

**Final Year B. Tech. (CSE) – I : 2021-22**  
**4CS462 : PE2 - Data Mining Lab**  
**Assignment No. 5**

**Group id: DM21G12**

**Group members:**

**Abhishek More(2018BTECS00037)**

**Sushil Wagh(2018BTECS00031)**

**Title :** Design the rule based classifier : Extract the rules from decision tree build in assignment no. 4.

**Objective/Aim :**

Tabulate the results and evaluate the performance of rules generated using following metrics :

- a. Coverage
- b. Accuracy
- C. Toughness (size)

Use the following categorical data sets from UCI machine learning repository :

- a. Balance Scale data set
- b. Car evaluation data set
- C. Breast-cancer data se

**Introduction:**

**Coverage of a rule:**

The percentage of instances that satisfy the antecedent of a rule (i.e., whose attribute values hold true for the rule's antecedent).

**Accuracy of a rule:**

Rule-Based Classification

The percentage of instances that satisfy both the antecedent and consequent of a rule

- **Rule accuracy and coverage:**

$$coverage(R) = \frac{n_{covers}}{|D|}$$

$$accuracy(R) = \frac{n_{correct}}{n_{covers}}$$

- *where*

- $D$ : class labeled data set
- $|D|$ : number of instances in  $D$
- $n_{covers}$ : number of instances covered by  $R$
- $n_{correct}$ : number of instances correctly classified by  $R$

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## Coverage and Accuracy

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- The rule  $R1$ :

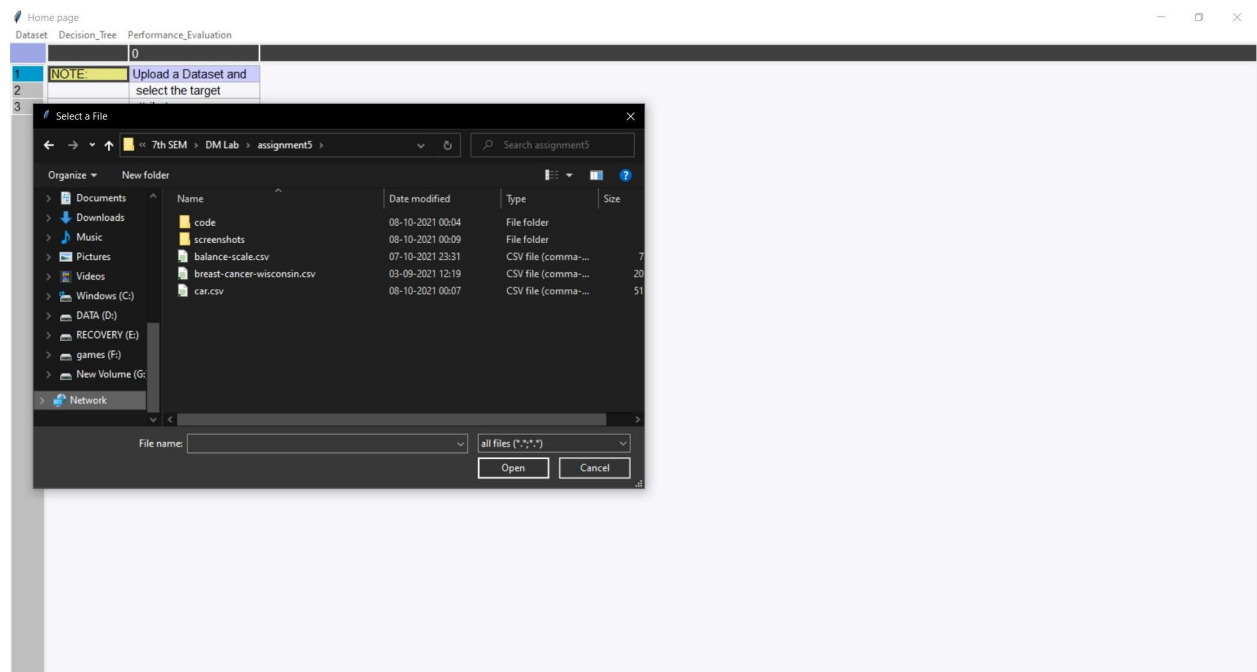
$R1$ : IF  $age = youth$  AND  $student = yes$  THEN  $buys\_computer = yes$

- $R1$  covers 2 of the 14 instances
- It can correctly classify both instances

- Therefore:

- $Coverage(R1) = 2/14 = 14.28\%$
- $Accuracy(R1) = 2/2 = 100\%$ .

**Result/Observations/Screenshots:**



|    |    | B | 1 | 1.1 | 1.2 | 1.3 |
|----|----|---|---|-----|-----|-----|
| 1  | 0  | R | 1 | 1   | 1   | 2   |
| 2  | 1  | R | 1 | 1   | 1   | 3   |
| 3  | 2  | R | 1 | 1   | 1   | 4   |
| 4  | 3  | R | 1 | 1   | 1   | 5   |
| 5  | 4  | R | 1 | 1   | 2   | 1   |
| 6  | 5  | R | 1 | 1   | 2   | 2   |
| 7  | 6  | R | 1 | 1   | 2   | 3   |
| 8  | 7  | R | 1 | 1   | 2   | 4   |
| 9  | 8  | R | 1 | 1   | 2   | 5   |
| 10 | 9  | R | 1 | 1   | 3   | 1   |
| 11 | 10 | R | 1 | 1   | 3   | 2   |
| 12 | 11 | R | 1 | 1   | 3   | 3   |
| 13 | 12 | R | 1 | 1   | 3   | 4   |
| 14 | 13 | R | 1 | 1   | 3   | 5   |
| 15 | 14 | R | 1 | 1   | 4   | 1   |
| 16 | 15 | R | 1 | 1   | 4   | 2   |
| 17 | 16 | R | 1 | 1   | 4   | 3   |
| 18 | 17 | R | 1 | 1   | 4   | 4   |
| 19 | 18 | R | 1 | 1   | 4   | 5   |
| 20 | 19 | R | 1 | 1   | 5   | 1   |
| 21 | 20 | R | 1 | 1   | 5   | 2   |
| 22 | 21 | R | 1 | 1   | 5   | 3   |
| 23 | 22 | R | 1 | 1   | 5   | 4   |
| 24 | 23 | R | 1 | 1   | 5   | 5   |
| 25 | 24 | L | 1 | 2   | 1   | 1   |
| 26 | 25 | B | 1 | 2   | 1   | 2   |
| 27 | 26 | R | 1 | 2   | 1   | 3   |
| 28 | 27 | R | 1 | 2   | 1   | 4   |
| 29 | 28 | R | 1 | 2   | 1   | 5   |
| 30 | 29 | B | 1 | 2   | 2   | 1   |
| 31 | 30 | R | 1 | 2   | 2   | 2   |
| 32 | 31 | R | 1 | 2   | 2   | 3   |
| 33 | 32 | R | 1 | 2   | 2   | 4   |
| 34 | 33 | R | 1 | 2   | 2   | 5   |
| 35 | 34 | R | 1 | 2   | 3   | 1   |
| 36 | 35 | R | 1 | 2   | 3   | 2   |
| 37 | 36 | R | 1 | 2   | 3   | 3   |
| 38 | 37 | R | 1 | 2   | 3   | 4   |

|    |    | B | 1 | 1.1 | 1.2 | 1.3 |
|----|----|---|---|-----|-----|-----|
| 1  | 0  | R | 1 | 1   | 1   | 2   |
| 2  | 1  | R | 1 | 1   | 1   | 3   |
| 3  | 2  | R | 1 | 1   | 1   | 4   |
| 4  | 3  | R | 1 | 1   | 1   | 5   |
| 5  | 4  | R | 1 | 1   | 2   | 1   |
| 6  | 5  | R | 1 | 1   | 2   | 2   |
| 7  | 6  | R | 1 | 1   | 2   | 3   |
| 8  | 7  | R | 1 | 1   | 2   | 4   |
| 9  | 8  | R | 1 | 1   | 2   | 5   |
| 10 | 9  | R | 1 | 1   | 3   | 1   |
| 11 | 10 | R | 1 | 1   | 3   | 2   |
| 12 | 11 | R | 1 | 1   | 3   | 3   |
| 13 | 12 | R | 1 | 1   | 3   | 4   |
| 14 | 13 | R | 1 | 1   | 3   | 5   |
| 15 | 14 | R | 1 | 1   | 4   | 1   |
| 16 | 15 | R | 1 | 1   | 4   | 2   |
| 17 | 16 | R | 1 | 1   | 4   | 3   |
| 18 | 17 | R | 1 | 1   | 4   | 4   |
| 19 | 18 | R | 1 | 1   | 4   | 5   |
| 20 | 19 | R | 1 | 1   | 5   | 1   |
| 21 | 20 | R | 1 | 1   | 5   | 2   |
| 22 | 21 | R | 1 | 1   | 5   | 3   |
| 23 | 22 | R | 1 | 1   | 5   | 4   |
| 24 | 23 | R | 1 | 1   | 5   | 5   |
| 25 | 24 | L | 1 | 2   | 1   | 1   |
| 26 | 25 | B | 1 | 2   | 1   | 2   |
| 27 | 26 | R | 1 | 2   | 1   | 3   |
| 28 | 27 | R | 1 | 2   | 1   | 4   |
| 29 | 28 | R | 1 | 2   | 1   | 5   |
| 30 | 29 | B | 1 | 2   | 2   | 1   |
| 31 | 30 | R | 1 | 2   | 2   | 2   |
| 32 | 31 | R | 1 | 2   | 2   | 3   |
| 33 | 32 | R | 1 | 2   | 2   | 4   |
| 34 | 33 | R | 1 | 2   | 2   | 5   |
| 35 | 34 | R | 1 | 2   | 3   | 1   |
| 36 | 35 | R | 1 | 2   | 3   | 2   |
| 37 | 36 | R | 1 | 2   | 3   | 3   |
| 38 | 37 | R | 1 | 2   | 3   | 4   |

| Dataset |    | Decision_Tree |   | Performance_Evaluation |     |     |
|---------|----|---------------|---|------------------------|-----|-----|
|         |    | Build         |   | Gini Index             |     |     |
|         |    | Display Tree  |   | Gain Ratio             |     |     |
|         |    | Display rules |   | Information Gain       |     |     |
|         |    |               |   | 1.1                    | 1.2 | 1.3 |
| 1       | 1  |               |   | 1                      | 1   | 2   |
| 2       | 1  |               |   | 1                      | 1   | 3   |
| 3       | 2  | R             | 1 | 1                      | 1   | 4   |
| 4       | 3  | R             | 1 | 1                      | 1   | 5   |
| 5       | 4  | R             | 1 | 1                      | 2   | 1   |
| 6       | 5  | R             | 1 | 1                      | 2   | 2   |
| 7       | 6  | R             | 1 | 1                      | 2   | 3   |
| 8       | 7  | R             | 1 | 1                      | 2   | 4   |
| 9       | 8  | R             | 1 | 1                      | 2   | 5   |
| 10      | 9  | R             | 1 | 1                      | 3   | 1   |
| 11      | 10 | R             | 1 | 1                      | 3   | 2   |
| 12      | 11 | R             | 1 | 1                      | 3   | 3   |
| 13      | 12 | R             | 1 | 1                      | 3   | 4   |
| 14      | 13 | R             | 1 | 1                      | 3   | 5   |
| 15      | 14 | R             | 1 | 1                      | 4   | 1   |
| 16      | 15 | R             | 1 | 1                      | 4   | 2   |
| 17      | 16 | R             | 1 | 1                      | 4   | 3   |
| 18      | 17 | R             | 1 | 1                      | 4   | 4   |
| 19      | 18 | R             | 1 | 1                      | 4   | 5   |
| 20      | 19 | R             | 1 | 1                      | 5   | 1   |
| 21      | 20 | R             | 1 | 1                      | 5   | 2   |
| 22      | 21 | R             | 1 | 1                      | 5   | 3   |
| 23      | 22 | R             | 1 | 1                      | 5   | 4   |
| 24      | 23 | R             | 1 | 1                      | 5   | 5   |
| 25      | 24 | L             | 1 | 2                      | 1   | 1   |
| 26      | 25 | B             | 1 | 2                      | 1   | 2   |
| 27      | 26 | R             | 1 | 2                      | 1   | 3   |
| 28      | 27 | R             | 1 | 2                      | 1   | 4   |
| 29      | 28 | R             | 1 | 2                      | 1   | 5   |
| 30      | 29 | B             | 1 | 2                      | 2   | 1   |
| 31      | 30 | R             | 1 | 2                      | 2   | 2   |
| 32      | 31 | R             | 1 | 2                      | 2   | 3   |
| 33      | 32 | R             | 1 | 2                      | 2   | 4   |
| 34      | 33 | R             | 1 | 2                      | 2   | 5   |
| 35      | 34 | R             | 1 | 2                      | 3   | 1   |
| 36      | 35 | R             | 1 | 2                      | 3   | 2   |
| 37      | 36 | R             | 1 | 2                      | 3   | 3   |

Home page

| Datas | Decision T | Performance | Evaluz   |          |          |          |          |          |           |          |         |       |  |
|-------|------------|-------------|----------|----------|----------|----------|----------|----------|-----------|----------|---------|-------|--|
|       |            | Sample c    | Clump Th | Uniformi | Uniformi | Marginal | Single E | Bare Nuc | Island Ch | Normal N | Mitoses | Class |  |
| 1     | 0          | 1000025     | 5        | 1        | 1        | 1        | 2        | 0        | 3         | 1        | 1       | 2     |  |
| 2     | 1          | 1002945     | 5        | 4        | 4        | 5        | 7        | 1        | 3         | 2        | 1       | 2     |  |
| 3     | 2          | 1015425     | 3        | 1        | 1        | 1        | 2        | 2        | 3         | 1        | 1       | 2     |  |
| 4     | 3          | 1016277     | 6        | 8        | 8        | 1        | 3        | 4        | 3         | 7        | 1       | 2     |  |
| 5     | 4          | 1017023     | 4        | 1        | 1        | 3        | 2        | 0        | 3         | 1        | 1       | 2     |  |
| 6     | 5          | 1017122     | 8        | 10       | 10       | 8        | 7        | 1        | 9         | 7        | 1       | 4     |  |
| 7     | 6          | 1018099     | 1        | 1        | 1        | 1        | 2        | 1        | 3         | 1        | 1       | 2     |  |
| 8     | 7          | 1018561     | 2        | 1        | 2        | 1        | 0        | 0        | 0         | 0        | 0       | 0     |  |
| 9     | 8          | 1033078     | 2        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 10    | 9          | 1033078     | 4        | 2        | 1        | 1        |          |          |           |          |         |       |  |
| 11    | 10         | 1035283     | 1        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 12    | 11         | 1036172     | 2        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 13    | 12         | 1041801     | 5        | 3        | 3        | 3        |          |          |           |          |         |       |  |
| 14    | 13         | 1043999     | 1        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 15    | 14         | 1044572     | 8        | 7        | 5        | 10       |          |          |           |          |         |       |  |
| 16    | 15         | 1047630     | 7        | 4        | 6        | 4        |          |          |           |          |         |       |  |
| 17    | 16         | 1048672     | 4        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 18    | 17         | 1049815     | 4        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 19    | 18         | 1050670     | 10       | 7        | 7        | 6        |          |          |           |          |         |       |  |
| 20    | 19         | 1050718     | 6        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 21    | 20         | 1054590     | 7        | 3        | 2        | 10       |          |          |           |          |         |       |  |
| 22    | 21         | 1054593     | 10       | 5        | 5        | 3        |          |          |           |          |         |       |  |
| 23    | 22         | 1056784     | 3        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 24    | 23         | 1057013     | 8        | 4        | 5        | 1        |          |          |           |          |         |       |  |
| 25    | 24         | 1059562     | 1        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 26    | 25         | 1065726     | 5        | 2        | 3        | 4        |          |          |           |          |         |       |  |
| 27    | 26         | 1068373     | 3        | 2        | 1        | 1        |          |          |           |          |         |       |  |
| 28    | 27         | 1068979     | 5        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 29    | 28         | 1067444     | 2        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 30    | 29         | 1070935     | 1        | 1        | 3        | 1        |          |          |           |          |         |       |  |
| 31    | 30         | 1070935     | 3        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 32    | 31         | 1071760     | 2        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 33    | 32         | 1072179     | 10       | 7        | 7        | 3        |          |          |           |          |         |       |  |
| 34    | 33         | 1074610     | 2        | 1        | 1        | 2        |          |          |           |          |         |       |  |
| 35    | 34         | 1075123     | 3        | 1        | 2        | 1        |          |          |           |          |         |       |  |
| 36    | 35         | 1079304     | 2        | 1        | 1        | 1        |          |          |           |          |         |       |  |
| 37    | 36         | 1080185     | 10       | 10       | 10       | 8        |          |          |           |          |         |       |  |
| 38    | 37         | 1081701     | 6        | 2        | 1        | 1        |          |          |           |          |         |       |  |

Figure 1

```
graph TD
    Root["Class <= 3.0  
gini = 0.825  
samples = 489  
class = 8  
value = [104, 9, 112, 115, 31, 27, 8, 56, 17, 10]"]
    Root --> Left["Normal Nucleoli <= 1.5  
gini = 0.699  
samples = 321  
class = 3  
value = [103, 0, 109, 92, 6, 3, 1, 7, 0, 0]"]
    Root --> Right["Uniformity of Cell Size <= 4.5  
gini = 0.835  
samples = 168  
class = 2  
value = [1, 9, 3, 23, 25, 24, 7, 49, 17, 10]"]
    Left --> LeftL["Sample code number <= 0.677  
gini = 0.677  
samples = 26  
class = 4  
value = [101, 97, 79, 0]"]
    Left --> LeftM["Bare Nuclei <= 0.744  
gini = 0.744  
samples = 3  
class = 8  
value = [2, 0, 12, 13, 6]"]
    Left --> LeftR["Bare Nuclei <= 0.818  
gini = 0.818  
samples = 122  
class = 2  
value = [1, 0, 1, 11, 10]"]
    LeftR --> LeftRL["Uniformity of Cell Shape <= 3.5  
gini = 0.818  
samples = 122  
class = 2  
value = [0, 9, 2, 12, 15, 12, 5, 42, 15, 10]"]
    LeftL --> LeftLL["Sample code <= 0.677  
gini = 0.677  
samples = 26  
class = 4  
value = [96]"]
    LeftL --> LeftLM["Clump <= 0.677  
gini = 0.677  
samples = 26  
class = 4  
value = [5]"]
    LeftLM --> LeftLM1["Sample code <= 0.677  
gini = 0.677  
samples = 26  
class = 4  
value = [2, 6]"]
    LeftLM1 --> LeftLM11["Sample code <= 0.677  
gini = 0.677  
samples = 26  
class = 4  
value = [0]"]
    LeftLM11 --> LeftLM111["Normal Nucleoli <= 1.5  
gini = 0.699  
samples = 321  
class = 3  
value = [1, 0]"]
    LeftLM111 --> LeftLM1111["Uniformity of Cell Size <= 4.5  
gini = 0.835  
samples = 168  
class = 2  
value = [1, 0]"]
    LeftLM1111 --> LeftLM11111["Normal Nucleoli <= 1.5  
gini = 0.699  
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    LeftLM1111111 --> LeftLM11111111["Uniformity of Cell Size <= 4.5  
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class = 2  
value = [1, 0]"]
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gini = 0.699  
samples = 321  
class = 3  
value = [1, 0]"]
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gini = 0.835  
samples = 168  
class = 2  
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gini = 0.699  
samples = 321  
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samples = 168  
class = 2  
value = [1, 0]"]
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gini = 0.699  
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gini = 0.835  
samples = 168  
class = 2  
value = [1, 0]"]
    LeftLM1111111111111111111111111111111111111111111111111111111111111111111 --> LeftLM1111111111111111111111111
```



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Dataset Decision\_Tree Performance\_Evaluation

|   | 0  | 1 | 2  | 3 | 4 | 5 | 6 | 7  | 8 | 9 |
|---|----|---|----|---|---|---|---|----|---|---|
| 0 | 14 | 0 | 31 | 3 | 0 | 0 | 0 | 0  | 0 | 0 |
| 1 | 0  | 0 | 0  | 0 | 0 | 1 | 0 | 10 | 0 | 0 |
| 2 | 10 | 0 | 36 | 5 | 0 | 2 | 0 | 1  | 0 | 0 |
| 3 | 1  | 0 | 30 | 6 | 1 | 1 | 0 | 6  | 0 | 5 |
| 4 | 0  | 0 | 1  | 2 | 0 | 2 | 0 | 4  | 0 | 0 |
| 5 | 0  | 0 | 0  | 1 | 0 | 1 | 0 | 5  | 0 | 0 |
| 6 | 0  | 0 | 0  | 0 | 0 | 0 | 0 | 2  | 0 | 0 |
| 7 | 0  | 0 | 0  | 2 | 0 | 3 | 0 | 12 | 0 | 0 |
| 8 | 0  | 0 | 0  | 1 | 0 | 1 | 0 | 8  | 0 | 1 |
| 9 | 0  | 0 | 0  | 0 | 0 | 0 | 0 | 1  | 0 | 0 |

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Dataset Decision\_Tree Performance\_Evaluation

|                 | precision           | recall              | f1-score            | support             |
|-----------------|---------------------|---------------------|---------------------|---------------------|
| 1               | 0.56                | 0.2916666666666667  | 0.3835616438356164  | 48.0                |
| 10              | 0.0                 | 0.0                 | 0.0                 | 11.0                |
| 2               | 0.3673469387755102  | 0.6666666666666666  | 0.47368421052631576 | 54.0                |
| 3               | 0.3                 | 0.12                | 0.17142857142857143 | 50.0                |
| 4               | 0.0                 | 0.0                 | 0.0                 | 9.0                 |
| 5               | 0.09090909090909091 | 0.14285714285714285 | 0.11111111111111111 | 7.0                 |
| 6               | 0.0                 | 0.0                 | 0.0                 | 2.0                 |
| 7               | 0.24489795918367346 | 0.7058823529411765  | 0.36363636363636365 | 17.0                |
| 8               | 0.0                 | 0.0                 | 0.0                 | 11.0                |
| 9               | 0.0                 | 0.0                 | 0.0                 | 1.0                 |
| 11 accuracy     | 0.32857142857142857 | 0.32857142857142857 | 0.32857142857142857 | 0.32857142857142857 |
| 12 macro avg    | 0.15631539886862747 | 0.19270728291316527 | 0.15034219005379784 | 210.0               |
| 13 weighted avg | 0.3167445887445887  | 0.32857142857142857 | 0.2834330038264532  | 210.0               |

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