

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

```
In [2]: #Loading the data set
df=pd.read_csv('iris.csv')
df.head()
```

```
Out[2]:
```

	sepal_length	sepal_width	petal_length	petal_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 [3]: #to display stats about data
df.describe()
```

```
Out[3]:
```

	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
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 [4]: #for basic information about dataset
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
 #   Column          Non-Null Count  Dtype
---  -
 0   sepal_length    150 non-null    float64
 1   sepal_width     150 non-null    float64
 2   petal_length    150 non-null    float64
 3   petal_width     150 non-null    float64
 4   species         150 non-null    object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

```
In [5]: df['species'].value_counts()
```

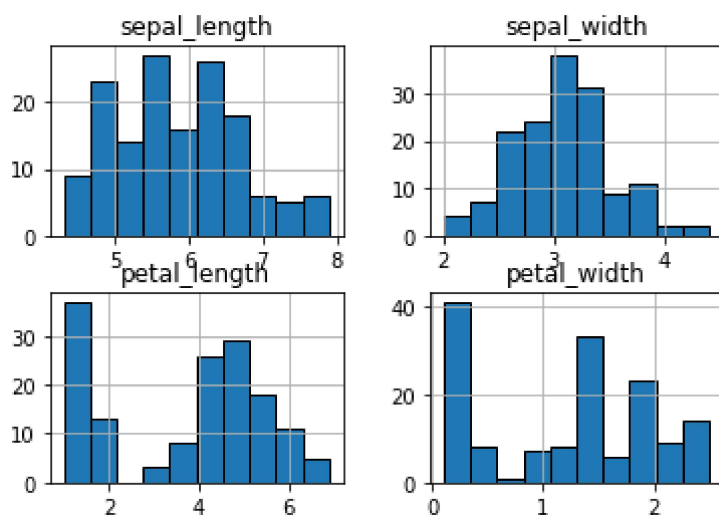
```
Out[5]: Iris-versicolor    50  
Iris-setosa               50  
Iris-virginica           50  
Name: species, dtype: int64
```

```
In [6]: #processing the dataset  
#check for null values  
df.isnull().sum()
```

```
Out[6]: sepal_length    0  
sepal_width           0  
petal_length          0  
petal_width           0  
species               0  
dtype: int64
```

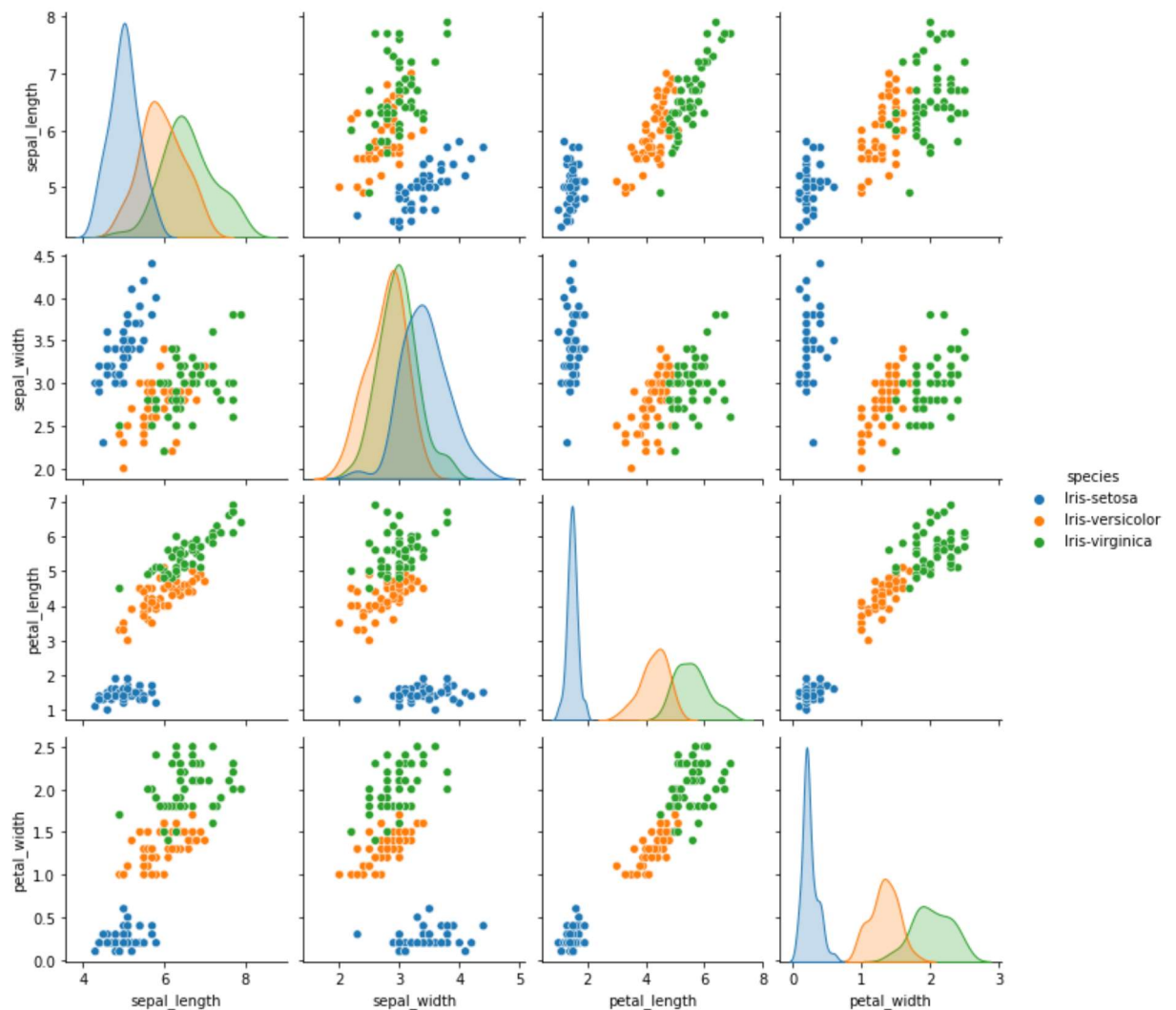
```
In [7]: #Extraordinary Data Analysis  
df.hist(edgecolor='black')
```

```
Out[7]: array([[<AxesSubplot:title={'center':'sepal_length'}>,  
               <AxesSubplot:title={'center':'sepal_width'}>],  
              [<AxesSubplot:title={'center':'petal_length'}>,  
               <AxesSubplot:title={'center':'petal_width'}>]], dtype=object)
```



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

```
Out[8]: <seaborn.axisgrid.PairGrid at 0x257005a7850>
```



```
In [9]: #observation
#All type of flowers are well separable for petal length and petal width
#also all type of bit of well seperable for petal width and sepal width
```

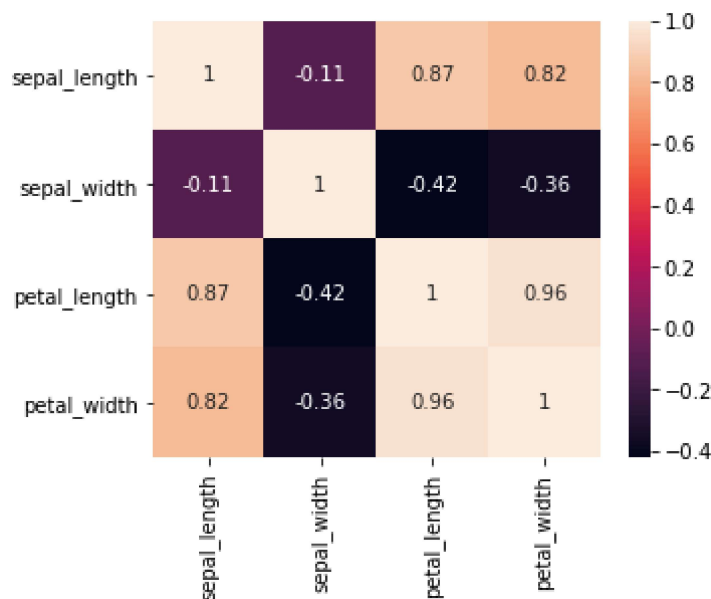
```
In [10]: #as this is classification problem, we will use classification algorithm for model
```

```
In [11]: from sklearn.linear_model import LogisticRegression
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn import svm
from sklearn import metrics
from sklearn.tree import DecisionTreeClassifier
```

```
In [12]: df.corr()

corr1=df.corr()
fig,ax=plt.subplots(figsize=(5,4))
sns.heatmap(corr1,annot=True,ax=ax)
```

Out[12]: <AxesSubplot:>



```
In [13]: df.corr()
```

```
Out[13]:
```

	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

```
In [14]: #observations
#petalLength and petalWidth:corr 0.96 :Having highest correlation
#petalLength and sepalLength:corr 0.87:Having second highest correlation
#petalWidth and sepalLength:corr 0.82 :Having fair enough correlation

#in the above figure,we can see that sepalLength and sepalWidth are not correlated
#while the petalLength and petalWidth are highly correlated
```

```
In [15]: #Iris-setosa :It's usually having smaller features except sepalwidth
#Iris-versicolor:It's having bigger features except sepalwidth
```

```
In [16]: #observation
```

```
In [17]: plt.figure(figsize=(10,10))

plt.subplot(2,2,1)
sns.violinplot(data=df, x='Species',y='SepalLength',palette='Set1')
plt.subplot(2,2,2)
sns.violinplot(data=df, x='Species',y='SepalWidth',palette='Set1')

plt.subplot(2,2,3)
sns.violinplot(data=df,x='Species', y='PetalLength',palette='Set1')
plt.subplot(2,2,4)
sns.violinplot(data=df, x='Species', y='PetalWidth',palette='Set1')
```

```
File "<ipython-input-17-243b53425a80>", line 1
    plt.figure(figsize=(10,10))
    ^
SyntaxError: invalid syntax
```

```
In [ ]: x_train, x_test, y_train, y_test= train X,y, test_size=0.33 ,random_state=4
```

```
In [ ]: model = LogisticRegression()
model.fit(X_train,y_train)
prediction=model.predict(X_test)
print('Logistic Regression accuracy= ',metrics.accuracy_score(prediction,y_test))
```

```
In [ ]: from sklearn.preprocessing import LabelEncoder
le=LabelEncoder() #LabelEncoder can be normalize Lable
```

```
In [ ]: df['species']=le.fit_transform(df['species']) #fit_transform:fit Label encoder ar
df.head()
```

```
In [ ]: X=df.drop(columns=['species']) #Drop Down
y=df['species']
X[:5] ## Return List from begining untel index 5
```

```
In [ ]: y[:5]
```

```
In [ ]: #splitting Dataset into Training set and Test set
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```
In [ ]: from sklearn.model_selection import train_test_split
X_train,X_test,y_train,y_test=train_test_split(X,y,test_size=0.3,random_state=1)
```

```
In [ ]: #selecting the model and Metrics
```

```
In [ ]: from sklearn.linear_model import LogisticRegression
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
from sklearn.naive_bayes import GaussianNB
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score
```

```
In [ ]: lr=LogisticRegression()
kmn=KNeighborsClassifier()
svm=SVC()
nb=GaussianNB()
dt=DecisionTreeClassifier()
rf=RandomForestClassifier()
```

```
models= [lr,kmn,svm,nb,dt,rf] scores=[]
for model in models: model.fit(X_train,y_train)
y_pred=model.predict(X_test)
scores.append(accuracy_score(y_test,y_pred))
print("Accuracy of "+type(model.name)+"is".accuracy_score(y_test,y_pred))
```

```
In [ ]: #Decision tree
model = DecisionTreeClassifier()
model.fit(X_train,y_train)

prediction = model.predict(X_test)
print('Decision Tree accuracy = ', metrics.accuracy_score(prediction,y_test))
```

```
In [ ]: #Support vector Machine (SVM)
model = svm.SVC()
model.fit(X_train,y_train)

prediction = model.predict(X_test)
print('SVM accuracy = ', metrics.accuracy_score(prediction,y_test))
```

```
In [ ]:
```