

Big Data Assignment Report

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COE17B010

Steps to RUN:

Pip Install Modules –

pickle, numpy, pandas, matplotlib seaborn, plotly, tqdm, functools, mlxtend, sklearn

Run - ***main.py***

Pre-processing

File: Preprocessing.py

Methods:

1. Reading Dataset – ReadCSVFile(path)
2. Frequency Distribution – FreqDist(Data)
3. Missing Count – MissingCount(Data)
4. Cleaning – MissingClean(Data)

Remove Technique

If missing data in field (Symbol or Scientific Name) – Remove Row as this is unique data

Ignore Technique

If missing data in Synonym field – no need to do anything as it is a optional field for data

Subset Technique

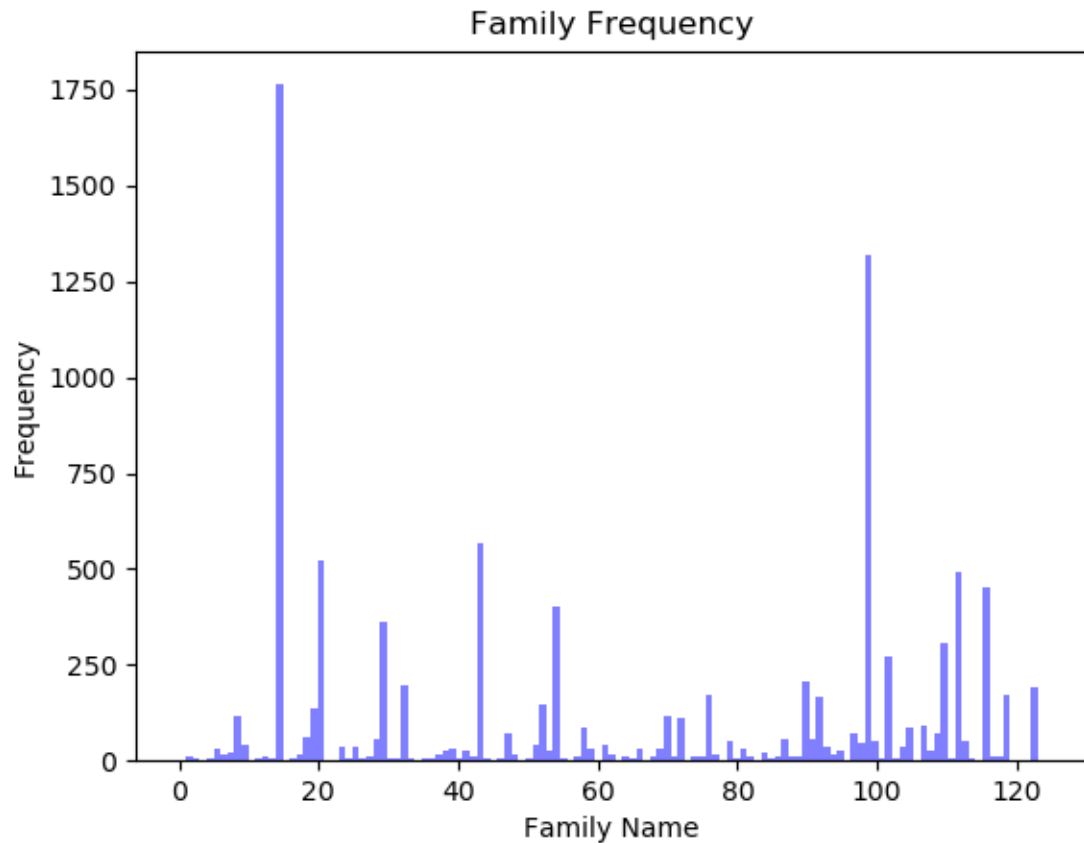
If missing data in Common Name or Family – replace smartly
Family of subspecies will be same as subspecies

So, if data missing, using Scientific Name, identify its species and use its Family to fill in missing family name (Subset Technique)

Redundant Technique – RedundantClean(Data)

If data rows are repeated, remove those redundant rows and keep only unique rows

5. Visualising – Histogram(Data) for Family Frequency



Part A

File: Algorithms.py

Algorithms:

Encoding – TransactionalEncoding(Data)

FIM

1. Apriori
2. FPGrowth

CFI

1. Charm
2. Apriori-Close

MFI

1. Pincer Search
2. Mafia

LFI

1. Apriori Based LFI
2. FPGrowth Based LFI

Part B

File: Algorithms.py

Function: RuleMining(FrequentItemsets)

Done for FPGrowth generated frequent itemsets

Confidence = 1.0

```
- Part B - Rules Mining -----
RuleSet:
<bound method NDFrame.head of
ction
2      (ALGR) (Alismataceae)      0.14      0.58      0.14      1.000000      1.724138      0.0588      inf
4      (ALTR7) (Alismataceae)      0.10      0.58      0.10      1.000000      1.724138      0.0420      inf
6      (SALA2) (Alismataceae)      0.26      0.58      0.26      1.000000      1.724138      0.1092      inf
1      (Aceraceae) (NULLVALUE)      0.20      0.38      0.10      0.500000      1.315789      0.0240      1.240000
7      (Alismataceae) (SALA2)      0.58      0.26      0.26      0.448276      1.724138      0.1092      1.341250
0      (NULLVALUE) (Aceraceae)      0.38      0.20      0.10      0.263158      1.315789      0.0240      1.085714
3      (Alismataceae) (ALGR)      0.58      0.14      0.14      0.241379      1.724138      0.0588      1.133636
5      (Alismataceae) (ALTR7)      0.58      0.10      0.10      0.172414      1.724138      0.0420      1.087500>
```

Part C

File: DecisionTree.py, BayesClassifier.py

Decision Tree done for balance-scale.csv dataset

```
Decision Tree on balance-scale dataset:

Dataset Length: 625
Dataset Shape: (625, 5)
Dataset:
class name left-weight left-distance right-weight right-distance
0      B      1      1      1      1
1      R      1      1      1      2
2      R      1      1      1      3
3      R      1      1      1      4
4      R      1      1      1      5
```

Results Using Gini Index:

Predicted values:

```
['R' 'L' 'R' 'R' 'R' 'L' 'R' 'L' 'L' 'L' 'R' 'L' 'L' 'L' 'R' 'L' 'R' 'L'
'L' 'R' 'L' 'R' 'L' 'L' 'R' 'L' 'L' 'L' 'R' 'L' 'L' 'L' 'R' 'L' 'L' 'L'
'L' 'R' 'L' 'L' 'R' 'L' 'R' 'L' 'R' 'R' 'L' 'L' 'L' 'L' 'R' 'R' 'L' 'R'
'R' 'L' 'R' 'R' 'L' 'L' 'R' 'R' 'L' 'L' 'L' 'L' 'L' 'L' 'R' 'R' 'L' 'L' 'R'
'R' 'R' 'L' 'L' 'L' 'R' 'R' 'L' 'L' 'L' 'L' 'L' 'L' 'R' 'R' 'L' 'R' 'L'
'R' 'L' 'R' 'L' 'R' 'R' 'L' 'R' 'R' 'R' 'R' 'L' 'R' 'R' 'R' 'R' 'R' 'R'
'L' 'L' 'L' 'R' 'R' 'R' 'R' 'L' 'R' 'R' 'R' 'L' 'L' 'R' 'L' 'R' 'L' 'R'
'L' 'L' 'R' 'L' 'L' 'R' 'L' 'R' 'L' 'R' 'R' 'R' 'L' 'R' 'R' 'R' 'R' 'R'
'L' 'L' 'R' 'R' 'R' 'R' 'L' 'R' 'R' 'R' 'L' 'R' 'L' 'L' 'L' 'L' 'R' 'R'
'L' 'R' 'R' 'L' 'L' 'R' 'R' 'R']
```

Confusion Matrix:

```
[[ 0  6  7]
```

```
[ 0 67 18]
```

```
[ 0 19 71]]
```

Accuracy:

73.40425531914893

C:\Users\Kausik N\AppData\Local\Programs\Python\Python38-32\lib\site-pack

Precision and F-score are ill-defined and being set to 0.0 in labels with

Report:

	precision	recall	f1-score	support
B	0.00	0.00	0.00	13
L	0.73	0.79	0.76	85
R	0.74	0.79	0.76	90
accuracy			0.73	188
macro avg	0.49	0.53	0.51	188
weighted avg	0.68	0.73	0.71	188

Results Using Entropy:

Predicted values:

```
['R' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'R' 'R' 'R' 'R' 'L' 'L' 'R' 'L' 'R' 'L'
'L' 'R' 'L' 'R' 'L' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'L' 'L'
'L' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'L' 'L' 'R' 'L' 'L' 'R' 'L' 'L'
'R' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'L' 'L' 'L' 'L' 'R' 'R' 'L' 'R' 'L'
'R' 'R' 'L' 'L' 'L' 'R' 'R' 'L' 'L' 'L' 'L' 'L' 'L' 'R' 'R' 'R' 'R' 'R'
'R' 'L' 'R' 'L' 'R' 'R' 'L' 'R' 'R' 'L' 'R' 'R' 'L' 'R' 'R' 'R' 'L' 'L'
'L' 'L' 'L' 'L' 'R' 'R' 'R' 'L' 'R' 'R' 'R' 'L' 'L' 'R' 'L' 'R' 'L' 'R'
'L' 'R' 'R' 'L' 'L' 'R' 'L' 'R' 'R' 'R' 'R' 'R' 'L' 'R' 'R' 'R' 'R' 'R'
'R' 'L' 'R' 'L' 'R' 'R' 'L' 'R' 'L' 'R' 'L' 'R' 'L' 'L' 'L' 'L' 'L' 'R'
'R' 'R' 'L' 'L' 'L' 'R' 'R' 'R']
```

Confusion Matrix:

```
[[ 0  6  7]
```

```
[ 0 63 22]
```

```
[ 0 20 70]]
```

Accuracy:

70.74468085106383

C:\Users\Kausik N\AppData\Local\Programs\Python\Python38-32\lib\site-pack

Precision and F-score are ill-defined and being set to 0.0 in labels with

Report:

	precision	recall	f1-score	support
B	0.00	0.00	0.00	13
L	0.71	0.74	0.72	85
R	0.71	0.78	0.74	90
accuracy			0.71	188
macro avg	0.47	0.51	0.49	188
weighted avg	0.66	0.71	0.68	188

Bayes Classifier done for Iris Dataset

Bayes Classifier on Iris dataset:

Results Using Bayes Classifier:

Predicted values:

```
[0 1 1 0 2 2 2 0 0 2 1 0 2 1 1 0 1 1 0 0 1 1 2 0 2 1 0 0 1 2 1 2 1 2 2 0 1
 0 1 2 2 0 1 2 1 2 0 0 0 1 0 0 2 2 2 2 2 1 2 1]
```

Confusion Matrix:

```
[[19  0  0]
 [ 0 19  2]
 [ 0  1 19]]
```

Accuracy:

95.0

Report:

	precision	recall	f1-score	support
0	1.00	1.00	1.00	19
1	0.95	0.90	0.93	21
2	0.90	0.95	0.93	20
accuracy			0.95	60
macro avg	0.95	0.95	0.95	60
weighted avg	0.95	0.95	0.95	60