OSDA Big Homework

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1 Link

Github code

https://github.com/codered-epidemy/OSDA

2 Data description

The dataset is taken from Kaggle (link). It was preprocessed by Prisha Sawhney. Original dataset is taken from UCL Library (link).

This data shows different features of mushrooms. Using this features we need to predict whether it is edible or not.

In original dataset from UCL there is 20 features. Kaggle dataset consists of 54035 rows with 8 features (all numerical). Target feature:

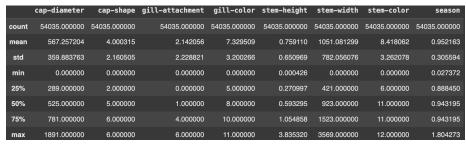
- 1 mushroom is poisonous
- 0 mushroom is edible

#	Column	Non-Null Count	Dtype	
0	cap-diameter	54035 non-null	int64	
1	cap-shape	54035 non-null	int64	
2	gill-attachment	54035 non-null	int64	
3	gill-color	54035 non-null	int64	
4	stem-height	54035 non-null	float64	
5	stem-width	54035 non-null	int64	
6	stem-color	54035 non-null	int64	
7	season	54035 non-null	float64	
dtypes: float64(2), int64(6)				

Features and their types

3 EDA

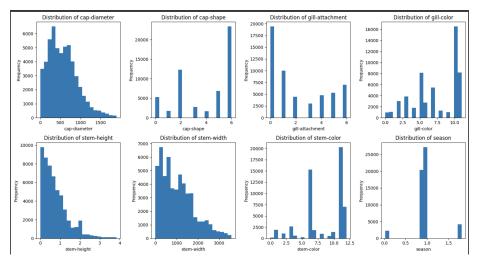
I checked NA values, get info about data (using .info()) and also used describe method.



Description of features

4 Visual DA

Here I used code from Alan's gitlab and get this distributions



Description of features

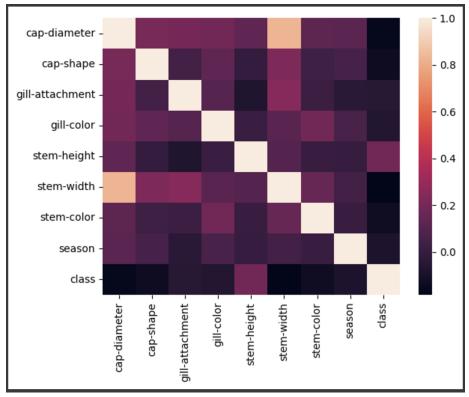
From this distributions I decided to use nominal scaling for cap-shape, gill-attachment, gill-color, stem-color, season. I also counted unique values in different features to check.

```
Feature: cap-diameter
More than 30 unique values
Feature: cap-shape
[2 6 4 0 1 5 3]
Feature: gill-attachment
[2 0 1 5 6 4 3]
Feature: gill-color
[10 5 7 9 0 3 11 8 1 6 4 2]
Feature: stem-height
More than 30 unique values
Feature: stem-width
More than 30 unique values
Feature: stem-color
[11 12 6 10 0 5 9 8 1 4 3 7 2]
Feature: season
[1.80427271 0.94319455 0.88845029 0.02737213]
```

Unique values in different features

For other features I decided to make 5 bins, divide values into them and use inter-ordinal scaling.

After that I checked correlation matrix:



Correlation matrix

Here we can see that most features do not correlate significantly with the target variable. Nevertheless, I would consider all the features.

5 Binarization

As I already mentioned, next step is binarization. For this, I applied interordinal scaling, where we apply rule "greater than or equal and less than or equal" to some threshold. I also used nominal scaling, because the amount of unique values was not so huge, so it would be appropriate for our methods to count it (taking into account time they spend on evaluations). For inter-ordinal, I took **min** and **max** values of features, divided into 5 bins and applied these bin rules to each object.

For nominal scaling I applied One Hot Encoding. After binarization I got 81 columns (features).

6 Metrics evaluation

For evaluation of the performance of the models I used:

• Accuracy (measures overall correctness of the model):

$$\frac{TP + TN}{TP + FP + FN + TN}$$

• **Precision** (measures how many of the predicted positive cases are actually positive):

$$\frac{TP}{TP+FP}$$

• **Recall** (measures the ability of the model to identify all relevant positive cases):

$$\frac{TP}{TP+FN}$$

• **F1-score** (balances precision and recall, providing a single metric when both are important.):

$$2*\frac{Precision*Recall}{Precision+Recall}$$

- **True positive** (indicates how many actual positives were correctly identified)
- False positive (indicates errors where the model predicted positive but the actual label was negative)
- False negative (indicates errors where the model missed positive cases)
- True negative (indicates how well the model identifies actual negatives)
- True Negative Rate (measures the ability of the model to correctly identify negatives):

$$\frac{TN}{TN+FP}$$

• False positive rate (measures how often the model falsely identifies negatives as positives):

$$\frac{FP}{TN+FP}$$

• False discovery rate (measures the trustworthiness of positive predictions):

$$\frac{FP}{TP+FP}$$

I also measured time of fitting and predicting.

7 Data split

I decided to split data into to equal parts and for this apply all models (Data split section in code). I suggest such variant because original data is not strongly imbalanced.

8 LazyFCA

LazyFCA is a hybrid machine learning method that combines the principles of Formal Concept Analysis with lazy learning techniques. It is designed for classification tasks where predictions are made based on the relationships between a test sample and subsets of training data, without building a full model beforehand.

LazyFCA avoids creating an exhaustive concept lattice (a hallmark of traditional FCA) and instead focuses on dynamically finding relevant patterns during the classification of each test sample. This approach leverages concepts (defined by shared attributes in the data) to classify samples in a transparent and interpretable way.

I modified Alan's code so it works faster now. The intersections (&) between the sample and X₋train subsets are computed in bulk, improving speed. Also I added verbose parameter to class.

Results:

Metric	Value
Accuracy	0.497436
Precision	0.497436
Recall	1.00000
F1	0.664384
TP	97
FP	98
FN	0
TN	0
TNR	0
FPR	1.000000
FDR	0.502564
Training time	0.000005
Prediciton time	0.490130

$9 \quad Modified Lazy FCA \\$

ModifiedLazyFCA incorporates weighted voting. This modification assigns weights to the classifiers based on their similarity to the sample being classified.

Definition of weight voting

Each classifier (positive or negative) contributes to the decision based on how

many features it shares with the sample. A higher number of overlapping features results in a higher weight, meaning that classifier is more relevant to the current sample.

How is weight computed?

For each classifier (positive or negative): Count the number of matching features with the test sample. Divide this count by the total number of features in the sample. This gives a normalized weight between 0 and 1, indicating the relevance of the classifier.

Results:

Metric	Value
Accuracy	0.656410
Precision	0.727273
Recall	0.494845
F1	0.588957
TP	48
FP	18
FN	49
TN	80
TNR	0.816327
FPR	0.183673
FDR	0.272727
Training time	0.000626
Prediciton time	0.437869

Difference between LazyFCA realizations

ModifiedLazyClassifierFCA achieves better overall performance by balancing precision and recall, leading to significantly improved accuracy and interpretability.

10 Comparison with other models

List of applied models:

- KNN
- Naive Bayes
- Logistic Regression
- SVM
- Decision Tree
- Random Forest
- XGBoost

• GBC

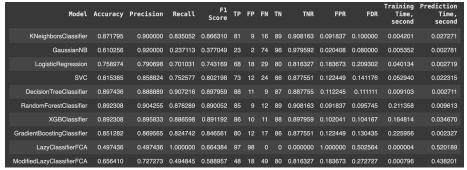
First I applied models to unbinarized data:



Metrics evaluation for unbinarized data

Here we can see that Random Forest is the best model.

Now let's apply same models to binarized data (including LazyFCA):



Metrics evaluation for binarized data

11 Conclusion

If we take a glance at the pictures above, we can find that state-of-the-art methods overcome the LazyFCA classificator (even after implementing modification). I suppose some other modification could improve LazyFCA efficiency and performance significantly. For this dataset, the best model is RandomForestClassifier (even without hyperparameter tuning).