

# Schema Evolution in Data Warehousing Environments — a schema transformation-based approach

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**Abstract.** In heterogeneous data warehousing environments, autonomous data sources are integrated into a materialised integrated database. The schemas of the data sources and the integrated database may be expressed in different modelling languages. It is possible for either the data source schemas or the warehouse schema to evolve. This evolution may include evolution of the schema, or evolution of the modelling language in which the schema is expressed, or both. In such scenarios, it is important for the integration framework to be evolvable, so that the previous integration effort can be reused as much as possible. This paper describes how the AutoMed heterogeneous data integration toolkit can be used to handle the problem of schema evolution in heterogeneous data warehousing environments. This problem has been addressed before for specific data models, but AutoMed has the ability to cater for multiple data models, and for changes to the data model.

## 1 Introduction

With the increasing use of the Internet in distributed applications, data warehouses may integrate data from remote, heterogeneous, autonomous data sources. The heterogeneity of these data sources has two aspects, heterogeneous data expressed in different data models, called *model heterogeneity* [10], and heterogeneous data within different data schemas expressed in the same data model, called *schema heterogeneity* [10, 18]. The common approach to handling model heterogeneity is to use a single conceptual data model (CDM) for the data transformation/integration. Each data source has a wrapper for translating its schema and data into the CDM. The warehouse schema is derived from these CDM schemas by means of view definitions, and is expressed in the same modelling language as them. With this approach, since they are both high-level conceptual data models, semantic mismatches may occur between the CDM and a source data model, and there may be a loss of information between them. Moreover, if a data source schema changes, it is not straightforward to evolve the view definitions of the warehouse schema.

Lakshmanan *et al* [11] argue that a uniform framework for schema integration and schema evolution is both desirable and possible, and this is our view

also. They define a higher-order logic language, SchemaSQL, which handles both data integration and schema evolution in relational multi-database systems. In contrast, our approach uses a simple set of schema transformation primitives, augmented with a functional query language, both of which are uniformly applicable to multiple data models. Other previous work on schema evolution, e.g. [1–4], has also presented approaches in terms of just one data model.

AutoMed is a heterogeneous data transformation and integration system which offers the capability to handle data integration across multiple data models<sup>1</sup>. In [7] we discussed how AutoMed metadata can be used to express the schemas and the cleansing, transformation and integration processes in heterogeneous data warehouse environments, supporting both schema heterogeneity and model heterogeneity. We discussed how this metadata can be used to populate and incrementally maintain the warehouse, and any data marts derived from it, and also to trace the lineage of data in the warehouse or the data marts. It is clearly advantageous to be able to reuse this kind of metadata if a schema evolves. In this paper we show how this can be achieved.

Earlier work [16] has shown how the AutoMed framework readily supports schema evolution in *virtual* data integration scenarios. In this paper we address the problem of schema evolution in *materialised* data integration scenarios, including both evolution of a source schema and of the warehouse schema, and also the impact on any data marts derived from the warehouse. This scenario is more complex than with virtual data integration, since both schemas and materialised data may be affected by an evolution.

The outline of the paper is as follows. Section 2 gives an overview of the AutoMed framework. Section 3 describes how AutoMed transformations can be used to express a schema evolution if either the schema changes, or the data model changes, or both. Section 4 describes the actions that are taken in order to evolve these transformations and the materialised data if the warehouse schema or a local schema evolves. Section 5 discusses the benefits of our approach and gives our concluding remarks.

## 2 Overview of AutoMed

AutoMed supports a low-level hypergraph-based data model (HDM). Higher-level modelling languages are defined in terms of this HDM. For example, previous work has shown how relational, ER, OO [15], XML [21], flat-file [5] and multidimensional [7] data models can be so defined. An HDM schema consists of a set of nodes, edges and constraints, and each modelling construct of a higher-level modelling language is specified as some combination of HDM nodes, edges and constraints. For any modelling language  $\mathcal{M}$  specified in this way (via the API of AutoMed’s Model Definitions Repository [5]), data source wrappers translate data source schemas expressed in  $\mathcal{M}$  into their AutoMed representation, without loss of information. AutoMed also provides a set of primitive schema transformations that can be applied to schema constructs expressed in  $\mathcal{M}$ . In particular, for

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<sup>1</sup> See <http://www.doc.ic.ac.uk/automed/>

every construct of  $\mathcal{M}$  there is an **add** and a **delete** primitive transformation which add to/delete from a schema an instance of that construct. For those constructs of  $\mathcal{M}$  which have textual names, there is also a **rename** primitive transformation.

In AutoMed, schemas are incrementally transformed by applying to them a sequence of primitive transformations  $t_1, \dots, t_r$ . Each primitive transformation adds, deletes or renames just one schema construct. Thus, intermediate schemas may contain constructs of more than one modelling language.

Each **add** or **delete** transformation is accompanied by a query specifying the extent of the new or deleted construct in terms of the rest of the constructs in the schema. This query is expressed in a functional query language IQL (see Section 2.1). Also available are **contract** and **extend** transformations which behave in the same way as **add** and **delete** except that they indicate that their accompanying query may only partially construct the extent of the new/removed schema construct. Moreover, their query may just be the constant **Void**, indicating that the extent of the new/removed construct cannot be derived even partially, in which case the query can be omitted.

We term a sequence of primitive transformations from one schema  $S_1$  to another schema  $S_2$  a transformation *pathway* from  $S_1$  to  $S_2$ , denoted  $S_1 \rightarrow S_2$ . All source, intermediate, and integrated schemas, and the pathways between them, are stored in AutoMed’s Schemas & Transformations Repository [5].

The queries present within transformations that add or delete schema constructs mean that each primitive transformation  $t$  has an automatically derivable *reverse transformation*,  $\bar{t}$ . In particular, each **add/extend** transformation is reversed by a **delete/contract** transformation with the same arguments, while each **rename** transformation is reversed by swapping its two arguments. Thus, AutoMed is a **both-as-view** (BAV) data integration system. As discussed in [17], BAV subsumes the global-as-view (GAV) and local-as-view (LAV) approaches [13], since it is possible to extract a definition of each global schema construct as a view over source schema constructs, and it is also possible to extract definitions of source schema constructs as views over the global schema. We refer the reader to [9] for details of AutoMed’s GAV and LAV view generation algorithms.

Figure 1 illustrates the general integration scenario with AutoMed. Each data source is described by a local schema  $LS_i$ . Each  $LS_i$  is first conformed into a schema  $CS_i$  (which may or may not be expressed in the same modelling language as  $LS_i$ ) by means of a transformation pathway  $T_i$ . Not all of the information within a local schema  $LS_i$  need be transferred into the global schema and this is asserted by means of **contract** transformation steps within  $T_i$ . Conversely, there may be information within the global schema which is not semantically derivable from  $LS_i$ , and this is asserted by the pathway from  $CS_i$  to a ‘union-schema’  $US_i$  which consists of the necessary **extend** transformations<sup>2</sup>.

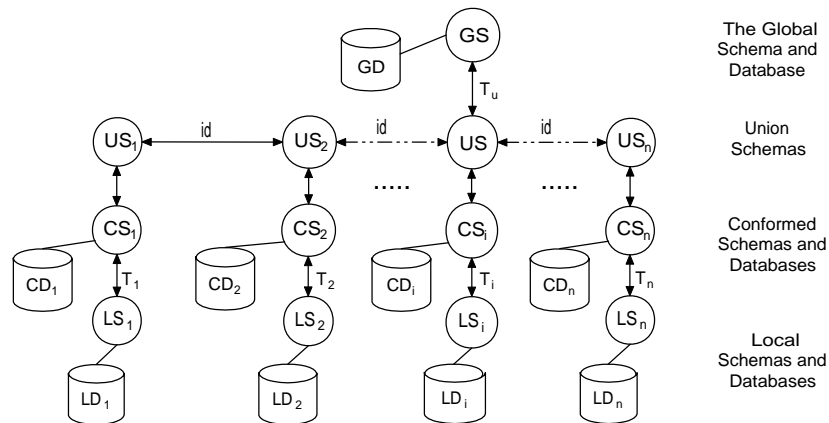
All the union schemas  $US_1, \dots, US_n$  are syntactically identical and this is asserted by creating a sequence of **id** transformations between each pair  $US_i$  and  $US_{i+1}$ , of the form **id**  $US_i : c \ US_{i+1} : c$  for each schema construct  $c$ . An **id** transformation signifies the semantic equivalence of syntactically identical

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<sup>2</sup> If there are none, then this pathway is empty and  $CS_i$  and  $US_i$  are the same schema

constructs in different schemas. The transformation pathways containing these *id* transformations are automatically generated by the AutoMed software. An arbitrary one of the  $US_i$  ( $US$  in Figure 1) can then be selected for further transformation into the global schema  $GS$  (by the pathway  $T_u$  in Figure 1). The extent of each construct  $c$  in a union schema  $US_i$  is equal to the bag-union of the extent of  $c$  in all union schemas  $US_1, \dots, US_n$ . That is, *id* is interpreted as *bag union* by AutoMed’s view generation functionality.

In a virtual data integration scenario, there is no materialised data associated with any of the schemas apart from the  $LS_i$ . In a data warehousing scenario, as illustrated in Figure 1, we assume that  $CS_1, \dots, CS_n$  are fully materialised and consist of the detailed data of the warehouse. This detailed data is further augmented with the necessary summary views by the transformations in the pathway  $T_u$ , and we assume that these summary views are materialised in the database  $GD$ . It would also be possible to partially or fully materialise more of the intermediate schemas in the network, or to not materialise  $CS_1, \dots, CS_n$  and to fully materialise  $GS$  instead. Our techniques in this paper easily generalise to these alternatives.



**Fig. 1.** Materialised Data Integration in AutoMed

For the purposes of this paper, we assume that all the  $LS_i$  and  $LD_i$  have been extracted from the original data sources and the data in the  $LD_i$  has been cleansed. The data cleansing process can also be expressed using AutoMed transformations — this is discussed in [7] and we do not consider it further here. See also that paper for some examples of how AutoMed transformations can express structural and representational changes to schemas and data.

We also assume here that there are no *contract* steps in the pathways  $T_i$ , i.e. that all the information in each  $LS_i$  will be transferred to  $CS_i$  and hence to  $US_i$ . This implies no loss of flexibility as each  $LS_i$  will be precisely that extract of the

original data source schema whose associated data is to be transferred into the warehouse.

## 2.1 The IQL query language

IQL is a comprehensions-based functional query language<sup>3</sup>. Such languages subsume query languages such as SQL and OQL in expressiveness [6]. IQL supports several primitive operators for manipulating lists. The list append operator, `++`, concatenates two lists together. The `distinct` operator removes duplicates from a list and the `sort` operator sorts a list. The `--` operator takes two lists and subtracts each member of the second list from the first e.g. `[1,2,3,2,4]--[4,4,2,1] = [3,2]`. The `fold` operator applies a given function `f` to each element of a list and then ‘folds’ a binary operator `op` into the resulting values. It is defined recursively as follows, where `(x:xs)` denotes a list with head `x` and tail `xs`:

```
fold f op e [] = e
fold f op e (x:xs) = (f x) op (fold f op e xs)
```

Other IQL list manipulation operators are defined using `fold` together with IQL’s set of built-in operators and its support of `lambda` abstractions. For example, the IQL functions `sum` and `count` are equivalent to SQL’s `SUM` and `COUNT` aggregation functions and are defined as

```
sum xs = fold (id) (+) 0 xs
count xs = fold (lambda x.1) (+) 0 xs
```

We also have

```
min xs = fold (id) lesser maxNum xs
max xs = fold (id) greater minNum xs
```

assuming constants `maxNum` and `minNum` and the following functions `lesser` and `greater`:

```
greater = lambda x.lambda y.if (x > y) then x else y
lesser = lambda x.lambda y.if (x < y) then x else y
```

The function `flatMap` applies a list-valued function `f` to each member of a list `xs` and is defined in terms of `fold`:

```
flatMap f xs = fold f (++) [] xs
```

`flatMap` can in turn be used to define selection, projection and join operators and, more generally, comprehensions. For example, the following comprehension iterates through a list of students and returns those students who are not members of staff:

```
[x | x <- <<student>>; not (member <<staff>> x)]
```

and it translates into:

```
flatMap (lambda x.if (not (member <<staff>> x))
        then [x] else []) <<student>>
```

Grouping operators are also definable in terms of `fold`. In particular, the operator `group` takes as an argument a list of pairs `xs` and groups them on their first component, while `gc aggFun xs` groups a list of pairs `xs` on their first component and then applies the aggregation function `aggFun` to the second component.

<sup>3</sup> We refer the reader to [8] for details of IQL.

There are several algebraic properties of IQL's operators that we can use in order to incrementally compute materialised data and to reason about IQL expressions, specifically for the purposes of this paper in a schema/data evolution context (note that the algebraic properties of `fold` below apply to all the operators defined in terms of `fold`):

- (a)  $e ++ [] = [] ++ e = e$ ,  $e -- [] = e$ ,  $[] -- e = []$ ,  
 $\text{distinct } [] = \text{sort } [] = []$   
 for any list-valued expression  $e$ . Since `Void` represents a construct for which no data is obtainable from a data source, it has the semantics of the empty list, and thus the above equivalences also hold if `Void` is substituted for `[]`.
- (b)  $\text{fold } f \text{ op } e [] = \text{fold } f \text{ op } e \text{ Void} = e$ , for any  $f$ ,  $\text{op}$ ,  $e$
- (c)  $\text{fold } f \text{ op } e (b1 ++ b2) = (\text{fold } f \text{ op } e b1) \text{ op } (\text{fold } f \text{ op } e b2)$   
 for any  $f$ ,  $\text{op}$ ,  $e$ ,  $b1$ ,  $b2$ . Thus, we can always incrementally compute the value of fold-based functions if collections expand.
- (d)  $\text{fold } f \text{ op } e (b1 -- b2) = (\text{fold } f \text{ op } e b1) \text{ op}' (\text{fold } f \text{ op } e b2)$   
 provided there is an operator  $\text{op}'$  which is the inverse of  $\text{op}$  i.e. such that  $(a \text{ op } b) \text{ op}' b = a$  for all  $a, b$ . For example, if  $\text{op} = +$  then  $\text{op}' = -$ , and thus we can always incrementally compute the value of aggregation functions such as `count`, `sum` and `avg` if collections contract. Note that this is not possible for `min` and `max` since `lesser` and `greater` have no inverses. Although IQL is list-based, if the ordering of elements within lists is ignored then its operators are faithful to the expected bag semantics, and within AutoMed we generally do assume bag semantics. Under this assumption,  
 $(xs ++ ys) -- ys = xs$   
 for all  $xs, ys$  and thus we can incrementally compute the value of `flatMap` and all its derivative operators if collections contract<sup>4</sup>.

## 2.2 An example

We will use schemas expressed in a simple relational data model and a simple XML data model to illustrate our techniques. However, we stress that these techniques are applicable to schemas defined in *any* data modelling language that has been specified within AutoMed's Model Definitions Repository.

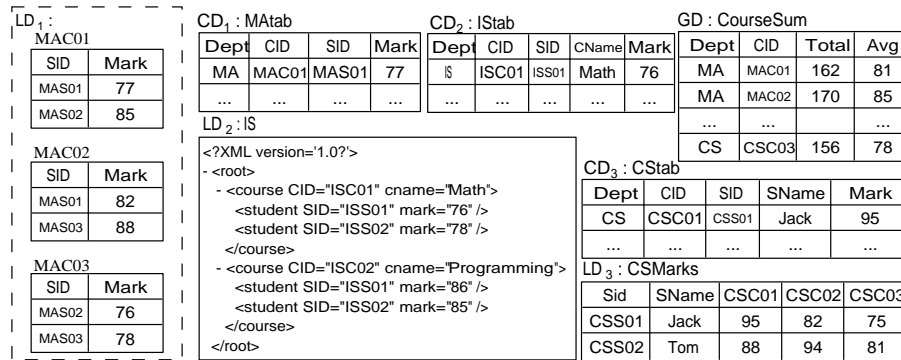
In the simple relational model, there are two kinds of schema construct: `Rel` and `Att`. The extent of a `Rel` construct  $\langle\langle R \rangle\rangle$  is the projection of the relation  $R$  onto its primary key attributes  $k_1, \dots, k_n$ . The extent of each `Att` construct  $\langle\langle R, a \rangle\rangle$  where  $a$  is an attribute (key or non-key) of  $R$  is the projection of relation  $R$  onto  $k_1, \dots, k_n, a$ . For example, the schema of table `MAtab` in Figure 2 consists of a `Rel` construct  $\langle\langle \text{MAtab} \rangle\rangle$ , and four `Att` constructs  $\langle\langle \text{MAtab}, \text{Dept} \rangle\rangle$ ,  $\langle\langle \text{MAtab}, \text{CID} \rangle\rangle$ ,  $\langle\langle \text{MAtab}, \text{SID} \rangle\rangle$ , and  $\langle\langle \text{MAtab}, \text{Mark} \rangle\rangle$ . We refer the reader to [15] for an encoding of a richer relational data model, including the modelling of constraints.

In the simple XML data model, there are three kinds of schema construct: `Element`, `Attribute` and `NestSet`. The extent of an `Element` construct  $\langle\langle e \rangle\rangle$  consists

<sup>4</sup> The `distinct` operator can also be used to obtain set semantics, if needed.

of all the elements with tag  $e$  in the XML document; the extent of each **Attribute** construct  $\langle\langle e, a \rangle\rangle$  consists of all pairs of elements and attributes  $x, y$  such that element  $x$  has tag  $e$  and has an attribute  $a$  with value  $y$ ; and the extent of each **NestSet** construct  $\langle\langle p, c \rangle\rangle$  consists of all pairs of elements  $x, y$  such that element  $x$  has tag  $p$  and has a child element  $y$  with tag  $c$ . We refer the reader to [21] for an encoding of a richer model for XML data sources, called XMLDSS, which also captures the ordering of children elements under parent elements and cardinality constraints. That paper gives an algorithm for generating the XMLDSS schema of an XML document. That paper also discusses a unique naming scheme for **Element** constructs so as to handle instances of the same element tag occurring at multiple positions in the XMLDSS tree.

Figure 2 illustrates the integration of three data sources  $LD_1$ ,  $LD_2$ , and  $LD_3$ , which respectively store students' marks for three departments MA, IS and CS.



**Fig. 2.** An example integration

Database  $LD_1$  for department MA has one table of students' marks for each course, where the relation name is the course ID. Database  $LD_2$  for department IS is an XML file containing information of course IDs, course names, student IDs and students' marks. Database  $LD_3$  for department CS has one table containing one row per student, giving the student's ID, name, and mark for the courses CSC01, CSC02 and CSC03.  $CD_1$ ,  $CD_2$ , and  $CD_3$  are the materialised conformal databases for each data source. Finally, the global database  $GD$  contains one table  $CourseSum(\underline{Dept}, \underline{CID}, Total, Avg)$  which gives the total and average mark for each course of each department. Note that the virtual union schema  $US$  (not shown) combines all the information from all the conformal schemas and consists of a virtual table  $Details(\underline{Dept}, \underline{CID}, \underline{SID}, CName, SName, Mark)$ .

The following transformation pathways express the schema transformation and integration processes in this example. Due to space limitations, we have not given the remaining steps for deleting/contracting the constructs in the source schema of each pathway (note that this 'growing' and 'shrinking' of schemas is characteristic of AutoMed schema transformation pathways):

```

T1 : LS1 → CS1
addRel ⟨⟨MAtab⟩⟩      [{ 'MA', 'MAC01', x } | x ← ⟨⟨MAC01⟩⟩ ] ++ [ { 'MA', 'MAC02', x } | x ← ⟨⟨MAC02⟩⟩ ]
                    ++ [ { 'MA', 'MAC03', x } | x ← ⟨⟨MAC03⟩⟩ ];
addAtt ⟨⟨MAtab, Dept⟩⟩ [ {k1, k2, k3, k1} | {k1, k2, k3} ← ⟨⟨MAtab⟩⟩ ];
addAtt ⟨⟨MAtab, CID⟩⟩ [ {k1, k2, k3, k2} | {k1, k2, k3} ← ⟨⟨MAtab⟩⟩ ];
addAtt ⟨⟨MAtab, SID⟩⟩ [ {k1, k2, k3, k3} | {k1, k2, k3} ← ⟨⟨MAtab⟩⟩ ];
addAtt ⟨⟨MAtab, Mark⟩⟩ [ { 'MA', 'MAC01', k, x } | {k, x} ← ⟨⟨MAC01, Mark⟩⟩ ]
                    ++ [ { 'MA', 'MAC02', k, x } | {k, x} ← ⟨⟨MAC02, Mark⟩⟩ ]
                    ++ [ { 'MA', 'MAC03', k, x } | {k, x} ← ⟨⟨MAC03, Mark⟩⟩ ];
delAtt ⟨⟨MAC01, Mark⟩⟩ [ {k3, x} | {k1, k2, k3, x} ← ⟨⟨MAtab, Mark⟩⟩; k2='MAC01' ];
delAtt ⟨⟨MAC01, SID⟩⟩ [ {k3, x} | {k1, k2, k3, x} ← ⟨⟨MAtab, SID⟩⟩; k2='MAC01' ];
delRel ⟨⟨MAC01⟩⟩      [ {k3} | {k1, k2, k3} ← ⟨⟨MAtab⟩⟩; k2='MAC01' ]
...

```

The removal of the other two tables in  $LS_1$  is similar.

```

T2 : LS2 → CS2
addRel ⟨⟨IStab⟩⟩      [ { 'IS', x, y } | {c, x} ← ⟨⟨course, CID⟩⟩; {s, y} ← ⟨⟨student, SID⟩⟩ ];
addAtt ⟨⟨IStab, Dept⟩⟩ [ {k1, k2, k3, k1} | {k1, k2, k3} ← ⟨⟨IStab⟩⟩ ];
addAtt ⟨⟨IStab, CID⟩⟩ [ {k1, k2, k3, k2} | {k1, k2, k3} ← ⟨⟨IStab⟩⟩ ];
addAtt ⟨⟨IStab, SID⟩⟩ [ {k1, k2, k3, k3} | {k1, k2, k3} ← ⟨⟨IStab⟩⟩ ];
addAtt ⟨⟨IStab, CName⟩⟩ [ { 'IS', x, y, n } | {c1, x} ← ⟨⟨course, CID⟩⟩; {c2, n} ← ⟨⟨course, cname⟩⟩; c1=c2;
                       {c3, s1} ← ⟨⟨course, student⟩⟩; c3=c2; {s2, y} ← ⟨⟨student, SID⟩⟩; s2=s1 ];
addAtt ⟨⟨IStab, Mark⟩⟩ [ { 'IS', x, y, m } | {c1, x} ← ⟨⟨course, CID⟩⟩; {c2, s1} ← ⟨⟨course, student⟩⟩; c1=c2;
                       {s2, y} ← ⟨⟨student, SID⟩⟩; s2=s1; {s3, m} ← ⟨⟨student, mark⟩⟩; s3=s2 ];
...

```

```

T3 : LS3 → CS3
addRel ⟨⟨CStab⟩⟩      [ { 'CS', x, y } | x ← [ 'CSC01', 'CSC02', 'CSC03' ]; y ← ⟨⟨CSMarks⟩⟩ ];
addAtt ⟨⟨CStab, Dept⟩⟩ [ {k1, k2, k3, k1} | {k1, k2, k3} ← ⟨⟨CStab⟩⟩ ];
addAtt ⟨⟨CStab, CID⟩⟩ [ {k1, k2, k3, k2} | {k1, k2, k3} ← ⟨⟨CStab⟩⟩ ];
addAtt ⟨⟨CStab, SID⟩⟩ [ {k1, k2, k3, k3} | {k1, k2, k3} ← ⟨⟨CStab⟩⟩ ];
addAtt ⟨⟨CStab, SName⟩⟩ [ { 'CS', x, k, s } | x ← [ 'CSC01', 'CSC02', 'CSC03' ]; {k, s} ← ⟨⟨CSMarks, SName⟩⟩ ];
addAtt ⟨⟨CStab, Mark⟩⟩ [ { 'CS', 'CSC01', k, x } | {k, x} ← ⟨⟨CSMarks, CSC01⟩⟩ ]
                    ++ [ { 'CS', 'CSC02', k, x } | {k, x} ← ⟨⟨CSMarks, CSC02⟩⟩ ]
                    ++ [ { 'CS', 'CSC03', k, x } | {k, x} ← ⟨⟨CSMarks, CSC03⟩⟩ ];
...

```

```

Tu : US → GS
addRel ⟨⟨CourseSum⟩⟩      distinct [ {k1, k3} | {k1, k2, k3} ← ⟨⟨Details⟩⟩ ];
addAtt ⟨⟨CourseSum, Dept⟩⟩ [ {k1, k2, k1} | {k1, k2} ← ⟨⟨CourseSum⟩⟩ ];
addAtt ⟨⟨CourseSum, CID⟩⟩ [ {k1, k2, k2} | {k1, k2} ← ⟨⟨CourseSum⟩⟩ ];
addAtt ⟨⟨CourseSum, Total⟩⟩ [ {x, y, z} | {x, y, z} ←
                          (gc sum [ {k1, k3, x} | {k1, k2, k3, x} ← ⟨⟨Details, Mark⟩⟩ ] ) ];
addAtt ⟨⟨CourseSum, Avg⟩⟩ [ {x, y, z} | {x, y, z} ←
                          (gc avg [ {k1, k3, x} | {k1, k2, k3, x} ← ⟨⟨Details, Mark⟩⟩ ] ) ];
...

```

### 3 Expressing Schema and Data Model Evolution

In a heterogeneous data warehousing environment, it is possible for either a data source schema or the integrated database schema to evolve. This schema evolution may be a change in the schema, or a change in the data model in which the schema is expressed, or both. AutoMed transformations can be used to express the schema evolution in all three cases:

- (a) Consider first a schema  $S$  expressed in a modelling language  $\mathcal{M}$ . We can express the evolution of  $S$  to  $S^{new}$ , also expressed in  $\mathcal{M}$ , as a series of primitive transformations that rename, add, extend, delete or contract constructs of  $\mathcal{M}$ . For example, suppose that the relational schema  $LS_1$  in the above example



evolves so its three tables become a single table with an extra column for the course ID. This evolution is captured by a pathway which is identical to the pathway  $LS_1 \rightarrow CS_1$  given above.

This kind of transformation that captures well-known equivalences between schemas can be defined in AutoMed by means of a parametrised transformation *template* which is schema- and data-independent. When invoked with specific schema constructs and their extents, a template generates the appropriate sequence of primitive transformations within the Schemas & Transformations Repository — see [5] for details.

- (b) Consider now a schema  $S$  expressed in a modelling language  $\mathcal{M}$  which evolves into an equivalent schema  $S^{new}$  expressed in a modelling language  $\mathcal{M}^{new}$ . We can express this translation by a series of **add** steps that define the constructs of  $S^{new}$  in  $\mathcal{M}^{new}$  in terms of the constructs of  $S$  in  $\mathcal{M}$ . At this stage, we have an intermediate schema that contains the constructs of both  $S$  and  $S^{new}$ . We then specify a series of **delete** steps that remove the constructs of  $\mathcal{M}$  (the queries within these transformations indicate that these are now redundant constructs since they can be derived from the new constructs).

For example, suppose that XML schema  $LS_2$  in the above example evolves into an equivalent relational schema consisting of single table with one column per attribute of  $LS_2$ . This evolution is captured by a pathway which is identical to the pathway  $LS_2 \rightarrow CS_2$  given above.

Again, such generic inter-model translations between one data model and another can be defined in AutoMed by means of transformation templates.

- (c) Considering finally to an evolution which is both a change in the schema and in the data model, this can be expressed by a combination of (a) and (b) above: either (a) followed by (b), or (b) followed by (a), or indeed by interleaving the two processes.

## 4 Handling Schema Evolution

In this section we consider how the general integration network illustrated in Figure 1 is evolvable in the face of evolution of a local schema or the warehouse schema. We have seen in the previous section how AutoMed transformations can be used to express the schema evolution if either the schema or the data model changes, or both. We can therefore treat schema and data model change in a uniform way for the purposes of handling schema evolution: both are expressed as a sequence of AutoMed primitive transformations, in the first case staying within the original data model, and in the second case transforming the original schema in the original data model into a new schema in a new data model.

In this section we describe the actions that are taken in order to evolve the integration network of Figure 1 if the global schema  $GS$  evolves (Section 4.1) or if a local schema  $LS_i$  evolves (Section 4.2). Given an evolution pathway from a schema  $S$  to a schema  $S^{new}$ , in both cases each successive primitive transformation within the pathway  $S \rightarrow S^{new}$  is treated one at a time. Thus, we describe in sections 4.1 and 4.2 the actions that are taken if  $S \rightarrow S^{new}$  consists of just

one primitive transformation. If  $S \rightarrow S^{new}$  is a composite transformation, then it is handled as a sequence of primitive transformations. Our discussion below assumes that the primitive transformation being handled is adding, removing or renaming a construct of  $S$  that has an underlying data extent. We do not discuss the addition or removal of constraints here as these do not impact on the materialised data, and we make the assumption that any constraints in the pathway  $S \rightarrow S^{new}$  have been verified as being valid.

#### 4.1 Evolution of the global schema

Suppose the global schema  $GS$  evolves by means of a primitive transformation  $t$  into  $GS^{new}$ . This is expressed by the step  $t$  being appended to the pathway  $T_u$  of Figure 1. The new global schema is  $GS^{new}$  and its associated extension is  $GD^{new}$ .  $GS$  is now an intermediate schema in the extended pathway  $T_u;t$  and it no longer has an extension associated with it.  $t$  may be a **rename**, **add**, **extend**, **delete** or **contract** transformation. The following actions are taken in each case:

1. If  $t$  is **rename**  $c$   $c'$ , then there is nothing further to do.  $GS$  is semantically equivalent to  $GS^{new}$  and  $GD^{new}$  is identical to  $GD$  except that the extent of  $c$  in  $GD$  is now the extent of  $c'$  in  $GD^{new}$ .
2. If  $t$  is **add**  $c$   $q$ , then there is nothing further to do at the schema level.  $GS$  is semantically equivalent to  $GS^{new}$ . However, the new construct  $c$  in  $GD^{new}$  must now be populated, and this is achieved by evaluating the query  $q$  over  $GD$ .
3. If  $t$  is **extend**  $c$ , then the new construct  $c$  in  $GD^{new}$  is populated by an empty extent. This new construct may subsequently be populated by an expansion in a data source (see Section 4.2).
4. If  $t$  is **delete**  $c$   $q$  or **contract**  $c$ , then the extent of  $c$  must be removed from  $GD$  in order to create  $GD^{new}$  (it is assumed that this a legal deletion/contraction, e.g if we wanted to delete/contract a table from a relational schema, then first the constraints and then the columns would be deleted/contracted and lastly the table itself; such syntactic correctness of transformation pathways is automatically verified by AutoMed). It may now be possible to simplify the transformation network, in that if  $T_u$  contains a matching transformation **add**  $c$   $q$  or **extend**  $c$ , then both this and the new transformation  $t$  can be removed from the pathway  $US \rightarrow GS^{new}$ . This is purely an optimization — it does not change the meaning of a pathway, nor its effect on view generation and query/data translation. We refer the reader to [19] for details of the algorithms that simplify AutoMed transformation pathways.

In cases **2** and **3** above, the new construct  $c$  will automatically be propagated into the schema  $DMS$  of any data mart derived from  $GS$ . To prevent this, a transformation **contract**  $c$  can be prefixed to the pathway  $GS \rightarrow DMS$ . Alternatively, the new construct  $c$  can be propagated to  $DMS$  if so desired, and materialised there. In cases **1** and **4** above, the change in  $GS$  and  $GD$  may impact on the data marts derived from  $GS$ , and we discuss this in Section 4.3.

## 4.2 Evolution of a local schema

Suppose a local schema  $LS_i$  evolves by means of a primitive transformation  $t$  into  $LS_i^{new}$ . As discussed in Section 2, there is automatically available a reverse transformation  $\bar{t}$  from  $LS_i^{new}$  to  $LS_i$  and hence a pathway  $\bar{t}; T_i$  from  $LS_i^{new}$  to  $CS_i$ . The new local schema is  $LS_i^{new}$  and its associated extension is  $LD_i^{new}$ .  $LS_i$  is now just an intermediate schema in the extended pathway  $\bar{t}; T_i$  and it no longer has an associated extension.

$t$  may be a **rename**, **add**, **delete**, **extend** or **contract** transformation. In 1–5 below we see what further actions are taken in each case for evolving the integration network and the downstream materialised data as necessary.

We first introduce some necessary terminology: If  $p$  is a pathway  $S \rightarrow S'$  and  $c$  is a construct in  $S$ , we denote by  $descendants(c, p)$  the constructs of  $S'$  which are directly or indirectly dependent on  $c$ , either because  $c$  itself appears in  $S'$  or because a construct  $c'$  of  $S'$  is created by a transformation **add**  $c' q$  within  $p$  where the query  $q$  directly or indirectly references  $c$ . The set  $descendants(c, p)$  can be straight-forwardly computed by traversing  $p$  and inspecting the query associated with each **add** transformation within in.

1. If  $t$  is **rename**  $c c'$ , then schema  $LS_i^{new}$  is semantically equivalent to  $LS_i$ . The new transformation pathway  $T_i^{new} : LS_i^{new} \rightarrow CS_i$  is  $\bar{t}; T_i = \text{rename } c' c; T_i$ . The new local database  $LD_i^{new}$  is identical to  $LD_i$  except that the extent of  $c$  in  $LD_i$  is now the extent of  $c'$  in  $LD_i^{new}$ .
2. If  $t$  is **add**  $c q$ , then  $LS_i$  has evolved to contain a new construct  $c$  whose extent is equivalent to the expression  $q$  over the other constructs of  $LS_i$ . The new transformation pathway  $T_i^{new} : LS_i^{new} \rightarrow CS_i$  is  $\bar{t}; T_i = \text{delete } c q; T_i$ .
3. If  $t$  is **delete**  $c q$ , this means that  $LS_i$  has evolved to not include a construct  $c$  whose extent is derivable from the expression  $q$  over the other constructs of  $LS_i$ , and the new local database  $LD_i^{new}$  no longer contains an extent for  $c$ . The new transformation pathway  $T_i^{new} : LS_i^{new} \rightarrow CS_i$  is  $\bar{t}; T_i = \text{add } c q; T_i$ .

In the above three cases, schema  $LS_i^{new}$  is semantically equivalent to  $LS_i$ , and nothing further needs to be done to any of the transformation pathways, schemas or databases  $CD_1, \dots, CD_n$  and  $GD$ . This may not be the case if  $t$  is a **contract** or **extend** transformation, which we consider next.

4. If  $t$  is **extend**  $c$ , then there will be a new construct available from  $LS_i^{new}$  that was not available before. That is,  $LS_i$  has evolved to contain the new construct  $c$  whose extent is not derivable from the other constructs of  $LS_i$ . If we left the transformation pathway  $T_i$  as it is, this would result in a pathway  $T_i^{new} = \text{contract } c; T_i$  from  $LS_i^{new}$  to  $CS_i$ , which would immediately drop the new construct  $c$  from the integration network. That is,  $T_i^{new}$  is consistent but it does not utilize the new data.

However, recall that we said earlier that we assume no **contract** steps in the pathways from local schemas to their union schemas, and that all the data in  $LS_i$  should be available to the integration network. In order to achieve this, there are four cases to consider:

- (a)  $c$  appears in  $US_i$  and has the same semantics as the newly added  $c$  in  $LS_i^{new}$ . Since  $c$  cannot be derived from the original  $LS_i$ , there must be a transformation **extend**  $c$ , in  $CS_i \rightarrow US_i$ . We remove from  $T_i^{new}$  the new **contract**  $c$  step and this matching **extend**  $c$  step. This propagates  $c$  into  $CS_i$ , and we populate its extent in the materialised database  $CD_i$  by replicating its extent from  $LD_i^{new}$ .
- (b)  $c$  does not appear in  $US_i$  but it can be derived from  $US_i$  by means of some transformation  $T$ . In this case, we remove from  $T_i^{new}$  the first **contract**  $c$  step, so that  $c$  is now present in  $CS_i$  and in  $US_i$ . We populate the extent of  $c$  in  $CD_i$  by replicating its extent from  $LD_i^{new}$ . To repair the other pathways  $T_j : LS_j \rightarrow CS_j$  and schemas  $US_j$  for  $j \neq i$ , we append  $T$  to the end of each  $T_j$ . As a result, the new construct  $c$  now appears in all the union schemas. To add the extent of this new construct to each materialised database  $CD_j$  for  $j \neq i$ , we compute it from the extents of the other constructs in  $CS_j$  using the queries within successive **add** steps in  $T$ . We finally append the necessary new **id** steps between pairs of union schemas to assert the semantic equivalence of the construct  $c$  within them.
- (c)  $c$  does not appear in  $US_i$  and cannot be derived from  $US_i$ . In this case, we again remove from  $T_i^{new}$  the first **contract**  $c$  step so that  $c$  is now present in schema  $CS_i$ . To repair the other pathways  $T_j : LS_j \rightarrow CS_j$  and schemas  $US_j$  for  $j \neq i$ , we append an **extend**  $c$  step to the end of each  $T_j$ . As a result, the new construct  $c$  now appears in all the conformed schemas  $CS_1, \dots, CS_n$ . The construct  $c$  may need further translation into the data model of the union schemas and this is done by appending the necessary sequence,  $T$ , of **add/delete/rename** steps to all the pathways  $LS_1 \rightarrow CS_1, \dots, LS_n \rightarrow CS_n$ . We compute the extent of  $c$  within the database  $CD_i$  from its extent within  $LD_i^{new}$  using the queries within successive **add** steps in  $T$ . We finally append the necessary new **id** steps between pairs of union schemas to assert the semantic equivalence of the new construct(s) within them.
- (d)  $c$  appears in  $US_i$  but has different semantics to the newly added  $c$  in  $LS_i^{new}$ . In this case, we rename  $c$  in  $LS_i^{new}$  to a new construct  $c'$ . The situation reverts to adding a new construct  $c'$  to  $LS_i^{new}$ , and one of (a)-(c) above applies.

We note that determining whether  $c$  can or cannot be derived from the existing constructs of the union schemas in (a)–(d) above requires domain or expert human knowledge. Thereafter, the remaining actions are fully automatic.

In cases (a) and (b), there is new data added to one or more of the conformed databases which needs to be propagated to  $GD$ . This is done by computing  $descendants(c, T_u)$  and using the algebraic equivalences of Section 2.1 to propagate changes in the extent of  $c$  to each of its descendant constructs  $gc$  in  $GS$ . Using these equivalences, we can in most cases incrementally recompute the extent of  $gc$ . If at any stage in  $T_u$  there is a transformation **add**  $c' q$  where no equivalence can be applied, then we have to recompute the whole extent of  $c'$ .

In cases (b) and (c), there is a new schema construct  $c$  appearing in the  $US_i$ . This construct will automatically appear in the schema  $GS$ . If this is not desired, a transformation contract  $c$  can be prefixed to  $T_u$ .

5. If  $t$  is contract  $c$ , then the construct  $c$  in  $LS_i$  will no longer be available from  $LS_i^{new}$ . That is,  $LS_i$  has evolved so as to not include a construct  $c$  whose extent is not derivable from the other constructs of  $LS_i$ . The new local database  $LD_i^{new}$  no longer contains an extent for  $c$ .

The new transformation pathway  $T_i^{new} : LS_i^{new} \rightarrow CS_i$  is  $\bar{t}; T_i = \text{extend } c; T_i$ . Since the extent of  $c$  is now **Void**, the materialised data in  $CD_i$  and  $GD$  must be modified so as to remove any data derived from the old extent of  $c$ .

In order to repair  $CD_i$ , we compute  $\text{descendants}(c, LS_i \rightarrow CS_i)$ . For each construct  $uc$  in  $\text{descendants}(c, LS_i \rightarrow CS_i)$ , we compute its new extent and replace its old extent in  $CD_i$  by the new extent. Again, the algebraic properties of IQL queries discussed in Section 2.1 can be used to propagate the new **Void** extent of construct  $c$  in  $LS_i^{new}$  to each of its descendant constructs  $uc$  in  $CS_i$ . Using these equivalences, we can in most cases incrementally recompute the extent of  $uc$  as we traverse the pathway  $T_i$ .

In order to repair  $GD$ , we similarly propagate changes in the extent of each  $uc$  along the pathway  $T_u$ .

Finally, it may also be necessary to amend the transformation pathways if there are one or more constructs in  $GD$  which now will always have an empty extent as a result of this contraction of  $LS_i$ . For any construct  $uc$  in  $US$  whose extent has become empty, we examine all pathways  $T_1, \dots, T_n$ . If all these pathways contain an **extend**  $uc$  transformation, or if using the equivalences of Section 2.1 we can deduce from them that the extent of  $uc$  will always be empty, then we can suffix a **contract**  $gc$  step to  $T_u$  for every  $gc$  in  $\text{descendants}(uc, T_u)$ , and then handle this case as paragraph 4 in Section 4.1.

### 4.3 Evolution of downstream data marts

We have discussed how evolutions to the global schema or to a source schema are handled. One remaining question is how to handle the impact of a change to the data warehouse schema, and possibly its data, on any data marts that have been derived from it.

In [7] we discuss how it is possible to express the derivation of a data marts from a data warehouse by means of an AutoMed transformation pathway. Such a pathway  $GS \rightarrow DMS$  expresses the relationship of a data mart schema  $DMS$  to the warehouse schema  $GS$ . As such, this scenario can be regarded as a special case of the general integration scenario of Figure 1, where  $GS$  now plays the role of the single source schema, databases  $CD_1, \dots, CD_n$  and  $GD$  collectively play the role of the data associated with this source schema and  $DMS$  plays the role of the global schema. Therefore, the same techniques as discussed in sections 4.1 and 4.2 can be applied.

## 5 Concluding Remarks

In this paper we have described how the AutoMed heterogeneous data integration toolkit can be used to handle the problem of schema evolution in heterogeneous data warehousing environments so that the previous transformation, integration and data materialisation effort can be reused. Our algorithms are mainly automatic, except for the aspects that require domain or expert human knowledge regarding the semantics of new schema constructs.

We have shown how AutoMed transformations can be used to express schema evolution within the same data model, or a change in the data model, or both, whereas other schema evolution literature has focussed on just one data model. Schema evolution within the relational data model has been discussed in previous work such as [11, 12, 18]. The approach in [18] uses a first-order schema in which all values in a schema of interest to a user are modelled as data, and other schemas can be expressed as a query over this first-order schema. The approach in [12] uses the notation of a *flat scheme*, and gives four operators UNITE, FOLD, UNFOLD and SPLIT to perform relational schema evolution using the SchemaSQL language. In contrast, with AutoMed the process of schema evolution is expressed using a simple set of primitive schema transformations augmented with a functional query language, both of which are applicable to multiple data models.

Our approach is complementary to work on mapping composition, e.g. [20, 14], in that in our case the new mappings are a composition of the original transformation pathway and the transformation pathway which expresses the schema evolution. Thus, the new mappings are, by definition, correct. There are two aspects to our approach: (i) handling the transformation pathways and (ii) handling the queries within them. In this paper we have in particular assumed that the queries are expressed in IQL. However, the AutoMed toolkit allows any query language syntax to be used within primitive transformations, and therefore this aspect of our approach could be extended to other query languages.

Materialised data warehouse views need to be maintained when the data sources change, and much previous work has addressed this problem at the data level. However, as we have discussed in this paper, materialised data warehouse views may also need to be modified if there is an evolution of a data source schema. Incremental maintenance of schema-restructuring views within the relational data model is discussed in [10], whereas our approach can handle this problem in a heterogeneous data warehousing environment with multiple data models and changes in data models. Our previous work [7] has discussed how AutoMed transformation pathways can also be used for incrementally maintaining materialised views at the data level. For future work, we are implementing our approach and evaluating it in the context of biological data warehousing.

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