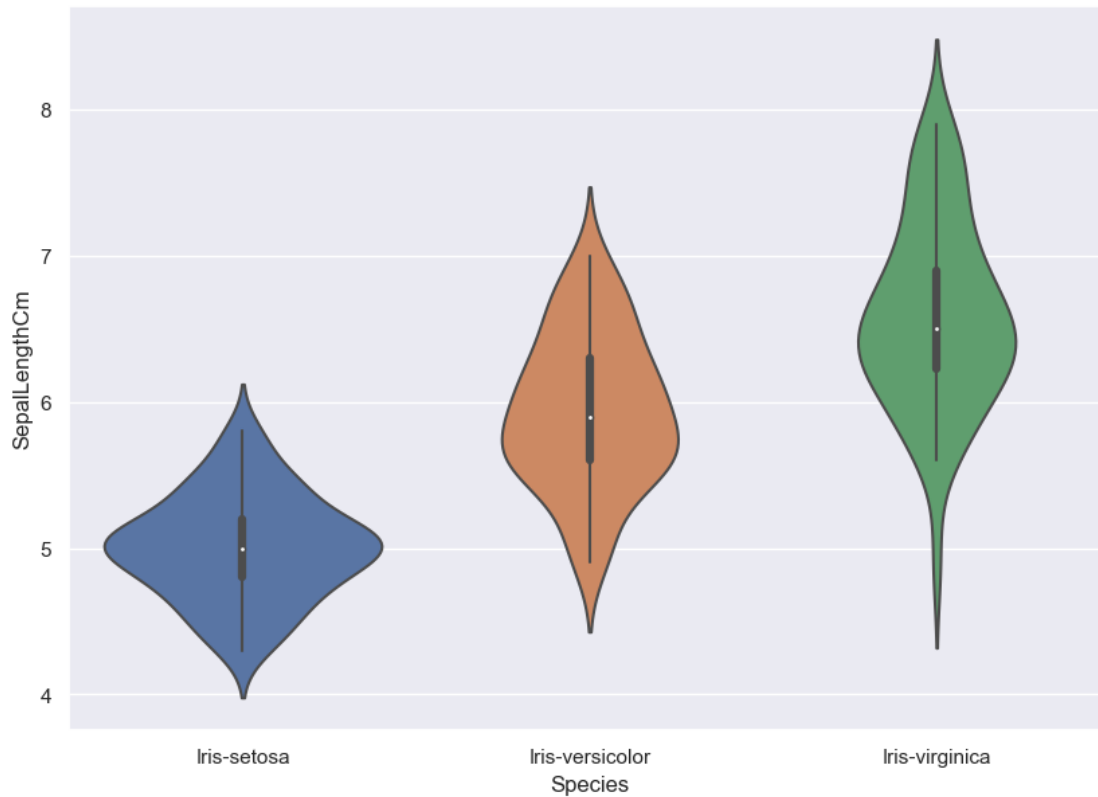
0.1.1 Combining Box and Strip Plots

```
[30]: fig=plt.gcf()
fig.set_size_inches(10,7)
fig=sns.boxplot(x='Species',y='SepalLengthCm',data=iris)
fig=sns.
↪stripplot(x='Species',y='SepalLengthCm',data=iris,jitter=True,edgecolor='gray')
```



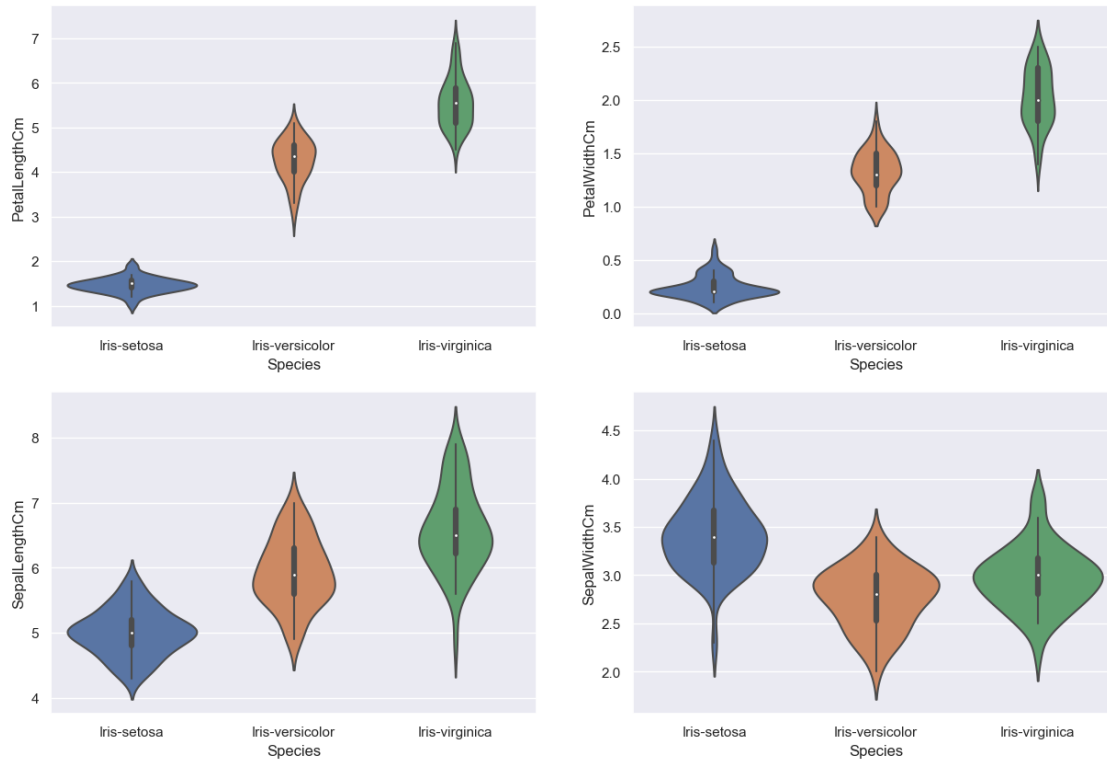
0.2 Violin Plot

```
[31]: fig=plt.gcf()
fig.set_size_inches(10,7)
fig=sns.violinplot(x='Species',y='SepalLengthCm',data=iris)
```



```
[32]: plt.figure(figsize=(15,10))
plt.subplot(2,2,1)
sns.violinplot(x='Species',y='PetalLengthCm',data=iris)
plt.subplot(2,2,2)
sns.violinplot(x='Species',y='PetalWidthCm',data=iris)
plt.subplot(2,2,3)
sns.violinplot(x='Species',y='SepalLengthCm',data=iris)
plt.subplot(2,2,4)
sns.violinplot(x='Species',y='SepalWidthCm',data=iris)
```

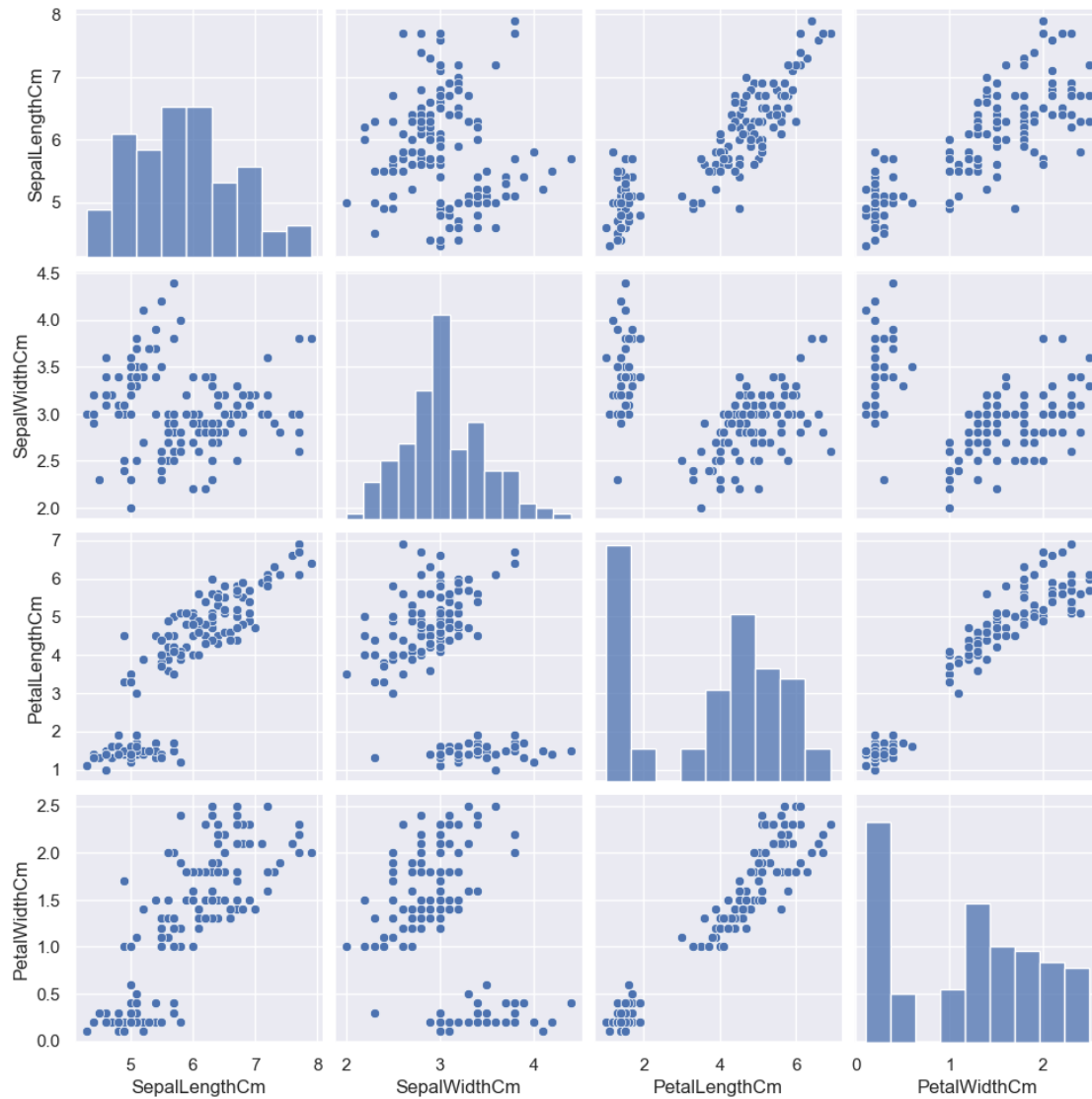
```
[32]: <Axes: xlabel='Species', ylabel='SepalWidthCm'>
```



0.3 Pair Plot

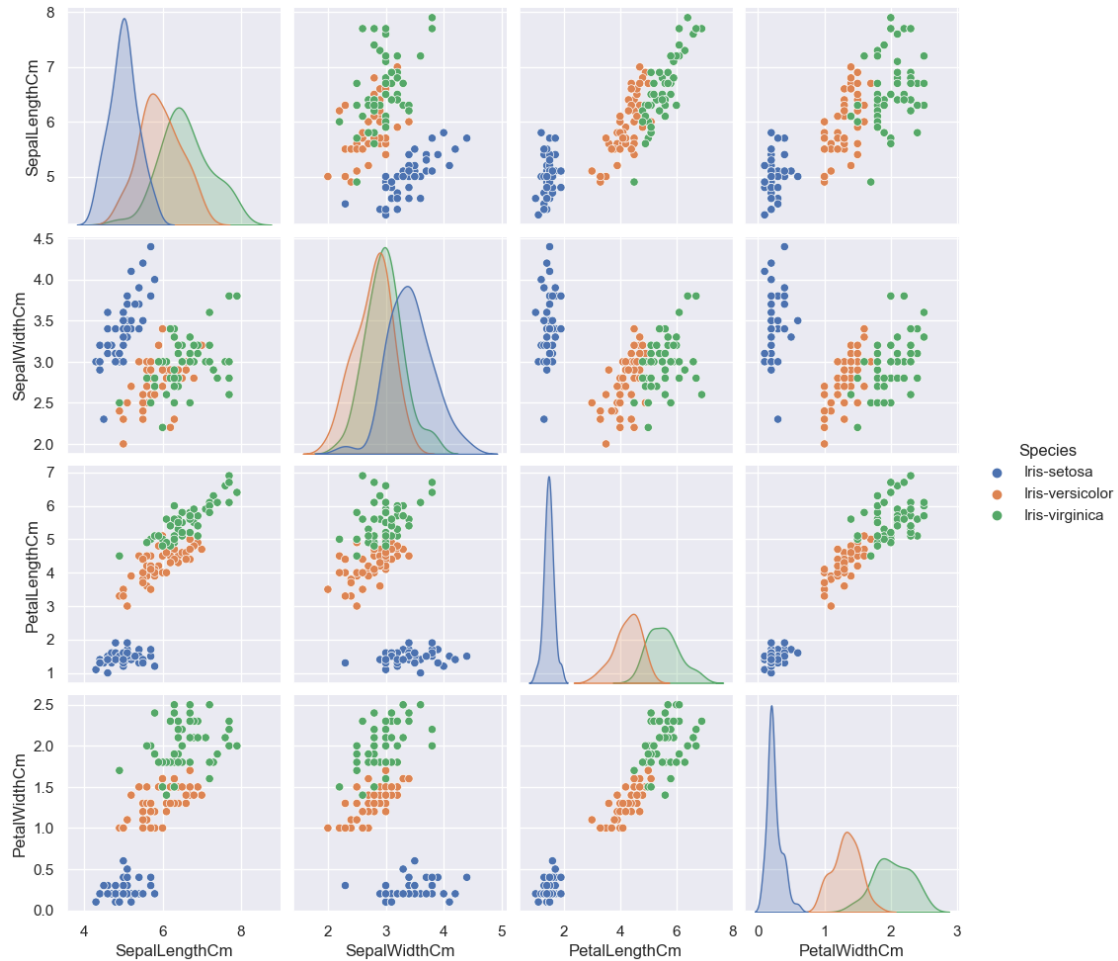
```
[33]: sns.pairplot(data=iris,kind='scatter')
```

```
[33]: <seaborn.axisgrid.PairGrid at 0x29e17104d90>
```



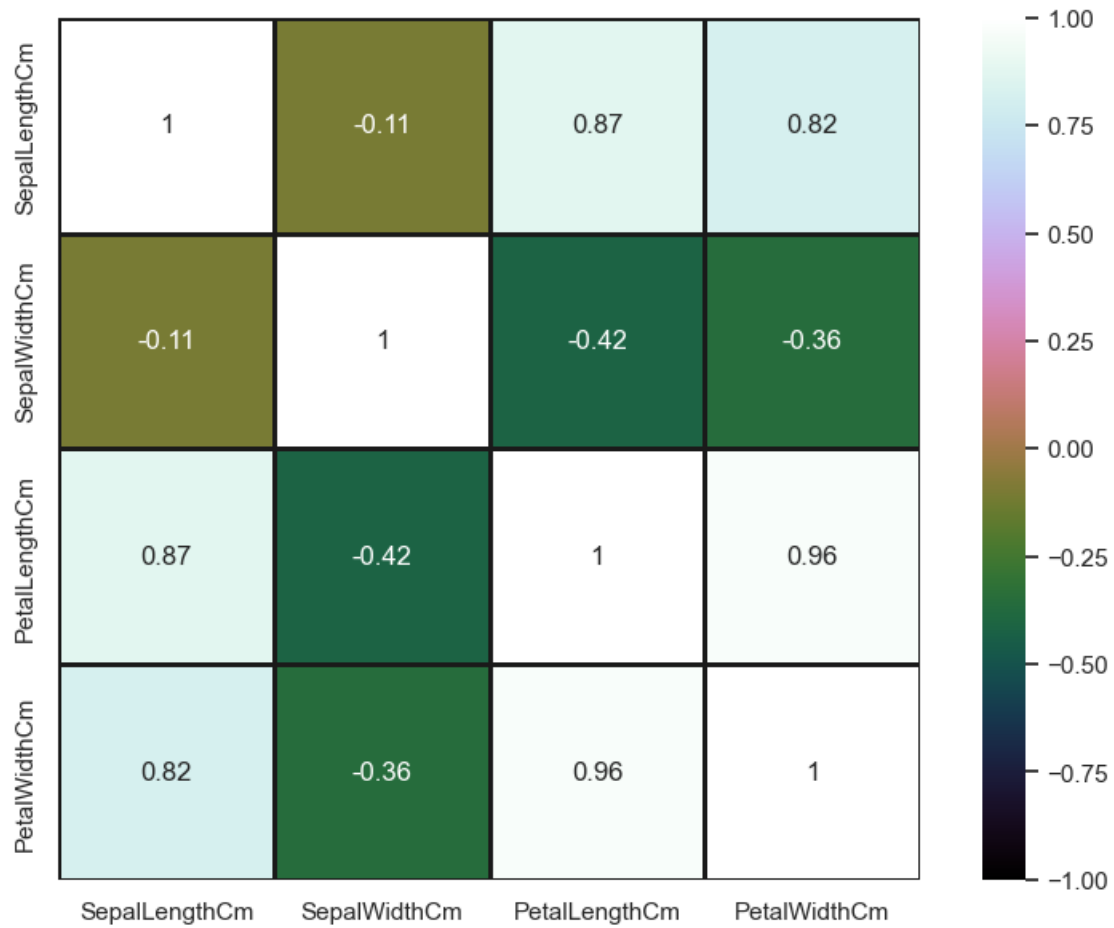
```
[34]: sns.pairplot(iris,hue='Species')
```

```
[34]: <seaborn.axisgrid.PairGrid at 0x29e17d68af0>
```



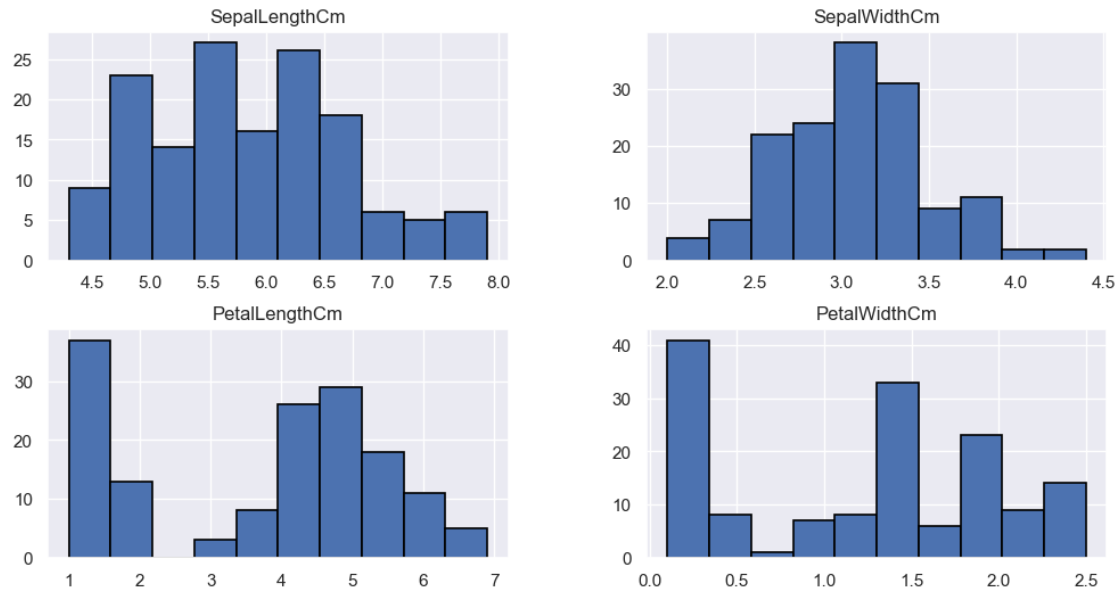
0.3.1 Heat map

```
[35]: fig=plt.gcf()
fig.set_size_inches(10,7)
fig=sns.heatmap(iris.
    ↪corr(),annot=True,cmap='cubehelix',linewidths=1,linecolor='k',square=True,mask=False,
    ↪vmin=-1, vmax=1,cbar_kws={"orientation": "vertical"},cbar=True)
```



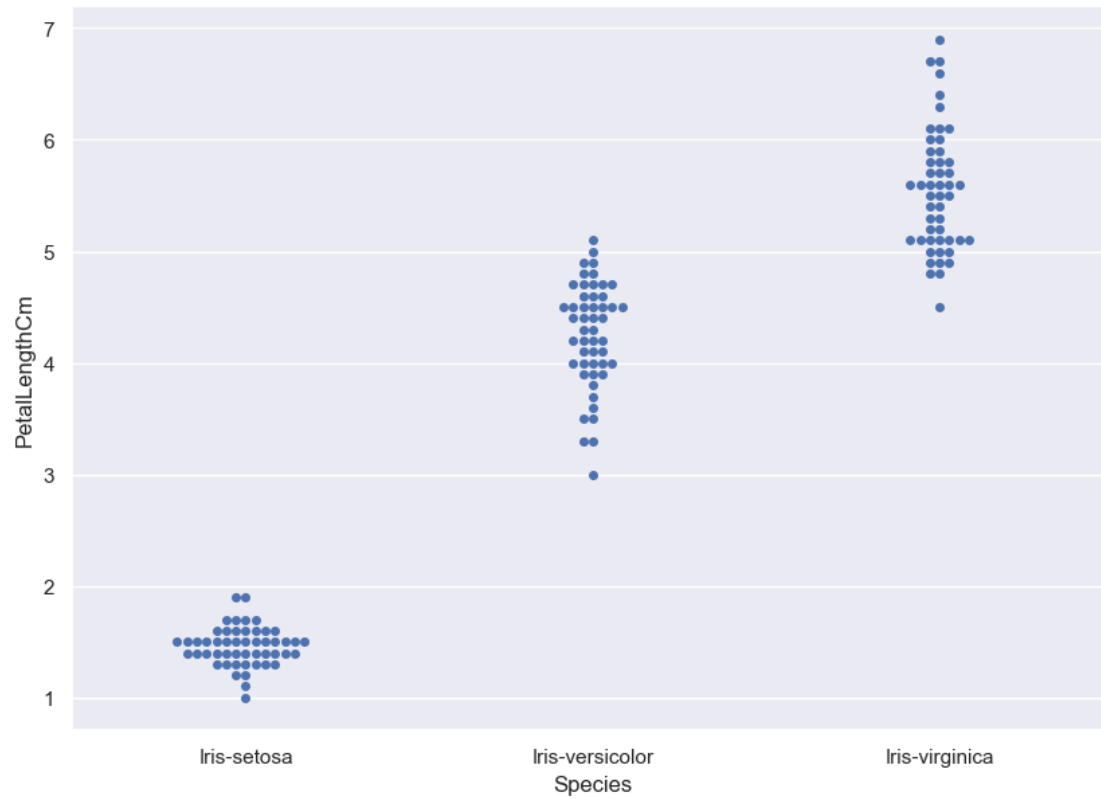
0.4 Distribution plot:

```
[36]: iris.hist(edgecolor='black', linewidth=1.2)
fig=plt.gcf()
fig.set_size_inches(12,6)
```

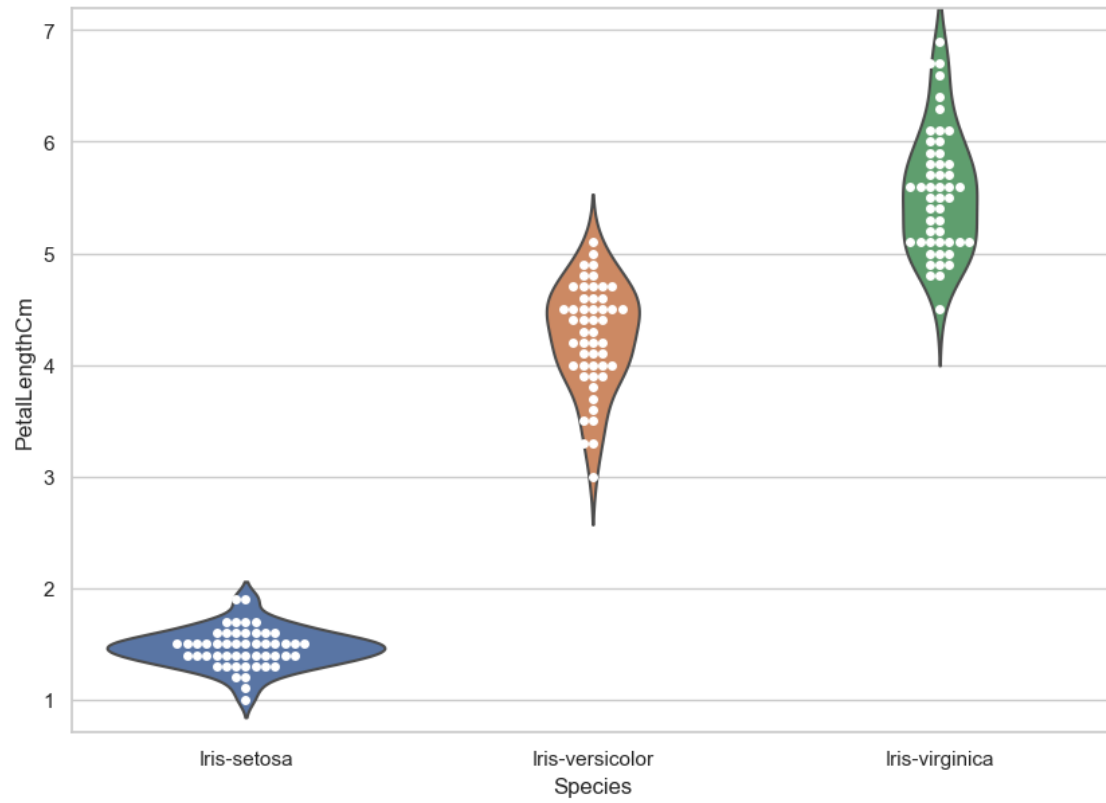


0.5 Swarm plot

```
[37]: sns.set(style="darkgrid")
fig=plt.gcf()
fig.set_size_inches(10,7)
fig = sns.swarmplot(x="Species", y="PetalLengthCm", data=iris)
```

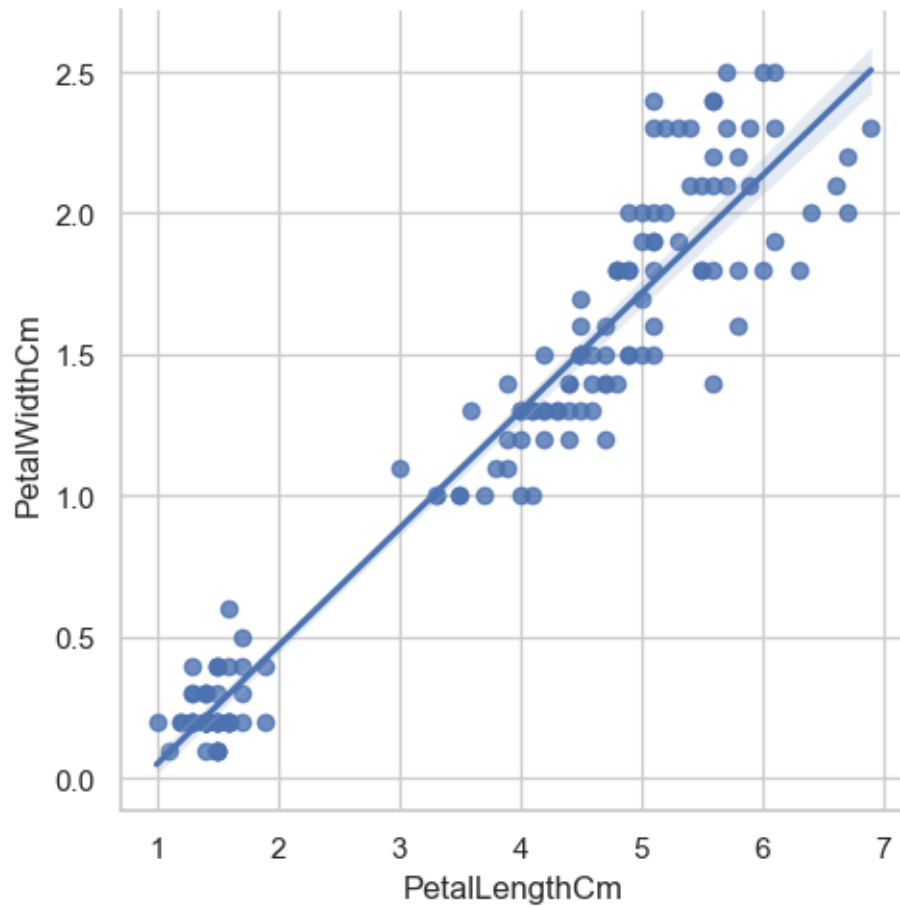



```
[38]: sns.set(style="whitegrid")
fig=plt.gcf()
fig.set_size_inches(10,7)
ax = sns.violinplot(x="Species", y="PetalLengthCm", data=iris, inner=None)
ax = sns.swarmplot(x="Species", y="PetalLengthCm", data=iris,color="white",
    edgecolor="black")
```



0.6 LM Plot

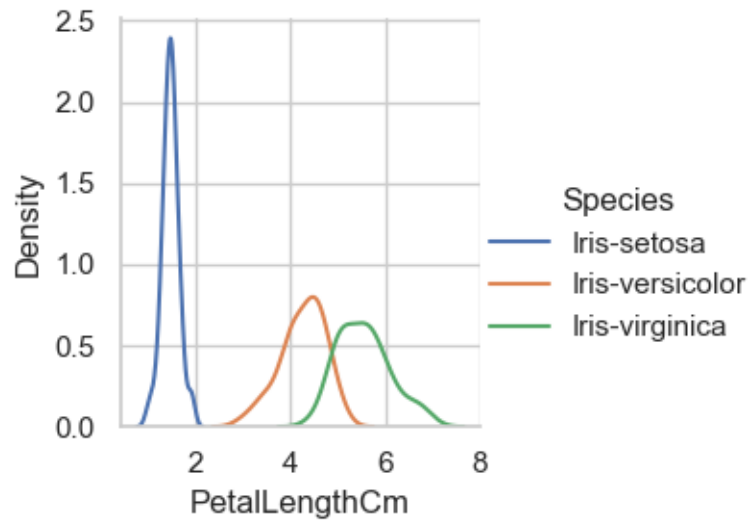
```
[39]: fig=sns.lmplot(x="PetalLengthCm", y="PetalWidthCm",data=iris)
```



0.7 Facegrid

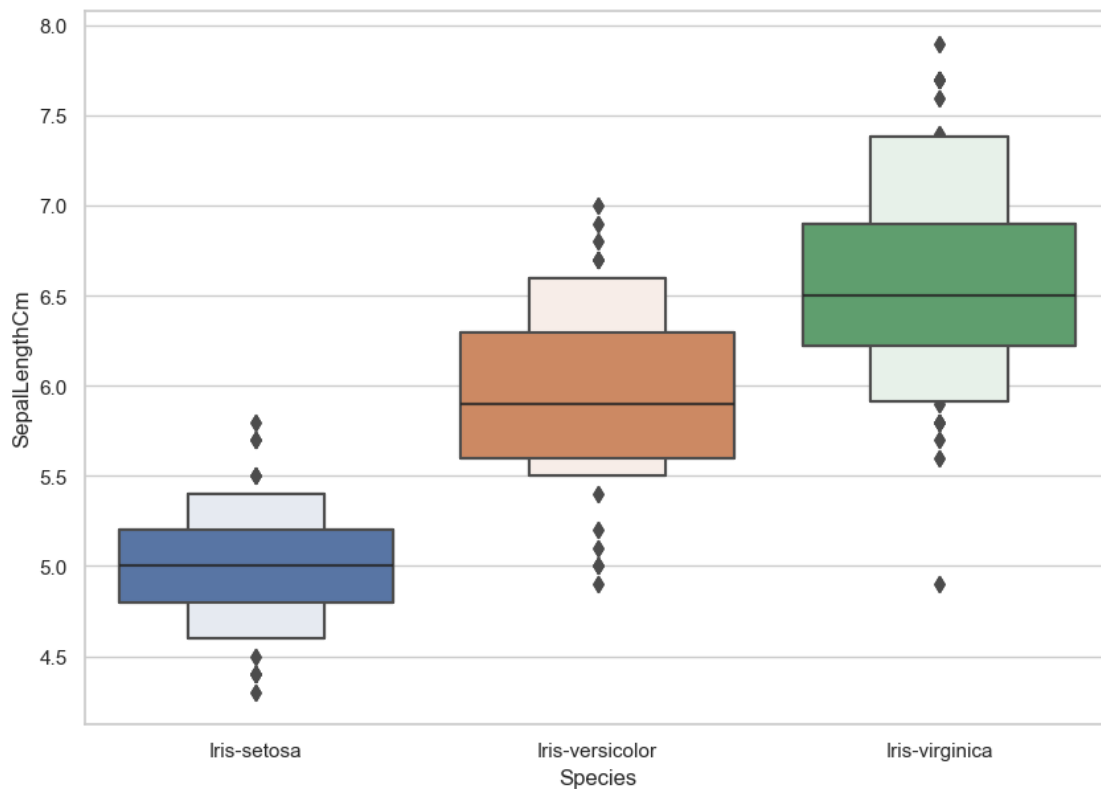
```
[41]: sns.FacetGrid(iris, hue="Species").map(sns.kdeplot, "PetalLengthCm").  
      ↪add_legend()  
      plt.ioff()
```

```
[41]: <contextlib.ExitStack at 0x29e17aaaa70>
```



0.8 Boxen Plot

```
[46]: fig=plt.gcf()
fig.set_size_inches(10,7)
fig=sns.boxenplot(x='Species',y='SepalLengthCm',data=iris)
plt.show()
```



0.9 Area Plot:

```
[53]: iris.plot.  
      ↪area(y=['SepalLengthCm','SepalWidthCm','PetalLengthCm','PetalWidthCm'],alpha=0.  
      ↪4,figsize=(12, 6));  
      plt.show()
```

