

Structural Matching and Discovery in Document Databases*

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1 Background

Structural matching and discovery in documents such as SGML and HTML is important for data warehousing [6], version management [7, 11], hypertext authoring, digital libraries [4] and Internet databases. As an example, a user of the World Wide Web may be interested in knowing changes in an HTML document [2, 5, 10]. Such changes can be detected by comparing the old and new version of the document (referred to as structural matching of documents). As another example, in hypertext authoring, a user may wish to find the common portions in the history list of a document or in a database of documents (referred to as structural discovery of documents). In SIGMOD 95 demo sessions, we exhibited a software package, called *TreeDiff* [13], for comparing two latex documents and showing their differences. Given two documents, the tool represents the documents as ordered labeled trees and finds an optimal sequence of edit operations to transform one document (tree) to the other. An edit operation could be an insert, delete, or change of a node in the trees. The tool is so named because documents are represented and compared using approximate tree matching techniques [9, 12, 14].

2 System Architecture and Operators

Here we present an extension of *TreeDiff* for querying, comparing and discovering structured documents. Our new system is equipped with a graphical interface and a powerful query language that allows the user to compare documents and to discover the (approximately) common portions of documents. The front end of the system is composed of a query processor and an SGML parser originally developed by James Clark. Given a document type definition (DTD), the SGML parser checks if an input document conforms to the DTD. If the syntax is correct, the parser translates the document to an ordered labeled tree. The query processor

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checks the syntax of a given query. When the query involves the comparison or structural discovery of documents, the query processor invokes the back end of the system, which performs approximate tree matching and discovery. Figure 1 illustrates the system architecture.

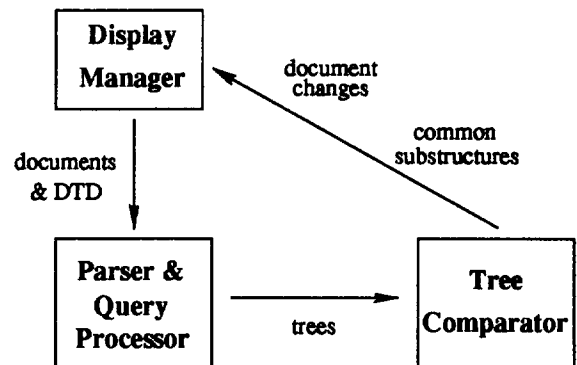


Fig. 1. System architecture.

The system provides several useful operators, which can be used alone or included in a query, for comparing two documents A and B:

- $\text{symdifference}(A, B)$ – to find all the differences between A and B (the structural elements in A not in B, the structural elements in B not in A, and the structural elements changed from A to B).
- $\text{difference}(A, B)$ – to find the structural elements in A not in B and the structural elements changed from A to B.
- $\text{intersection}(A, B)$ – to find the structural elements in common between A and B (cf. Figures 2 – 4).
- $\text{union}(A, B)$ – to find the structural elements in common, the structural elements in A not in B, the structural elements in B not in A, and the structural elements changed from A to B.
- $\text{mergable}(O, A, B)$ – to determine whether A and B can be merged where A, B are the documents obtained by modifying document O; this function is true if B is obtained by modifying different portions of O than A.

- merge(O, A, B) – to perform the merge of A and B; whenever some portion has changes from both A and B, then A's changes will occur, but not B's.

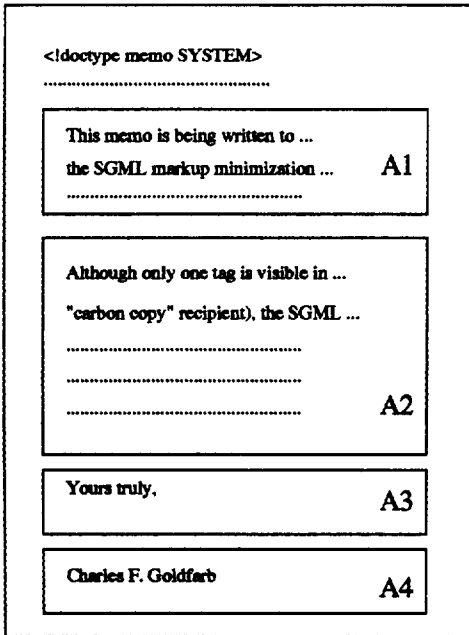


Fig. 2. An SGML memo document A.

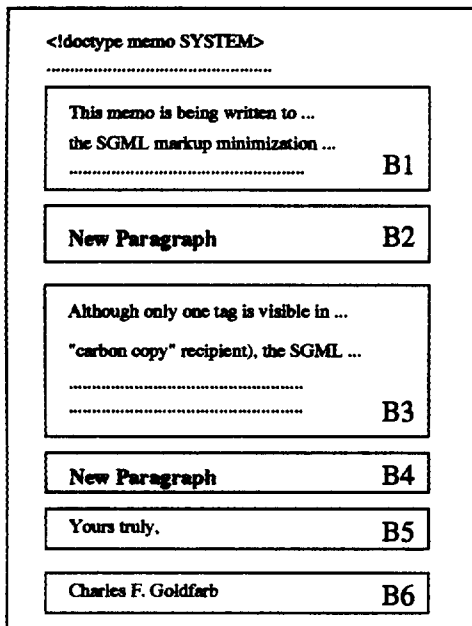


Fig. 3. An SGML memo document B.

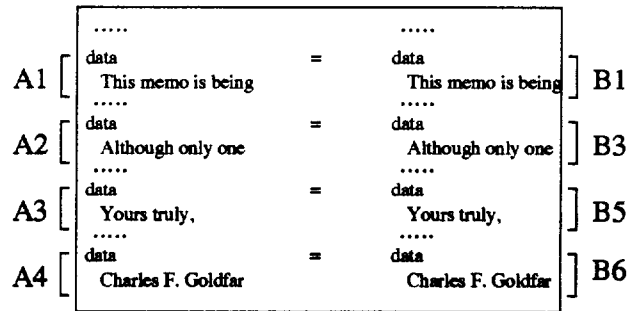


Fig. 4. The result of intersection(A, B). For each document element (e.g. a paragraph), only the first 18 characters are displayed.

3 Related Work

Whereas much database research has been conducted to manage structured documents such as SGML and HTML (e.g., [1, 3, 8]), very few systems have been built to support their comparison or the discovery of patterns in their structures. As far as we know, only the *LaDiff* program developed in [2] supports structural matching. Both our system and *LaDiff* represent hierarchically structured documents as trees and identify changes in them. In contrast to other systems dealing with flat information, the two systems are concerned with not only the "nodes" in the data, but also their relationships. For example, if a node (and its children) is moved from one location to another, both systems represent it as a "move" operation in the data. The two systems differ, however, in some respects. First, they use different notions of editing. We use a generalization of edit distance as used in the Unix utility *diff*. *LaDiff* approximates this edit distance in order to run faster. Second, we include both querying and structural pattern discovery.

4 The Demonstration

In the demonstration of SIGMOD 97, we will show new queries for performing:

- document matching: aligning two documents and detecting changes or differences of the documents;
- document discovery: aligning two documents and identifying the largest (approximately) common portions of the documents;
- multiple documents discovery: finding the (approximately) common portions of a database of documents;
- similarity search: finding the document in a database that is most similar to a given document;
- substructure search: finding the document in a database that contains a given document;
- superstructure search: finding the document in a database that is the same as a substructure of a given document.

The output of a query can be displayed alone or combined with that of other utilities such as *diff*. For example, Figure 5 illustrates document matching, showing the result of *syndifference*(C, D) of two memos C and D. "=" indicates that the two document elements (e.g. two section titles or two paragraphs) are the same, while "|" indicates that the

document elements are changed, ">" indicates that the corresponding document element is inserted, and "<" indicates the document element is deleted. The detailed comparisons of the changed paragraphs using *diff* are displayed at the end.

```

MEMO | MEMO
<MEMO FINAL SEC=" | <MEMO FINAL SEC="
... | ...
data | <
  Here is more infor | ...
... | > data
... | This paragraph has
... | ...
data | data
  Although only one | Although only one
... | ...
data | data
  recipient), the SG | recipient
... | ...
... | ...
... | ...
**** EDITED TEXT ****
:::: memo_C.sgm line 13-20      memo_D.sgm: line 14-21 ::::
3,5c3,5
< carbon copy recipient), the SGML parser recognizes 15 .....
< element types. This allows each element to be formatted ....
< for example, the start and end of a quotation can have .....
...
> carbon copy recipient), the SGML parser recognizes 14 .....
> different element types. This allows each element to be .....
> differently; for example, the start and end of a quotation .....

```

Fig. 5. The result of *syndifference*(C, D).

Document changes or common portions are highlighted in a colorful fashion through the graphical interface of our system. The system is implemented using C, Perl and Tcl-Tk on SUN SPARC workstations. Figure 6 shows a screen shot of the system. We have prepared three versions: one for SunOS, one for Solaris and one for other operating systems. The software is available for research purposes and can be obtained from the authors (please visit the Web site <http://www.cis.njit.edu/~jason/demo.html> for details).

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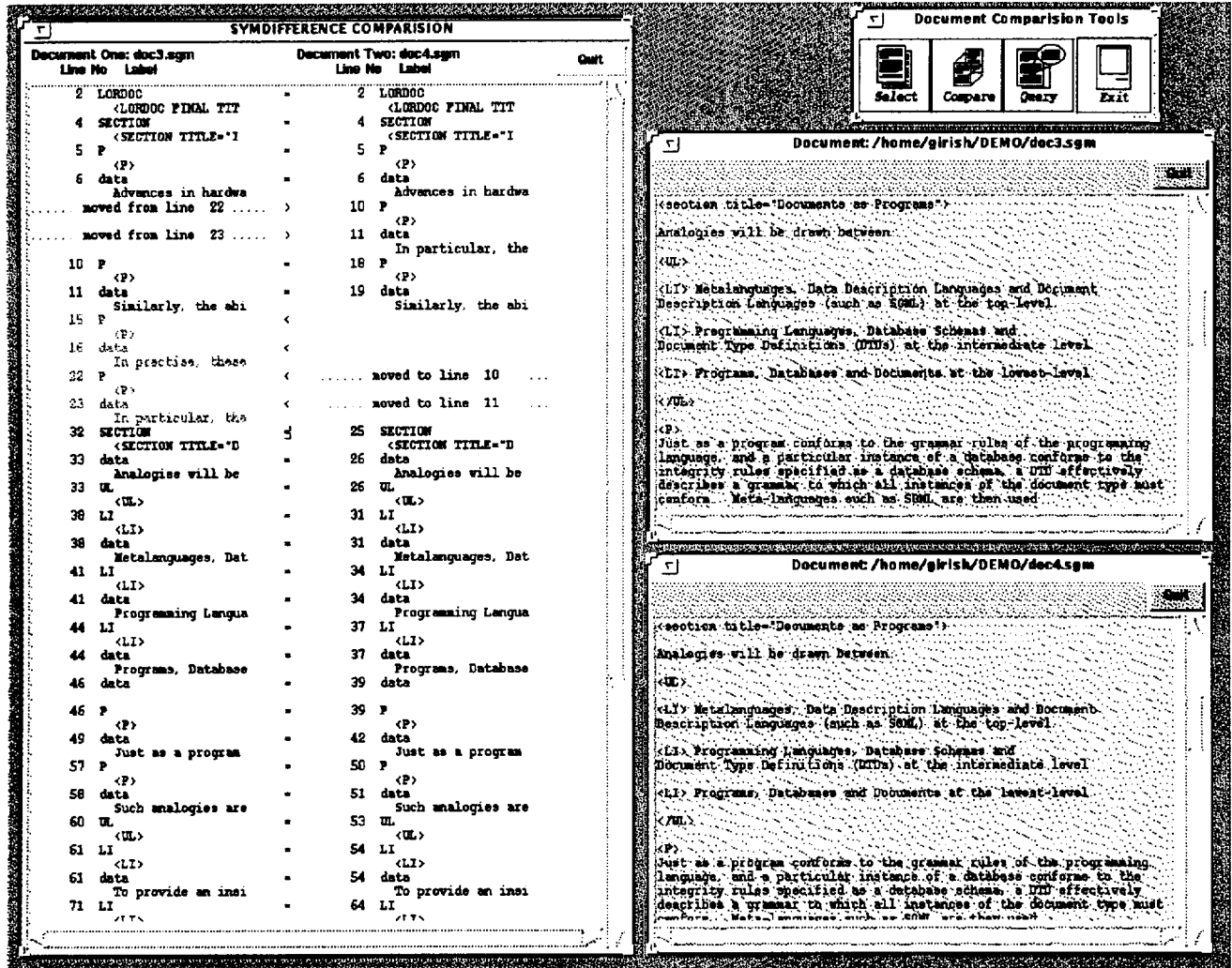


Fig. 6. The result of comparing two SGML articles. The two windows on the right show the articles. The left window displays the output of the comparison. In addition to showing the inserts, deletes and changes of paragraphs, the system can also show their movement (lines 10 and 22). When the user clicks on the particular document element of interest, that portion of the document is scrolled to the top in the right windows. If the user is interested in section changes only, rather than paragraph changes, the system displays a size parameter for each section, where "size = n" means that the total number of inserts, deletes and changes of paragraphs in that section is n. Moreover, the section with a larger change size is displayed in a deeper color.