

Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a set of training data samples. Read the following data from a .csv file.

```

from pandas import DataFrame
data = DataFrame.from_csv('c:/Users/hp/Desktop/4MT17CS
                           043 Harshitha / lab1.csv')
columnLength = data.shape[1]
print(data)
h = ['?'] * (columnLength - 1)
hp = []
hn = []
for trainingExample in data.values:
    if trainingExample[-1] == 'no':
        hp.append(list(trainingExample))
    else:
        hn.append(list(trainingExample))
for i in range(len(hp)):
    for j in range(columnLength - 1):
        if (h[j] == '?'):
            h[j] = hp[i][j]
        if (h[j] != hp[i][j]):
            h[j] = '?'
else:
    h[j] = hp[i][j]

```

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Expt. No. \_\_\_\_\_

print ("In The positive hypothesis are : ", hp)

print ("In The negative hypothesis are : ", hn)

print ("In The maximally specific hypothesis is : ", L)

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Sl. No	Sky	A <sup>o</sup> temp	Humidity	wind	water	Forecast	Enjoy
1	Sunny	Warm	normal	Strong	Warm	Same	Yes
2	Sunny	Warm	high	Strong	Warm	Same	Yes
3	Rainy	Cold	high	Strong	Warm	Change	No
4	Bunny	Warm	high	Strong	Cool	Change	Yes

The positive hypothesis are :

- [ ['sunny', 'warm', 'normal', 'strong', 'warm', 'same', 'yes'] ]
- [ ['sunny', 'warm', 'high', 'strong', 'warm', 'same', 'yes'] ]
- [ ['sunny', 'warm', 'high', 'strong', 'cool', 'change', 'yes'] ]

The negative hypothesis are :

- [ ['rainy', 'cold', 'high', 'strong', 'warm', 'change', 'no'] ]

The maximally specific hypothesis is :

- [ ['sunny', 'warm', '?', 'strong', '?', '?'] ]

For a given set of training data examples stored in a .csv file, implement and demonstrate the candidate elimination algorithm to output a description of the set of all hypothesis consistent with the training examples.

```
import csv
```

```
with open ('c:/User/hp/Desktop/4MT17CS043/Harshitha/  
lab2.csv') as f:
```

```
    CSV-file = csv.reader (f)  
    data = list (CSV-file)
```

```
print (data)
```

```
s = data [1] [: - 1]
```

```
print (s)
```

```
g = [[? for i in range (len (s))] for j in range  
(len (s))]
```

```
for i in data :
```

```
    if i [-1] == "Yes":
```

```
        for j in range (len (s)):
```

```
            if i [j] != s [j]:
```

```
                s [j] = ?
```

```
                g [i] [j] = ?
```

```
    elif i [-1] == "No":
```

```
        for j in range (len (s)):
```

```
            if i [j] != s [j]:
```

```
                g [i] [j] = s [j]
```

```
            else :
```

```
                g [i] [j] = ?
```

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print ("steps of candidate elimination algorithm", data[i],  
(i)+1)

print (s)

print (g)

gh = [ ]

for i in g:

    for j in i:

        if j != '?':

            gh.append (i)

        break

print ("In Final Specific hypothesis : ", s)

print ("In Final general hypothesis : ", gh)

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[['sunny', 'warm', 'normal', 'strong', 'warm', 'same', 'yes'],  
[['sunny', 'warm', 'high', 'strong', 'warm', 'same', 'yes'],  
[['rainy', 'cold', 'high', 'strong', 'warm', 'change', 'no'],  
[['sunny', 'warm', 'high', 'strong', 'cool', 'change', 'yes']]  
[['sunny', 'warm', 'high', 'strong', 'warm', 'same']]

Steps of candidate elimination algorithm 1:

('sunny', 'warm', '?', 'strong', 'warm', 'same')  
[('?', '?', '?', '?', '?', '?'), ('?', '?', '?', '?', '?', '?'),  
[('?', '?', '?', '?', '?', '?'), ('?', '?', '?', '?', '?', '?'),  
[('?', '?', '?', '?', '?', '?'), ('?', '?', '?', '?', '?', '?')]])

Steps of candidate elimination algorithm 2:

('sunny', 'warm', '?', 'strong', 'warm', 'same')  
[('?', '?', '?', '?', '?', '?'), ('?', '?', '?', '?', '?', '?'),  
[('?', '?', '?', '?', '?', '?'), ('?', '?', '?', '?', '?', '?'),  
[('?', '?', '?', '?', '?', '?'), ('?', '?', '?', '?', '?', '?')]])

Steps of candidate elimination algorithm 3:

{'sunny', 'warm', '?', 'strong', 'warm', 'same']}

[{'sunny', '?', '?', '?', '?', '?'], {'?', 'warm', '?', '?', '?', '?']}

[{'?', '?', '?', '?', '?', '?'], {'?', '?', '?', '?', '?', '?']}

[{'?', '?', '?', '?', '?', '?'], {'?', '?', '?', '?', '?', 'same']}

Steps of candidate elimination algorithm 4:

{'sunny', 'warm', '?', 'strong', '?', '?')}

[{'sunny', '?', '?', '?', '?', '?'], {'?', 'warm', '?', '?', '?', '?']}

[{'?', '?', '?', '?', '?', '?'], {'?', '?', '?', '?', '?', '?']}

[{'?', '?', '?', '?', '?', '?'], {'?', '?', '?', '?', '?', '?'}]

Final general hypothesis:

[{'sunny', '?', '?', '?', '?', '?'], {'?', 'warm', '?', '?', '?', '?'}]

Write a program to demonstrate the working of the decision tree based ID3 algorithm. Use an appropriate dataset for building the decision tree and apply this knowledge to classify a new sample.

```
import pandas as pd
from pandas import DataFrame
df_tennis = pd.read_csv('C:/Users/hp/Desktop/4MT17CS043_Harshitha/play tennis.csv')
attribute_names = list(df_tennis.columns)
attribute_names.remove('Play Tennis')
print(attribute_names)
```

```
def entropy_of_list(lst):
    from collections import Counter
    count = Counter(x for x in lst)
    num_instances = len(lst) * 1
    probs = [x / num_instances for x in count.values()]
    return entropy(probs)
```

```
def entropy(probs):
    import math
    return sum([-prob * math.log(prob, 2) for prob in probs])
```

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total\_entropy = entropy\_of\_list (df['tennis']('Play Tennis'))

def information\_gain (df, split\_attribute\_name, target\_attribute\_name, trace=0):

df\_split = df.groupby(split\_attribute\_name)

nobs = len(df.index) + 1

df\_agg\_ent = df\_split.agg([target\_attribute\_name:])

 (entropy\_of\_list, lambda x: len(x)/nobs)]))

df\_agg\_ent.columns = ['Entropy', 'proportion']

new\_entropy = sum(df\_agg\_ent['Entropy'] \* df\_agg\_ent['proportion'])

old\_entropy = entropy\_of\_list (df[target\_attribute\_name])

print(split\_attribute\_name, 'IG:', old\_entropy - new\_entropy)

return old\_entropy - new\_entropy

def id3 (df, target\_attribute\_name, attribute\_names, default\_class = None):

from collections import Counter

count = Counter (x for x in df[target\_attribute\_name])

if len(count) == 1:

return next(iter(count))

elif df.empty or (not attribute\_names):

return default\_class

else :

default\_class = max(count.keys())

gain = [

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information-gain(df, attr, target-attribute-name) for  
attr in attribute-names)

index-of-max = gain\_index(max(gain))

best\_attr = attribute\_names[index-of-max]

tree = {best\_attr: {}}

remaining\_attribute\_names = [i for i in attribute\_names  
if i != best\_attr]

for attr\_val, data\_subset in df.groupby(best\_attr):

subtree = id3(data\_subset, target-attribute-name,

remaining\_attribute\_names, default\_class)

tree[best\_attr][attr\_val] = subtree

return tree.

from pprint import pprint

tree = id3(df\_tennis, 'PlayTennis', attribute\_names)

print("The resultant Decision tree is :")

pprint(tree)

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[ 'outlook', 'Temperature', 'Humidity', 'wind' ]

Outlook IG: 0.2467498197744391

Temperature IG: 0.0299222565658954647

Humidity IG: 0.15183550136234136

Wind IG: 0.04818703040826927

Temperature IG : 0.01997309402197489

Humidity IG : 0.01997309402197489

Wind IG : 0.9709505944546686

Temperature IG: 0.5709505944546686

Humidity IG: 0.9709505944546686

Wind IG: 0.01997309402197489

The result decision tree is :

```
{'outlook': { 'overcast': 'yes',
    'Rain': { 'wind': { 'strong': 'no', 'weak': 'yes' },
        'sunny': { 'Humidity': { 'high': 'no',
            'Normal': 'yes' } } } }}
```

Build an artificial neural network by implementing the backpropagation algorithm and test the same using appropriate dataset.

```
from math import exp
from random import seed
from random import random
```

```
def initialize_networks (n_inputs, n_hidden, n_outputs):
    network = list()
    hidden_layer = [{`weights`:[random() for i in range
        (n_inputs+1)]} for i in range (n_hidden)]
    network.append (hidden_layer)
    output_layer = [{`weights`:[random() for i in range
        (n_hidden+1)]} for i in range (n_outputs)]
    network.append (output_layer)
    return network.
```

```
def activate (weights, inputs):
    activation = weights [-1]
    for i in range (len (weights) - 1):
        activation += weights [i] * inputs [i]
    return activation
```

```
def transfer (activation):
    return 1.0 / (1.0 + exp (-activation))
```

```
def forward_propagate (network, row):
```

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```

inputs - row
for layer in network:
    new_inputs = []
    for neuron in layer:
        activation = activate(neuron['weights'].dot(inputs))
        neuron['output'] = transfer(activation)
        new_inputs.append(neuron['output'])
    inputs = new_inputs
return inputs

def transfer_derivative(output):
    return output * (1.0 - output)

def backward_propagate_error(network, expected):
    for i in reversed(range(len(network))):
        layer = network[i]
        errors = list()
        if i == len(network) - 1:
            for j in range(len(layer)):
                error = 0.0
                for neuron in network[i+1]:
                    error += (neuron['weights'][j] * neuron['delta'])
                errors.append(error)
        else:
            for j in range(len(layer)):
                neuron = layer[j]
                errors.append(expected[j] - neuron['output'])
    return errors

```

```

for j in range (len (layer)):
    neuron = layer [j]
    neuron ['delta'] = errors [j] * transfer - derivative
        (neuron ['output'])

def update - weights (network , row , l - rate):
    for i in range (len (network)):
        inputs = row [: - 1]
        if i != 0:
            inputs = [neuron ['output'] for neuron in network
                      [i - 1]]
        for neuron in network [i]:
            for j in range (len (inputs)):
                neuron ['weights'] [j] += l - rate * neuron
                    ['delta'] * inputs [j]
                neuron ['weights'] [- 1] += l - rate * neuron
                    ['delta']

def train - network (network , train , l - rate , n - epoch , n - output):
    for epoch in range (n - epoch):
        sum - error = 0
        for row in train:
            outputs = forward - propagate (network , row)
            expected [row (- 1)] = 1
            sum - error += sum ((expected [i] - output [i])
                                ** 2 for i in range (len (expected)))
            backward propagate error (network , expected)
            update - weights (network , row , l - rate)

```

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```
print ('>epoch = %.d, l-rate = %.3f, error = %.3f' % (epoch,
    l-rate, sum_error))
```

seed(1)

dataset = [

[2.7810836, 2.550537003, 0],

[1.465489372, 2.362125076, 0],

[3.396561688, 4.400293529, 0],

[1.38807019, 1.850220317, 0],

[3.06407232, 3.005305973, 0],

[7.6127531214, 2.759262235, 1],

[5.332441248, 2.088626775, 1],

[6.922596716, 1.77106367, 1],

[8.675418651, -0.242088655, 1],

[7.673756466, 3.508563011, 1]]

n\_inputs = len(dataset[0]) - 1

n\_outputs = len(set([row[-1] for row in dataset]))

network = initialize\_networks(n\_inputs, 2, n\_outputs)

train\_network(network, dataset, 0.5, 20, n\_outputs)

for layer in network:

print(layer)

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> epoch = 0 , lrate = 0.500 , error = 6.350  
> epoch = 1 , lrate = 0.500 , error = 5.531  
> epoch = 2 , lrate = 0.500 , error = 5.221  
> epoch = 3 , lrate = 0.500 , error = 4.951  
> epoch = 4 , lrate = 0.500 , error = 4.519  
> epoch = 5 , lrate = 0.500 , error = 4.173  
> epoch = 6 , lrate = 0.500 , error = 3.835  
> epoch = 7 , lrate = 0.500 , error = 3.501  
> epoch = 8 , lrate = 0.500 , error = 3.192  
> epoch = 9 , lrate = 0.500 , error = 2.898  
> epoch = 10 , lrate = 0.500 , error = 2.628  
> epoch = 11 , lrate = 0.500 , error = 2.377  
> epoch = 12 , lrate = 0.500 , error = 2.153  
> epoch = 13 , lrate = 0.500 , error = 1.953  
> epoch = 14 , lrate = 0.500 , error = 1.774  
> epoch = 15 , lrate = 0.500 , error = 1.614  
> epoch = 16 , lrate = 0.500 , error = 1.472  
> epoch = 17 , lrate = 0.500 , error = 1.346  
> epoch = 18 , lrate = 0.500 , error = 1.233  
> epoch = 19 , lrate = 0.500 , error = 1.182

{'weights': [-1.4688375095432327, 1.85088732543951]

, 1.0858178629550297], 'output': 0.029980305604012

'delta': -0.005956604162323625}, ;

{'weights': [0.37711098142462157, -0.0625909894552

987, 0.2765123702642716], 'output': 0.9456290002113 'delta': 0.0026274652850563837}]

{'weights': [2.515394649397849, -0.3391929502445985,

-0.9671565426390275], 'output': 0.2344679420957587, 'delta': -0.04270059278364587},

{'weights': [-2.5584149848484263, 1.0036422106209202, 0.42383086467582715], 'output': 0.779053520243836, 'delta': 0.038031325964373543}]

Write a program to implement the naive Bayesian classifier for a sample training dataset stored as a .csv file. Compute the accuracy of the classifier, considering few test data sets.

```

import csv, random, math
import statistics as st
def loadcsv(filename):
    lines = csv.reader(open(filename, "r"))
    dataset = list(lines)
    for i in range(len(dataset)):
        dataset[i] = [float(x) for x in dataset[i]]
    return dataset

def splitDataset(dataset, splitRatio):
    testSize = int(len(dataset) * splitRatio)
    trainset = list(dataset)
    testset = []
    while len(testset) < testSize:
        index = random.randrange(len(trainset))
        testset.append(trainset.pop(index))
    return (trainset, testset)

def separateByClass(dataset):
    separated = {}
    for i in range(len(dataset)):
        x = dataset[i]
        if (x[-1] not in separated):
            separated[x[-1]] = []
        separated[x[-1]].append(x)
    return separated

```

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```

separated[x[-1]] = []
separated[x[-1]].append(x)
return separated

```

```

def Compute_mean_std(dataset):
    mean_std = [st.mean(attribute), st.stdev(attribute)]
    for attribute in zip(*dataset):
        del mean_std[-1]
    return mean_std

```

```

def Summarize_By_Class(dataset):
    separated = separateByClass(dataset)
    summary = {}
    for classValue, instances in separated.items():
        summary[classValue] = compute_mean_std(
            instances)
    return summary

```

```

def estimateProbability(x, mean, stdDev):
    exponent = math.exp(-(math.pow(x - mean, 2) /
                           (2 * math.pow(stdDev, 2))))
    return (1 / (math.sqrt(2 * math.pi) * stdDev)) *
           exponent.

```

```

def calculateClassProbabilities(summaries, testVector):
    p = {}
    for classValue, classSummaries in summaries.items():

```

```
P [classValue] = 1
for i in range (len(classSummaries)):
    mean, stdDev = classSummaries[i]
    x = testVector[i]
    P [classValue] = estimateProbability (x, mean,
                                           stdDev)
return p
```

```
def predict (summaries, testVector):
    allP = calculateClassProbabilities (summaries, testVector)
    bestLabel, bestProb = None, -1
    for (lbl, p) in allP.items():
        if bestLabel is None or p > bestProb:
            bestProb = p
            bestLabel = lbl
    return bestLabel
```

```
def performClassification (summaries, testSet):
    predictions = []
    for i in range (len(testSet)):
        result = predict (summaries, testSet[i])
        predictions.append (result)
    return predictions
```

```
def getAccuracy (testSet, predictions):
    correct = 0
    for i in range (len(testSet)):
        if testSet[i][-1] == predictions[i]:
            correct += 1
    return correct / len(testSet)
```

```
correct += 1
return (correct / float(len(testset))) * 100.0

dataset = loadcsv('c:/users/hp/Desktop/4MT17CS043  
Harskrita/diabetes.csv'):
print ('Pima Indian Diabetes Dataset loaded ...')
print ('Total instances available: ', len(dataset))
print ('Total attributes present: ', len(dataset[0])-1)
print ('First Five instances of dataset: ')
for i in range(5):
    print (i+1, '::', dataset[i])
splitRatio = 0.2
trainingset, testset = splitDataset(dataset, splitratio)
print ('Dataset is split into training and testing set.')
print ("Training examples = {} \n Testing examples = {}".format(len(trainingset), len(testset)))
summaries = summariseByClass (trainingset)
predictions = performClassification (summaries, testset)
accuracy = getAccuracy (testset, predictions)
print ("The Accuracy of the Naive Bayesian classifier is : ", accuracy)
```

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Pima Indian Diabetes Dataset loaded.  
Total instances available : 768  
Total attributes present : 8

First five instances of dataset :

1: [6.0, 148.0, 72.0, 35.0, 0.0, 33.6, 0.627, 50.0,  
1.0]

2: [1.0, 85.0, 66.0, 290, 0.0, 26.6, 0.351, 31.0,  
0.0]

3: [8.0, 183.0, 64.0, 0.0, 0.0, 23.3, 0.672, 32.0,  
1.0]

4: [1.0, 89.0, 66.0, 23.0, 94.0, 28.1, 0.167, 21.0,  
0.0]

5: [0.0, 131.0, 40.0, 35.0, 168.0, 43.1, 2.288,  
33.0, 1.0]

Dataset is split into training and testing set

Training examples = 615

Testing examples = 153

Accuracy of the Naive Bayesian classifier is :  
75.16339869281046.

Assuming a set of documents that need to be classified, use the naive bayesian classifier model to perform this task. Built-in Java classes/API can be used to write the program. Calculate the accuracy, precision and recall for your dataset.

```
import pandas as pd
msg = pd.read_csv('C:/Users/HP/Desktop/4MT17CS043_Harshitha/lab6.csv') name='message',
'label')
print ("Total instances in the dataset : ", msg.shape[0])
msg['labelnum'] = msg.label.map({'pos': 1, 'neg': 0})
x = msg.message
y = msg.labelnum
print ("The message and its label of first 5
instances are listed below")
x5, y5 = x[0:5], msg.label[0:5]
for x, y in zip(x5, y5):
    print (x, ',', y)
```

```
from sklearn.model_selection import train_test_split
xtrain, xtest, ytrain, ytest = train_test_split(x, y)
print ("Dataset is split into training and testing
samples")
```

```
print ("Total training instances : ", xtrain.shape[0])
print ("Total testing instances : ", xtest.shape[0])
```

```
from sklearn.feature_extraction.text import CountVectorizer
```

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```

count_vect = CountVectorizer()
xtrain_dtm = count_vect.fit_transform(xtrain)
xtest_dtm = count_vect.transform(xtest)
print ("Total features extracted using Count Vectorizer : ", xtrain_dtm.shape[1])
print ("The Features for first 5 training instances are listed below")
df = pd.DataFrame(xtrain_dtm.toarray(), columns= count_vect.get_feature_names())
print (df[0:5])

```

```

from sklearn.naive_bayes import MultinomialNB
df = MultinomialNB().fit(xtrain_dtm, ytrain)
predicted = df.predict(xtest_dtm)
print ("Classification results of testing samples are given below")
for doc, p in zip(xtest, predicted):
    pred = 'pos' if p == 1 else 'neg'
    print ("%s -> %s" % (doc, pred))

```

```

from sklearn import metrics
print ("Accuracy metrics")
print ("Accuracy of the classifier is :",
      metrics.accuracy_score(ytest, predicted))
print ("Recall :",
      metrics.recall_score(ytest, predicted))
print ("Confusion matrix")
print (metrics.confusion_matrix(ytest, predicted))

```

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Total instances in the dataset: 18

The message and its label of first 5 instances are listed below:

I love this sandwich, pos

This is an amazing place, pos

I feel very good about these beers, pos

This is my best work, pos

What an awesome view, pos.

Dataset is split into Training & Testing Samples

Total training instances: 13

Total testing instances: 5

Total features extracted using countvectorizer: 46

Features for first 5 training instances are listed below.

about am an awesome beers best beers can ded do do

0	0	0	0	0	0	0	0	0	0	1	-0
1	0	0	0	0	0	0	0	0	0	0	0
2	0	0	0	0	0	0	0	0	1	1	0
3	0	0	1	1	0	0	0	0	0	0	0
4	0	0	0	0	0	0	0	0	0	0	0

Write a program to implement k-nearest neighbor algorithm to classify the iris dataset. Point both correct and wrong predictions. Java / Python ML library classes can be used for this problem.

```

from sklearn.datasets import load_iris
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification_report
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix
from sklearn.metrics import accuracy_score
iris_dataset = load_iris()

print ("In IRIS FEATURES) TARGET NAMES: \n", iris_
dataset, target_names)
for i in range (len(iris_dataset.target_names)):
    print ("In [{}]: [{}]\n".format (i, iris_dataset.
target_names[i]))
# print ("In Iris data: ", iris_dataset["data"])

x_train, x_test, y_train, y_test = train_test_split(iris_
dataset["data"], iris_dataset["target"]
random_state=0)

classifier = KNeighborsClassifier(n_neighbors=8, p=3,
metric="euclidean")

```

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```
classifier.fit(x_train, y_train)
y_pred = classifier.predict(x_test)
cm = confusion_matrix(y_test, y_pred)
print("confusion matrix is as follows (n,cm)")
print("Accuracy metrics")
print(classification_report(y_test, y_pred))
print("correct prediction", accuracy_score(y_test,
                                             y_pred))
print("wrong prediction", (1 - accuracy_score(y_test,
                                              y_pred)))
```

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## IRIS FEATURES | TARGET NAMES:

['setosa', 'versicolor', 'virginica']

- [0] : [setosa]
- [1] : [versicolor]
- [2] : [virginica]

kNeighbors Classifier (algorithm = 'auto', leaf\_size = 30, metric = 'euclidean' metric\_params = None, n\_jobs = None, n\_neighbors = 5, p = 3, weights = 'uniform')

Confusion matrix is as follows:

$$\begin{bmatrix} (13 & 0 & 0) \\ (0 & 15 & 1) \\ (0 & 0 & 9) \end{bmatrix}$$

Accuracy matrices

		prediction	recall	f1-score	support
	0	1.00	1.00	1.00	13
	1	1.00	0.94	0.97	6
	2	0.90	1.00	0.95	9
accuracy				0.97	38
macro avg		0.97	0.98	0.97	38
weighted avg		0.98	0.97	0.97	38

Correct prediction : 0.9736842105263158

wrong prediction : 0.02631578947368418