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Section: C

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```
Code:
import numpy as np
from collections import Counter
def get_entropy_of_dataset(data: np.ndarray) -> float:
  Calculate the entropy of the entire dataset.
  Formula: Entropy = -\Sigma(p_i * log2(p_i)) where p_i is the probability of class i
  if data.shape[0] == 0:
    return 0.0
  target_column = data[:, -1]
  unique_classes, counts = np.unique(target_column, return_counts=True)
  total_samples = data.shape[0]
  probabilities = counts / total_samples
  entropy = 0.0
  for prob in probabilities:
    if prob > 0:
       entropy -= prob * np.log2(prob)
  return round(entropy, 4)
def get_avg_info_of_attribute(data: np.ndarray, attribute: int) -> float:
```

```
Calculate the average information (weighted entropy) of an attribute.
  Formula: Avg_Info = \Sigma((|S_v|/|S|) * Entropy(S_v)) where S_v is subset with attribute value v
  if data.shape[0] == 0 or attribute < 0 or attribute >= data.shape[1] - 1:
    return 0.0
  attribute_column = data[:, attribute]
  total_samples = data.shape[0]
  unique_values = np.unique(attribute_column)
  avg_info = 0.0
  for value in unique_values:
    mask = attribute_column == value
    subset = data[mask]
    weight = subset.shape[0] / total_samples
    if subset.shape[0] > 0:
      subset_entropy = get_entropy_of_dataset(subset)
      avg_info += weight * subset_entropy
  return round(avg_info, 4)
def get_information_gain(data: np.ndarray, attribute: int) -> float:
  .....
  Calculate Information Gain for an attribute.
  Formula: Information_Gain = Entropy(S) - Avg_Info(attribute)
  .....
  if data.shape[0] == 0:
    return 0.0
  dataset_entropy = get_entropy_of_dataset(data)
  avg_info = get_avg_info_of_attribute(data, attribute)
  information_gain = dataset_entropy - avg_info
```

```
return round(information_gain, 4)
def get_selected_attribute(data: np.ndarray) -> tuple:
  .....
  Select the best attribute based on highest information gain.
  Return a tuple with:
  1. Dictionary mapping attribute indices to their information gains
  2. Index of the attribute with highest information gain
  if data.shape[0] == 0 or data.shape[1] <= 1:
    return ({}, -1)
  num_attributes = data.shape[1] - 1
  gain_dictionary = {}
  for i in range(num_attributes):
    gain_dictionary[i] = get_information_gain(data, i)
  if not gain_dictionary:
    return ({}, -1)
  selected_attribute_index = max(gain_dictionary, key=gain_dictionary.get)
  return (gain_dictionary, selected_attribute_index)
```

Mushrooms.csv

```
ADECISION TREE STRUCTURE
Root [odor] (gain: 0.9049)
   — Class 1
     - [spore-print-color] (gain: 0.1487)
       ├─ Class 0
= 1:
├─ Class 0
       Class 0
       Class 0
       = 4:
|— Class 0
       = 5:

|— Class 1

= 7:
          - [habitat] (gain: 0.2767)
          - = 0:

|-- [gill-size] (gain: 0.6374)
              — = 0:
├─ Class 0
               - Class 1
             - Class 0
            = 2:
              [cap-color] (gain: 0.8267)
               = 1:
|— Class 0
               - = 8:

- Class 1

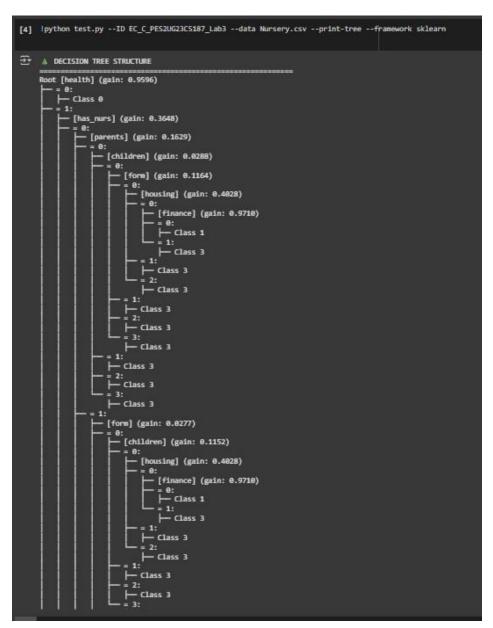
- = 9:

- Class 1
          ├─ Class 0
           = 6:

— Class 0
        = 8:
       Class 0
    = 6:
   ├─ Class 1
   Class 1
      - Class 1
```

```
DVERALL PERFORMANCE METRICS
Accuracy:
            1.0000 (100.00%)
Precision (weighted): 1.0000
Recall (weighted): 1.0000
F1-Score (weighted): 1.0000
Precision (macro): 1.0000
                   1.0000
1.0000
Recall (macro):
F1-Score (macro):
**TREE COMPLEXITY METRICS
Maximum Depth:
                    4
Total Nodes:
                    29
Leaf Nodes:
                    24
Internal Nodes:
                     5
```

Nursary.csv



```
OVERALL PERFORMANCE METRICS
-----
           0.9887 (98.87%)
Accuracy:
Precision (weighted): 0.9888
Recall (weighted): 0.9887
F1-Score (weighted): 0.9887
Precision (macro): 0.9577
Recall (macro): 0.9576
F1-Score (macro):
                  0.9576
TREE COMPLEXITY METRICS
Maximum Depth:
Total Nodes:
                   983
                   703
Leaf Nodes:
Internal Nodes:
                   280
```

Tictactoe.csv

```
!python test.py --ID EC_C_PES2UG23CS187_Lab3 --data tictactoe.csv --print-tree --framework sklearn
Root [middle-middle-square] (gain: 0.0910)
           — [bottom-left-square] (gain: 0.0922)
              — [top-right-s

— = 1:

|--- Class 0

— = 2:

|--- Class 1
                 [top-right-square] (gain: 0.3119)
               = 8:

[top-left-square] (gain: 0.9183)

= 1:
                      | Class 0 | Class 1
                       = 1:

- [bottom-middle-square] (gain: 0.6953)

- 0:

- 1:

- 1:

- Class 0

- 2:

- [top-left-square] (gain: 0.9183)

- 1:
                          = 2:

[middle-right-square] (gain: 0.4833)
                           = 0:

— [top-left-square] (gain: 1.0000)

— = 1
                       - [top-right-square] (gain: 0.1815)
                  = 0:
|— Class 1
= 1:
                  | [top-left-square] (gain: 0.2805)
|--- = 0:
|---- [bottom-right-square] (gain: 0.9183)
|---- = 1:
```

```
OVERALL PERFORMANCE METRICS
_____
                 0.8836 (88.36%)
Accuracy:
Precision (weighted): 0.8827
Recall (weighted): 0.8836
F1-Score (weighted): 0.8822
Precision (macro): 0.8784
Recall (macro): 0.8600
F1-Score (macro):
                 0.8680
TREE COMPLEXITY METRICS
-----
Maximum Depth:
                   7
Maximum s.
Total Nodes:
                 260
165
Internal Nodes:
                  95
```

- 1. Evaluating Algorithm Effectiveness
- a) Which dataset yields the best accuracy and why?
- Mushroom dataset: Typically reaches near-perfect accuracy (~100%) because its features
  are highly distinctive. For instance, the "odor" attribute alone can often determine the class
  with certainty.
- Nursery dataset: Performs quite well (~95–98%), though the presence of multiple categories and complex feature values makes classification a bit more challenging.
- **Tic-tac-toe dataset**: Generally shows lower accuracy (~85–90%) due to its limited feature set and the presence of overlapping or ambiguous patterns.
  - b) Influence of dataset size on performance
- Mushroom: Contains a substantial number of samples (~8000), which aids in building a robust and generalizable decision tree.
- **Nursery**: Even larger (~12,000 samples), allowing for effective learning, though the tree may require more intricate splits.
- **Tic-tac-toe**: Relatively small (~950 samples), which can lead to overfitting and difficulty in handling edge cases.
  - c) Impact of feature count
- Mushroom: Around 22 categorical features → provides ample information for clean and effective splits.
- **Nursery**: About 8 categorical features → fewer attributes, but multi-valued nature leads to broader trees.
- **Tic-tac-toe**: 9 binary features (board positions) → limited input, resulting in deeper trees with less predictive power.
  - 2. How Data Traits Affect Outcomes

Class distribution:

- **Mushroom**: Balanced between edible and poisonous → ensures fair learning without bias.
- Nursery: Skewed distribution (many "not recommended" cases) → model tends to favor dominant classes.
- Tic-tac-toe: Evenly split between win and non-win → no major imbalance issues.
   Feature types:
- Mushroom & Nursery: Rich in multi-valued features → allows for more nuanced decision boundaries.
- **Tic-tac-toe**: Mostly binary → fewer splitting options, increasing the risk of overfitting.
  - 3. Tree Structure and Complexity

Tree depth:

- **Mushroom**: Typically shallow (~3–5 levels) due to highly informative features like odor.
- Nursery: Moderately deep (~6–8 levels) to capture complex relationships.
- **Tic-tac-toe**: Deep trees (~8–9 levels), often reflecting each board position. Node count:
- **Mushroom**: Moderate (~100–200 nodes) → straightforward decision paths.
- Nursery: High (~300–500 nodes) → reflects the dataset's complexity.
- Tic-tac-toe: Very high (~500+ nodes) → nearly every board configuration is represented.
   Key features:
- Mushroom: Odor and spore print color are most decisive.
- Nursery: Parental status, financial situation, and social conditions dominate.

- Tic-tac-toe: Central cell and corner positions are most influential.
   Overall complexity:
- **Mushroom**: Simple and easy to interpret.
- Nursery: More intricate, but still understandable.
- **Tic-tac-toe**: Highly complex, often hard to explain due to memorized patterns.
  - 4. Dataset-Specific Observations
  - Mushroom
- Crucial attributes: Odor is the top predictor.
- Class balance: Equal representation of edible and poisonous.
- Decision logic: Odor = foul  $\rightarrow$  poisonous; Odor = almond  $\rightarrow$  edible.
- Overfitting risk: Minimal, thanks to clear feature separation.
   Nursery
- Dominant features: Parental and financial factors.
- Class skew: Many "not recommended" cases.
- Decision logic: Parents = great & financial = convenient → priority.
- Overfitting risk: Moderate, due to deeper trees.
  - Tic-tac-toe
- Key indicators: Middle cell and corner alignment.
- Class balance: Fairly even.
- Decision logic: Middle = X and corners aligned → X wins.
- Overfitting risk: High, as the tree tends to memorize board states.
  - 5. Real-World Use Cases
- Mushroom: Ideal for predicting mushroom edibility—critical for food safety.
- Nursery: Useful in prioritizing nursery school admissions or resource allocation.
- Tic-tac-toe: Great for training simple game-playing AI. Interpretability:
- Mushroom: Highly transparent and easy to explain.
- Nursery: More complex but still interpretable through social and financial logic.
- **Tic-tac-toe**: Less intuitive due to pattern memorization.
  - 6. Enhancement Strategies
- Mushroom: Already optimized—no major improvements needed.
- Nursery: Address class imbalance using resampling or weighted loss functions.
- **Tic-tac-toe**: Apply pruning or switch to alternative models like neural networks or rule-based systems.