```
In [1]: #importing libraries
          import numpy as np
          import pandas as pd
          import matplotlib.pyplot as plt
          import seaborn as sns
          import random
In [2]: | df = pd.read_csv('iris.csv',header = None)
          df.columns = ['separ_length','separ_width','petal_length','petal_width','clas
In [3]: df.head(10)
Out[3]:
             separ_length separ_width petal_length petal_width
                                                                     class
          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
                                                            0.2 Iris-setosa
                                   3.1
                                                1.5
                      4.6
                      5.0
                                   3.6
                                                1.4
                                                            0.2 Iris-setosa
           5
                      5.4
                                   3.9
                                                1.7
                                                            0.4 Iris-setosa
                                                            0.3 Iris-setosa
           6
                      4.6
                                   3.4
                                                1.4
          7
                      5.0
                                   3.4
                                                1.5
                                                            0.2 Iris-setosa
          8
                      4.4
                                   2.9
                                                1.4
                                                            0.2 Iris-setosa
           9
                      4.9
                                   3.1
                                                1.5
                                                            0.1 Iris-setosa
In [4]: | df.shape
```

Out[4]: (150, 5)

In [5]: df.describe()

Out[5]:

	separ_length	separ_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
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%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

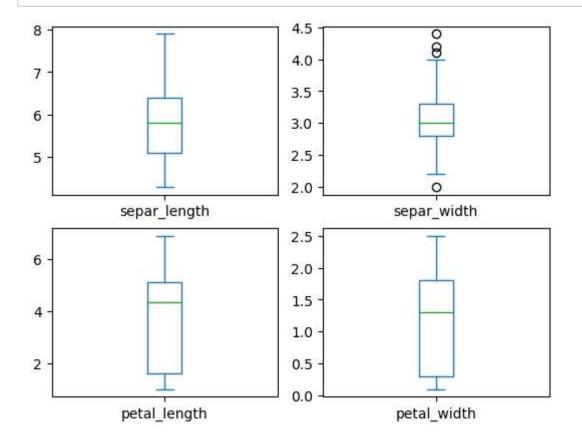
In [6]: df['class'].value_counts()

Out[6]: Iris-setosa 50
Iris-versicolor 50
Iris-virginica 50

Name: class, dtype: int64

In [7]: #ploting box plot

df.plot(kind='box',subplots=True,layout=(2,2),sharex=False,sharey=False)
plt.show()



```
In [8]: #removing outliers from the dataset

Q1 = df.separ_width.quantile(0.25)
Q3 = df.separ_width.quantile(0.75)
IQR = Q3-Q1
df = df[(df.separ_width>=Q1-1.5*IQR) & (df.separ_width<=Q3+1.5*IQR)]</pre>
```

In [9]: df

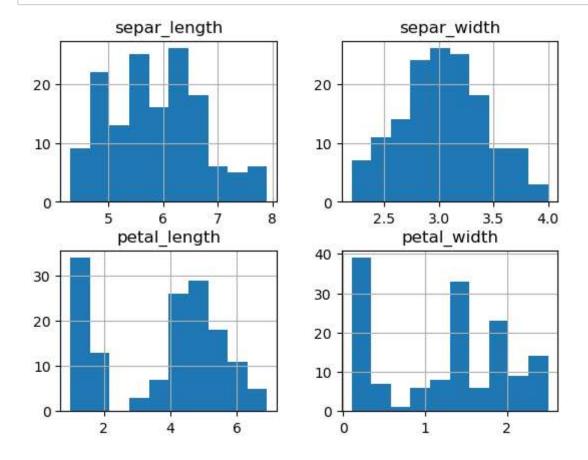
Out[9]:

	separ_length	separ_width	petal_length	petal_width	class
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
145	6.7	3.0	5.2	2.3	Iris-virginica
146	6.3	2.5	5.0	1.9	Iris-virginica
147	6.5	3.0	5.2	2.0	Iris-virginica
148	6.2	3.4	5.4	2.3	Iris-virginica
149	5.9	3.0	5.1	1.8	Iris-virginica

146 rows × 5 columns

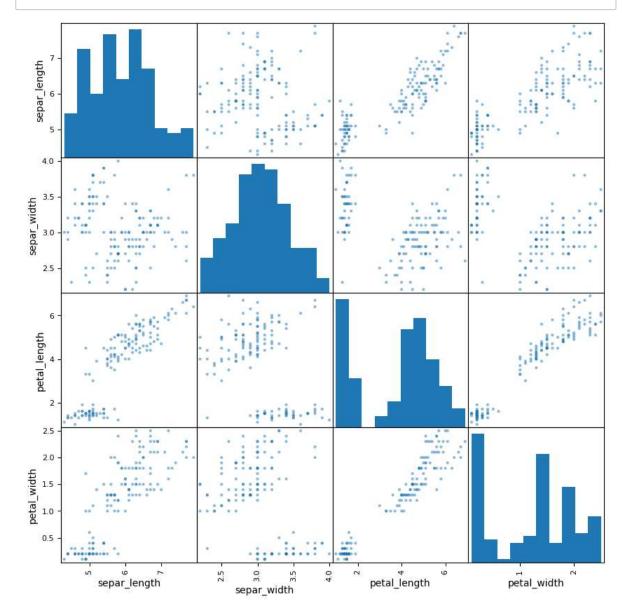
In [10]: #ploting histogram

df.hist()
 plt.show()

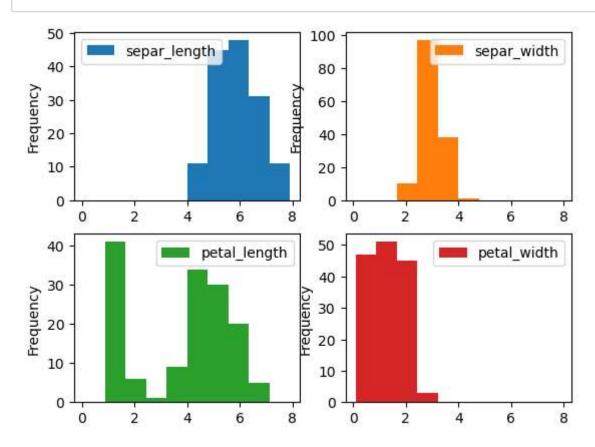


In [11]: #show scatter matrics

pd.plotting.scatter_matrix(df,figsize=(10,10))
plt.show()



In [12]: df.plot(kind='hist',subplots=True,layout=(2,2),sharex=False,sharey=False)
plt.show()



```
In [13]: #spitting dataset

arr = df.values
x = arr[:,:4]
y = arr[:,4]
```

- In [14]: #changing string labels to number values/labels with the help of LabelEncoder
 from sklearn.preprocessing import LabelEncoder

 y = LabelEncoder().fit_transform(y)
- In [15]: #importint libraries
 from sklearn.model_selection import train_test_split
 from sklearn.model_selection import cross_validate
- In [16]: x_train,x_test,y_train,y_test = train_test_split(x,y,test_size = 0.2,random_s

```
In [17]: #determining shape of splitting dataset
         print("shape of x_train = ",x_train.shape)
         print("shape of x_test = ",x_test.shape)
         print("shape of y_train = ",y_train.shape)
         print("shape of y_test = ",y_test.shape)
         shape of x_{train} = (116, 4)
         shape of x test = (30, 4)
         shape of y_{train} = (116,)
         shape of y_{test} = (30,)
In [18]: #import multiple models for comparison
         from sklearn.linear model import LogisticRegression
         from sklearn.tree import DecisionTreeRegressor
         from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
         from sklearn.neighbors import KNeighborsClassifier
         from sklearn.naive bayes import GaussianNB
         from sklearn.svm import SVC
         from sklearn.model selection import cross val score
         from sklearn.model_selection import KFold
In [19]: #creating set of models
         models = \{\}
         models["LR"] = LogisticRegression()
         models["DTR"] = DecisionTreeRegressor()
         models["LDA"] = LinearDiscriminantAnalysis()
         models["KNC"] = KNeighborsClassifier()
         models["GNB"] = GaussianNB()
         models["SVM"] = SVC()
```

```
In [20]: result = []
    for key in models:
        kfold = KFold(n_splits=10,random_state=1,shuffle=True)
        cv_results = cross_val_score(models[key],x_train,y_train,cv=kfold,scoring result.append(cv_results)
        print('%s : %f (%f)',(key,cv_results.mean(),cv_results.std()))
```

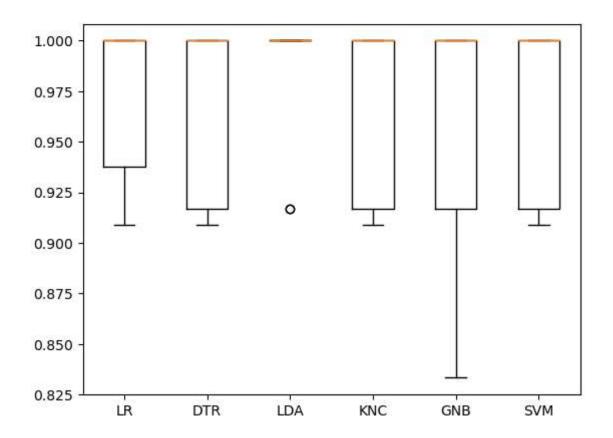
```
%s : %f (%f) ('LR', 0.97424242424242, 0.0393939393939393941)
%s : %f (%f) ('DTR', 0.96590909090908, 0.041804180791883035)
%s : %f (%f) ('LDA', 0.9833333333333, 0.0333333333333335)
%s : %f (%f) ('KNC', 0.9651515151515151, 0.04274768478686633)
%s : %f (%f) ('GNB', 0.957575757576, 0.05650926120900459)
%s : %f (%f) ('SVM', 0.96515151515151, 0.04274768478686633)
```

```
C:\ProgramData\anaconda3\Lib\site-packages\sklearn\linear_model\_logistic.p
y:460: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max_iter) or scale the data as shown in:
    https://scikit-learn.org/stable/modules/preprocessing.html (https://scik
it-learn.org/stable/modules/preprocessing.html)
Please also refer to the documentation for alternative solver options:
    https://scikit-learn.org/stable/modules/linear model.html#logistic-regre
ssion (https://scikit-learn.org/stable/modules/linear model.html#logistic-re
gression)
  n iter i = check optimize result(
C:\ProgramData\anaconda3\Lib\site-packages\sklearn\linear model\ logistic.p
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ssion (https://scikit-learn.org/stable/modules/linear_model.html#logistic-re
gression)
  n_iter_i = _check_optimize_result(
C:\ProgramData\anaconda3\Lib\site-packages\sklearn\linear_model\_logistic.p
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```

```
ssion (https://scikit-learn.org/stable/modules/linear_model.html#logistic-re
gression)
 n_iter_i = _check_optimize_result(
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ssion (https://scikit-learn.org/stable/modules/linear_model.html#logistic-re
gression)
  n_iter_i = _check_optimize_result(
```

```
In [21]: fig = plt.figure()
    fig.suptitle("Algorithm comparison0")
    ax = fig.add_subplot(111)
    plt.boxplot(result)
    ax.set_xticklabels(models.keys())
    plt.show()
```

Algorithm comparison0



In [22]: #importing libraries

```
from sklearn.metrics import classification_report
from sklearn.metrics import confusion_matrix
from sklearn.metrics import accuracy_score
```

```
In [23]: svm = SVC()
         svm.fit(x_train,y_train)
         predictions = svm.predict(x_test)
         print("accuracy : ",accuracy_score(y_test,predictions))
         print("confusion matrix:\n",confusion_matrix(y_test,predictions))
         print("classification_report:\n",classification_report(y_test,predictions))
         accuracy: 0.966666666666667
         confusion matrix:
          [[ 9 0 0]
          [ 0 10 1]
          [ 0 0 10]]
         classification_report:
                                      recall f1-score
                        precision
                                                         support
                    0
                            1.00
                                       1.00
                                                 1.00
                                                              9
                    1
                            1.00
                                       0.91
                                                 0.95
                                                             11
                    2
                            0.91
                                       1.00
                                                 0.95
                                                             10
                                                 0.97
                                                             30
             accuracy
                            0.97
                                       0.97
                                                 0.97
                                                             30
            macro avg
                            0.97
                                                 0.97
         weighted avg
                                       0.97
                                                             30
```

```
In [ ]:
```