# MTL782 Assignment

Data Mining - Multiclass Classification

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#### Data Set Used: Maternal Health Risk Data Set

#### (https://archive.ics.uci.edu/ml/datasets/Maternal+Health+Risk+Data+Set)

1) A Data description

The data is about the health risk of females during pregnancy based on their blood pressure, heart rate, and blood glucose level. Data has been collected from different hospitals, community clinics, maternal health cares in the rural areas of Bangladesh through the IoT-based risk monitoring system.

2) Benefits of using Data Mining

The data can help in pre-diagnosing the pregnant females that their health condition is good enough for giving birth or they need any kind of premedication to make their vitals normal before going into labor. This can help the patient in giving safe birth and reassure the doctor that the health of the patient will be alright after giving birth.

- 3) Further information about data
- i) There were some data that did not align with our predictions this could be because of some noise present that is an error in collecting the data. The other reason is that every human body is different and it is very difficult to accurately predict the risk level by using a limited amount of dataset.
- ii) We can try to fix the error by taking the data from different parts in India and different hospitals (this will improve our test and training sets), we can also introduce new attributes using the current attributes or introduce some new attributes that might be useful (this will improve our algorithm).

# **Code for Decision Tree**

```
import pandas as pd
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
from sklearn import metrics

data = pd.read_csv("Maternal Health Risk Data Set.csv")
X = pd.read_csv("Maternal Health Risk Data Set.csv", usecols=[i for i in data.columns if i != 'RiskLevel' ])
y = data['RiskLevel'].values.reshape((-1, 1))
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=1)
clf = DecisionTreeClassifier()
clf = clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
print("Accuracy:", metrics.accuracy_score(y_test, y_pred))
```

#### **Code for Random Forest**

```
import pandas as pd
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import train_test_split
from sklearn import metrics

data = pd.read_csv("Maternal Health Risk Data Set.csv")
X = pd.read_csv("Maternal Health Risk Data Set.csv", usecols=[i for i in
data.columns if i != 'RiskLevel' ])
y = data['RiskLevel'].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=1)
clf = RandomForestClassifier()
clf = clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
print("Accuracy:", metrics.accuracy score(y test, y pred))
```

### **Code for KNN Classifier**

```
import pandas as pd
from sklearn.neighbors import KNeighborsClassifier
from sklearn.model_selection import train_test_split
from sklearn import metrics

data = pd.read_csv("Maternal Health Risk Data Set.csv")
X = pd.read_csv("Maternal Health Risk Data Set.csv", usecols=[i for i in
data.columns if i != 'RiskLevel' ])
y = data['RiskLevel'].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,
random_state=1)
clf = KNeighborsClassifier()
clf = clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
print("Accuracy:", metrics.accuracy score(y test, y pred))
```

# **Code for Naïve Bayes Classifier**

```
import pandas as pd
from sklearn.naive_bayes import GaussianNB
from sklearn.model_selection import train_test_split
from sklearn import metrics

data = pd.read_csv("Maternal Health Risk Data Set.csv")
X = pd.read_csv("Maternal Health Risk Data Set.csv", usecols=[i for i in data.columns if i != 'RiskLevel'])
y = data['RiskLevel'].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=1)
clf = GaussianNB()
clf = clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
print("Accuracy:", metrics.accuracy score(y test, y pred))
```

# Code for comparing performances using K-fold Cross Validation

```
from sklearn.naive bayes import GaussianNB
warnings.filterwarnings('ignore')
class kFoldCV:
    def crossValSplit(self, dataset, numFolds):
                fold.append(dataCopy.pop(index))
    def kFCVEvaluate(self, dataset, numFolds):
        scores = list()
```

```
testSet.append(rowCopy)
trainLabels = [row[-1] for row in trainSet]
trainSet = [train[:-1] for train in trainSet]
scores.append(accuracy)
accuracy = printMetrics(actual, predicted)
scores.append(accuracy)
```

```
trainSet = list(folds)
testSet = list()
    testSet.append(rowCopy)
trainLabels = [row[-1] for row in trainSet]
trainSet = [train[:-1] for train in trainSet]
RF.fit(trainSet, trainLabels)
predicted = RF.predict(testSet)
scores.append(accuracy)
trainSet = [train[:-1] for train in trainSet]
NB.fit(trainSet, trainLabels)
testSet = [test[:-1] for test in testSet]
predicted = NB.predict(testSet)
scores.append(accuracy)
```

```
def readData(fileName):
    data = []
    labels = []

with open(fileName, "r") as file:
        lines = file.readlines()
    for line in lines:
        splitline = line.strip().split(',')
        data.append(splitline)
        labels.append(splitline[-1])
    return data, labels

File = 'Maternal Health Risk Data Set.csv'

Data, Label = readData(File)
df = pd.DataFrame(Data)
df = df.apply(preprocessing.LabelEncoder().fit_transform)
Features = df.values.tolist()
Labels = [car[-1] for car in Features]
kfov = kFoldCV()
print('*' * 20)
print('Maternal Health Risk Data Set')
print('\n')
kfcv.kFCVEvaluate(Features, 10)
```

# **Code for Apriori Algorithm**

```
support = float(ssCnt[key] / numItem)
         if support >= minSupport:
         supportData[key] = support # Update support dictionary
    return retList, supportData
         for j in range(i + 1, lenLK):
    L1 = list(LK[i])[:k - 2]
    L2 = list(LK[j])[:k - 2]
                  retList.append(LK[i] | LK[j])
    return retList
def aprioriGen(LK, k): # Create candidate item set CK, where k is the
             if L1 == L2:
    return retList
def apriori(dataSet, minSupport):
    C1 = DataSet(dataSet)
         D.add(tid)
         supportData.update(supK)
         L.append(LK)
minSupport = 0.2
minconfidence = 0.5
def createData (name): # Preprocess the data and output the dataset
```

```
D = D.fillna(0)
    dataset, dic = apriori(dataset, minSupport)
    Rsupport = []
                if A.issubset(AB):
                        Rconfidence.append(conf)
                        Rsupport.append(dic.get(AB))
    return Rname, Rsupport, Rconfidence
def outputdata(Rname, Rsupport, Rconfidence):
dataset = createData(name)
R1, R2, R3 = calculate(dataset)
df = outputdata(R1, R2, R3)
df.to csv('Report.txt')
```

#### TXT file used above: menu\_orders.txt

```
a,c,e
b,d
b,c
a,b,c,d
b,c
a,b
b,c
```

```
a,b,c,e
a,b,c
a,c,e
```

#### Code for FP-Growth

```
import pandas as pd
      self.parent = parent
self.link = link
      self.children = {}
      output.append(str(vocabdic[self.word]) + " " +
               output.append(self.children[i].visittree())
      self.wordlinesort = []
      self.wordsortdic = []
```

```
elf.construct(data)
    def construct(self, data):
x[1], x[0])
            self.nodetable.append(wordinfo)
self.wordorderdic[k])
                for i in sortsupword:
                    if i in R.children.keys():
                        R.children[i].word count += 1
                        R = R.children[i]
                        R = R.children[i]
```

```
else:
                line.append(PN.word)
                condtreeline.append(line)
    def findfqt(self, parentnode=None):
        if len(list(self.root.children.keys())) == 0:
            conwords = contree.findfqt(fqset)
                     result.append(words)
    def checkheight(self):
min sup = 2
```

```
import numpy as np
import pandas as pd

dataset = pd.read_csv('menu_orders.csv', sep='delimeter', header=None)
test_data = []
for sublist in dataset.values.tolist():
    clean_sublist = [item for item in sublist if item is not np.nan]
    test_data.append(clean_sublist)

fp_tree = fptree(test_data, min_sup) # create FP tree on data

# print ("\n========== Printing Frequent Word Set on " + i +" =========")
frequentwordset = fp_tree.findfqt() # mining frequent patt
frequentwordset = sorted(frequentwordset, key=lambda k: -k[1])

# print frequent patt
for word in frequentwordset:
    count = (str(word[1]) + "\t")
    words = ''
    for val in word[0]:
        words += (str([val]) + " ")
    print(count + words)

for i in fp_tree.nodetable[::-1]:
    lines = fp_tree.condtreetran(i['linknode'])
    condtree = fptree(lines, min_sup)
    if (condtree.checkheight()):
        print('Condtional FPTree Root on ' + (vocabdic[i['wordn']]))
        print('Condtional FPTree Root on ' + (vocabdic[i['wordn']]))
        print('Condtree.root.visittree())
```

## **Modified Algorithm for Apriori**

Optimization in Apriori algorithm

- 1.It is possible to delete infrequent itemsets directly in the transaction database to avoid multiple scans and reduce I/O overhead
- 2. For frequent K itemsets, if a single item i appears less than k times, the itemset containing i cannot appear in the frequent k+1 itemset. In the frequent K itemset, items containing one i should be deleted and linked together

```
ssCnt[can] = 1
            dataSet.remove(tid)
for tid in retList:
                tids.append(tid)
```

```
for j in range(i + 1, lenLK):
    L1 = list(LK[i])[:k - 2]
    L2 = list(LK[j])[:k - 2]
              L1.sort()
                   retList.append(LK[i] | LK[j])
     return retList
def apriori(dataSet, min support 1):
         supportData.update(supK)
         L.append(LK)
name = 'menu orders.txt'
minSupport = 0.2
minconfidence = 0.5
def createData(name): # Preprocess the data and output the dataset
    D = pd.read csv(name, header=None, index col=False, names=['1', '2',
    D = D.fillna(0)
```