

Applying distances between terms to both flat and hierarchical data

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- 2. Distances over terms
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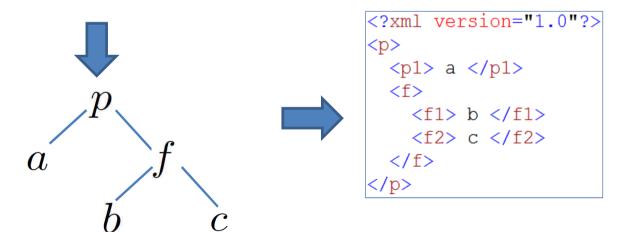
1. Introduction

- Distances are measures of dissimilarity with some special properties, such as symmetry and triangle inequality, which make them more advantageous for many algorithms.
- There are distances for virtually any kind of object, including complex or highly structured ones, such as tuples, sets, lists, trees, graphs, etc.
- Distances between atoms may not only be useful in ILP, but also in other areas where structured (hierarchical) information is involved, such as learning from ontologies or XML documents.
- The advantage of term distances is that they are able to consider context and, some of them, repetitions.

1. Introduction

- Tree structures and functional terms have strong similarities.
- Some popular languages for information representation are based on trees or hierarchies, such as XML.

$$t = p(a, f(b, c))$$



1. Introduction

Objectives:

- Define a transformation procedure to convert semi-structured data into a term-based representation in XML.
- Define several mechanisms to extract hierarchies from flat data, using the attribute names or their correlation.
- Experimentally analyse the performance of term distances over the transformed data.

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- In order to use with a general inductive method, such as k-NN, we worked with the following distances over terms:
 - Nienhuys-Cheng's distance
 - Estruch et al.'s distance

Nienhuys-Cheng distance:

Given two ground expressions:
$$s=s_0(s_1,\ldots,s_n)$$
 $t=t_0(t_1,\ldots,t_n)$

distance between them is recursively defined as:

$$d_N(s,t) = \begin{cases} 0, & \text{if } s = t \\ 1, & \text{if } \neg Compatible(s,t) \\ \frac{1}{2n} \sum_{i=1}^n d(s_i, t_i), & \text{otherwise} \end{cases}$$

It takes into account the depth of the symbol occurrences.

Nienhuys-Cheng distance example:

$$d_N(s,t) = \begin{cases} 0, & \text{if } s = t \\ 1, & \text{if } \neg Compatible(s,t) \\ \frac{1}{2n} \sum_{i=1}^n d(s_i, t_i), & \text{otherwise} \end{cases}$$

Given two ground expressions: s = p(a, a) t = p(f(b), f(b))

$$d_N(s,t) = \frac{1}{4} \cdot \left(d(a, f(b) + d(a, f(b))) \right)$$

$$d_N(s,t) = \frac{1}{4} \cdot (1+1) = \frac{1}{2}$$

Estruch et al. distance:

Given two ground expressions: $s = s_0(s_1, \ldots, s_n)$ $t = t_0(t_1, \ldots, t_n)$

distance between them is defined as:

$$d_E(s,t) = \sum_{o \in O^*(s,t)} \frac{w(o)}{C(o)} \left(Size'(s|_o) + Size'(t|_o) \right)$$

It takes into account:

- Context of the differences C(o).
- Syntactical complexity through size of the expression Size'(s|_{o'})
- The weight w(o) associated with the repetitions, using equivalence relations.

Estruch et al. distance example:

$$d_E(s,t) = \sum_{o \in O^*(s,t)} \frac{w(o)}{C(o)} \left(Size'(s|_o) + Size'(t|_o) \right)$$

Given two ground expressions: s = p(a, a) t = p(f(b), f(b))

$$C(1) = C(2) = 2 \cdot (2+1) = 6$$

$$Size'(a) = 1/4 \text{ and } Size'(f(b)) = 5/16$$

$$w(1) = 1$$
 and $w(2) = 7/8$

$$d_E(s,t) = \frac{1}{6} \left(\frac{1}{4} + \frac{5}{16} \right) + \frac{7}{48} \left(\frac{1}{4} + \frac{5}{16} \right)$$

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- A common language for representing several datatypes is XML.
- We distinguished two situations:
 - Flat data: are given as a table of attribute-value pairs.
 - Hierarchical data at source: consist of data with a hierarchical structure which is represented by a tree or a functional term.

Schema definition:

- We need to make some decisions in order to use XML to represent functional terms, and also to adapt the previous types of data to a common representation.
- It is necessary to ensure that distance calculations are not affected by the absence of features or their order.
- This difficulty requires the creation of a general schema that allows each instance, with its own features, to be properly adjusted, without losing any element or content, and in a defined order.

Schema definition:

Deriving hierarchical XML schemas from flat data

This work considered three possible sources of structures from flat representations:

- 1. Value equality
- 2. Name-induced hierarchies
- 3. Attribute-similarity hierarchy

1. Value equality

A detailed inspection shows that some attributes are related by the values they take.

For instance:

Two variables X1 and X2 can take the values **East**, **West**, **North** and **South**; there is clearly a connection between them that can be exploited, especially through the use of equalities.

2. Name-induced hierarchies

```
1. cap-shape
 2. cap-surface
 3. cap-color
 4. bruises
 5. odor
 6. gill-attachment
 7. gill-spacing
 8. gill-size
 9. gill-color
10. stalk-shape
11. stalk-root
12. stalk-surface-above-ring
13. stalk-surface-below-ring
14. stalk-color-above-ring
15. stalk-color-below-ring
16. veil-type
17. veil-color
18. ring-number
19. ring-type
20. spore-print-color
21. population
22. habitat
```

```
(Mushroom)
 <cap>
   <shape>CONVEX</shape>
   <surface>SMOOTH</surface>
   <color>WHITE</color>
 </cap>
  <bruises>BRUISES
 <odor>ALMOND</odor>
 <qi11>
   <attachment>FREE</attachment>
   <spacing>CROWDED</spacing>
   <size>NARROW</size>
   <color>WHITE</color>
 </gill>
 <stalk>
   <shape>TAPERING</shape>
   <root>BULBOUS</root>
   <surface>
     <above ring>SMOOTH</above ring>
     <below ring>SMOOTH</below ring>
   </surface>
   (color)
     <above ring>WHITE</above ring>
     <below ring>WHITE</below ring>
   </color>
 </stalk>
 <veil>
   <type>PARTIAL</type>
   <color>WHITE</color>
 </veil>
```

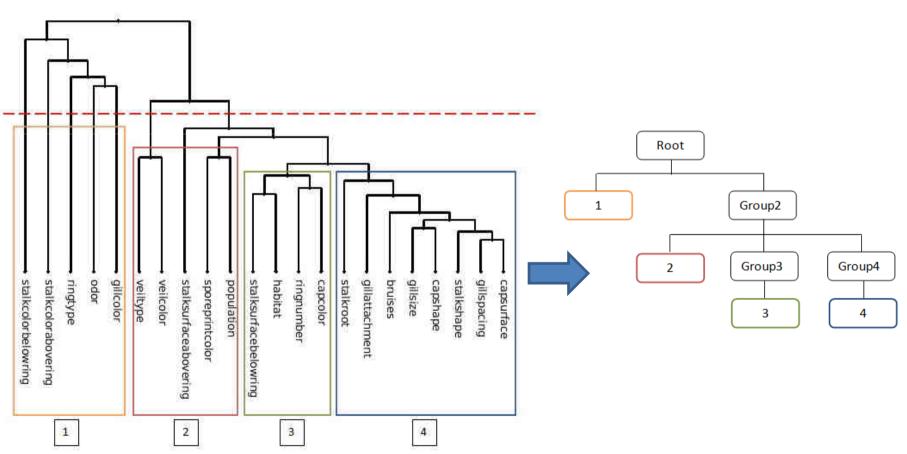
2. Name-induced hierarchies

```
Mushroom (
cap (CONVEX, SMOOTH, WHITE),
BRUISES, ALMOND,
gill (FREE, CROWDED, NARROW, WHITE),
stalk (TAPERING, BULBOUS,
surface(SMOOTH, SMOOTH),
color (WHITE, WHITE)),
veil (PARTIAL, WHITE),
ring(ONE, PENDANT),
spore (print (BROWN)),
SEVERAL, WOODS
)
```



```
(Mushroom)
 <cap>
   <shape>CONVEX</shape>
   <surface>SMOOTH</surface>
   <color>WHITE</color>
 </cap>
 <bruises>BRUISES
 <odor>ALMOND</odor>
 <qill>
   <attachment>FREE</attachment>
   <spacing>CROWDED</spacing>
   <size>NARROW</size>
   <color>WHITE</color>
  </gill>
  <stalk>
   <shape>TAPERING</shape>
   <root>BULBOUS</root>
   <surface>
     <above ring>SMOOTH</above ring>
     <below ring>SMOOTH</below ring>
   </surface>
    <color>
     <above ring>WHITE</above ring>
     <below ring>WHITE</below ring>
   </color>
 </stalk>
 <veil>
   <type>PARTIAL</type>
   <color>WHITE</color>
 </veil>
```

3. Attribute-similarity hierarchy



Deriving hierarchical XML schemas from hierarchical data

It is necessary to determine whether order, repetitions and labels are relevant, in order to determine the features and place them correctly in the XML document.

location mediterranean ecologicalecological **Deriving hierarchical XML** schemas from hierarchical data none <BODY-SIZE>MEDIUM</BODY-SIZE> <CONSISTENCY>STONY</CONSISTENCY> colour <HAND>TOUGH</HAND> external-<TOUCH>VELVETY</TOUCH> <SURFACE>BRISTLY</SURFACE> <GROWING> <DEF>GROWING</DEF> description sponge-220 <GROW>ERECT</GROW> <LINE-FORM> <DEF>LINE-FORM</DEF> <FORM>BRANCHING</FORM> uniform-length no-acanthos <PEDUNCLE>YES</PEDUNCLE> </LINE-FORM> </GROWING> fusiform <BORING>NO</BORING> smooth-ornamentation <COLOUR> two-or-more-lengths <DEF>COLOUR</DEF> <EXTERNAL>YELLOW</EXTERNAL> smooth-</COLOUR> mamentation <CRIBLE>NO</CRIBLE> <HOLLOW>NO</HOLLOW> <BRUSH>NO</BRUSH> <BRIOZOA>NO</BRIOZOA> <MACRO-DEBRIS>NONE</MACRO-DEBRIS> 23

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- k-nearest neighbor (k-NN) algorithm.
- Weighted k-NN variant, using an attraction function which gives more or less weight to each of the k-most nearest examples, defined as

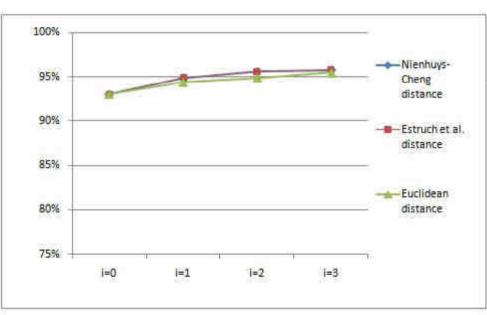
$$\frac{1}{d^i}$$
 (*i* varying from 0 to 3).

- Three distances: the Nienhuys-Cheng distance, the Estruch et al. distance and the Euclidean distance.
- Datasets:
 - Flat: Mushroom (1000 examples) and Soybean:
 - 10-fold cross validation experiment.
 - A paired t-test between the methods (confidence 95%).
 - Hierarchy: Demospongiae (sponge) dataset.
 - 60% for training examples and a 40% for test examples.

Mushroom dataset

Distances without hierarchy data.

	i=0 %	i=1 %	i=2 %	i=3 %
d_N	93.0	94.9	95.6	95.8
d_{E}	93.0	94.9	95.6	95.8
d _U	93.0	94.4	94.9	95.5

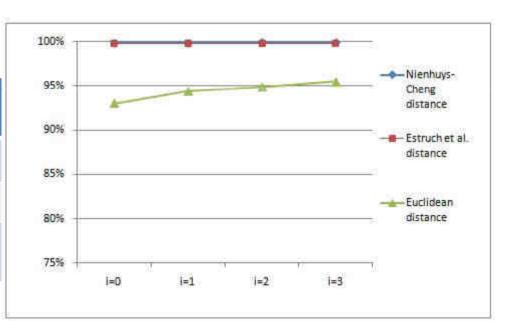


The differences between distances were not significant.

Mushroom dataset

Distances with hierarchy induced from the attributes names.

	i=0 %	i=1 %	i=2 %	i=3 %
d _N	99.8	99.8	99.9	99.9
d_{E}	99.8	99.8	99.8	99.8
d _U	93.0	94.4	94.9	95.5

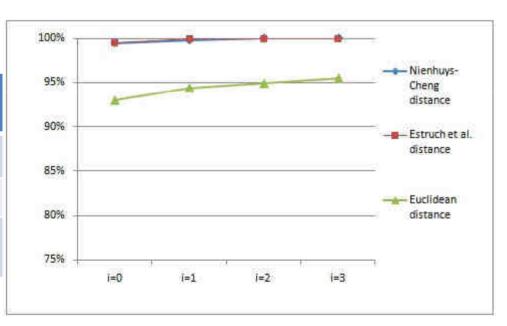


 The performance of Estruch et al's and Nienhuys-Cheng's distances was optimum.

Mushroom dataset

Distances with hierarchy using similarities between attributes.

	i=0 %	i=1 %	i=2 %	i=3 %
d _N	99.5	99.8	100	100
d_E	99.5	100	100	100
d _U	93.0	94.4	94.9	95.5

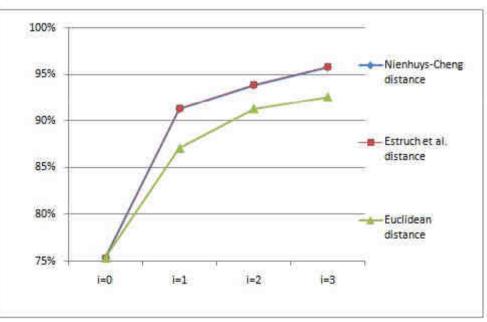


 The distances between terms exploited this structure and got much better results than the Euclidean distance.

SoyBean dataset

Distances without hierarchy data.

	i=0 %	i=1 %	i=2 %	i=3 %
d_N	75.3	91.3	93.9	95.8
d_E	75.3	91.3	93.9	95.8
d _U	75.3	87.1	91.3	92.6

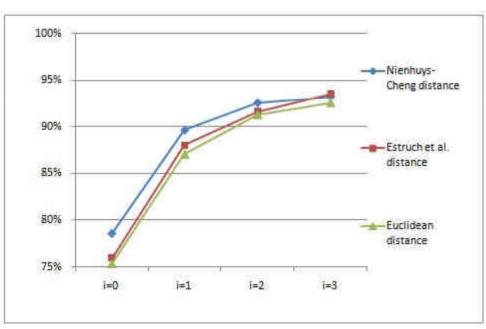


The Euclidean distance obtained less favorable results than the two term distances.

SoyBean dataset

Distances with hierarchy induced from the attributes names.

	i=0 %	i=1 %	i=2 %	i=3 %
d _N	78.6	89.6	92.6	93.2
d_E	76.0	88.0	91.6	93.5
d _U	75.3	87.1	91.3	92.6

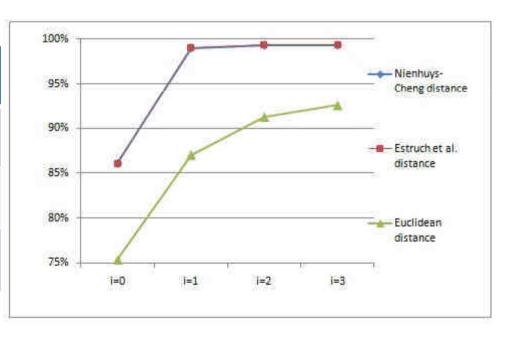


 The behavior of the distances was similar and the differences were not statistically significant.

SoyBean dataset

Distances with hierarchy using similarities between attributes.

	i=0	i=1	i=2	i=3
	%	%	%	%
d _N	86.1 <i>d</i> _U	99.6 <i>d_U</i>	99.4 <i>d</i> _U	99.4 d _U
d _E	86.1	99.0	99.4	99.4
	<i>d</i> _U	<i>d</i> _U	<i>d</i> _U	<i>d</i> _U
d _U	75.3	87.1	91.3	92.6
	d _N d _E			



The term distances obtained much better results than the Euclidean distance.

Demospongiae dataset

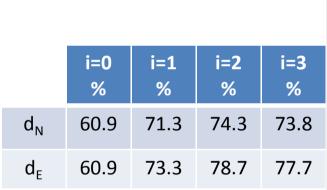
- It is a hierarchical dataset represented as a tree using terms in Lisp language.
- Each tree has a depth between 5 and 8 levels and its number of leaves varies between 17 and 51.
- In order to do the transformations from Lisp, each line in Lisp was converted into one or several well-formed XML elements.

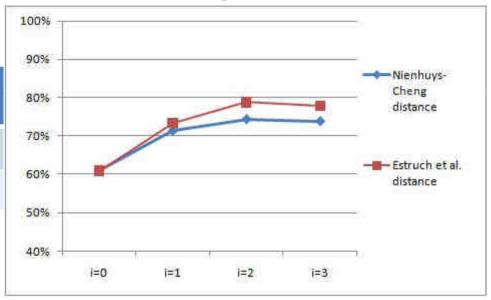
For instance:

Each simple feature such as: (BODY-SIZE SMALL)
 was converted into an element <BODY-SIZE>SMALL</BODY-SIZE>

Demospongiae dataset

Differences between the term-based distances using semi-structured data.





 The Estruch et al's distance showed a better accuracy than Nienhuys-Cheng's distance.

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5. Conclusions

- Regarding the three transformations to adapt different degrees of structures and hierarchical data to be used with term distances:
 - The method for constructing the hierarchical from the attribute names does not always seem to provide good results.
 - Using the similarity between the attributes to construct a dendrogram from which a hierarchy is constructed improves the results of the flat dataset.
 - Different transformations take place when the original dataset already has a hierarchy, here, the relevance of using repetitions or not can be seen.
- We have seen a promising application of term distances to different types of datasets, which suggests that the use of term distances can be broader than it is now.

Thank you for your attention!