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```
from PIL import Image
 In [1]:
          irf=Image.open('irisimages.jpg')
 In [4]:
 In [5]:
          print(irf)
          <PIL.JpegImagePlugin.JpegImageFile image mode=RGB size=367x137 at 0x26F664170A0>
          irf
 In [6]:
 Out[6]:
               Iris Versicolor
                                  Iris Setosa
                                                 Iris Virginica
          #importing the libraries
 In [8]:
          import pandas as pd
          import numpy as np
          import seaborn as sns
          from matplotlib import pyplot as plt
          #load and read the iris dataset
 In [9]:
          data=pd.read_csv('IRIS.csv')
          print(data)
In [10]:
               sepal_length sepal_width petal_length petal_width
                                                                              species
         0
                        5.1
                                      3.5
                                                    1.4
                                                                  0.2
                                                                          Iris-setosa
                        4.9
         1
                                      3.0
                                                                  0.2
                                                                          Iris-setosa
                                                    1.4
         2
                        4.7
                                      3.2
                                                    1.3
                                                                  0.2
                                                                          Iris-setosa
         3
                        4.6
                                      3.1
                                                    1.5
                                                                  0.2
                                                                          Iris-setosa
                        5.0
                                                                  0.2
                                                                          Iris-setosa
         4
                                      3.6
                                                    1.4
                        . . .
                                      . . .
                                                    . . .
                                                                  . . .
          . .
                        6.7
                                                                  2.3 Iris-virginica
         145
                                      3.0
                                                    5.2
                        6.3
                                                                  1.9 Iris-virginica
         146
                                      2.5
                                                    5.0
         147
                        6.5
                                      3.0
                                                    5.2
                                                                  2.0 Iris-virginica
         148
                                                                  2.3 Iris-virginica
                        6.2
                                      3.4
                                                    5.4
         149
                        5.9
                                      3.0
                                                    5.1
                                                                  1.8 Iris-virginica
         [150 rows x 5 columns]
In [11]:
          data.head()
```

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Out[11]:	sepal_	length se	pal_width p	oetal_length p	etal_width	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 [12]:	data.tai	il()						
Out[12]:	sepa	al_length	sepal_width	petal_length	petal_widtl	n spe	cies	
	145	6.7	3.0	5.2	2.3	3 Iris-virgii	nica	
	146	6.3	2.5	5.0	1.9	9 Iris-virgii	nica	
	147	6.5	3.0	5.2	2.0	O Iris-virgii	nica	
	148	6.2	3.4	5.4	2.3	3 Iris-virgii	nica	
	149	5.9	3.0	5.1	1.8	8 Iris-virgii	nica	
In [13]:	data.sha	ape						
Out[13]:	(150, 5)							
In [14]:	data.va]	lue_count	s()					
Out[14]:	sepal_le 4.9 5.8 5.9 6.2	ength se 3. 2. 4. 3.	7 0 0	petal_lengt 1.5 5.1 1.2 4.2 5.4	h petal_w 0.1 1.9 0.2 1.5 2.3	In: In: In: In:	ecies is-setosa is-virginica is-setosa is-versicolor is-virginica	3 2 1 1
	5.57.9Length:	2. 2. 3. 147, dty	4 5	4.0 3.7 3.8 4.0 6.4	1.3 1.0 1.1 1.3 2.0	Ir: Ir: Ir:	is-versicolor is-versicolor is-versicolor is-versicolor is-virginica	1 1 1 1 1

In [15]: data.isnull()

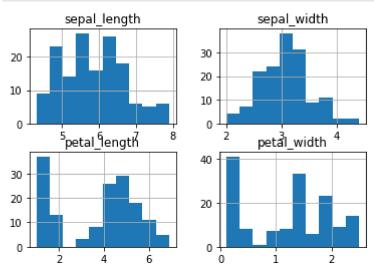
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Out[15]:

sepal_length sepal_width petal_length petal_width species 0 **False False** False False False 1 False False False False False 2 False False False False False 3 False False False False False 4 False **False** False False False ••• 145 **False False** False False **False** 146 False False False False False 147 **False False** False False **False** 148 False False False False **False** 149 **False False** False False **False**

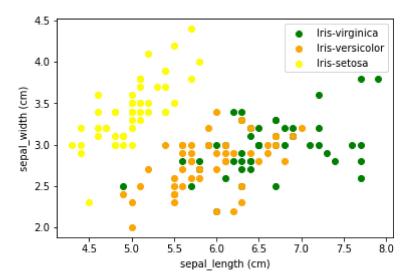
150 rows × 5 columns

```
data.isnull().sum()
In [16]:
         sepal_length
                          0
Out[16]:
          sepal width
                          0
         petal length
                          0
         petal_width
                          0
          species
         dtype: int64
          data.hist()
In [17]:
          plt.show()
```

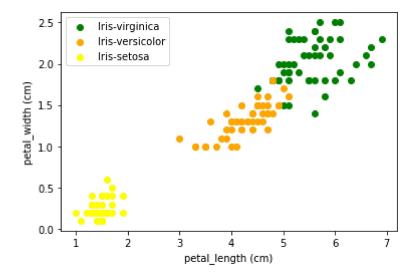


```
plt.ylabel('sepal_width (cm)')
plt.legend()
```

Out[21]: <matplotlib.legend.Legend at 0x26f6e4890a0>

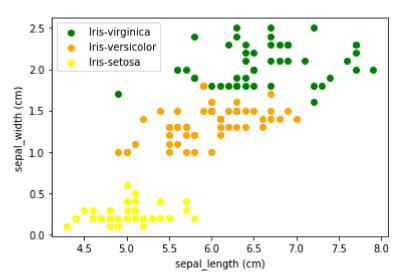


Out[22]: <matplotlib.legend.Legend at 0x26f6e56aca0>



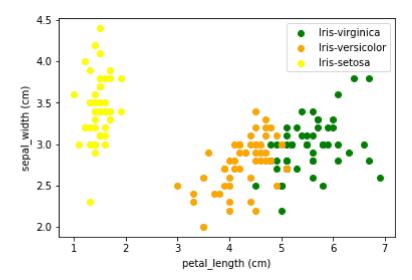
```
In [24]:
    c=['green','orange','yellow']
    f=['Iris-virginica','Iris-versicolor','Iris-setosa']
    for i in range(3):
        x=data[data['species'] == f[i]]
        plt.scatter(x['sepal_length'],x['petal_width'],c=c[i],label=f[i])
    plt.xlabel('sepal_length (cm)')
    plt.ylabel('petal_width (cm)')
    plt.legend()
```

Out[24]: <matplotlib.legend.Legend at 0x26f6e4f8a30>



```
In [25]:
    c=['green','orange','yellow']
    f=['Iris-virginica','Iris-versicolor','Iris-setosa']
    for i in range(3):
        x=data[data['species'] == f[i]]
        plt.scatter(x['petal_length'],x['sepal_width'],c=c[i],label=f[i])
    plt.xlabel('petal_length (cm)')
    plt.ylabel('sepal_width (cm)')
    plt.legend()
```

Out[25]: <matplotlib.legend.Legend at 0x26f6acf8d00>



In [26]: data.corr()

Out[26]:

	sepal_length	sepal_width	petal_length	petal_width
sepal_length	1.000000	-0.109369	0.871754	0.817954
sepal_width	-0.109369	1.000000	-0.420516	-0.356544
petal_length	0.871754	-0.420516	1.000000	0.962757
petal_width	0.817954	-0.356544	0.962757	1.000000

```
iris data analysis
           sns.heatmap(data.corr())
In [27]:
           <AxesSubplot:>
Out[27]:
                                                                       -1.0
           sepal_length -
                                                                       - 0.8
                                                                       - 0.6
            sepal_width
                                                                       - 0.4
                                                                       - 0.2
           petal length
                                                                       - 0.0
                                                                       -0.2
            petal_width
                                                                        -0.4
                       sepal_length sepal_width petal_length petal_width
In [29]:
           from sklearn.linear model import LinearRegression,LogisticRegression
           from sklearn.model selection import train test split
           from sklearn.tree import DecisionTreeClassifier
           from sklearn.preprocessing import LabelEncoder
In [31]:
           e=LabelEncoder()
           data['species']=e.fit transform(data['species'])
In [32]:
           data.head(10)
Out[32]:
              sepal_length
                           sepal_width petal_length petal_width species
           0
                       5.1
                                                 1.4
                                                              0.2
                                                                        0
                                    3.5
           1
                       4.9
                                    3.0
                                                 1.4
                                                              0.2
                                                                        0
           2
                       4.7
                                    3.2
                                                 1.3
                                                              0.2
                                                                        0
           3
                                                 1.5
                                                              0.2
                                                                        0
                       4.6
                                    3.1
           4
                       5.0
                                    3.6
                                                 1.4
                                                              0.2
                                                                        0
           5
                       5.4
                                    3.9
                                                 1.7
                                                              0.4
                                                                        0
           6
                       4.6
                                    3.4
                                                 1.4
                                                              0.3
                                                                        0
                                                                        0
           7
                       5.0
                                    3.4
                                                 1.5
                                                              0.2
           8
                                    2.9
                                                 1.4
                                                              0.2
                                                                        0
                       4.4
```

data.tail()

In [33]:

4.9

3.1

1.5

0.1

0

```
sepal_length sepal_width petal_length petal_width species
Out[33]:
           145
                          6.7
                                       3.0
                                                     5.2
                                                                  2.3
                                                                             2
           146
                          6.3
                                       2.5
                                                     5.0
                                                                  1.9
                                                                             2
           147
                          6.5
                                       3.0
                                                     5.2
                                                                  2.0
                                                                             2
           148
                          6.2
                                                     5.4
                                                                  2.3
                                                                             2
                                       3.4
           149
                                                                             2
                          5.9
                                       3.0
                                                     5.1
                                                                  1.8
```

```
In [34]: x=data.drop(columns=['species'])
    y=data['species']
```

In [35]:

Out[35]:

	sepal_length	sepal_width	petal_length	petal_width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
•••		•••		***
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

150 rows × 4 columns

```
In [36]:
                 0
Out[36]:
                 0
                 0
          2
         3
                 0
                 0
         145
                 2
         146
                 2
                 2
         147
         148
                 2
         149
         Name: species, Length: 150, dtype: int32
         x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.30)
In [37]:
         lg=LogisticRegression()
In [38]:
```

```
In [39]:
        lg.fit(x_train,y_train)
        LogisticRegression()
Out[39]:
In [40]:
         s=lg.score(x_test,y_test)*100
In [41]:
         print(s)
        97.7777777777777
         print("the accuracy of logistic regerssion is",s)
In [43]:
        the accuracy of logistic regerssion is 97.7777777777777
        y_predict=lg.predict(x_test)
In [44]:
In [45]:
        print(y_predict)
        2 2 2 0 0 0 2 2]
        dt=DecisionTreeClassifier()
In [46]:
         dt.fit(x_train,y_train)
In [47]:
        DecisionTreeClassifier()
Out[47]:
In [48]:
        dscore=dt.score(x_test,y_test)*100
In [49]:
         print(dscore)
        91.11111111111111
In [50]: print("the accuracy of logistic regerssion is",dscore)
        the accuracy of logistic regerssion is 91.111111111111111
        from sklearn.metrics import confusion matrix
In [51]:
         cm=confusion_matrix(y_test,y_predict)
         print(cm)
        [[17 0 0]
         [ 0 14 1]
         [ 0 0 13]]
In [ ]:
```

In []:	
Tn [].	
In []:	
In []:	
In []:	
In []:	