#### **Artificial Intelligence Lab Evaluation Assignment 2**

#### COE 9

**Q1.** Create a dataset (.csv) of students having following attributes: Written score (out of 100), Personal interview Score (out of 50), Group discussion score (out of 150), Qualified for IIM (for qualified students mention the Name of IIM; otherwise mention "Not qualified").

- Implement KNN to predict whether following students are qualified for an IIM or not?
   Mention IIM name in case of Qualified otherwise predicted result should display not qualified
  - 80% scored in Written test, 70% scored in Personal interview, and 70% scored in Group discussion.
  - 50% scored in Written test, 80% scored in Personal interview, and 40% scored in Group discussion.
- 2. Compare the accuracy of prediction of KNN with Bayesian learning model and show it using plots.
- 3. Run your program for different values of K and find its best value. Also observe the effect on train test ratio on the best value of K.

**CODE Cells with output:** 

### # Creating a dataset

```
In [31]:
# Kulpreet_q1_30/11/20
import random as r
# campus=['IIM-DL','IIM-GJ'] # possible IIM campuses
fp=open('data_1k.csv','w') # Open the file in writing mode
fp.write('written(100),pi(50),gd(150),qualified\n')
for i in range(10000):
 written=r.randint(0,100)
 pi=r.randint(0,50)
 gd=r.randint(0,150)
 tot=0.33*written+0.17*pi+0.50*gd # merit qualification factor
 if tot>90:
    college= r.randint(1,2) # r.choice(campus)
    college= 0 # 'not qualified'
 stu='%d,%d,%d,%d\n'%(written,pi,gd,college)
 fp.write(stu) # Writing to the file line by line
fp.close()
print ('Done! \n Open the file to view the dataset.')
Done!
 Open the file to view the dataset.
```

```
In [32]:
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
In [33]:
df = pd.read_csv('data_1k.csv')
df.head()
Out [33]:
```

	written(100)	pi(50)	gd(150)	qualified
0	14	27	96	0
1	49	18	97	0
2	96	10	71	0
3	45	47	61	0
4	86	18	130	2

```
In [34]:
x = df[['written(100)', 'pi(50)', 'gd(150)']]
y = df['qualified']
print(x.shape, y.shape)

(10000, 3) (10000,)
```

## # Scaling the Input Data

```
In [35]:
stu11= pd.Series(data={'written(100)': 50, 'pi(50)' : 40, 'gd(150)' : 60})
x = x.append(stu11, ignore_index=True)
stu22= pd.Series(data={'written(100)': 80, 'pi(50)' : 35, 'gd(150)' : 105})
x = x.append(stu22, ignore_index=True)

In [36]:
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
x = sc.fit_transform(x)
np.set_printoptions(suppress = True)
x
```

### # Splitting the training and testing data

```
In [38]:
from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split(x, y, test_size = 0.20, random_state = 0)
print(x_train.shape, x_test.shape)
print(x_test[0])

(8000, 3) (2000, 3)
[-1.7166161 -1.33470476 1.70089931]
```

### # Applying the KNN model for n=1 initially

```
from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n_neighbors=1)
knn.fit(x_train,y_train)
kpred = knn.predict(x_test)

In [40]:
from sklearn.metrics import classification_report,confusion_matrix
print('WITH K = 1 (initially)\n')
print('Confusion Matrix:')
print(confusion_matrix(y_test,kpred))
print('\nClassification_report(y_test,kpred))

WITH K = 1 (initially)
```

In [39]:

```
Confusion Matrix:
      5
[[1769
[ 4
       50
            481
    8
       50
           59]]
Classification Report:
            precision
                      recall f1-score support
                0.99
                     0.99
                                 0.99
                                          1781
         1
                0.48
                        0.49
                                 0.48
                                          102
                0.52
                        0.50
                                 0.51
                                           117
   accuracy
                                 0.94
                                          2000
```

0.66

0.94

macro avg

weighted avg

## # Checking for the best value of n in range(1,40) for KNN

0.66

0.94

2000

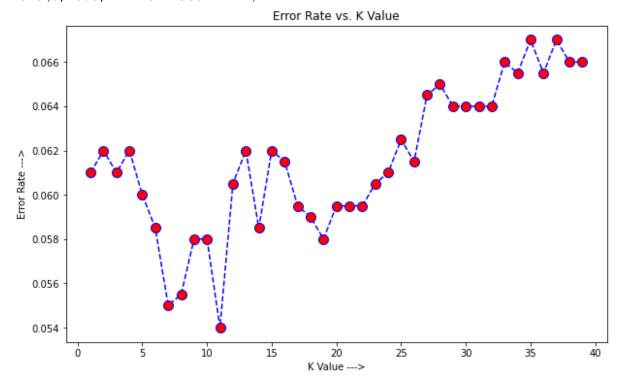
2000

0.66

0.94

```
In [41]:
error_rate = []
for i in range(1,40):
 knn = KNeighborsClassifier(n_neighbors=i)
 knn.fit(x_train,y_train)
 kpred_i = knn.predict(x_test)
 error_rate.append(np.mean(kpred_i != y_test))
 if (error_rate[i-1]==min(error_rate)):
   dip = i
print('min error=', dip, min(error_rate))
print(error_rate)
min error= 11 0.054
[0.061, 0.062, 0.061, 0.062, 0.06, 0.0585, 0.055, 0.0555, 0.058, 0.058, 0.0
54, 0.0605, 0.062, 0.0585, 0.062, 0.0615, 0.0595, 0.059, 0.058, 0.0595, 0.0
595, 0.0595, 0.0605, 0.061, 0.0625, 0.0615, 0.0645, 0.065, 0.064, 0.064, 0.
064, 0.064, 0.066, 0.0655, 0.067, 0.0655, 0.067, 0.066, 0.066]
In [58]:
plt.figure(figsize=(10,6))
plt.plot(range(1,40),error_rate,color='blue', linestyle='dashed', marker='o',markerfacecolor='red', marke
rsize=10)
plt.title('Error Rate vs. K Value')
plt.xlabel('K Value --->')
plt.ylabel('Error Rate --->')
```

## Out[58]: Text(0, 0.5, 'Error Rate --->')



# # Again running KNN with best value of n i.e. 11 in our case

```
In [55]:
knn = KNeighborsClassifier(n_neighbors= 11)
knn.fit(x_train,y_train)
kpred = knn.predict(x_test)
In [44]:
print('FINALLY WITH K = 11 (error rate minima)\n')
print('Confusion Matrix:')
print(confusion_matrix(y_test,kpred))
print('\nClassification Report:')
print(classification_report(y_test,kpred))
FINALLY WITH K = 11 (error rate minima)
Confusion Matrix:
[[1780
            0
                  1]
      9
           49
                 44]
    17
           37
                 63]]
Classification Report:
                 precision
                                recall f1-score
                                                         support
```

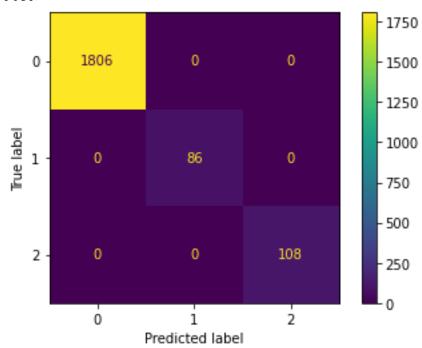
0	0.99	1.00	0.99	1781
1	0.57	0.48	0.52	102
2	0.58	0.54	0.56	117
accuracy			0.95	2000
macro avg	0.71	0.67	0.69	2000
weighted avg	0.94	0.95	0.94	2000

In [45]:

from sklearn.metrics import plot\_confusion\_matrix
plot\_confusion\_matrix(knn, x\_test, kpred)

#### Out[45]:

<sklearn.metrics.\_plot.confusion\_matrix.ConfusionMatrixDisplay at 0x21d7090
96c8>



### # Predicting the result for the given two Inputs

```
In [46]:
campus=['Not Qualified','Selected in IIM-DL','Selected in IIM-GJ']
def kpredictfor(testcase):
    college = knn.predict(testcase)
    return college

print('Custom Cases:')
# testA = [[50,40,60]]
collegeA = kpredictfor([stu1])
print('KNN: Student A was ---> ',campus[int(collegeA)])
```

```
#testB = [[80,35,105]]
collegeB = kpredictfor([stu2])
print('KNN: Student B was ---> ',campus[int(collegeB)])

Custom Cases:
KNN: Student A was ---> Not Qualified
KNN: Student B was ---> Not Qualified
```

## # Applying Bayesian Learning Model on the same dataset

```
In [47]:
from sklearn.naive_bayes import GaussianNB
gnb = GaussianNB()
gnb.fit(x_train,y_train)
bpred = gnb.predict(x_test)
In [48]:
from sklearn.metrics import classification_report,confusion_matrix
print('Confusion Matrix:')
print(confusion_matrix(y_test,bpred))
print('\nClassification Report:')
print(classification_report(y_test,bpred))
Confusion Matrix:
[[1781 0 0]
[ 31 22
               49]
[ 43 31 43]]
Classification Report:
               precision recall f1-score support
           0
                   0.96
                              1.00
                                         0.98
                                                    1781
                    0.42
           1
                              0.22
                                          0.28
                                                      102
                   0.47 0.37
                                         0.41
                                                     117
                                          0.92
                                                     2000
    accuracy
```

#### In [49]:

macro avg

weighted avg

from sklearn.metrics import plot\_confusion\_matrix
plot\_confusion\_matrix(gnb, x\_test, bpred)

0.61

0.90

0.53

0.92

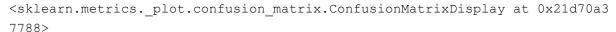
0.56

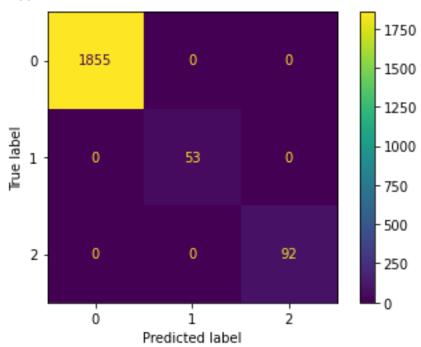
0.91

2000

2000

```
Out[49]:
```





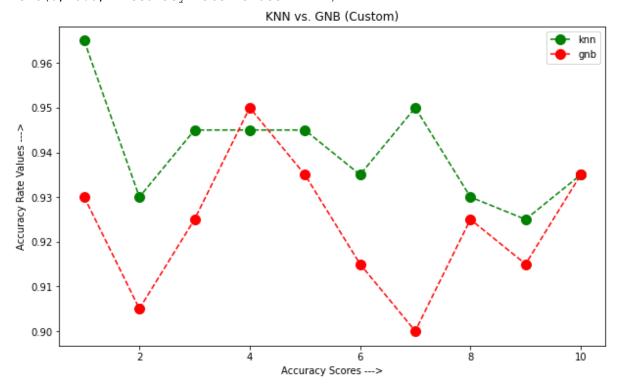
## # Comparing the KNN with Bayesian learning model

```
In [52]:
from sklearn.naive_bayes import GaussianNB
from sklearn.neighbors import KNeighborsClassifier
from sklearn.model_selection import cross_val_score
print('Comparison KNN vs GNB \n')
print('knn', cross_val_score(knn, x_test, y_test, scoring='accuracy', cv=10).mean())
print('gnb', cross_val_score(gnb, x_test, y_test, scoring='accuracy', cv=10).mean())
Comparison KNN vs GNB (Custom)
knn 0.9405000000000001
gnb 0.9235000000000001
In [53]:
plt.figure(figsize=(10,6))
plt.plot(range(1,11),cross_val_score(knn, x_test, y_test, scoring='accuracy', cv=10),color='green', linestyle
='dashed', marker='o',markerfacecolor='green', markersize=10, label='knn')
plt.plot(range(1,11),cross_val_score(gnb, x_test, y_test, scoring='accuracy', cv=10),color='red', linestyle='
dashed', marker='o',markerfacecolor='red', markersize=10, label='gnb')
plt.legend()
plt.title('KNN vs. GNB')
```

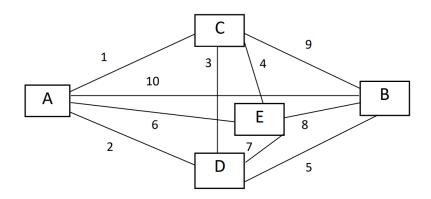
```
plt.xlabel('Accuracy Scores --->')
plt.ylabel('Accuracy Rate Values --->')
```

#### Out[53]:

Text(0, 0.5, 'Accuracy Rate Values --->')



**Q2.** Solve following TSP problem using prolog. Consider source node= 'A' and Goal Node = 'B'



#### CODE:

edge(a,b,10).

edge(a,c,1).

edge(a,d,2).

edge(a,e,6).

edge(b,a,10).

edge(b,c,9).

```
edge(b,d,5).
edge(b,e,8).
edge(c,a,1).
edge(c,b,9).
edge(c,d,3).
edge(c,e,4).
edge(d,a,2).
edge(d,b,5).
edge(d,c,3).
edge(d,e,7).
edge(e,a,6).
edge(e,b,8).
edge(e,c,4).
edge(e,d,7).
len([], 0).
len([H|T], N):-
len(T, X),
N is X+1.
best_path(Visited, Total):- path(a, b, Visited, Total).
path(Start, Fin, Visited, Total) :-
path(Start, Fin, [Start], Visited, 0, Total).
path(Start, Fin, CurrentLoc, Visited, Costn, Total) :-
edge(Start, StopLoc, Distance),
NewCostn is Costn + Distance,
\+ member(StopLoc, CurrentLoc),
path(StopLoc, Fin, [StopLoc|CurrentLoc], Visited, NewCostn, Total).
path(Start, Fin, CurrentLoc, Visited, Costn, Total) :-
edge(Start, Fin, Distance), reverse([Fin|CurrentLoc], Visited),
len(Visited, Q),
(Q)=5 \rightarrow Total is 100000; Total is Costn + Distance).
shortest_path(Path):-setof(Cost-Path, best_path(Path,Cost), Holder),pick(Holder,Path).
best(Cost-Holder,Bcost-_,Cost-Holder):-
Cost<Bcost,!.
best(\_,X,X).
pick([Cost-Holder|R],X):-
pick(R,Bcost-Bholder),
best(Cost-Holder,Bcost-Bholder,X),
!.
pick([X],X).
```

#### **OUTPUT:**

```
SWI-Prolog (AMD64, Multi-threaded, version 8.2.3)

File Edit Settings Run Debug Help

Welcome to SWI-Prolog (threaded, 64 bits, version 8.2.3)

SWI-Prolog comes with ABSOLUTELY NO WARRANTY. This is free software.

Please run ?- license. for legal details.

For online help and background, visit https://www.swi-prolog.org

For built-in help, use ?- help(Topic). or ?- apropos(Word).

?-

Warning: c:/users/kulpr/documents/prolog/tsp.pl:23:
Warning: Singleton variables: [H]
?- shortest_path(X).
X = 17-[a, d, c, e, b].
?- ■
```

**Q3.** Write a prolog program to implement Medical Diagnosis Expert System (MDES). Create your own facts and rules and apply it in MDES implementation.

#### CODE:

```
:- dynamic known answers/3.
:- dynamic patient record/2.
:- dynamic illness/2.
%-----
% symptom
% if you want to add more symptoms, just add more predicates below
%-----
symptom(fever).
symptom(cough).
symptom(shivering).
symptom(runny nose).
%-----
% new patient/1
% starts a new session
% It must be initiated with a patent name.
% This goal clears all known_answers,
% and starts the examine subgoal followed by diagnosis subgoal.
%-----
new_patient(PatientName) :- not(patient_already_exists(PatientName)),
retractall(known\_answer(\_,\_,\_)), write('\nStarting
examination...\n'),examine,confirmed_symptoms(S), write('\nConfirmed symptoms are '),
write_term(S, []), write('\n'), assert(patient_record(PatientName,S)), !,
write('Determining illness...\n'), diagnose(S,I),!, write('AI diagnosed that you have '),
```

```
write_term(I, []), !.
%-----
% symptoms/1
% returns a list of all symptoms61
%-----
symptoms(L):-findall(X, symptom(X), L).
%-----
% examine/0
% starts the examination process
% by asking for a yes/ no question against each symptom
examine:-symptoms(L), check symptoms(L).
% -----
% diagnose/2
% starts the diagnosis process by checking PatientSymptoms and unifying the Illness
% This works by checking whether an Ilness exists with symptoms being
% subset of PatientSymptoms -----
diagnose(PS, I):-length(PS, MustMatchCount), diagnose(PS, MustMatchCount, I).
%-----
% diagnose/2
% The following predicate is expected to match when no other illness is identified.
%-----
diagnose(_, unknown_disease).
%-----
% diagnose/3
% recursively matches the illness symptoms and patient symptoms with
% decreasing number of matches PS: Patient Symptoms (expected to be
% passed as a parameter) I: Illness (expected to be unified)
% MustMatchCount: The number of symptoms that should exist in Illness
% -----
diagnose(PS, MustMatchCount, I) :- (illness(I,S), length(S, MustMatchCount),
subset(S,PS),!); (MustMatchCount > 1, NewCount is MustMatchCount-1,
diagnose(PS, NewCount, I)).
%-----
% check symptoms/1
% given a list of symptoms, ask quetions
%-----
check_symptoms([]) :- !.
check_symptoms([H|T]) :- ask(symptom,H), check_symptoms(T).
%-----
% confirmed_symptoms/1
% returns a list of symptoms for which the answer is yes
confirmed_symptoms(C) :- findall(X,known_answer(yes,symptom,X),C).
%-----
% ask/2
% given an attribute and a value, gets a yes/no answer from the user
% It works by writing a prompt and having a subgoal to assert the answer
```

```
ask(Attr,Val):-write(Attr:Val), write('?'), read(Y), asserta(known answer(Y,Attr,Val)).
%-----
% fix_diagnosis/2
% learns "actual" illness of a "patient" and improves the learning process
fix_diagnosis(PatientName, ActualIllness) :- patient_record(PatientName, PS),
write('Confirmed patient symptoms '), write_term(PS, []),
write(' will be related to '), write_term(ActualIllness, []), !, update_definition(ActualIllness, PS, FS),!,
write('\nNew definition is '), write_term(FS,[]).
%-----
% update definition/3
% given an illness and new symptoms, returns the updated symptoms for that illness
% Following case is when the illness is not already defined.
update_definition(Illness, RelateSymptoms, RelateSymptoms) :- not(illness(Illness, _)),
write('\nThere was no earlier definition of '), write_term(Illness, []),assert(illness,
RelateSymptoms)),!.
%-----
% update_definition/3
% given an illness and new symptoms, returns the updated symptoms for that illness
% Following case is when the illness is already defined, and hence takes
% an intsection of old and new symtpoms -----
update definition(Illness, RelateSymptoms, FinalSymptoms):- illness(Illness, OldSymptoms),
write('\nEarlier definition of '),write(Illness),write(' was '),
illness(Illness, OldSymptoms), write_term(OldSymptoms,[]),
intersection(OldSymptoms, RelateSymptoms, FinalSymptoms), retractall(illness(Illness, )),
assert(illness(Illness, FinalSymptoms)).
%-----
% rediagnose/1
% given a patient name, rechecks diagnosis based on existing symptoms
% This goal could be requested, for example, when illness predicates are updated
%-----
rediagnose(PatientName):-patient_record(PatientName, C), write('\nConfirmed symptoms were
write term(C, []), write('\nRediagnosing...'), diagnose(C, NewIllness),!,
write('\nUpdated diagnosis is that Patient '), write_term(PatientName, []),
write(' is having '),write_term(NewIllness, []).
%-----
% patient_already_exists/1
% true if given patient name is already in the patient records
%-----
patient_already_exists(PatientName) :-patient_record(PatientName,_),
write('Patient'), write_term(PatientName,[]), write('already exists.\n').
%-----
% show_patient_records/0
% shows all patient records
%-----
```

```
show_patient_records :- findall((P,S), patient_record(P,S), L), !,show_records(L).
%-----
% show records/1
% Calls show_record for each (PatientName, Symptom) pair
%-----
show records([]).
show_records([(P,S)|T]):-show_record(P,S),show_records(T).
show_record(P, S):- diagnose(S, I),!, write_term(P, []), write(' has symptoms '),
write term(S, []), write(' and diagnosed '),
write_term(I, []), write('\n').
%-----
% change diagnosis/2
% Associate the symptoms from one illness to another for a given patient
change diagnosis(Patient, NewIllness):-patient record(Patient, Symptoms),!.
write_term(Patient, []),write(' has symptoms '),
write term(Symptoms, []), diagnose(Symptoms, OldIllness),
write(' and was diagnosed '), write term(OldIllness, []),
write('\n'),!,write('Changing it to '),
write_term(NewIllness, []), write('\n'),
retractall(illness(NewIllness, _)), retractall(illness(OldIllness, _)),
assert(illness(NewIllness, Symptoms)).
OUTPUT:
                                                                                          X
 SWI-Prolog (AMD64, Multi-threaded, version 8.2.3)
File Edit Settings Run Debug Help
% c:/users/kulpr/documents/prolog/mdes compiled 0.00 sec, 0 clauses
Warning: [Thread poe] The predicates below are not defined. If these are defined Warning: [Thread poe] at runtime using assert/1, use :- dynamic Name/Arity.
           [Thread pce]
[Thread pce]
Warning:
Warning:
                         known_answer/3, which is referenced by
Warning: [Thread pce]
                            c:/users/kulpr/documents/prolog/mdes.pl:67:35: 1-st clause
 of confirmed_symptoms/1
?- new_patient(abc).
Starting
examination...
symptom:fever? no
symptom:cough? |: no.
symptom:shivering? |: no.
symptom:runny_nose? |: yes.
Confirmed symptoms are [runny_nose]
Determining illness.
AI diagnosed that you have unknown_disease
true.
 ?- fix_diagnosis(abc,cold).
Confirmed patient symptoms [runny_nose] will be related to cold
 There was no earlier definition of cold
```

New definition is [runny\_nose]

true.

```
SWI-Prolog (AMD64, Multi-threaded, version 8.2.3)
                                                                                                                         X
File Edit Settings Run Debug Help
?- rediagnose(abc).
Confirmed symptoms were
[runny_nose]
Rediagnosing
Updated diagnosis is that Patient abc is having cold
true.
?- new_patient(def).
Starting
examination..
symptom:fever? no.
symptom:cough? |: no.
symptom:shivering? |: no.
symptom:runny_nose? |: yes.
Confirmed symptoms are [runny_nose]
Determining illness...
AI diagnosed that you have cold
true.
?- new_patient(ghi).
Starting
examination.
svmptom:fever? ves.
SWI-Prolog (AMD64, Multi-threaded, version 8.2.3)
                                                                                                                         ×
File Edit Settings Run Debug Help
true.
?- new_patient(ghi).
Starting
examination . .
examination...
symptom:fever? yes.
symptom:cough? |: no.
symptom:shivering? |: no.
symptom:runny_nose? |: yes.
Confirmed symptoms are [runny_nose,fever]
Determining illness...
AI diagnosed that you have cold
true.
?- fix_diagnosis(ghi,flu).
Confirmed patient symptoms [runny_nose, fever] will be related to flu
There was no earlier definition of flu
New definition is [runny_nose, fever]
true.
?- ■
```