# USING IRIS DATA SET FOR ANALYSIS

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The data set consists of 50 samples from each of three species of *Iris* (*Iris setosa*, *Iris virginica* and *Iris versicolor*).

Four <u>features</u> were measured from each sample: the length and the width of the <u>sepals</u> and <u>petals</u>, in centimetres.

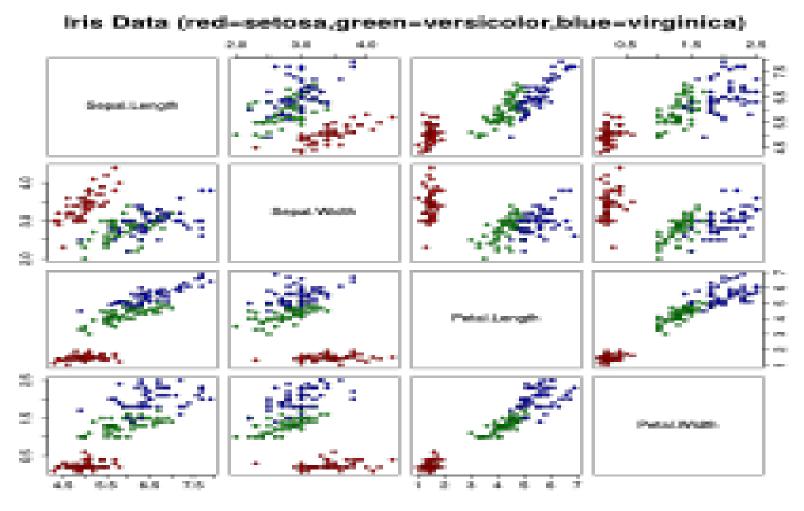
Based on the combination of these four features, Fisher developed a linear discriminant model to distinguish the species from each other.

# Case Study and Practice in Python: IRIS Data Set

This *data sets* consists of 3 different types of *irises*' (Setosa, Versicolour, and Virginica) petal and sepal length, stored in a 150x4 **numpy.ndarray**.

The rows being the samples and the columns being: Sepal Length, Sepal Width, Petal Length and Petal Width.

The below plot uses the first two features.



```
In [15]: import numpy as np
   import seaborn as sns
   import matplotlib.pyplot as plt
   sns.set(color_codes=True)
   import pandas as pd
   %matplotlib inline
   dataset = pd.read_csv('D:/python/IRIS.csv')
   print(dataset)
```

# Download Iris data and see the output

	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	
5	6	5.4	3.9	1.7	0.4	
6	7	4.6	3.4	1.4	0.3	
7	8	5.0	3.4	1.5	0.2	
8	9	4.4	2.9	1.4	0.2	
9	10	4.9	3.1	1.5	0.1	
10	11	5.4	3.7	1.5	0.2	
11	12	4.8	3.4	1.6	0.2	
12	13	4.8	3.0	1.4	0.1	
13	14	4.3	3.0	1.1	0.1	
14	15	5.8	4.0	1.2	0.2	
15	16	5.7	4.4	1.5	0.4	
16	17	5.4	3.9	1.3	0.4	
17	18	5.1	3.5	1.4	0.3	
	4.5	F 7	^ ^	1 7	^ ^	

```
In [15]: import numpy as no
         import seaborn as sns
         import matplotlib.pyplot as plt
         sns.set(color codes=True)
         import pandas as pd
         %matplotlib inline
         dataset = pd.read csv('D:/python/IRIS.csv')
         print (dataset)
         133 Iris-virginica
         134 Iris-virginica
         135 Iris-virginica
         136 Iris-virginica
         137 Iris-virginica
         138 Iris-virginica
         139 Iris-virginica
         140 Iris-virginica
         141 Iris-virginica
         142 Iris-virginica
         143 Iris-virginica
         144 Iris-virginica
         145 Iris-virginica
         146 Iris-virginica
         147 Iris-virginica
         148 Iris-virginica
         149 Iris-virginica
         [150 rows x 6 columns]
```

In [16]: dataset.head()

## Out[16]:

	ld	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

In [17]: #drop Id column dataset = dataset.drop('Id',axis=1) dataset.head()

# DROP ID COLUMN

## Out[17]:

	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

```
In [18]: # shape - SUMMARY OF DATA SET
         print(dataset.shape)
         (150, 5)
In [19]: # more information about iris data
         print(dataset.info())
         <class 'pandas.core.frame.DataFrame'>
         RangeIndex: 150 entries, 0 to 149
         Data columns (total 5 columns):
         SepalLengthCm 150 non-null float64
         SepalWidthCm 150 non-null float64
         PetalLengthCm 150 non-null float64
         PetalWidthCm 150 non-null float64
         Species 150 non-null object
         dtypes: float64(4), object(1)
         memory usage: 5.9+ KB
         None
```

```
In [20]:
         # descriptions
         print(dataset.describe())
```

	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
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
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.350000	1.300000
75 <del>%</del>	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

```
In [21]: # class distribution
         print(dataset.groupby('Species').size())
```

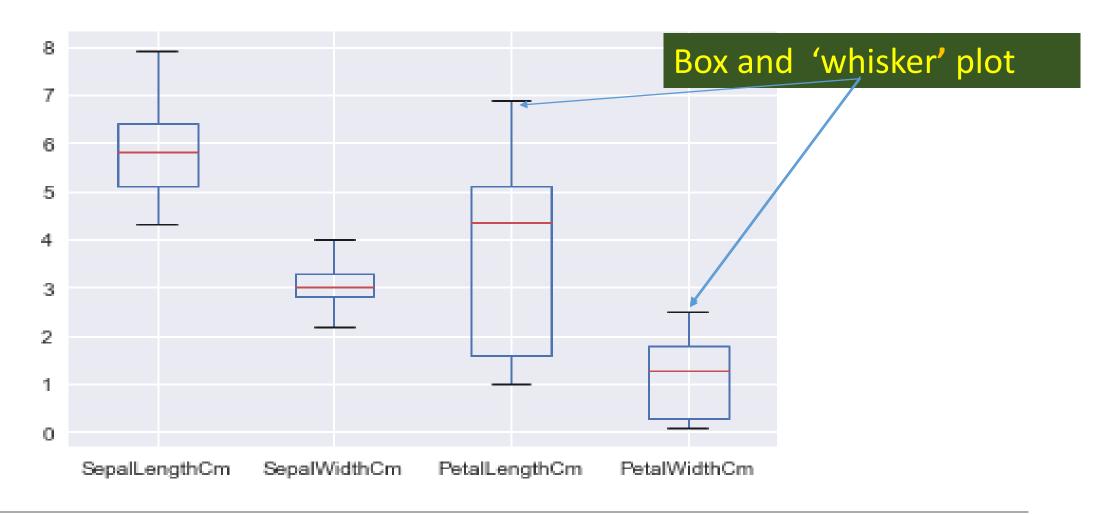
Use 'groupby' to combine

# Species

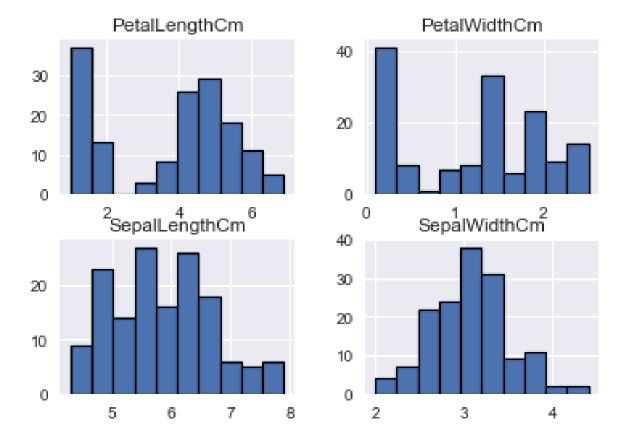
50 Iris-setosa Iris-versicolor 50 50 Iris-virginica dtype: int64

```
In [22]: # box and whisker plots
dataset.plot(kind='box', sharex=False, sharey=False)
```

Out[22]: <matplotlib.axes.\_subplots.AxesSubplot at 0xb0e6668>

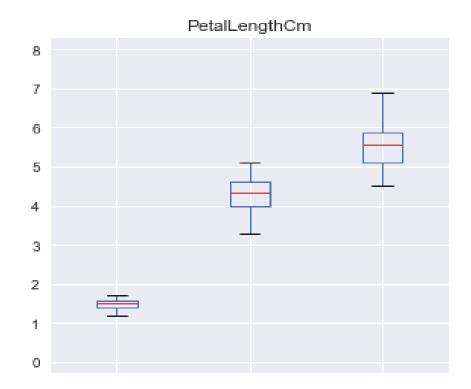


```
In [23]: # histograms
dataset.hist(edgecolor='black', linewidth=1.2)
```

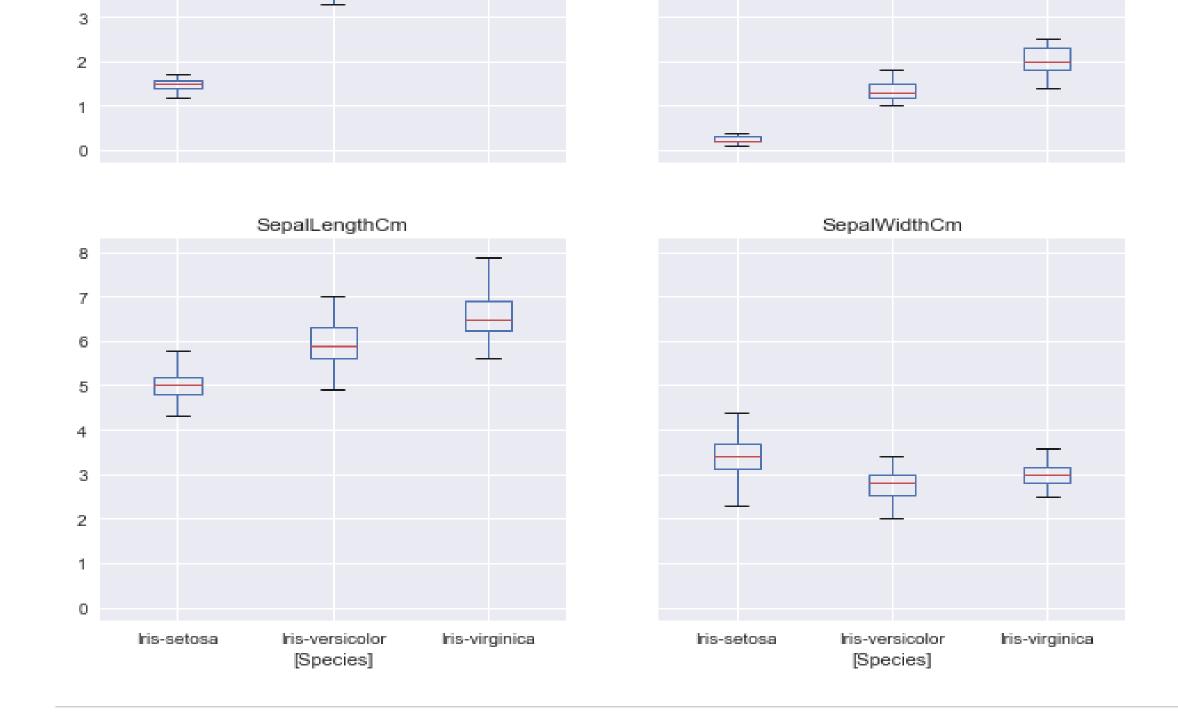


```
In [24]: # boxplot on each feature split out by species
dataset.boxplot(by="Species", figsize=(10,10))
```

Boxplot grouped by Species

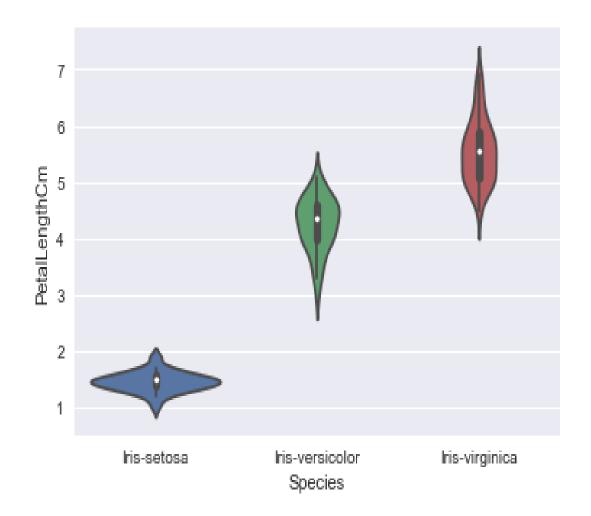






In [25]: # violinplots on petal-length for each species sns.violinplot(data=dataset, x="Species", y="PetalLengthCm")

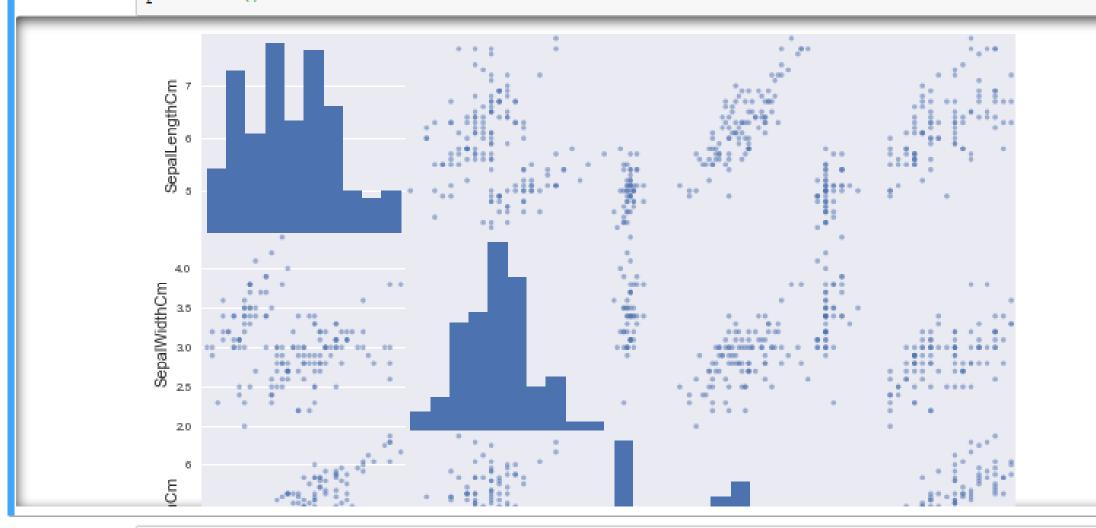
Out[25]: <matplotlib.axes. subplots.AxesSubplot at 0xb48b8d0>





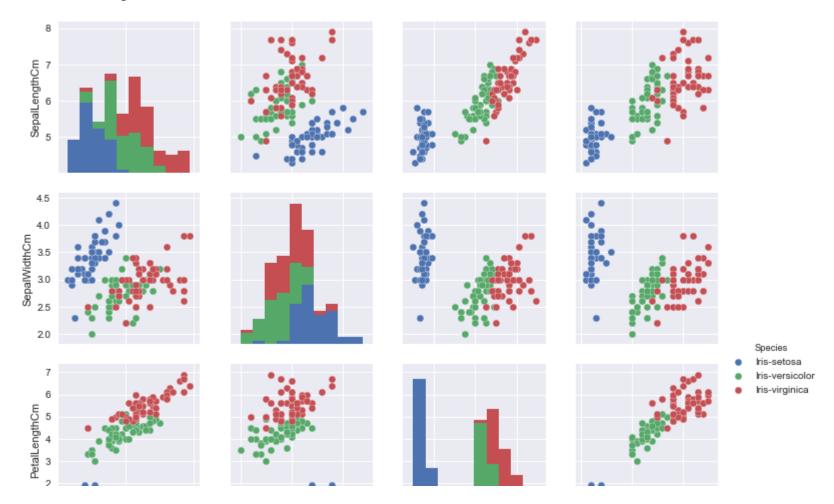
**VOILIN PLOTS** 

```
In [26]: from pandas.plotting import scatter_matrix
     # scatter plot matrix
     scatter_matrix(dataset, figsize=(10,10))
     plt.show()
```



In [27]: # Using seaborn pairplot to see the bivariate relation between each pair of features
 sns.pairplot(dataset, hue="Species")

Out[27]: <seaborn.axisgrid.PairGrid at 0xbc2f550>



```
In [28]: # Importing metrics for evaluation
         from sklearn.metrics import confusion matrix
         from sklearn.metrics import classification report
In [29]: # Seperating the data into dependent and independent variables
         X = dataset.iloc[:, :-1].values
         y = dataset.iloc[:, -1].values
         # Splitting the dataset into the Training set and Test set
         from sklearn.cross validation import train test split
         X train, X test, y train, y test = train test split(X, y, test size = 0.2, random state = 0)
         C:\Users\LENOVO\Anaconda1\lib\site-packages\sklearn\cross validation.py:41: DeprecationWarning: This module was deprecated
         in version 0.18 in favor of the model selection module into which all the refactored classes and functions are moved. Also
```

note that the interface of the new CV iterators are different from that of this module. This module will be removed in 0.20

"This module will be removed in 0.20.", DeprecationWarning)

```
In [31]: # LogisticRegression
    from sklearn.linear_model import LogisticRegression
    classifier = LogisticRegression()
    classifier.fit(X_train, y_train)
    y_pred = classifier.predict(X_test)

# Summary of the predictions made by the classifier
    print(classification_report(y_test, y_pred))
    print(confusion_matrix(y_test, y_pred))
# Accuracy score
    from sklearn.metrics import accuracy_score
    print('accuracy is',accuracy_score(y_pred,y_test))
```

	precision	recall	f1-score	support
Iris-setosa	1.00	1.00	1.00	11
Iris-versicolor	1.00	0.92	0.96	13
Iris-virginica	0.86	1.00	0.92	6
avg / total	0.97	0.97	0.97	30

[[11 0 0] [ 0 12 1] [ 0 0 6]] accuracy is 0.96666666666666667

```
In [32]: # Decision Tree's
    from sklearn.tree import DecisionTreeClassifier

    classifier = DecisionTreeClassifier()

    classifier.fit(X_train, y_train)

    y_pred = classifier.predict(X_test)

# Summary of the predictions made by the classifier
    print(classification_report(y_test, y_pred))
    print(confusion_matrix(y_test, y_pred))

# Accuracy score
    from sklearn.metrics import accuracy_score
    print('accuracy is',accuracy_score(y_pred,y_test))
```

	precision	recall	f1-score	support
Iris-setosa Iris-versicolor	1.00 1.00	1.00 1.00	1.00 1.00	11 13
Iris-virginica	1.00	1.00	1.00	6
avg / total	1.00	1.00	1.00	30
[[11 0 0] [ 0 13 0] [ 0 0 6]] accuracy is 1.0				