

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

```
In [2]: columns = ['Sepal length', 'Sepal width', 'Petal length', 'Petal width', 'Class_labels']
```

```
In [5]: df = pd.read_csv('iris.data', names=columns)
```

```
In [6]: df.head()
```

Out[6]:

	Sepal length	Sepal width	Petal length	Petal width	Class_labels
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

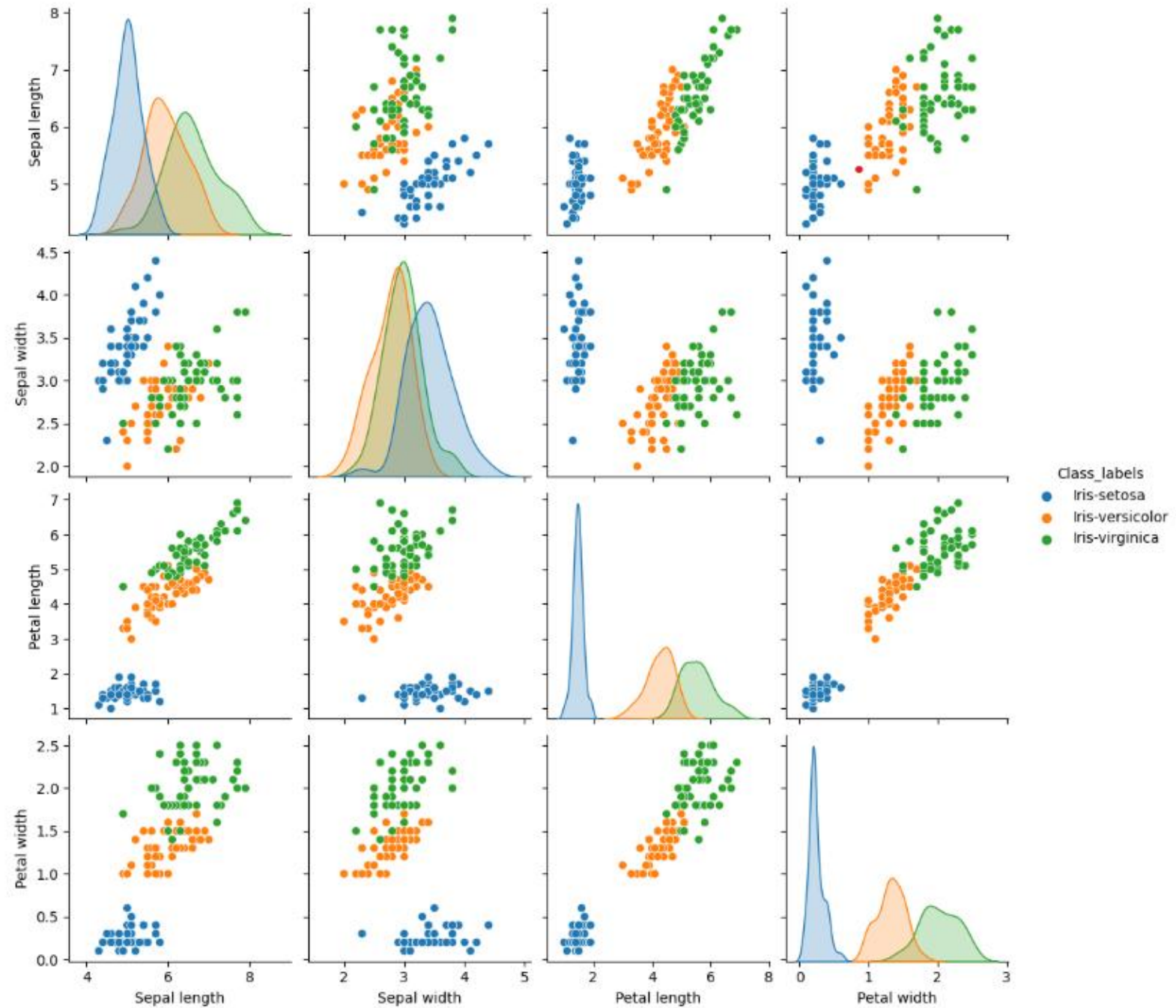
```
In [7]: df.describe()
```

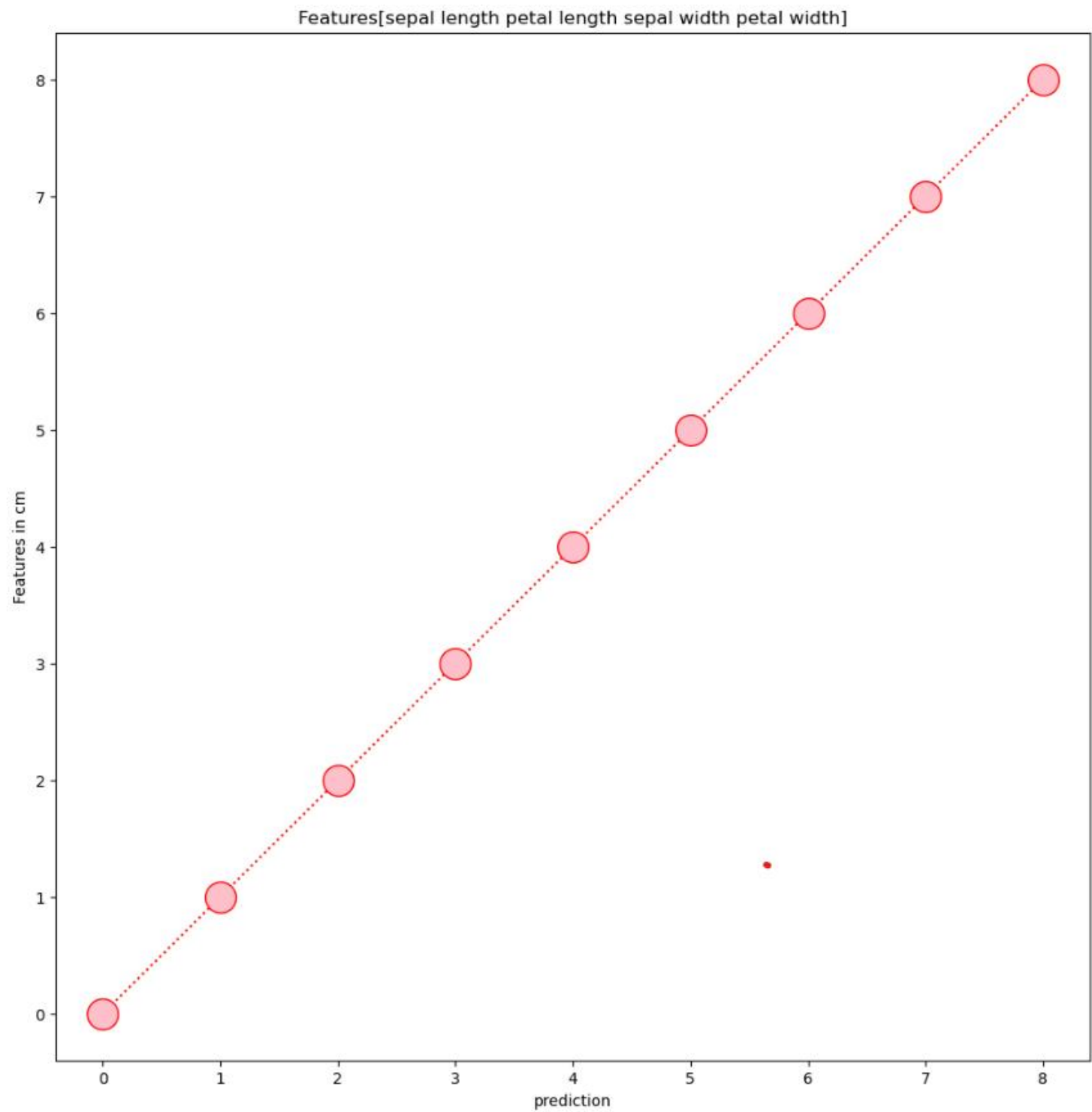
Out[7]:

	Sepal length	Sepal width	Petal length	Petal width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.054000	3.758667	1.198667
std	0.828066	0.433594	1.764420	0.763161

```
In [8]: sns.pairplot(df, hue='Class_labels')
```

```
Out[8]: <seaborn.axisgrid.PairGrid at 0x2b9bf06bd00>
```





```
In [12]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, Y, test_size=0.2)
```

```
In [13]: from sklearn.svm import SVC
svn = SVC()
svn.fit(X_train, y_train)
```

```
Out[13]: ▾ SVC
SVC()
```

```
In [14]: predictions = svn.predict(X_test)
```

```
In [15]: from sklearn.metrics import accuracy_score
accuracy_score(y_test, predictions)
```

```
Out[15]: 0.9666666666666667
```

```
In [18]: from sklearn.metrics import classification_report
print(classification_report(y_test, predictions))
```

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	8
Iris-versicolor	0.92	1.00	0.96	12
Iris-virginica	1.00	0.90	0.95	10
accuracy			0.97	30
macro avg	0.97	0.97	0.97	30
weighted avg	0.97	0.97	0.97	30

```
In [19]: X_new = np.array([[3, 2, 1, 0.2], [ 4.9, 2.2, 3.8, 1.1 ], [ 5.3, 2.5, 4.6, 1.9 ]])
#Prediction of the species from the input vector
prediction = svn.predict(X_new)
print("Prediction of Species: {}".format(prediction))
```

```
Prediction of Species: ['Iris-setosa' 'Iris-versicolor' 'Iris-versicolor']
```

```
In [18]: from sklearn.metrics import classification_report
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```

	precision	recall	f1-score	support
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In [19]: X_new = np.array([[3, 2, 1, 0.2], [ 4.9, 2.2, 3.8, 1.1 ], [ 5.3, 2.5, 4.6, 1.9 ]])
#Prediction of the species from the input vector
prediction = svm.predict(X_new)
print("Prediction of Species: {}".format(prediction))
```

Prediction of Species: ['Iris-setosa' 'Iris-versicolor' 'Iris-versicolor']

```
In [20]: import pickle
with open('SVM.pickle', 'wb') as f:
    pickle.dump(svm, f)
```

```
In [21]: with open('SVM.pickle', 'rb') as f:
    model = pickle.load(f)
```

```
In [22]: model.predict(X_new)
```

```
Out[22]: array(['Iris-setosa', 'Iris-versicolor', 'Iris-versicolor'], dtype=object)
```