1. Write a program to perform arithmetic operations in python.

Program:

```
a=int(input("Enter a value:"))
b=int(input("Enter b value:"))
print("add:",a+b)
print("Sub:",a-b)
print("Mul:",a*b)
print("Div:",a/b)
print("Mod:",a%b)
a=float(input("Enter a value:"))
b=float(input("Enter b value:"))
print("add:",a+b)
print("Sub:",a-b)
print("Mul:",a*b)
print("Div:",a/b)
print("Mod:",a%b)
```

Output:

```
Enter a value:10
Enter b value:22
add: 32
Sub: -12
Mul: 220
Div: 0.45454545454545453
Mod: 10
Enter a value:1.2
Enter b value:3.5
add: 4.7
Sub: -2.3
Mul: 4.2
Div: 0.34285714285714286
Mod: 1.2
```

Machine Learning Lab III B.Tech II Sem CSE-B 20255A0507 2021-2022 2. Write a program to check whether the string is symmetrical/palindrome. Program:

```
st=input("Enter a string:")
def palindrome(st):
  m=(len(st)-1)//2
  1=0
  h=len(st)-1
  flag=0
  while(l<=m):</pre>
    if st[1]==st[h]:
       1=1+1
       h=h-1
    else:
       flag=1
       break;
  if flag==0:
    print(st,"is palindrome")
  else:
    print(st,"is not palindrome")
def symmetrical(st):
  n=len(st)
  flag=0
  if n%2==0:
    mid=n//2
  else:
    mid=n//2+1
  1=0
  m=mid
  while(l<mid and m<n):
    if(st[1]==st[m]):
      1=1+1
       m=m+1
    else:
       flag=1
       break
  if flag==0:
    print(st,"is symmetrical")
  else:
    print(st,"is not symmetrical")
palindrome(st)
symmetrical(st)
Output:
         Enter a string:khokho
         khokho is not palindrome
          khokho is symmetrical
```

3. Write a program to print the multiplication table of a given number.

Program:

```
n=int(input("enter a number:"))
for i in range(1,11):
    print(n,"*",i,"=",n*i)
```

Output:

```
enter a number:5

5 * 1 = 5

5 * 2 = 10

5 * 3 = 15

5 * 4 = 20

5 * 5 = 25

5 * 6 = 30

5 * 7 = 35

5 * 8 = 40

5 * 9 = 45

5 * 10 = 50
```

4. Write a program to print largest of three numbers using conditional statements.

Program:

```
a=int(input("Enter a value:"))
b=int(input("Enter b value:"))
c=int(input("Enter c value:"))
if a>=b and a>=c:
    print(a,"is largest")
elif b>=a and b>=c:
    print(b,"is largest")
else:
    print(c,"is largest")
```

Output:

Enter a value:12 Enter b value:13 Enter c value:45 45 is largest 5. Write a python program to count the occurrences of each word in a text file.

Program:

```
fd=open("demo.txt",'r')
d=fd.read()
count=dict()
words=d.split()
for i in words:
    if i in count:
        count[i]+=1
    else:
        count[i]=1
print(count)
```

Output:

{'python': 1, 'is': 2, 'a': 1, 'simple': 2, 'programming': 1, 'language.': 1, 'it': 1, 'very': 1, 'and': 1, 'easier': 1, 'to': 1, 'understand.': 1}

6. Write a program to perform the different basic operations on a tuple.

Program:

```
t1=('akhila','aakanksha','sai')
t2=(10,22,3,1)
print(len(t1))
print(max(t2))
print(min(t2))
t=t1+t2
print(t)
print(t1[-2])
print(t2[::-1])
print(list(t))
Output:
3
22
('akhila', 'aakanksha', 'sai', 10, 22, 3, 1)
aakanksha
(1, 3, 22, 10)
['akhila', 'aakanksha', 'sai', 10, 22, 3, 1]
```

7. Write a program to get the largest number from a list.

Program:

```
l=[22,56,23,45,87,10,108,46,67,82]
mx=l[1]
for i in range(0,len(1)-1):
    if l[i]>l[i+1]:
        mx=l[i]
print("Largest number from list is",mx)
```

Output:

Largest number from list is 108

8. Write a program to get the maximum and minimum value in a dictionary.

Program:

```
d={'akhila':92,'anusha':96,'pavani':85}
mx=max(d.keys(),key=(lambda k:d[k]))
mn=min(d.keys(),key=(lambda k:d[k]))
print("Maximum value:",d[mx])
print("Minimum value:",d[mn])

Output:

Maximum value: 96
Minimum value: 85
```

9. Explore python classes, packages related to Machine Learning (numpy, pandas, matplotlib)

Numpy

```
import numpy as np
a=np.array([5,10,22,15,13,60])
  array([ 5, 10, 22, 15, 13, 60])
print(a[0:3])
  [ 5 10 22]
print(a[:5])
  [ 5 10 22 15 13]
print(a[3:])
  [15 13 60]
print(a[::-1])
  [60 13 15 22 10 5]
print(np.sort(a))
 [ 5 10 13 15 22 60]
type(a)
  numpy.ndarray
b=np.array([[1,2,3],[4,5,6],[7,8,9]])
b
  array([[1, 2, 3],
[4, 5, 6],
[7, 8, 9]])
type(b)
  numpy.ndarray
print(a.dtype)
  int64
a.shape
  (6,)
b.shape
  (3, 3)
a.ndim
  1
```

```
b=np.array([[1,2,3],[4,5,6],[7,8,9]])
array([[1, 2, 3],
[4, 5, 6],
[7, 8, 9]])
type(b)
numpy.ndarray
print(a.dtype)
int64
a.shape
(6,)
b.shape
(3, 3)
a.ndim
b.ndim
2
c=np.array([[(1,2,3),(3,3,3)],
            [(7,8,9),(4,4,4)],
[(1,1,1),(3,3,3)]
c.shape
(3, 2, 3)
np.zeros((4,3))
array([[0., 0., 0.],
       [0., 0., 0.],
[0., 0., 0.],
[0., 0., 0.]])
np.zeros((4,3),dtype=np.int32)
d=np.arange(1,20,2)
```

```
c=np.array([[(1,2,3),(3,3,3)],
                [(7,8,9),(4,4,4)],
               [(1,1,1),(3,3,3)]
               ])
[ ] c.shape
    (3, 2, 3)
[ ] np.zeros((4,3))
    array([[0., 0., 0.],
           [0., 0., 0.],
           [0., 0., 0.],
           [0., 0., 0.]])
[ ] np.zeros((4,3),dtype=np.int32)
    array([[0, 0, 0],
           [0, 0, 0],
           [0, 0, 0],
           [0, 0, 0]], dtype=int32)
[ ] d=np.arange(1,20,2)
[ ] d
    array([ 1, 3, 5, 7, 9, 11, 13, 15, 17, 19])
[ ] e=np.full((4,3),10)
[ ] e
    array([[10, 10, 10],
           [10, 10, 10],
           [10, 10, 10],
           [10, 10, 10]])
[ ] np.linspace(1,5,10)
                 , 1.44444444, 1.88888889, 2.33333333, 2.77777778,
    array([1.
```

Pandas

```
[ ] import pandas as pd
Creating data frame
[ ] df=pd.DataFrame([[4,5,6],[3,4,5],[12,13,14]],index=[1,2,3],columns=["a","b","c"])
[ ] df
           a b c
       1 4 5 6
       2 3 4 5
      3 12 13 14
[ ] df = pd.DataFrame(
    {"a" : [4 ,5 ,6],
    "b" : [7, 8, 9],
    "c" : [18, 11, 12]},
    index = pd.MultIndex.from_tuples(
    [('d',1),('d',2),('e',2)],
    names=['n','v']))
[ ] df
       d 1 4 7 10
        2 5 8 11
       e 2 6 9 12
[ ] dfn = pd.DataFrame(
                        "Braund, Mr. Owen Harris",
"Allen, Mr. William Henry",
"Bonnell, Miss. Elizabeth",
                  ],
"Age": [22, 35, 58],
"Sex": ["male", "male", "female"],
[ ] dfn
                            Name Age Sex
       0 Braund, Mr. Owen Harris 22 male
       1 Allen, Mr. William Henry 35 male
       2 Bonnell, Miss. Elizabeth 58 female
```

printing one column of dataframe

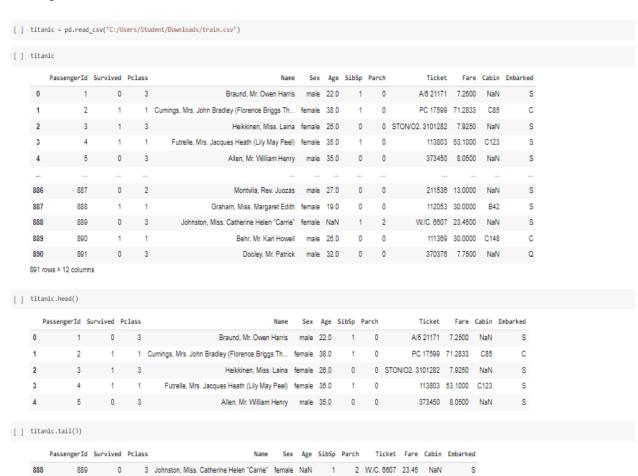
```
[ ] dfn["Age"]

0 22
1 35
2 58
Name: Age, dtype: int64
```

Series is one dimensional data

```
[ ] ages = pd.Series([22, 35, 58], name="Age")
[ ] ages
    0 22
1 35
2 58
    Name: Age, dtype: int64
[ ] dfn["Age"].max()
[ ] ages.max()
    58
[ ] dfn.describe()
     count 3.000000
     mean 38.333333
      std 18.230012
      min 22.000000
     25% 28.500000
      50% 35.000000
     75% 46.500000
      max 58.000000
```

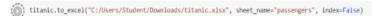
Reading csv file into dataframe



Dooley, Mr. Patrick male 32.0 0 0 370376 7.75 NaN Q

writng dataframe to excel sheet

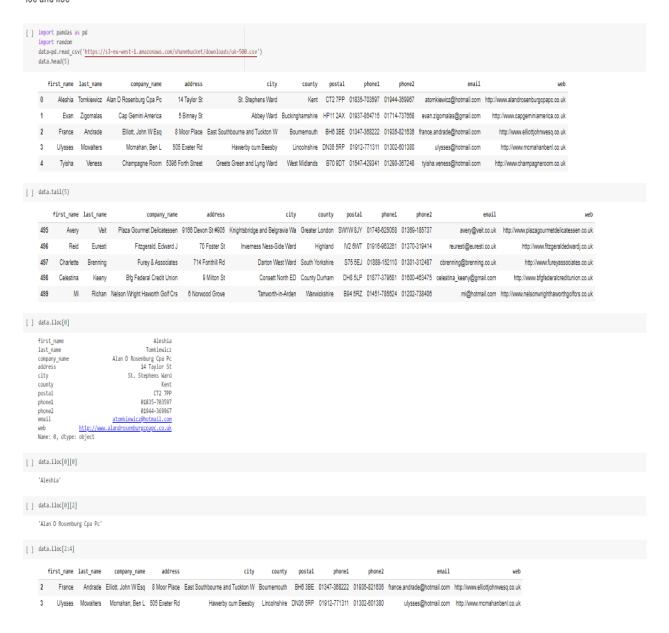
890 891 0 3



(titanic.info() #Gives summary of dataframe

Iloc and loc

loc and iloc

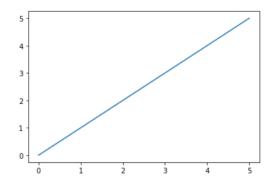


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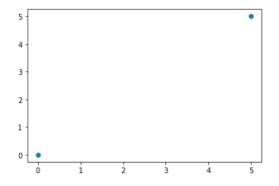


Matplotlib

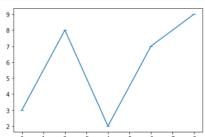
```
import matplotlib.pyplot as plt
import numpy as np
x=np.array([0,5])
y=np.array([0,5])
plt.plot(x,y)
plt.show()
```



```
x=np.array([0,5])
y=np.array([0,5])
plt.plot(x,y,'o')
plt.show()
```

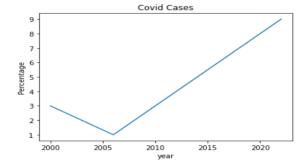


```
x=np.array([0,2,4,6,8])
y=np.array([3,8,2,7,9])
plt.plot(x,y,marker="_")
plt.show()
```



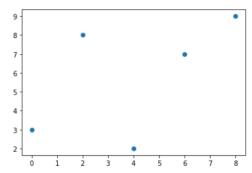
```
y=np.array([3,8,2,7,9])
plt.plot(y,marker="_")
plt.show()
   5
               1.5
                   2.0
                      2.5
y=np.array([3,8,2,7,9])
plt.plot(y, 'o:g')
plt.show()
    8
    6
    5
    3
                    2.0
             1.0
                1.5
                        2.5
                           3.0
                               3.5
y=np.array([3,8,2,7,9])
plt.plot(y,marker="*",ms=10,mec='g',mfc='y',color='y')
plt.show()
               1.5
                      2.5 3.0
            1.0
                  2.0
x=np.array([2000,2006,2008,2022])
y=np.array([3,1,2,9])
plt.plot(x,y)
```

```
plt.title("Covid Cases")
plt.xlabel("year")
plt.ylabel("Percentage")
plt.show()
```



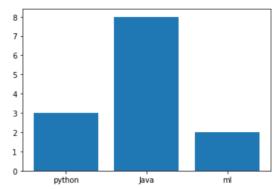
Scatter

```
x=np.array([0,2,4,6,8])
y=np.array([3,8,2,7,9])
plt.scatter(x,y)
plt.show()
```

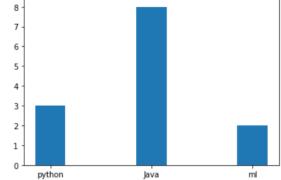


Bar graph

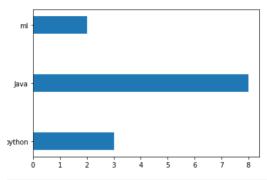
```
x=np.array(['python','Java','ml'])
y=np.array([3,8,2])
plt.bar(x,y)
plt.show()
```



```
x=np.array(['python','Java','ml'])
y=np.array([3,8,2])
plt.bar(x,y,width=0.3)
plt.show()
```

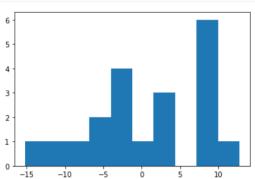


```
x=np.array(['python','Java','ml'])
y=np.array([3,8,2])
plt.barh(x,y,height=0.3)
plt.show()
```



Histogram:

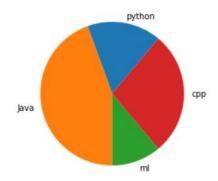
```
y=np.random.normal(3,8,20)
plt.hist(y)
plt.show()
```



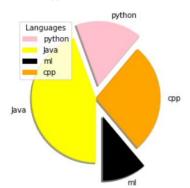
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Pie chart

```
x=np.array([3,8,2,5])
label=['python','Java','ml','cpp']
plt.pie(x,labels=label,startangle=50)
plt.show()
```



```
x=np.array([3,8,2,5])
label=['python','Java','ml','cpp']
exp=[0.2,0,0.3,0]
color=['pink','yellow','k','orange']
plt.pie(x,labels=label,startangle=50,explode=exp,shadow=True,colors=color)
plt.legend(title="Languages")
plt.show()
```



10. Implement FIND-S algorithm for finding the most specific hypothesis based on a given set of training data sample.

```
Program:
```

```
import pandas as pd
import random
data=pd.read csv("enjoySport.csv")
print(data)
#making an array of all the attributes
d = np.array(data)[:,:-1]
print("The attributes are:\n",d)
#segragating the target that has positive and negative examples
target = np.array(data)[:,-1]
print("The target is: ",target)
#training function to implement find-s algorithm
def train(c,t):
    for i, val in enumerate(t):
         if val == "Y":
             specific hypothesis = c[i].copy()
             break
    for i, val in enumerate(c):
         if t[i] == "Y":
              for x in range(len(specific hypothesis)):
                  if val[x] != specific hypothesis[x]:
                       specific hypothesis[x] = '?'
                  else:
                      pass
    return specific hypothesis
Output:
          Sky AirTemp Humidity
                               Wind Water Forecast EnjoySport
     0 Sunny
                Warm Normal Strong Warm
                                                Same
                                                            Yes
     1 Sunny
                 Warm
                        High Strong Warm
                                                Same
                                                            Yes
     2 Rainy
                 Cold
                         High Strong Warm
                                              Change
                                                            No
     3 Sunny
                 Warm
                         High Strong Cool
                                              Change
                                                            Yes
     The attributes are:
      [['Sunny' 'Warm' 'Normal' 'Strong' 'Warm' 'Same']
      ['Sunny' 'Warm' 'High' 'Strong' 'Warm' 'Same']
['Rainy' 'Cold' 'High' 'Strong' 'Warm' 'Change']
['Sunny' 'Warm' 'High' 'Strong' 'Cool' 'Change']]
     The target is: ['Yes' 'Yes' 'No' 'Yes']
```

11. For a given set of training data examples, implement Candidate-Elimination algorithm to output a description of the set of all consistent hypothesis.

```
Program:
import random
import csv
def g_0(n):
  return ("?",)*n
def s_0(n):
  return ('0',)*n
def more_general(h1, h2):
  more_general_parts = []
  for x, y in zip(h1, h2):
     mg = x == "?" \text{ or } (x != "0" \text{ and } (x == y \text{ or } y == "0"))
     more_general_parts.append(mg)
  return all(more general parts)
def fulfills(example, hypothesis):
  ### the implementation is the same as for hypotheses:
  return more_general(hypothesis, example)
def min_generalizations(h, x):
  h_new = list(h)
  for i in range(len(h)):
     if not fulfills(x[i:i+1], h[i:i+1]):
       h_{new[i]} = '?' \text{ if } h[i] != '0' \text{ else } x[i]
  return [tuple(h_new)]
def min_specializations(h, domains, x):
  results = []
  for i in range(len(h)):
     if h[i] == "?":
       for val in domains[i]:
          if x[i] != val:
            h_new = h[:i] + (val_n) + h[i+1:]
            results.append(h_new)
     elif h[i] != "0":
       h_new = h[:i] + ('0',) + h[i+1:]
       results.append(h_new)
  return results
with open('enjoySport.csv') as csvFile:
     examples = [tuple(line) for line in csv.reader(csvFile)]
examples
def get_domains(examples):
  d = [set() \text{ for i in examples}[0]]
  for x in examples:
     for i, xi in enumerate(x):
        d[i].add(xi)
  return [list(sorted(x)) for x in d]
get_domains(examples)
def candidate elimination(examples):
  domains = get_domains(examples)[:-1]
```

 $G = set([g_0(len(domains))])$

```
S = set([s_0(len(domains))])
  print("\n G[\{0\}]:".format(i),G)
  print("\n S[\{0\}]:".format(i),S)
  for xcx in examples:
     i=i+1
     x, cx = xcx[:-1], xcx[-1] # Splitting data into attributes and decisions
     if cx=='Y': # x is positive example
        G = \{g \text{ for } g \text{ in } G \text{ if fulfills}(x, g)\}
        S = generalize\_S(x, G, S)
     else: # x is negative example
        S = \{s \text{ for } s \text{ in } S \text{ if not fulfills}(x, s)\}
        G = \text{specialize } G(x, \text{domains, } G, S)
     print("\n G[\{0\}]:".format(i),G)
     print("\n S[\{0\}]:".format(i),S)
  return
def generalize S(x, G, S):
  S_prev = list(S)
  for s in S_prev:
     if s not in S:
        continue
     if not fulfills(x, s):
        S.remove(s)
        Splus = min\_generalizations(s, x)
        ## keep only generalizations that have a counterpart in G
        S.update([h for h in Splus if any([more_general(g,h)
                                for g in G])])
        ## remove hypotheses less specific than any other in S
        S.difference_update([h for h in S if
                       any([more_general(h, h1)
                          for h1 in S if h != h1])])
  return S
def specialize_G(x, domains, G, S):
  G_prev = list(G)
  for g in G prev:
     if g not in G:
        continue
     if fulfills(x, g):
        G.remove(g)
        Gminus = min_specializations(g, domains, x)
        ## keep only specializations that have a conuterpart in S
        G.update([h for h in Gminus if any([more_general(h, s)
                                 for s in S])])
        ## remove hypotheses less general than any other in G
        G.difference_update([h for h in G if
                       any([more_general(g1, h)
                          for g1 in G if h != g1])])
  return G
candidate_elimination(examples)
Output:
[('Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same', 'Y'),
  ('Sunny', 'Warm', 'High', 'Strong', 'Warm', 'Same', 'Y'),
  ('Rainy', 'Cold', 'High', 'Strong', 'Warm', 'Change', 'N'),
  ('Sunny', 'Warm', 'High', 'Strong', 'Cool', 'Change', 'Y')]
```

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2021-2022

1.Implement Decision tree for classification of any given data

Program:

```
set.import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
#%matplotlib inline#for encoding
from sklearn.preprocessing import LabelEncoder#for train test splitting
from sklearn.model selection import train test split#for decision tree object
from sklearn.tree import DecisionTreeClassifier#for checking testing results
from sklearn.metrics import classification report, confusion matrix#for visualiz
ing tree s
from sklearn.tree import plot tree
#reading the data
df = sns.load dataset('iris')
df.head()
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa

```
#getting information of dataset
df.info()
<class 'pandas.core.frame.DataFrame'>
```

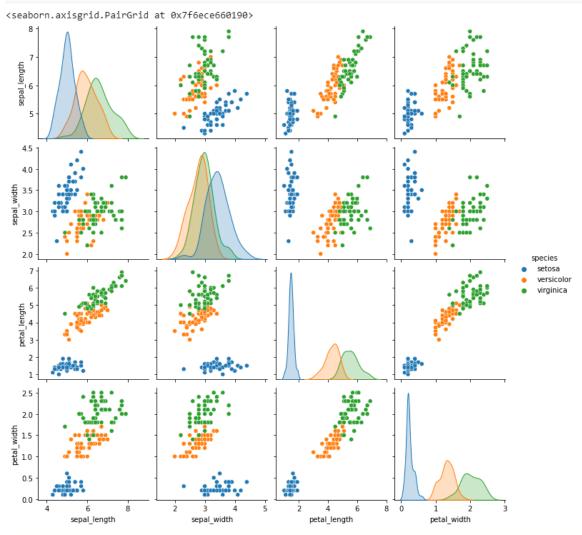
```
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
             Non-Null Count Dtype
# Column
               -----
--- -----
0 sepal_length 150 non-null float64
1 sepal_width 150 non-null float64
2 petal_length 150 non-null float64
3 petal width 150 non-null float64
4 species
               150 non-null
                             object
dtypes: float64(4), object(1)
```

```
df.shape
(150, 5)
df.isnull().any()
```

memory usage: 6.0+ KB

sepal_length False sepal_width False petal_length False petal_width False species False dtype: bool

let's plot pair plot to visualise the attributes all at once
sns.pairplot(data=df, hue = 'species')



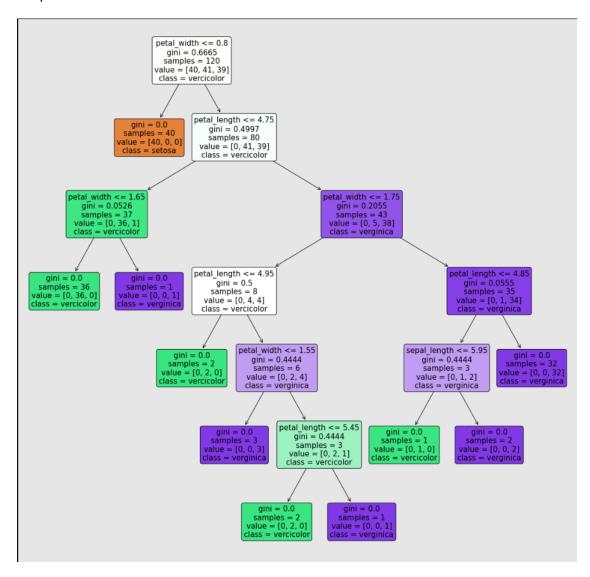
correlation matrix
sns.heatmap(df.corr())

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```
<matplotlib.axes._subplots.AxesSubplot at 0x7f6ed5505e90>
                               -1.0
 sepal length -
                                - 0.8
                                0.6
 sepal_width
                                - 04
                                0.2
 petal_length
                                0.0
                                -0.2
 petal width
                                -0.4
       sepal_length sepal_width petal_length petal_width
target=df['species']
df1=df.copy()
df1=df1.drop('species',axis=1) #removing whole species column
X = df1
target=df['species']
target
       setosa
1
       setosa
2
       setosa
3
       setosa
4
       setosa
145
     virginica
146
     virginica
147
     virginica
148
     virginica
149
     virginica
Name: species, Length: 150, dtype: object
#label encoding
le = LabelEncoder()
target = le.fit transform(target)
target
  1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
       y=target
# Splitting the data - 80:20 ratio
X train, X test, y train, y test = train test split(X , y, test size = 0.2, rand
om state = 42)
print("Training split input- ", X train.shape)
print("Testing split input- ", X test.shape)
Training split input- (120, 4)
  Testing split input- (30, 4)
```

```
# Defining the decision tree algorithm
dtree=DecisionTreeClassifier()
dtree.fit(X train,y train)
print('Decision Tree Classifier Created')
# Predicting the values of test data
y pred = dtree.predict(X test)
print("Classification report - \n", classification report(y test, y pred))
   Classification report -
               precision
                          recall f1-score
                                         support
            0
                  1.00
                          1.00
                                  1.00
                                            10
            1
                  1.00
                          1.00
                                  1.00
                                             9
                  1.00
                          1.00
                                  1.00
                                            11
                                  1.00
                                            30
      accuracy
                  1.00
                          1.00
                                  1.00
                                            30
     macro avg
   weighted avg
                  1.00
                          1.00
                                  1.00
                                            30
cm = confusion_matrix(y_test, y_pred)
plt.figure(figsize=(5,5))
sns.heatmap(data=cm,linewidths=.5, annot=True,square = True, cmap = 'Blues')
plt.ylabel('Actual label')
plt.xlabel('Predicted label')
all_sample_title = 'Accuracy Score: {0}'.format(dtree.score(X_test, y_test))
plt.title(all sample title, size = 15)
   Text(0.5, 1.0, 'Accuracy Score: 1.0')
           Accuracy Score: 1.0
                          0
          10
                          0
                  0
           0
                                 - 0
               Predicted label
# Visualising the graph without the use of graph
vizplt.figure(figsize = (20,20))
dec_tree = plot_tree(decision_tree=dtree, feature_names = dfl.columns,
                       class names =["setosa", "vercicolor", "verginica"] , filled
 = True , precision = 4, rounded = True)
```

Output:



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