

ML LAB WEEK 3

REPORT:

PES2UG23CS387

TICTACTOE DATASET

| 🇮🇹 OVERALL PERFORMANCE METRICS | |
|--------------------------------|-----------------|
| ===== | |
| Accuracy: | 0.8730 (87.30%) |
| Precision (weighted): | 0.8741 |
| Recall (weighted): | 0.8730 |
| F1-Score (weighted): | 0.8734 |
| Precision (macro): | 0.8590 |
| Recall (macro): | 0.8638 |
| F1-Score (macro): | 0.8613 |
| | |
| 🌳 TREE COMPLEXITY METRICS | |
| ===== | |
| Maximum Depth: | 7 |
| Total Nodes: | 281 |
| Leaf Nodes: | 180 |
| Internal Nodes: | 101 |

Roots and early splits are usually the middle squares because they decide winning conditions.

Class distribution is balanced between win (positive) and loss(negative)

Decision patterns: Tree first checks the critical winning lines

Overfitting indicators: Highly complex tree but still generalizes (87% accuracy)

NURSERY DATASET

| OVERALL PERFORMANCE METRICS | |
|-----------------------------|-----------------|
| ===== | |
| Accuracy: | 0.9867 (98.67%) |
| Precision (weighted): | 0.9876 |
| Recall (weighted): | 0.9867 |
| F1-Score (weighted): | 0.9872 |
| Precision (macro): | 0.7604 |
| Recall (macro): | 0.7654 |
| F1-Score (macro): | 0.7628 |
| | |
| TREE COMPLEXITY METRICS | |
| ===== | |
| Maximum Depth: | 7 |
| Total Nodes: | 952 |
| Leaf Nodes: | 680 |
| Internal Nodes: | 272 |

Root splits have attributes like *parents*, *has_nurs*

Class Distribution is very imbalanced because most are **not_recom**, while the other values are rare.

Decision Patterns: Longer and more complex because of many classes and imbalance of values.

Overfitting indicators: Accuracy very high, but recall is lower meaning tree struggles with minority classes. Therefore, it is *overfitting* to the **not_recom** class. (Deeper branches to cover rare cases)

MUSHROOM DATASET

| 📊 OVERALL PERFORMANCE METRICS | | |
|-------------------------------|--------|-----------|
| ===== | | |
| Accuracy: | 1.0000 | (100.00%) |
| Precision (weighted): | 1.0000 | |
| Recall (weighted): | 1.0000 | |
| F1-Score (weighted): | 1.0000 | |
| Precision (macro): | 1.0000 | |
| Recall (macro): | 1.0000 | |
| F1-Score (macro): | 1.0000 | |
| | | |
| 🌳 TREE COMPLEXITY METRICS | | |
| ===== | | |
| Maximum Depth: | 4 | |
| Total Nodes: | 29 | |
| Leaf Nodes: | 24 | |
| Internal Nodes: | 5 | |

Odor and spore-print-colour dominate the root and early splits.

Class Distribution: Fairly balanced

Decision Patterns: Very simple rules, single feature splits

Overfitting indicators: Achieves 100% accuracy, not overfitting because it is separable by a few categorical features.

Algorithm Performance:

Highest Accuracy: Mushroom dataset (100%) because features like odor are perfect predictors.

Effect of Dataset Size: Larger datasets (Nursery: 12960, Mushroom: 8124) are more stable and accurate models. Smaller dataset (Tictactoe: 958) leads to lower accuracy, more variance.

Role of Features: More features and multi-valued attributes (Nursery, Mushroom) give richer splits. Tictactoe has only 9 simple attributes, so tree is more complex to capture patterns.

Data Characteristics Impact

Class Imbalance: Nursery has imbalance because tree tends to predict not_recom often; affects macro precision/recall because of bias in predictions. Mushrooms is mostly balanced, so no bias.

Binary vs Multi-valued: Multi-valued categorical features (Mushroom, Nursery) help trees split cleanly and improve accuracy. Binary (Tictactoe outcomes) makes patterns harder to learn.

Practical Applications

Tictactoe: Game AI, strategy modelling, rule-based systems. Advantage: interpretable strategies (which moves lead to wins).

Nursery: Admission or resource allocation systems. Advantage: clear decision rules for fairness/priority.

Mushroom: Food safety and toxicology. Advantage: easily interpretable Eg. If odor = foul, poisonous.

Improving Performance:

Tictactoe: Use pruning or Random Forests to reduce overfitting and capture general patterns.

Nursery: Handle class imbalance (resampling, weighted loss).

Mushroom: Already perfect no improvements needed

CONCLUSION

Mushrooms dataset was **clean** and **separable**

simple, perfect model

Nursery dataset was **large** and **imbalanced**

very accurate but careful handling of minority classes

Tictactoe dataset was **small** and **binary**

very complex tree lower accuracy