## Computer Science 477

## Continuous Attributes

Lecture 9

### TIDIDT Constraints

- TIDIDT Requires categorical attributes
- Can take individual values 6.3, 7.2, 8.3, 9.2 as categorical
- (For reasons discussed), over splits the training data □ Large number of subsets, each with few instances.
- **Nore common to separate into non-overlapping subsets:**

- Discretization Further Examples<br>
Convert age to infant, child, young adu ■ Convert age to infant, child, young adult, adult, middle-aged, old
- Convert height to very short, short, medium, tall, very\_tall
- **Length, ranging from 0.3 to 6.6, inclusive:**
- **Divide into equal width intervals** 
	- $\Box$  0.3  $\leq length$   $<$  2.4
	- $\Box$  2.4  $\leq$  length  $<$  4.5
	- $\Box$  4.5  $\leq$  length  $\leq$  6.6

### Equal Width Intervals

- Number of ranges arbitrary
	- □ 3, not 4, not 12
- **Perhaps, many, of the values are in a narrow range such** as 2.35 to 2.45
	- $\Box$  A rule involving a test on length  $\leq$  2.4 would include instances where length is say 2.39999 and exclude those where length is 2.40001.
	- □ Unlikely that there is any real difference between those values, especially if they were all measured imprecisely by different people at different times.
	- $\Box$  If there were no values between say 2.3 and 2.5, a test such as  $length < 2.4$  reasonable.

Equal Frequency Intervals

- **Divide length into three ranges,** 
	- Same number of instances in each of the three ranges:
	- $\Box$  0.3  $\leq$  length  $<$  2.385
	- $\Box$  2.385  $\leq length < 3.0$
	- $\Box$  3.0  $\leq$  length  $\leq$  6.6

### ■ Same problem at cut points,

 length of 2.99999 treated differently from one of 3.00001

### **Oversensitivity**

- Whichever cut points are chosen there will always be a potential problems with values that fall just below a cut point being treated differently from those that fall just above for no principled reason.
- I Ideally, find 'gaps' in the range of values.
- If in the *length* there are many values from 0.3 to 0.4 with the next smallest value being 2.2, a test such as length < 1.0 would avoid problems around the cut point
	- $\Box$  No instances (in the training set) with values close to 1.0.
	- $\Box$  The value 1.0 is y arbitrary and a different cut point, e.g. 1.5 could be chosen
	- □ Unfortunately the same gaps may not occur in unseen test data. If there were values such as 0.99, 1.05, 1.49 and 1.51 in the test data
	- □ Choice of 1.0 or 1.5 could be of critical importance.

### Problems

**Equal width intervals and the equal frequency** intervals take no account of the classifications when determining where to place the cut points

**One solution:** 

Local versus Global Discretization

### Adding Local Discretization to TDIDT

- Convert continuous attributes to a categorical ones at each stage of the process □ (e.g. at each node of the decision tree).
- Approach 1: convert at each step attributes according to previously noted methods, namely,
	- □ Equal width
	- □ Equal frequency

## Approach 2

- At each node to convert each continuous attribute to a number of alternative categorical attributes.
- **Example: if continuous attribute A has values** −12.4, −2.4, 3.5, 6.7 and 8.5
	- □ (each possibly occurring several times)
- Test  $A < 3.5$  splits the training data into two parts
- Equivalent to a kind of categorical attribute with two possible values
	- **D** True and false.
- **Pseudo-attribute**

### Pseudo-attributes

**Attribute A with n distinct values**  $v_1, v_2, \ldots, v_n$  **there are**  $n-1$  possible corresponding pseudo-attributes

 $\Box A \leq v_2, A \leq v_3, \ldots, A \leq v_n$ 

- If one of the pseudo-attributes,  $Age < 27.3$ , is selected at a node, we can consider the continuous attribute Age as having been discretized into two intervals with cut point 27.3.
- **This is a local discretization which does not lead to the** continuous attribute itself being discarded.
- Hence there may be a further test such as  $Age < 14.1$ at a lower level in the 'yes' branch descending from the test  $Age < 27.3$ .

### Algorithmically

- **For each continuous attribute A** 
	- a) Sort the instances into ascending numerical order.
	- $\Box$  b) If there are n distinct values  $v_{1}$ ,  $v_{2}$ , ...,  $v_{n}$ , calculate the values of information gain (or of GINI index or other measure) for each of the  $n-1$  corresponding pseudo-attributes  $A \leq$  $v_2, A < v_3, ..., A < v_n$ . .
	- c) Find which of the  $n-1$  attribute values gives the largest value of information gain (or optimizes some other measure).
	- If this is  $v_i$  return the pseudo-attribute  $A < vi$ , and the value of the corresponding measure.
- Calculate the value of information gain (or other measure) for any categorical attributes.
- Select the attribute or pseudo-attribute with the largest value of information gain (or which optimizes some other measure).

# Pseudo-attributes - Information Gain<br>
Finne Stages

### **Three stages**

- **First: Count the number of instances with** each of the possible classifications in the part of the training set under consideration at the node.
- Values do not depend on which attribute is subsequently processed and so only have to be counted once at each node of the tree.

- Pseudo-attributes Stage Two<br>• Work through the continuous attributes one by c ■ Work through the continuous attributes one by one.  $\Box$  Call attribute  $Var$
- **Consider all possible pseudo-attributes**  $Var <$ X where X is one of the values of  $Var$ 
	- $\Box$  In the part of the training set under consideration at the given node.
- Call the values of attribute  $Var$  candidate cut points.
- Call the largest value of measure maxmeasure and the value of  $X$  that gives that largest value the cut point for attribute  $Var$ .

- Pseudo-Attributes Stage Three<br> **Example: Having found the value of** *maxmeasure* **(a Having found the value of maxmeasure (and** the corresponding cut points)
- Find the largest and then compare it with the values of the measure obtained for any categorical attributes to determine which attribute or pseudo-attribute to split on at the node.

Example – Golf Data Set<br>
Count the number of instances with each of the po ■ Count the number of instances with each of the possible classifications.



9 play and 5 don't play, a total of 14.

- **Process each of the continuous attributes in turn (Stage 2).** 
	- □ Two: temperature and humidity.
	- □ Illustrate Stage 2 using attribute temperature

- Example Stage 2<br>
Sort attribute value in ■ Sort attribute value in ascending order
	- □ Construct a two-column table
	- Attribute value and classification
	- □ Sorted instances table
- **Twelve distinct values**



### Processing Sorted Instance Table

- $n$  instances and rows in sorted instances numbered 1 to  $n$
- **Nork through the table from bottom to top** 
	- □ Accumulate a count of the number of instances of each classification
- **As each row is processed it attribute value is compared with** the value for the row below **□** Accumulate a count of the number of instances of each<br>classification<br>As each row is processed it attribute value is compared with<br>the value for the row below<br>**□** If larger, treat as candidate cut point<br>**□** Value of me
	- □ If larger, treat as candidate cut point
	- □ Value of measure is computed using the "frequency table method"
- **Algorithm returns** *maxmeasure* and *cutvalue* 
	- whatever).
	- □ Cutvalue is the value is the attribute value that currently maximizes the maxmeasure

# Algorithm - Processing Sorted Instance Table<br>Algorithm for Processing a Sorted Instances Table<br>Set count of all classes to zero

Set maxmeasure to a value less than the smallest possible value of the measure used

```
for i=1 to n-1 {
  increase count of class(i) by 1
  if value(i) < value(i + 1)(a) Construct a frequency table for pseudo-attribute
        Var < value(i + 1)(b) Calculate the value of measure
    (c) If measure > maxmeasure {
         maxmeasure=measure
         cutvalue = value(i + 1)
```
### Processing the sorted instance table

- golf training set and continuous attribute temperature
- Temperature 64 and class play.
- Increase the count for class play to 1.
- Count for class *don't play* is zero.
- Temperature is less than that for the next instance
- So proceed to construct a frequency table for the pseudo-attribute temperature < 65





Processing second row of sorted instance table

- **Temperature of 68, class don't play**
- **Create new frequency table** 
	- □ Update class count(s)
	- □ Column totals





- Sort items according to continuous attribute values into ascending numerical order
- Construct a frequency table giving the number of occurrences of each distinct value of the attribute for each possible classification



- Interpret each row not just as a single attribute
	- □ As an *interval*, i.e. a range of values
	- □ starting at the value, continuing up to but excluding the value given in the row below.
	- □ Row labelled 1.3 corresponds to the interval  $1.3 \leq A \leq 1.4$ . indicate the lowest number in the range of values included in that interval. The



**Figuency table could be augmented by an** additional column showing the interval corresponding to each classification



- ChiMerge systematically applies statistical tests to combine pairs
- **Does not merge intervals that are statistically different**
- Implicitly, if a pair is merged if it doesn't modify outcome, classification
- $\blacksquare$  For each pair, tests the hypothesis

**Hypothesis** 

The class is independent of which of the two adjacent intervals an instance belongs to.

If the hypothesis is confirmed, intervals are merged

## Statistical test:  $\chi^2$

- 
- $\int \chi^2$  test for independence
- For each pair of adjacent rows, construct a contingency table:



**The 'row sum' (right-hand column) and the 'column sum'** 

### ■ Correspond (respectively) to

- **Q** Number of instances for each value of A (i.e. with their value of attribute in the corresponding interval)
- **□** Number of instances in each class for both intervals combined.

### Use of the  $\chi^2$  statistic

- $\rightarrow \chi^2$  value is then compared with a threshold value T
	- **Depends on the number of classes and**
	- □ The level of statistical significance required.
- **•** (Here) use a significance level of 90%
	- Gives a threshold value of 4.61.
- If we assume that the classification is independent of which of the two adjacent intervals an instance belongs to, there is a 90% probability that  $\chi^2$  will be less than 4.61.
- If  $\chi^2$  is less than 4.61 the hypothesis of independence is supported at the 90% significance level
	- □ The two intervals are merged.

### Calculating the Expected Values and  $\chi^2$

- **For a given pair of adjacent rows (intervals) the value of**  $\chi^2$  is calculated using
	- The 'observed' and 'expected' frequency values
	- □ For each combination of class and row.
- There are three classes so there are six such combinations.
- Observed frequency value, denoted by O, is the frequency that actually occurred.
- **Expected value E** is the frequency value that would be expected to occur by chance
	- Given the assumption of independence

### Calculating the Expected Values and  $\chi^2$

- Row is *i* and the class is *j*, then let the total number of instances in row *i* be rowsum, and let the total number of occurrences of class  $j$  be colsum<sub>i</sub>.
- The grand total number of instances for the two rows combined be sum.
- **Assuming the hypothesis that the class is independent of** which of the two rows an instance belongs, calculate the expected number of instances in row *i* for class *j* thus: follows.
- **There are a total of colsum** occurrences of class *j* in the two intervals combined,
- So class *j* occurs a proportion of  $\frac{cousum_j}{cous_j}$  the time.

### Calculating the Expected Values and  $\chi^2$

- There are a total of colsum<sub>i</sub> occurrences of class *j* in the two intervals combined,
- So class *j* occurs a proportion of  $\frac{cotsum_j}{c}$  the time.
- There are rowsum, instances in row  $i$ ; expect pected Values and  $\chi^2$ <br>colsum<sub>j</sub> occurrences of class *j* in the<br>ned,<br>proportion of  $\frac{colsum_j}{sum}$  the time.<br>instances in row *i* ; expect<br>currences of class *j* in row *i*. rowsum<sub>i</sub>  $\frac{colsum_j}{sum}$  occurrences of class *j* in row *i*.
- To calculate this value for any combination of row and class,
	- □ Take the product of the corresponding row sum and column sum
	- □ Divide by the grand total of the observed values for the two rows.

# Expected Values and  $\chi^2$  - Example<br>
To calculate expected value for any combination of row and class,<br>
Take the product of the corresponding row sum and column

- To calculate expected value for any combination of row and class,
	- □ Take the product of the corresponding row sum and column sum
	- □ Divided by the grand total of the observed values for the two rows.
- For the adjacent intervals labelled 8.7 and 12.1 the six values of O and E are:
	- **Expected value for C1 at value** 8.7 is  $\frac{7\times13}{10} = 4.789 \sim 4.79$  $19 \t\t \ldots \t\t \ldots$





Finally calculating  $\chi^2$ 

- Using observed and expected values, calculate  $\frac{(0-E)}{E}$  for మ for each of the six combinations
- Value of  $\chi^2$  is the sum of the six values for  $\frac{(0-E)^2}{E}$ మ
- If the independence hypothesis is correct O and E values would be the same and  $\chi^2$  is zero
	- **E** Small value for  $\chi^2$  supports hypothesis

□ Larger value militates against it

- When  $\chi^2$  exceeds threshold, hypothesis is rejected
- Important adjustment, when  $E < 0.5$  replace it with 0.5

### Final Calculation

- $x^2$  at each row is the value for the pair of adjacent row □ That row
	- The row below
- **Original table has 11 rows, so**  $10 \times \chi^2$  **calculations,** values
- **Each value represents** is the value for that row and the one below it



### Final Step

- Select the smallest  $\chi^2$  value
- Compare it to the threshold
- If it falls below the threshold, merge it with the row immediately below it
	- $\Box$  The independence of which the  $\chi^2$  represents
- $\blacksquare$  Smallest value is 1.08 for row

1.4

**New resulting interval is** 

 $1.4 \leq x < 2.4$ 



### New Table

### Revised frequency table



### Final Step

- $\rightarrow$   $\chi^2$  values calculated for the new frequency table
- Only need to do this for rows adjacent to the recently **merged one**  $\sqrt{\frac{Value of A}{Value of A}}$  Frequency for class  $\sqrt{\frac{Value of \chi^2}{Value of \chi^2}}$



- Smallest  $\chi^2$  is 1.20
	- □ Below the threshold
- **Intervals 87.1 and 89.0 merged**
- Continue until one reaches a fixed point:
	- **Example 1** Smallest  $\chi^2$  is above the threshold

### Final Table

### **All possible merging complete**



- **1.3**  $\leq$   $x$   $\leq$  56.2
- $\blacksquare$  56.2  $\leq x < 87.1$
- $x \geq 87.1$

# minIntervals and maxIntervals<br>Two extrema:<br>a Large number of intervals

- Two extrema:
	- □ Large number of intervals
	- Just one interval
- **Large numbers of intervals does little to solve the problem of** discretization
- **Just one interval cannot contribute to a decision making** process
	- □ Attribute value is independent of classification.
- **Two solutions**
- □ Modify significance level hypothesis of independence must pass, triggering interval merge. discretization<br>
Just one interval cannot contribute to a decision making<br>
process<br>  $\Box$  Attribute value is independent of classification.<br>
Two solutions<br>  $\Box$  Modify significance level hypothesis of independence must<br>
pas
	- **□** Set a minimum and a maximum number of intervals
- theory