Shourya Jindal 2020336 ML Assignment 2

Section A

(Q1)

(RF)

(a) In random forests, correlation regers to the degree to which individual decision trees, in the RF are similar to each other.

> High Correlation = Trees make similer error and decision on somedota pts

= Make dig. devisions on Barne data pts.

Diversity regers to differences among the trees in the RF. () diff. Jecture selections, theresholds, Tules. high aircroits => Reduces orugiting, reduces variance.

So, tradeby blue these 2 is that

- · if we increase correlation too much, it is similer to having Single decision tree. leads to Overyithing => Higher Bass Low Bies (= Irligh Variona aloes not perform well
- on unseen data . But Andig increese diversits too much, it leads to lack of consensus in the predictions. Reduces overall accuracy. It reduces overfitting, they reducing verience.
- (b) In 'naive Bayes', the "curse of dimensionality" becomes an issue, when , 'no. of features' i.e the dimensionalis of data becomes very large. It leads to very high computational complexits, overfitting. In hisher dimensions deta becomes spenese making it difficult to colculate probabilities in 'Naive Bias' also leading to incressed Competational complexity. Also, leads to overyithing, thus poor performence on unseen det a.

Strategies + Mitigate this:

- · Feature selection: Use only relevant deatures and discend redundent ".
- · Dimensionality red": Use methods like PCA, TSNE reduce dimensionality
- · Regularization: Use Li, lz methodo to prevent overfiting
- · ensemble Methods. Use ensemble methods like boosting, random forest => Mor effective in hisher dimensions.

(c) you, if some value of attributes which was not present in the training set is encountered, it will affect the injurence repults.

Problem:

- · when unsur value of attri. is encountered, it will assign it prob. 0.
- · This can lead to worong predictions
- . Thus leading to high variance inc low a courses to on uns een det q.
- · Accuracy reduces
- Solm: , Laplace smoothing: Add a small value (3) severally 1, to court of each attribute value pair. = No zero prob.
 - · Use prior prob: Top If we have access to prior prob. for altri volves, these can be siver to the model to help estimate unseen values prob.

Training: < mail 1 be lotters of free 1 St

Smail	1 to Lottery	· free	Spem Not
- Liver	909	yes	900
2-	yes	No	949
3	No	99	NO
4	No	No	1 No

Test sample: anails: "Lottery free hello". = X her In tuis, p("hello" | "spam") = 0 =) p (spem | xnew) = 0 although it contains both "lotter's our "dree".

but single "hello" is (= unseen, it is but if leplace smoothing declared as sporm. will be used it will prob \$00 >0.

So, prob. are:

Possible outcomes our prob:

Vom lest part (c)

* Cerdio:

P[Cerdio] = (0.42) (0.8). (0.5) + (0.18) (0.4) (0.5)

= 0.168 + 0.36 = 0.528

P[weigner] = Sem or Cerdio

= 0.528

= 0.084 + 0.108 = 0.084 + 0.108

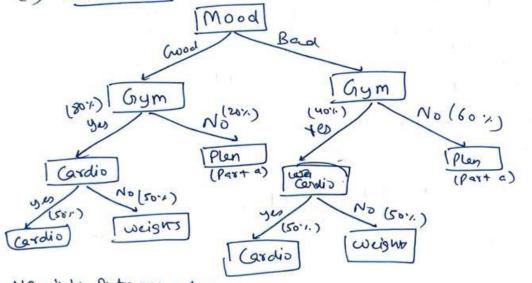
Sea Toto calculate exacts which sport he will play is info. is not available.

So, He will visit sym and would either do

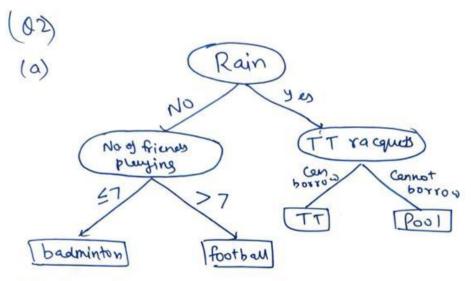
Cardio or weights with equal probability.

-0.528)

(c) Decision Tree:



Affossible Outcomes and prob ..



Possible Outcomes & their Probabilities:

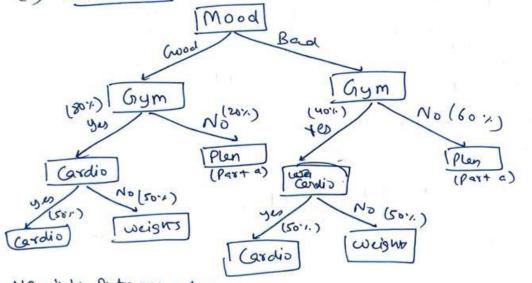
Now, first we will calculate p[Rain]

for this we will use total probability theorem:

P[Predicts Rainy] = P[Rain] + P[Predicts Rains | Rains | + P[Clean] + P[Preak cts Rain | Clean)

(Using siven a and e, n 1) 0.3 = 0.8 P[Rain] + 0.1 P[(Len)

(c) Decision Tree:



Affossible Outcomes and prob ..

(d) Yes, if some attri have more condinality, there can be bias in splitting so decision tree mode using Info. Gain.

This is because cardinalib means that no. of distinct values an attribute can take.

If an attribute was high cardinality of they can be split into more subset (more by work of the work of the tree)

() Thus, blasedness:

* Other criterion that can be used:

- · Crimi Index: Gaini (s)= 1- Epi2
 it is less sensitive to condinations
- · Crain Retio: It takes into cendinality and penelizes attribtes with high 11.

Example:

Suppose, we are building a decision tree to predict, which specy was ice- Green Sales.

Consider 3 attributes.

- (1) Age Group: (2) { 2000 < 13, 18-435, 36-60, >60}
- (2) Weather conditions: { Rainy, hot, cold}
- 3 Flavour = ¿ Chocolote, butterscotch, herry, -- 3 hish cerdinality

If we use I'm as an criterion, it can favour flavour attribute (due to its high cerdinality).

But, this may not be was optimal approach, as

Solut of ice-cream are generally to effected most by the

factors like weather condition.

* Rest of the outcomes ere some just use previous formula and multiply them with an do following: P(Event) = P(Plen) . P(Event) 20.2 where P(Plan): P(Good Mood) . P(No asm) Good Mood) +P (Bad Mood). P (" " | Bad Mood) -0.6 (d) P (Gross word) = 0. 6 Assuming the selft 7 hrs P(Bad Mood) = U.y one night begon P(R=7)Gm)=0.7 50, P[F=7]=1 P(F=7/BM)=0.45 we are assuming that be stop p (Gm) = 0. 4 Solon east last now P(Com) P(F=>16m) + PEgm) P FF=>16m) Now, for next day: P[GM]F=7] = P[F=7]Gm].P[Gm]
P[F=7] - (0.7)(0.6)=0.42 P[BM|F=]= P[F=7|BM]-P[BM] = (0.45) (0.4) = 0.18

Section B

PART A: Preprocessing and EDA

Read the data and store it in the pandas' data frame. In total, 13 features and 1 target column. Toral rows: 303.

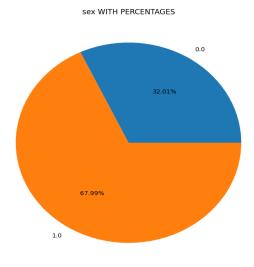
Preprocessing:

- Converted the classification to binary classification.
 Target values with value > 0 were converted to 1
- In the data, missing values were marked as '?'. Replaced them with the mode value of their respective columns, as most values were categorical data.

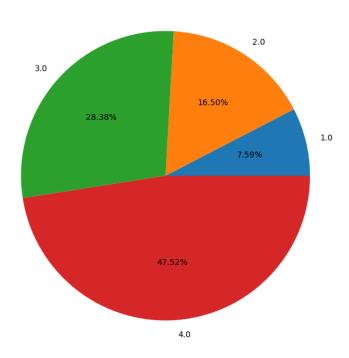
Columns: {0: 'age', 1: 'sex', 2: 'cp', 3: 'trestbps', 4: 'chol', 5: 'fbs', 6: 'restecg', 7: 'thalach', 8: 'exang', 9: 'oldpeak', 10: 'slope', 11: 'ca', 12: 'thal', 13: 'num'}

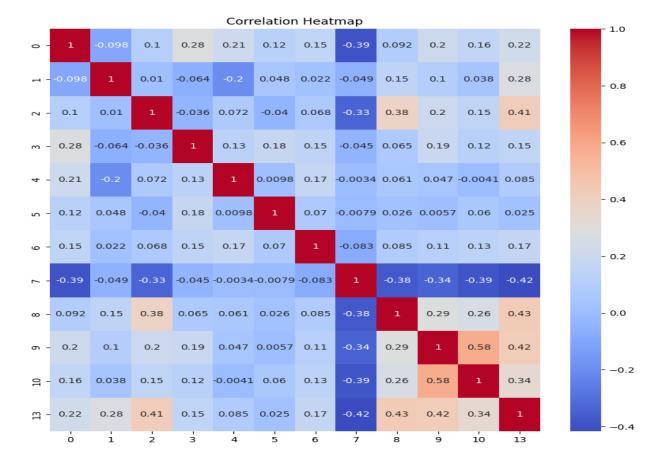
Where the target variable is column no. 13 or num.

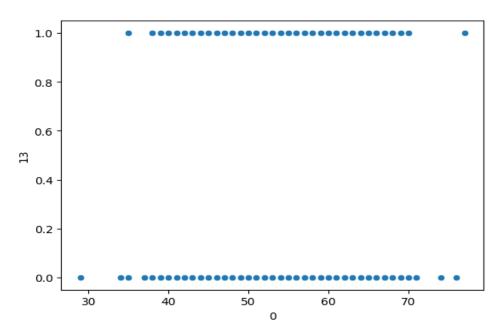
EDA: I visualised the data by creating several graphs and plots. Scatter plot, heatmap, pairplots, piecharts. Some of these are:



cp WITH PERCENTAGES







X-axis: age. Y-axis: target.

Some insights are:

- From the correlation heatmap, we can see that most of the features are not correlated, except features no. 9 and 10. Also, target variable no. 13 somewhat correlates with features 2, 8, and 9.
- Made scatter plots and pair plots to find outliers. But there was not much of any outlier.
- On a surprising, from the scatter plot above, we can see that age does not much affect the outcome of heart disease.
- Data has an unequal distribution of sex. One gender data is almost double the other, but it is not specified.

Part B:

Split the data into training and testing in the ratio of 80:20. Done using sklearn library:

```
x_train, x_test,y_train,y_test = train_test_split(X,Y,test_size = 0.2)
```

Part C:

Trained the decision tree using <u>entropy</u> as the splitting criterion. Got the following accuracy:

```
#Entropy
Entropy = DecisionTreeClassifier(criterion='entropy')
Entropy.fit(x_train, y_train)
ypredEntropy = Entropy.predict(x_test)
accuracyEntropy = accuracy_score(y_test, ypredEntropy)
print(accuracyEntropy)

0.7049180327868853
```

When trained using 'gini impurity' as the splitting criterion, it got the following accuracy:

```
#Gini
Gini = DecisionTreeClassifier(criterion='gini')
Gini.fit(x_train, y_train)
ypredGini = Gini.predict(x_test)
accuracyGini = accuracy_score(y_test, ypredGini)
print(accuracyGini)

0.6557377049180327
```

So, based on the accuracy score, 'Entropy' was the better criterion for node splitting.

Part D:

Performed hyperparameter search for the parameters min_samples_split and max_features using Grid search, with the help sklearn.

Values passed in the grid were:

```
'min_samples_split': [2, 3, 5, 7, 9, 10],
'max_features': ['sqrt', 'log2', None]
```

The criterion used for splitting was: 'Entropy' (from part c).

After running this and using the best parameters that came out were:

```
Hyperparameters Values:
min_samples_split:9
max_features:log2
Accuracy:0.7213114754098361
```

With an accuracy of 0.72.

Part E: Random Forests

Performed hyperparameter search for the parameters n_estimators, max_depth and min_samples_split using Grid search, with the help of sklearn.

Values passed in the grid were:

```
'n_estimators': [50, 100, 150],
'max_depth': [None, 10, 20, 30],
'min_samples_split': [2, 3, 5, 7, 9, 10]
```

After running a grid search, the following were the results:

```
Hyperparameters Values:
n_estimators:50
max_depth:20
min samples split:9
Classification Report:
            precision recall f1-score
                                        support
                0.72
                         0.91
                                  0.81
         0
                                             32
         1
                0.86
                                  0.72
                                             29
                         0.62
   accuracy
                                  0.77
                                             61
                                  0.76
  macro avg 0.79
                         0.76
                                             61
weighted avg
               0.79
                         0.77
                                  0.76
                                             61
Accuracy:0.7213114754098361
```

This shows the combination of best hyperparameters, classification reports, and accuracy.