Introduction to Artificial Intelligence



COMP307/AIML420 Decision Tree Learning Methods

Dr Andrew Lensen
Andrew.Lensen@vuw.ac.nz



How do **humans** make decisions?

- Have a think about the last decision you made...
- What factors did you take into account?
- Can you make these into a set of rules?



How can we make this into an algorithm?

• It would be really useful to teach computers to "think" in this way

Allows computers to automatically make decisions for us

• For example, "Should this credit card purchase be approved?"

Today we will discuss decision trees, which do just this!

A real-world example:

 Which habitat is best for raising capybara offspring?



Table 1. Decision rules for allocating values of forage, shelter and rest and protection for offspring for different categories in which plant species were grouped according to criteria of Quintana and Bolkovic (unpublished data 2014).

| Ecological requirements | Plant category | | | Mean size | Value | |
|-------------------------|----------------------|--|--------|-----------------------|--------|---|
| | Aquatic | Oplimenopsis Luziola peruv Leersia hexandr | | All | High | 3 |
| | | All other | | All | Low | 1 |
| Forage | Broadleaf herbs | | | All | Low | 1 |
| | Sedges | | | Shorter than 1 m | High | 3 |
| | | | | Taller than 1 m | Medium | 2 |
| | Terrestrial grasses | | | All | High | 3 |
| | Shrubs | | | All | Null | 0 |
| | Trees | | | All | Null | 0 |
| | Aquatic | | | All | Medium | 2 |
| | | | | Shorter than 0.4 m | Null | 0 |
| | Herbaceous | (broadleaf h | nerbs, | Between 0.4 and 0.8 m | Low | 1 |
| Shelter | sedges, and grasses) | | | Between 0.8 and 1.2 m | Medium | 2 |
| | | | | Taller than 1.2 m | High | 3 |
| | Shrubs | | | All | High | 3 |
| | Trees | | | All | Medium | 2 |
| | Aquatic | | | All | High | 3 |
| | Herbaceous | (broadleaf h | nerbs, | Shorter than 1 m | Low | 1 |
| Rest and | sedges, and g | rasses) | | Taller than 1 m | High | 3 |
| protection for | | | | Shorter than 0.5 m | Low | 1 |
| offspring | Shrubs | | | Between 0.5 and 1 m | Medium | 2 |
| | | | | Taller than 1 m | High | 3 |
| | Trees | | | All | High | 3 |

Schivo, Facundo, et al. "A Habitat Suitability Model for Capybara (Hydrochoerus Hydrochaeris) at Its Core Area in Argentina." Tropical Conservation Science, Mar. 2015, pp. 150–168, doi:10.1177/194008291500800113.

Outline

What is a decision tree?

How do we create a decision tree given a set of instances (inputs)?

How can we use impurity to measure a decision tree node's quality?

Using DTs for multi-class classification or with other purity measures

What is a Decision Tree?

- A symbolic classifier which uses a series of decisions/rules to make a classification
 - One "big" rule to make an overall decision
 - Easy to interpret?
- Decision tree learning is a method for creating decision trees
 - One of the first classification learning methods in AI!

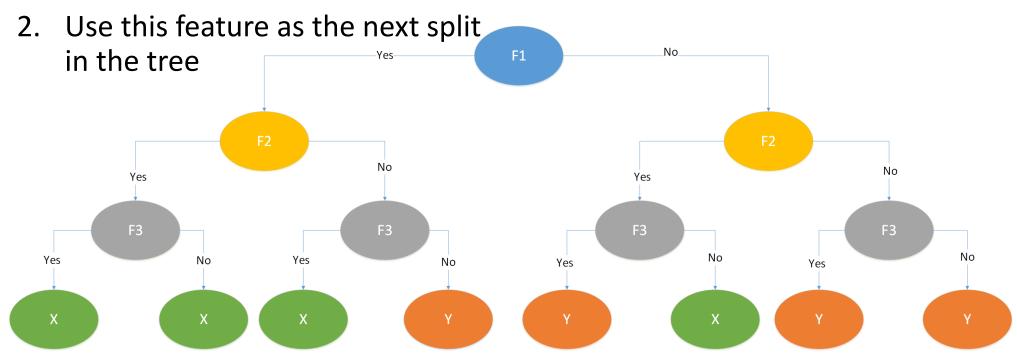
An example decision tree

 Decide whether or not a habitat Foliage is good for capybara breeding -Large plants Little/none Location Location Seashore Seashore Inland Inland Shrubs? Good Good Bad No Yes Bad Good

Constructing Decision Trees

- Easy to build a naïve (full) decision tree:
 - 1. Pick one feature at a time

- Repeat until all features used
- 4. Label each leaf with the corresponding class.



What are the problems with this?

- No learning is taking place! Just remembering the inputs
 - Too specific...not likely to work well on future (unlabelled) inputs

How can we do better?

A smaller decision tree:

- Capture the common patterns across the inputs
- Discover some underlying rules which are more generic

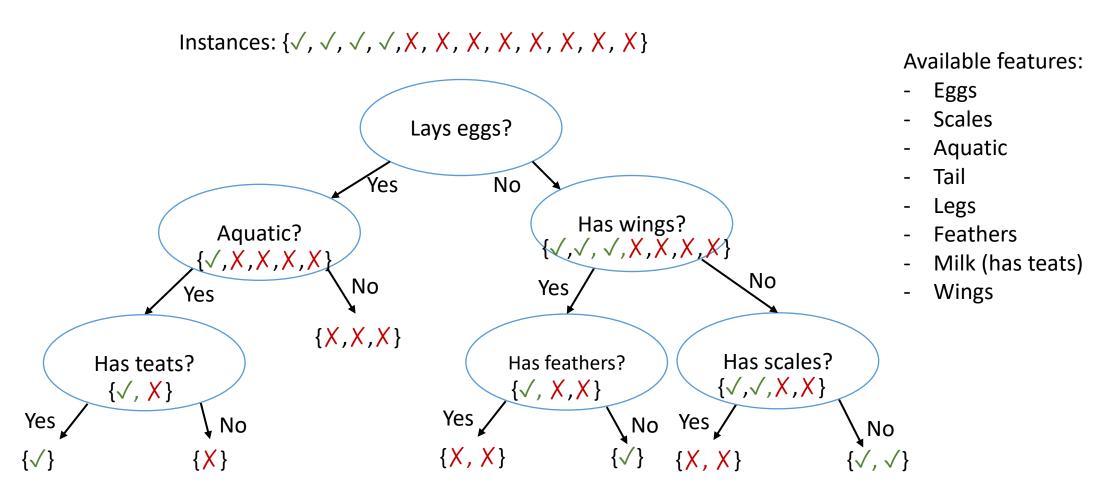
A better algorithm

Input: a set of instances defined by their feature values

Output: a decision tree classifier which performs classification

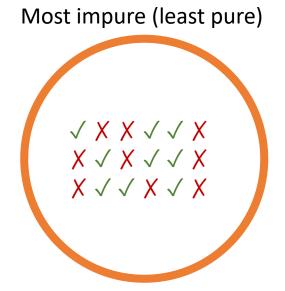
- 1. Compute if the set of instances is *pure* enough if so, then stop!
- 2. Select the best (unused) feature to use as the next node the one which would give the highest *purity*
- 3. Split the dataset into two sub-sets according to the chosen node
- 4. Recurse on each of the two sub-sets

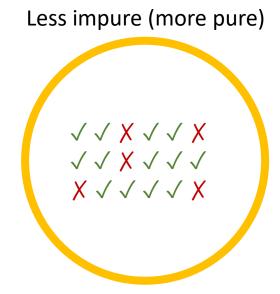
Building a DT: is the animal a mammal?

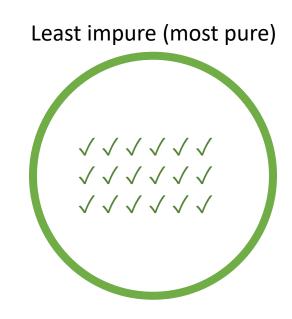


Measuring purity

- How do we measure the purity of a node?
 - Many different ways! Probability-based, information theory-based, ...
 - More pure = can predict the class more confidently
 - Often easier to talk about impurity







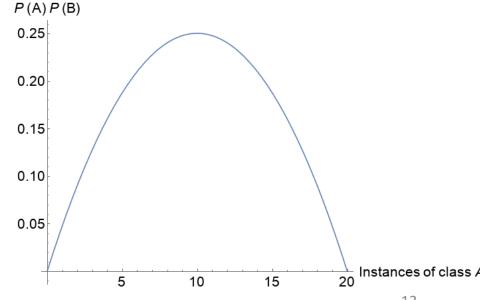
A probability-based impurity measure

If we have class A and B, then we measure a node's impurity using P(A)P(B), where there are m instances of class A and n of class B:

Impurity =
$$P(A)P(B) = \frac{m}{m+n} \times \frac{n}{m+n} = \frac{mn}{(m+n)^2}$$

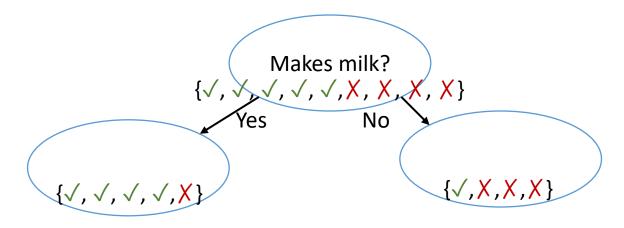
• For example, if we have 20 instances:

This measure is smooth



Choosing the next feature split

- When we choose a feature to be the next node, we create two children nodes, which each have their own impurity
- How do we combine these two impurity values?



Impurity =
$$\frac{4}{1+4} \times \frac{1}{1+4}$$
$$= 0.16$$

Impurity =
$$\frac{1}{1+3} \times \frac{3}{1+3}$$

= 0.1875

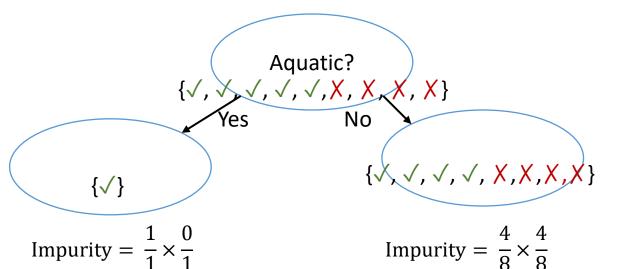
The simplest way: take the average of the two!

Average Impurity =
$$\frac{0.16+0.1875}{2}$$

= 0.17375

What is wrong with this?

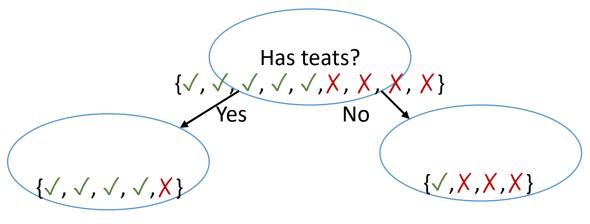
Consider this example:



Average Impurity =
$$\frac{0+0.25}{2}$$

= 0.125

But wasn't this a better split?:



Impurity =
$$\frac{4}{1+4} \times \frac{1}{1+4}$$
$$= 0.16$$

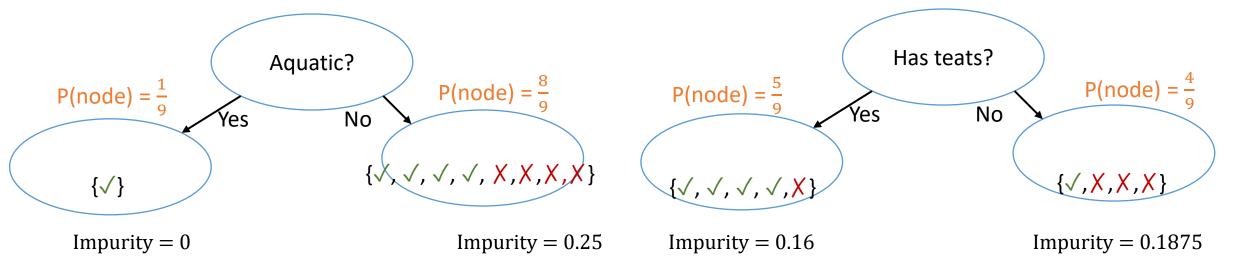
Impurity =
$$\frac{1}{1+3} \times \frac{3}{1+3}$$

= 0.1875

Average Impurity = 0.17375!

The solution: weighted average impurity

• If we weight a child's impurity by how many instances it has, then the measure is not biased to uneven splits



Weighted average impurity =
$$\frac{1}{9} \times 0 + \frac{8}{9} \times 0.25$$

 ≈ 0.222

Weighted average impurity =
$$\frac{5}{9} \times 0.16 + \frac{4}{9} \times 0.1875$$

 ≈ 0.172

16

Weighted average impurity =
$$\sum_{i} P(node_i) \times Impurity(node_i)$$

Advanced topics: More than two classes?

- Impurity = P(A)P(B) works well for two classes
 - What about three? Impurity = P(A)P(B)P(C)
 - 10? Impurity = P(A)P(B)P(C)P(D)P(E) ...
 - Any problem?

Alternatives:

- Gini impurity: $Gini(X) = \sum_{i=1}^{C} P(i) \sum_{j \neq i} P(j) = 1 \sum_{i=1}^{C} P(i)^2$
- Entropy: $H(X) = -\sum_{i=1}^{C} P(i) \log_2 P(i)$
- These often give very similar results!

Advanced topics: Numerical features (1)

- Many features are numerical, e.g. height, tail length, ...
- How can our DT algorithm cope with this?

| Instance # | Height (cm) | Tail length (cm) | Can fly? | Class |
|------------|-------------|------------------|----------|---------|
| 1 | 170 | 0 | No | Mammal |
| 2 | 19.5 | 1.1 | Yes | Bird |
| 3 | 90 | 4 | No | Bird |
| 4 | 15 | 30 | Yes | Fish |
| 5 | 4 | 70 | No | Reptile |

Advanced topics: Numerical features (2)

- Sort the feature we're considering
- Choose split points based on class boundaries
- Compute impurity for each split, choose best one

| Instance # | Height (cm) | Tail length (cm) | Can fly? | Class | |
|------------|-------------|------------------|----------|---------|---------|
| 5 | 4 | 70 | No | Reptile | 115 000 |
| 4 | 15 | 30 | Yes | Fish | <15cm |
| 2 | 19.5 | 1.1 | Yes | Bird | <19.5cm |
| 3 | 90 | 4 | No | Bird | |
| 1 | 170 | 0 | No | Mammal | <170cm |

Advanced topics: Overfitting

• The larger our tree, the more it will overfit, as it becomes too specific to the training set

How can we avoid this? Stop splitting earlier

 Need to be careful we don't stop too soon – otherwise nodes will not be pure enough → worse performance!

Some approaches:

- Stop at a given impurity threshold
- Stop at a pre-defined max depth
- Use pruning to simplify a completed tree

