iris-flower-classification-oasis

March 4, 2024

1 IRIS FLOWER CLASSIFICATION (OASIS)

Problem Statement:

Iris flower has three species; setosa, versicolor, and virginica, which differs according to their measurements. Now assume that you have the measurements of the iris flowers according to their species, and here your task is to train a machine learning model that can learn from the measurements of the iris species and classify them.

[1]:
```
# Import Necessary Libraries
```

[2]:
```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import plotly.express as px

from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier

import warnings
warnings.filterwarnings('ignore')
```

[3]:
```
# Loading Dataset
```

[4]:
```
Data = pd.read_csv('Iris.csv')
```

[5]:
```
# Exploratory Data Analysis
```

[6]:
```
Data.head()
```

[6]:

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa

[7]:
```
Data.tail()
```

[7]:

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
145	146	6.7	3.0	5.2	2.3	Iris-virginica
146	147	6.3	2.5	5.0	1.9	Iris-virginica
147	148	6.5	3.0	5.2	2.0	Iris-virginica
148	149	6.2	3.4	5.4	2.3	Iris-virginica
149	150	5.9	3.0	5.1	1.8	Iris-virginica

[8]:
```
Data.columns
```

[8]:
```
Index(['Id', 'SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm',
       'Species'],
      dtype='object')
```

[9]:
```
Data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 6 columns):
 #   Column         Non-Null Count  Dtype
---  ------         --------------  -----
 0   Id             150 non-null    int64
 1   SepalLengthCm  150 non-null    float64
 2   SepalWidthCm   150 non-null    float64
 3   PetalLengthCm  150 non-null    float64
 4   PetalWidthCm   150 non-null    float64
 5   Species        150 non-null    object
dtypes: float64(4), int64(1), object(1)
memory usage: 7.2+ KB
```

[10]:
```
Data.describe()
```

[10]:

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
count	150.000000	150.000000	150.000000	150.000000	150.000000
mean	75.500000	5.843333	3.054000	3.758667	1.198667
std	43.445368	0.828066	0.433594	1.764420	0.763161
min	1.000000	4.300000	2.000000	1.000000	0.100000
25%	38.250000	5.100000	2.800000	1.600000	0.300000
50%	75.500000	5.800000	3.000000	4.350000	1.300000

```
75%     112.750000       6.400000       3.300000       5.100000       1.800000
max     150.000000       7.900000       4.400000       6.900000       2.500000
```

[11]:
```
Data.isnull().sum()
```

[11]:
```
Id               0
SepalLengthCm    0
SepalWidthCm     0
PetalLengthCm    0
PetalWidthCm     0
Species          0
dtype: int64
```

[12]:
```
Data.isnull().sum().sum()
```

[12]:
```
0
```

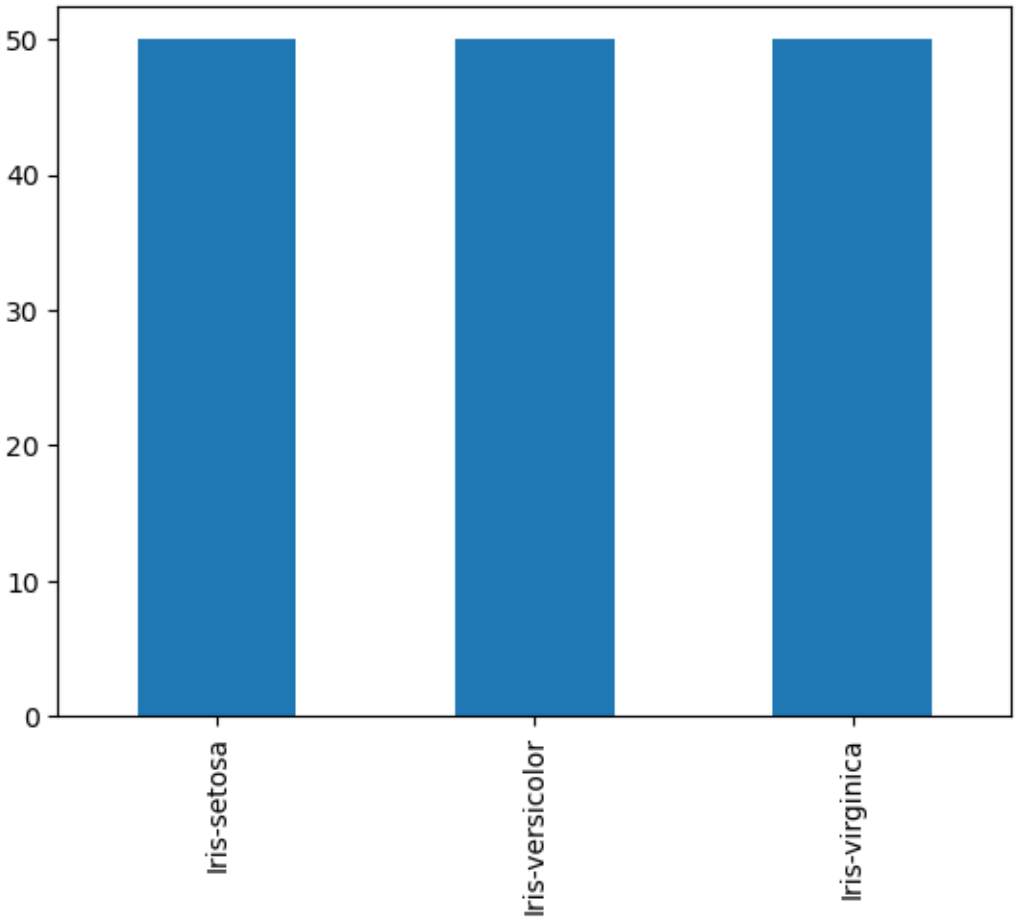
[13]:
```
Data[Data.duplicated()]
```

[13]:
```
Empty DataFrame
Columns: [Id, SepalLengthCm, SepalWidthCm, PetalLengthCm, PetalWidthCm, Species]
Index: []
```

[14]:
```
# Data Visualization
```

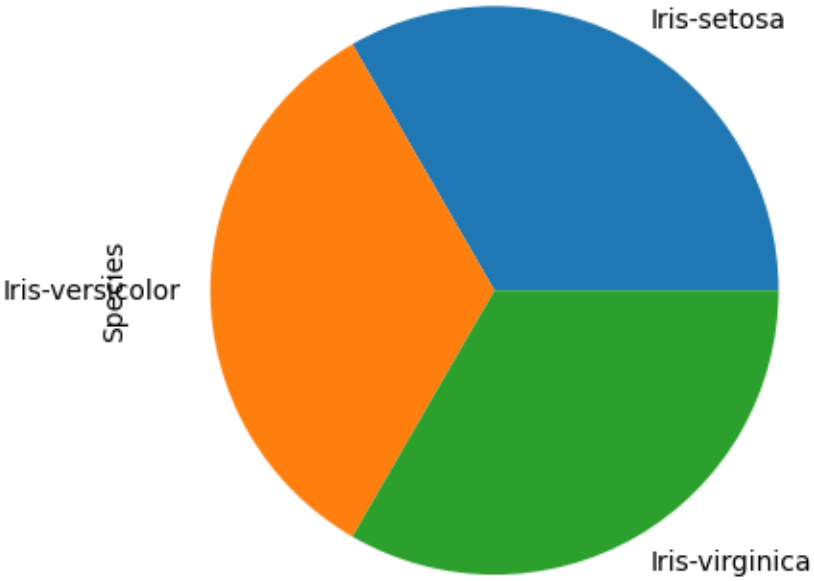
[15]:
```
Data['Species'].value_counts().plot(kind='bar')
```

[15]:
```
<Axes: >
```



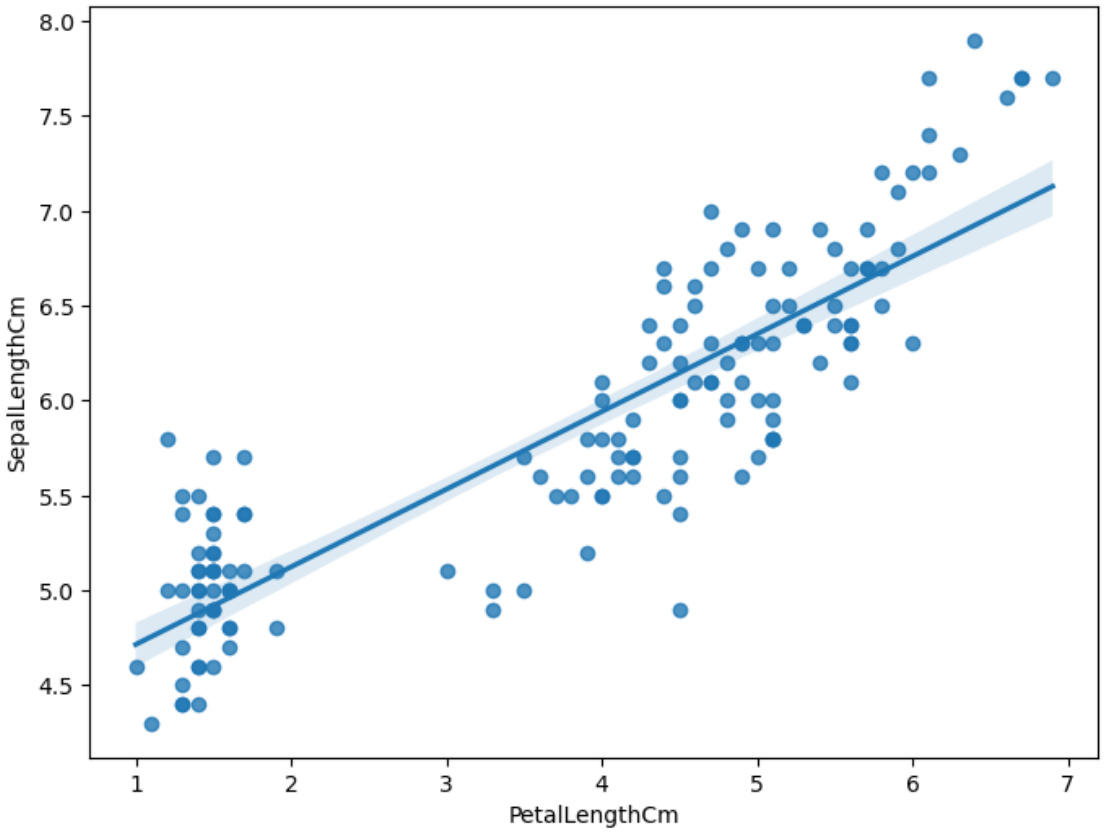
[16]:
```
Data['Species'].value_counts().plot.pie()
```

[16]: <Axes: ylabel='Species'>



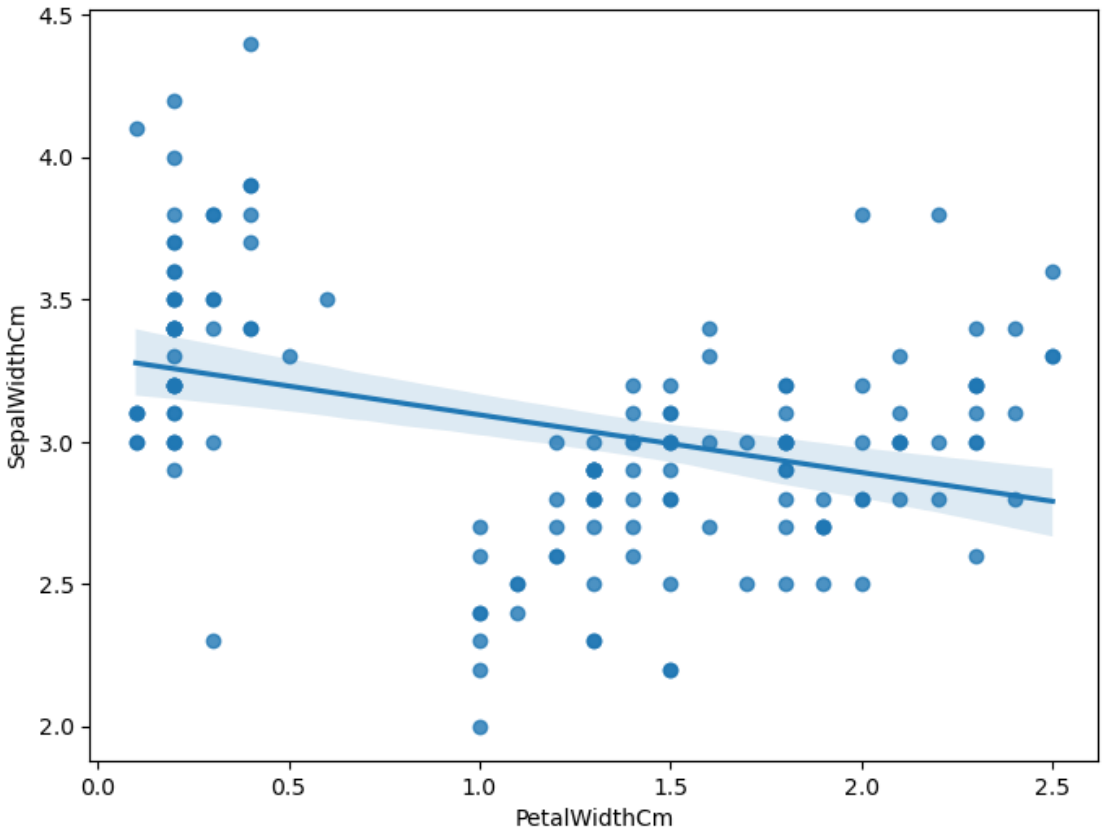
[17]:
```
plt.figure(figsize=(8,6))
sns.regplot(x='PetalLengthCm',y='SepalLengthCm',data=Data)
```

[17]: <Axes: xlabel='PetalLengthCm', ylabel='SepalLengthCm'>



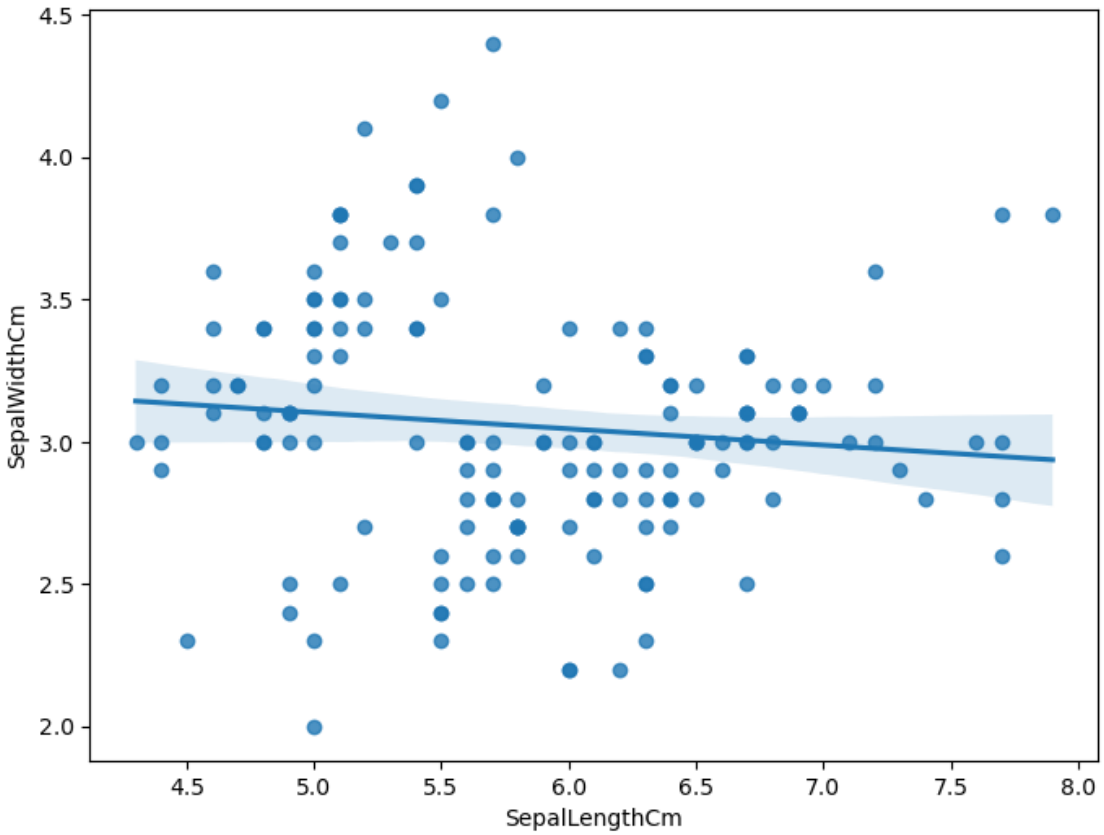
[18]:
```
plt.figure(figsize=(8,6))
sns.regplot(x='PetalWidthCm',y='SepalWidthCm',data=Data)
```

[18]: <Axes: xlabel='PetalWidthCm', ylabel='SepalWidthCm'>



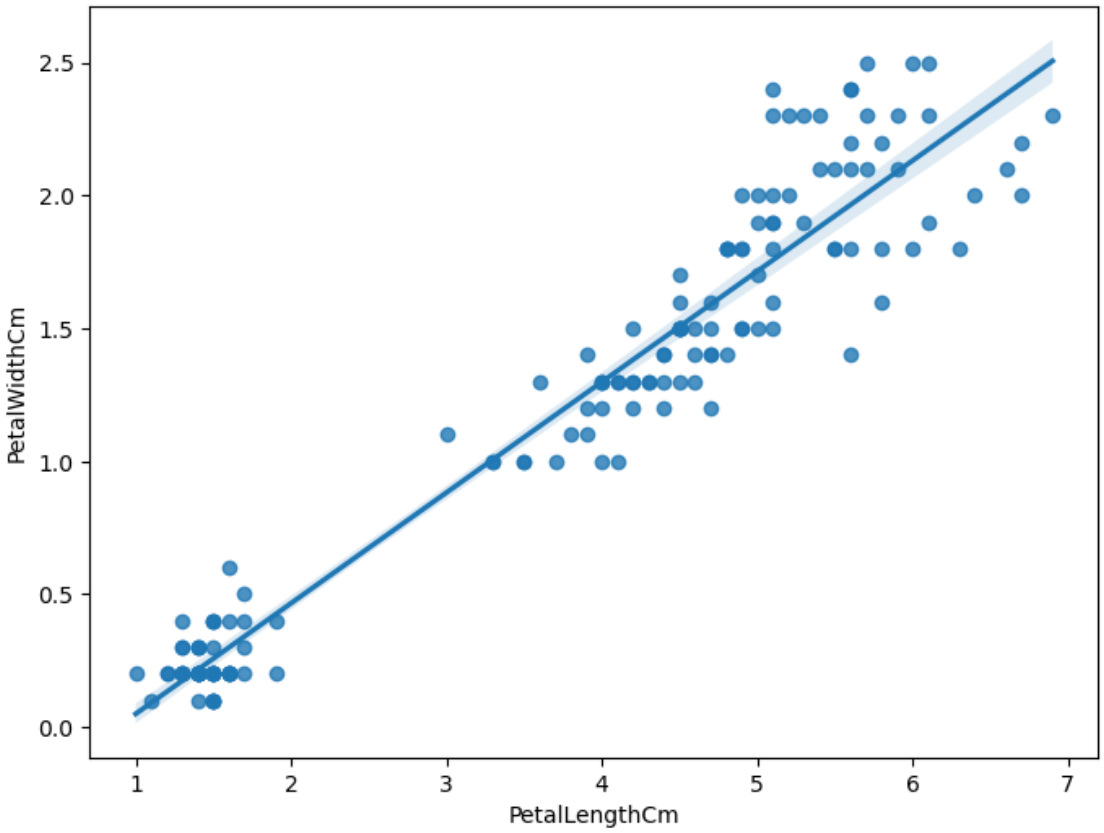
[19]:
```
plt.figure(figsize=(8,6))
sns.regplot(x='SepalLengthCm',y='SepalWidthCm',data=Data)
```

[19]: <Axes: xlabel='SepalLengthCm', ylabel='SepalWidthCm'>



[20]:
```
plt.figure(figsize=(8,6))
sns.regplot(x='PetalLengthCm',y='PetalWidthCm',data=Data)
```

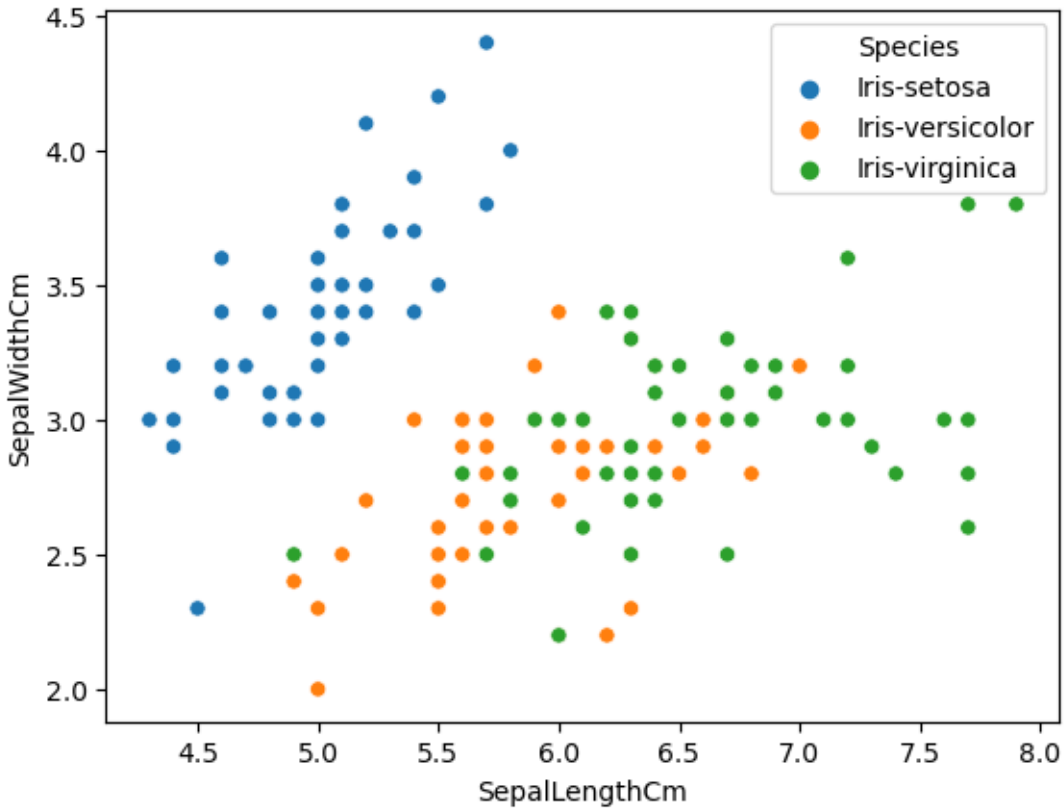
[20]: <Axes: xlabel='PetalLengthCm', ylabel='PetalWidthCm'>



[21]:
```
# Bivariate analysis
```

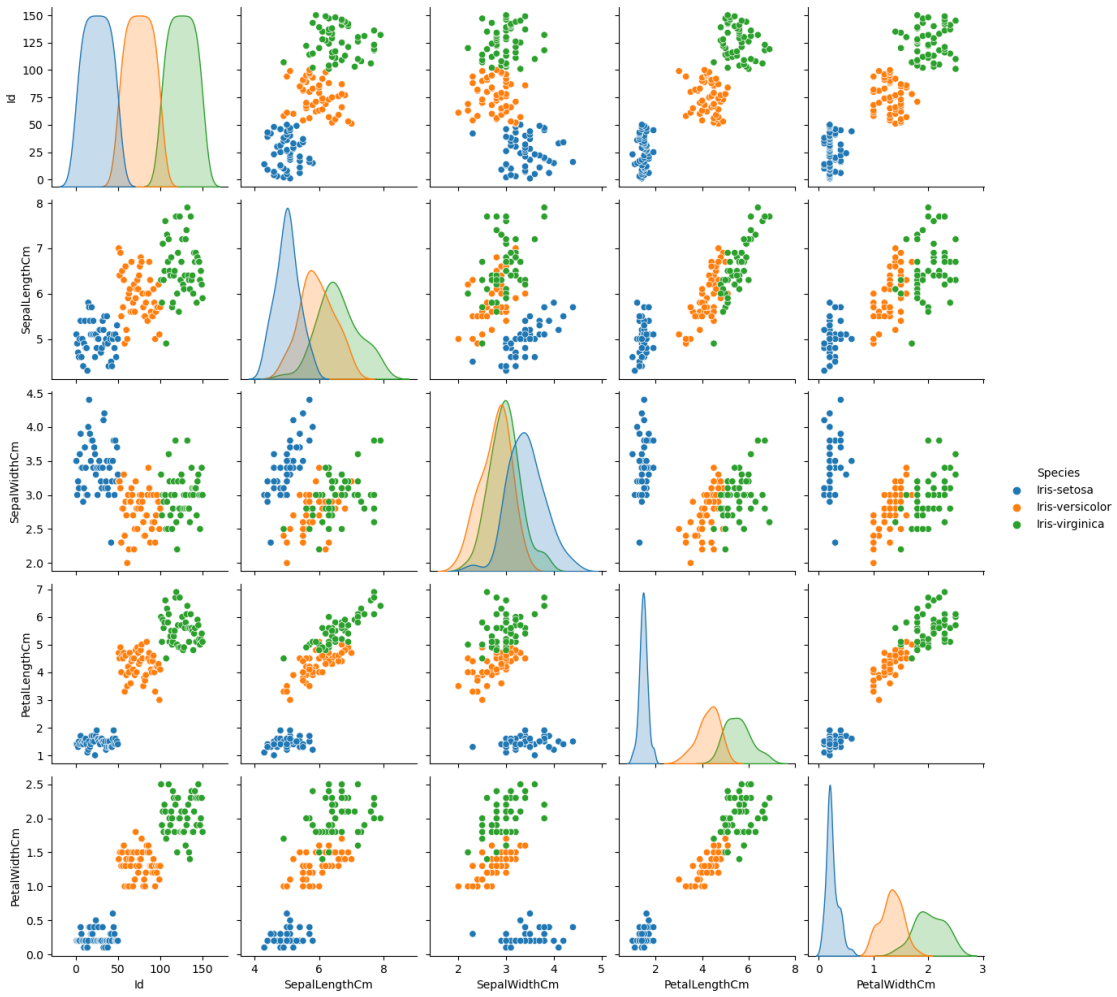
[22]:
```
sns.
↪scatterplot(x=Data["SepalLengthCm"],y=Data["SepalWidthCm"],hue=Data['Species'])
```

[22]:
```
<Axes: xlabel='SepalLengthCm', ylabel='SepalWidthCm'>
```



[23]:
```
sns.pairplot(data = Data , hue = 'Species')
```

[23]: <seaborn.axisgrid.PairGrid at 0x18a39f8ce20>



[24]:
```
#  Machine Learning Models
```

[25]:
```
x = Data.iloc[: , 0:4].values
y = Data.iloc[:,4].values
```

[26]:
```
y
```

[26]:
```
array([0.2, 0.2, 0.2, 0.2, 0.2, 0.4, 0.3, 0.2, 0.2, 0.1, 0.2, 0.2, 0.1,
       0.1, 0.2, 0.4, 0.4, 0.3, 0.3, 0.3, 0.2, 0.4, 0.2, 0.5, 0.2, 0.2,
       0.4, 0.2, 0.2, 0.2, 0.2, 0.4, 0.1, 0.2, 0.1, 0.2, 0.2, 0.1, 0.2,
       0.2, 0.3, 0.3, 0.2, 0.6, 0.4, 0.3, 0.2, 0.2, 0.2, 0.2, 1.4, 1.5,
       1.5, 1.3, 1.5, 1.3, 1.6, 1. , 1.3, 1.4, 1. , 1.5, 1. , 1.4, 1.3,
       1.4, 1.5, 1. , 1.5, 1.1, 1.8, 1.3, 1.5, 1.2, 1.3, 1.4, 1.4, 1.7,
       1.5, 1. , 1.1, 1. , 1.2, 1.6, 1.5, 1.6, 1.5, 1.3, 1.3, 1.3, 1.2,
       1.4, 1.2, 1. , 1.3, 1.2, 1.3, 1.3, 1.1, 1.3, 2.5, 1.9, 2.1, 1.8,
       2.2, 2.1, 1.7, 1.8, 1.8, 2.5, 2. , 1.9, 2.1, 2. , 2.4, 2.3, 1.8,
```

```
       2.2, 2.3, 1.5, 2.3, 2. , 2. , 1.8, 2.1, 1.8, 1.8, 1.8, 2.1, 1.6,
       1.9, 2. , 2.2, 1.5, 1.4, 2.3, 2.4, 1.8, 1.8, 2.1, 2.4, 2.3, 1.9,
       2.3, 2.5, 2.3, 1.9, 2. , 2.3, 1.8])
```

[27]:
```
from sklearn.preprocessing import LabelEncoder
```

[28]:
```
encode = LabelEncoder()
y = encode.fit_transform(y)
y
```

[28]:
```
array([ 1,  1,  1,  1,  1,  3,  2,  1,  1,  0,  1,  1,  0,  0,  1,  3,  3,
        2,  2,  2,  1,  3,  1,  4,  1,  1,  3,  1,  1,  1,  1,  3,  0,  1,
        0,  1,  1,  0,  1,  1,  2,  2,  1,  5,  3,  2,  1,  1,  1,  1, 10,
       11, 11,  9, 11,  9, 12,  6,  9, 10,  6, 11,  6, 10,  9, 10, 11,  6,
       11,  7, 14,  9, 11,  8,  9, 10, 10, 13, 11,  6,  7,  6,  8, 12, 11,
       12, 11,  9,  9,  9,  8, 10,  8,  6,  9,  8,  9,  9,  7,  9, 21, 15,
       17, 14, 18, 17, 13, 14, 14, 21, 16, 15, 17, 16, 20, 19, 14, 18, 19,
       11, 19, 16, 16, 14, 17, 14, 14, 14, 17, 12, 15, 16, 18, 11, 10, 19,
       20, 14, 14, 17, 20, 19, 15, 19, 21, 19, 15, 16, 19, 14],
      dtype=int64)
```

[29]:
```
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score

# Encode the species column
label_encoder = LabelEncoder()
Data['Species'] = label_encoder.fit_transform(Data['Species'])

# Split the data into features and target
X = Data.drop(['Species', 'Id'], axis=1)
y = Data['Species']

# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,␣
 ↪random_state=42)

# Train a Random Forest Classifier
rf_classifier = RandomForestClassifier()
rf_classifier.fit(X_train, y_train)

# Predict on the test set
y_pred = rf_classifier.predict(X_test)

# Calculate the accuracy of the model
accuracy = accuracy_score(y_test, y_pred)
```

```
accuracy
```

[29]:
```
1.0
```

[30]:
```
X_train , X_test , y_train , y_test = train_test_split(x,y,test_size=.20)
```

[31]:
```
model = KNeighborsClassifier(n_neighbors=4)
model.fit(X_train , y_train)
```

[31]:
```
KNeighborsClassifier(n_neighbors=4)
```

[32]:
```
y_pred = model.predict(X_test)
y_pred
```

[32]:
```
array([0, 2, 1, 0, 1, 2, 1, 2, 1, 2, 1, 0, 1, 1, 2, 2, 2, 0, 1, 1, 1, 0,
       0, 1, 2, 0, 0, 1, 0, 0])
```

[33]:
```
y_test
```

[33]:
```
15     0
111    2
61     1
29     0
52     1
144    2
87     1
108    2
59     1
129    2
75     1
8      0
65     1
74     1
142    2
100    2
106    2
40     0
60     1
72     1
71     1
36     0
1      0
70     1
137    2
31     0
20     0
92     1
```

```
7      0
27     0
Name: Species, dtype: int32
```

[34]:
```
from sklearn.metrics import accuracy_score

accuracy_score(y_test , y_pred)
```

[34]:
```
1.0
```

[35]:
```
test_errors = []

for k in range(1,10):
    model = KNeighborsClassifier(n_neighbors=k)
    model.fit(X_train , y_train)


    y_pred_test = model.predict(X_test)
    error = 1- accuracy_score(y_test , y_pred_test)
    test_errors.append(error)

test_errors
```

[35]:
```
[0.0, 0.0, 0.0, 0.0, 0.0, 0.033333333333333326, 0.0, 0.033333333333333326, 0.0]
```