# Keyword Search over Relational Tables and Streams

ALEXANDER MARKOWETZ
University of Bonn
YIN YANG and DIMITRIS PAPADIAS
Hong Kong University of Science and Technology

Relational Keyword Search (R-KWS) provides an intuitive way to query relational data without requiring SQL, or knowledge of the underlying schema. In this article we describe a comprehensive framework for R-KWS covering snapshot queries on conventional tables and continuous queries on relational streams. Our contributions are summarized as follows: (i) We provide formal semantics, addressing the temporal validity and order of results, spanning uniformly over tables and streams; (ii) we investigate two general methodologies for query processing, graph based and operator based, that resolve several problems of previous approaches; and (iii) we develop a range of algorithms and optimizations covering both methodologies. We demonstrate the effectiveness of R-KWS, as well as the significant performance benefits of the proposed techniques, through extensive experiments with static and streaming datasets.

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## 1. INTRODUCTION

With the rise of the Web, the vast majority of users have adapted *KeyWord Search* (KWS) as a primary tool to access information. In conventional KWS,

Authors' addresses: A. Markowetz, Department of Computer Science, University of Bonn, Germany; email: alex@iai.uni-bonn.de; Y. Yang, D. Papadias, Department of Computer Science and Engineering, Hong Kong University of Science and Technology, Hong Kong, email: {yini, dimitris}@cse.ust.hk.

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each document/Web page constitutes one *unit of information*, and is considered a result if it contains a subset of the query's keywords. Recently, KWS has also been applied to relational DBMS, allowing data retrieval without SQL. In *Relational KeyWord Search* (R-KWS), the basic unit of information is a record/tuple. In contrast to KWS on documents, R-KWS queries cannot be answered by inspecting records individually. Instead, results have to be *constructed* by joining tuples. R-KWS has several benefits over SQL queries. First, it liberates users from studying a (possibly messy) database schema. Queries can instead be issued without knowledge of tables, their attributes, or join conditions. Second, R-KWS allows querying for terms in unknown locations (tables/attributes). Finally, a keyword query replaces numerous complex SQL statements, whose number (frequently in the thousands) prohibits hand-coded SQL on any database with a nontrivial schema. For such broad queries, R-KWS poses the only practical solution.

The external simplicity of R-KWS hides a great internal complexity, involving a vast search space. Specifically, an R-KWS system must explore all possible keyword occurrences (in every table and attribute), as well as their interactions. There are two general methodologies for processing R-KWS queries: *Graph Based* (GB) and *Operator Based* (OB). The former maintains an in-memory *data graph*, where data tuples are represented by nodes, connected through edges iff they can be joined. Results (subgraphs) are retrieved by means of graph traversal. In contrast, OB R-KWS enumerates and executes a set of operator trees, similar to hand-coded SQL. These trees are generated exhaustively, such that *any* combination of keyword occurrences is found and returned to the user.

In this article we propose a general framework covering both snapshot R-KWS on static tables and continuous queries on relational streams. Our contributions are summarized as follows.

- —We propose uniform semantics that accommodate general join conditions, and take into account temporal validity and order of results.
- —We present GB and OB query processing methodologies that are applicable to both streams and static databases. The methodologies resolve problems of previous approaches related to duplicate elimination.
- —We devise a streamlined graph traversal that significantly accelerates GB query processing. Keyword labeling schemes further improve performance for continuous queries, by indicating which keywords can be reached from a given node.
- —We design a highly efficient algorithm for OB systems that integrates operator trees into a *mesh* which can grow and shrink dynamically, adapting to data characteristics.
- —We experimentally compare our methods using static and streaming data, and investigate their effectiveness under different settings.

The remainder of the article is structured as follows. Section 2 outlines related work on R-KWS and data streams. Sections 3 and 4 present generalized R-KWS semantics and processing methodologies for static databases and

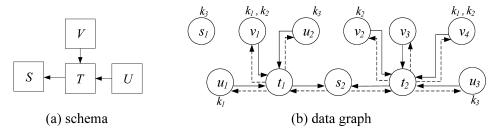


Fig. 1. Database and graph representation.

data streams, respectively. Section 5 enhances the efficiency of GB using keyword labels, and Section 6 proposes optimizations for OB. Section 7 evaluates the benefits of our algorithms experimentally. Finally, Section 8 concludes the work.

#### 2. RELATED WORK

Section 2.1 outlines GB, and Section 2.2 OB systems. For each approach we discuss one representative system, namely *Banks* and *Discover*, respectively. Section 2.3 presents existing work that falls outside these categories. Section 2.4 provides an overview on data streams.

#### 2.1 Graph-Based R-KWS

Graph-based R-KWS query processing was introduced in Banks [Bhalotia et al. 2002], the first system for relational keyword search. Banks defines R-KWS semantics on a graph representation of the database. Each node in the data graph G corresponds to a tuple, and edges connect nodes/tuples that can be joined. Figure 1(b) illustrates the data graph for a database consisting of four tables (S, T, U, V), whose schema is shown in Figure 1(a). In our notation,  $s_i$  signifies a tuple of  $S, t_i$  one of T, etc. Keywords  $\{k_1, k_2, k_3\}$  are noted next to the tuples in which they occur, for example,  $k_1$  and  $k_2$  exist in  $v_1$ . Two tuples (e.g.,  $s_2, t_1$ ) are connected in G by a (solid) edge, iff: (i) their corresponding relations (S, T) are connected in the schema, and (ii) the tuples satisfy the corresponding join conditions. Banks limits join conditions to foreign to primary key relationships, and uses a pair of directed edges for each connection. Specifically, a  $forward\ edge$  (shown in solid lines) points from the tuple containing a foreign key to its primary key partner (e.g.,  $v_1$  has a foreign key referencing  $t_1$ ), and a  $backward\ edge$  (in dotted lines) runs in the opposite direction.

The result of an R-KWS query  $q = \{k_1, \ldots, k_m\}$  is the set of trees in G satisfying the following conditions: (i) Edges (forward or backward) point from the root towards the leafs, (ii) each leaf node contains at least one keyword in q, and (iii) the nodes jointly cover all terms of q. Assuming the data graph of Figure 1 and the query  $q = \{k_1, k_2, k_3\}$ , Figure 2 depicts several result trees. Observe that the two trees in Figures 2(a) and 2(b) differ only in the direction of the edge between  $t_1$  and  $t_2$ . From a user's perspective, these are defacto duplicates, since both edges  $t_1$  and  $t_2$  and  $t_3$  and  $t_4$  and  $t_4$  represent the same

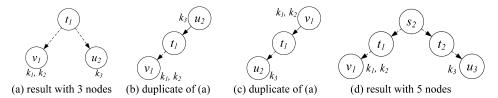


Fig. 2. Example outputs of *Banks* for  $q = \{k_1, k_2, k_3\}$ .

association between  $t_1$  and  $v_1$ . Similarly, reversing the edge between  $t_1$  and  $v_1$  in Figure 2(a) yields another duplicate (Figure 2(c)). In general, for every result r with |r| nodes, Banks produces |r| copies, each of which has one of the |r| nodes as the root.

Banks maintains the data graph in main memory. In order to reduce the space consumption, the graph only consists of tuple identifiers; concrete attribute values are stored on disk. Bhalotia et al. [2002] report 100MB of memory for the graph of the DBLP dataset, containing roughly 120,000 nodes and 310,000 edges. For an incoming query, the system first identifies all tuples with keywords, using a disk-resident inverted index. Next, it traverses G for result trees. Specifically, it initiates a graph traversal at each node containing a keyword, following edges in their backward direction. Visited tuples record the source nodes and their keywords. Once a node has been visited by all keywords, it generates a result by following the reverse paths to the sources. Assuming  $q = \{k_1, k_2, k_3\}$  in Figure 1, traversals are initiated at  $v_1$  and  $u_3$  (among others), since these nodes contain keywords. When the traversals meet (e.g., at  $s_2$ ) a new result is created, consisting of  $s_2$  and the paths to  $v_1$  and  $u_3$  (Figure 2(d)). In contrast to this simplified description, Banks weighs nodes and edges to rank results, and attempts an early generation of high scoring results, using Dijkstra's algorithm.

Kacholia et al. [2005] accelerate GB query processing by bidirectional expansion that traverses the data graph both backwards (from nodes that can reach all keywords, as in the original system) and forwards (from nodes containing at least one keyword) simultaneously. From a theoretical perspective, Kimelfeld and Sagiv [2006, 2005] and Golenberg et al. [2008] investigate the complexity of relational keyword search, and propose algorithms that produce top-k results with the guarantee of polynomial delay. Blinks [He et al. 2007] reduces the search space through a precomputed two-level reachability index of the data graph. Reachability indexing is further explored by Markowetz et al. [2009].

## 2.2 Operator-Based R-KWS

Discover [Hristidis and Papakonstantinou 2002] translates an R-KWS query into a series of SQL statements, executed directly on the underlying DBMS. The union of results answers the query. Although this methodology differs radically from GB, its semantics are also defined on a data graph G, which remains conceptual and is never materialized. Similarly to Banks, the system

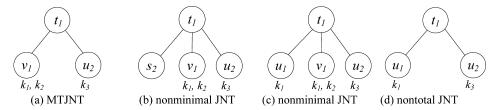


Fig. 3. Join networks of tuples.

is restricted to foreign-to-primary key relationships. However, Hristidis and Papakonstantinou [2002] avoid de facto duplicates, by considering a data graph without back-edges. Given the schema and data of Figure 1, the graph remains the same, but the dotted edges are removed. Consequently, the trees of Figures 2(b) and 2(c) do not exist in this graph, and hence do not form results. Tuples  $t_1$ ,  $v_1$  and  $u_2$  (Figure 3(a)) are still connected and intuitively constitute a result that is equivalent to those of Figures 2(b) and 2(c). However, the structure of Figure 3(a) does not have a distinct root node. Accordingly, *Discover* evades the concept of trees through *Join Networks of Tuples* (JNT), which are connected acyclic components of G.

Figure 3 depicts several JNT for the data of Figure 1. JNT lack a root node: Depending on the structure's orientation, every node can serve as root. Likewise, there are no distinct leaf nodes. We hence refer to nodes with degree ≤ 1 (in the JNT) as terminal nodes. Given a query, a JNT is called total iff it contains all keywords. The JNT in Figures 3(a) through 3(c) are total for q= $\{k_1,k_3,k_3\}$ ; the one in Figure 3(d) lacks  $k_2$ . A JNT is called *Minimal Total JNT* (MTJNT) iff it is impossible to remove any node and find the remainder to be total. In particular, minimalism is satisfied iff every terminal node contains at least one unique keyword (that is not contained in any other node of the JNT). The JNT in Figure 3(b) (respectively 3(c)) is not minimal because after node s<sub>2</sub> (respectively  $u_1$ ) is removed, the remainder still constitutes a total JNT. Discover answers an R-KWS query q with the set of MTJNT. In contrast, Banks only requires terminal nodes to contain some keyword. Consequently, its results form a superset of *Discover's*. For instance, the JNT of Figure 3(c) is not a result in Discover, but the equivalent tree is a result of Banks. Finally, Discover imposes an upper limit of  $T_{max}$  nodes per MTJNT, in order to avoid long chains of joins, which usually lead to uninteresting results.

Discover's query processing relies on an expanded schema<sup>1</sup> and Candidate Networks (CN). Specifically, given a query  $q = \{k_1, \ldots, k_m\}$ , its corresponding expanded schema EG(q) is a graph, constructed as follows. For each table S in the database and every set of keywords  $K \subseteq q$ , the expanded schema contains a node  $S\{K\}$ , where K may be empty. A pair of nodes  $S\{K\}$  and  $T\{K'\} \in EG(q)$  is connected by an edge iff their base relations S and T can be joined through a foreign-key to primary-key relationship. For example, assuming  $q = \{k_1, k_2, k_3\}$ , there are eight nodes in EG(q) for each relation S, namely  $S\{\}$ ,  $S\{k_1\}$ ,

<sup>&</sup>lt;sup>1</sup>The *expanded schema* is proposed, but not named, in Hristidis and Papakonstantinou [2002]. We introduce the term for easier reference.

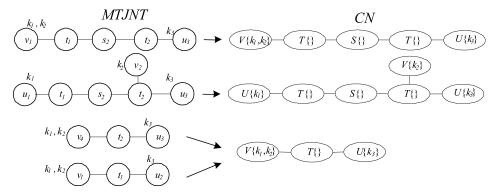


Fig. 4. Examples of MTJNT and CN.

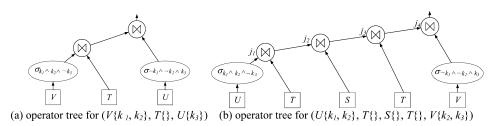


Fig. 5. Example operator trees in Discover.

 $S\{k_2\}$ ,  $S\{k_3\}$ ,  $S\{k_1, k_2\}$ ,  $S\{k_1, k_3\}$ ,  $S\{k_2, k_3\}$  and  $S\{k_1, k_2, k_3\}$ . Every  $S\{K\}$  is connected with each  $T\{K'\}$  since S and T are related in the schema of Figure 1. Candidate Networks (CN) are projections of MTJNT onto the expanded schema. In particular, a tuple s of relation S maps to node  $S\{K\} \in EG(q)$ , iff s contains all keywords in K, but does not contain any other term in  $q \setminus K$ . Figure 4 illustrates several MTJNT and their corresponding CN. For instance, the MTJNT  $\{v_1, t_1, u_2\}$  maps to the CN  $\{v_1, v_2\}$ ,  $\{v_3\}$ . Observe that an MTJNT projects to a unique CN, while it is possible for multiple MTJNT, for example,  $\{v_1, t_1, u_2\}$  and  $\{v_4, v_2, v_3\}$ , to map to the same CN. In contrast to MTJNT, a CN potentially contains multiple copies of the same node in EG(q), such as  $T\{\}$ .

Discover answers a query by: (i) generating all possible CN, (ii) converting CN to operator trees, and (iii) executing operator trees to produce results. We first clarify the translation from CN to operator trees. Given a CN (e.g., at the bottom of Figure 4), Discover generates the corresponding operator tree, shown in Figure 5(a) as follows. Each node with an nonempty keyword set  $(V\{k_1, k_2\})$  and  $U\{k_3\}$  translates into a selection over the base table  $(\sigma_{k1} \wedge_{k2} \wedge \neg_{k3} V)$  and  $(\sigma_{k1} \wedge_{k2} \wedge_{k3} U)$ , reporting tuples that contain precisely those keywords  $(\{k_1, k_2\})$  and  $\{k_3\}$ . The remaining CN nodes with an empty keyword set (e.g.,  $T\{\}\}$ ) map to the base relation (e.g., T) without selection, referred to as free tuple set [Hristidis and Papakonstantinou 2002]. On top of the selections and free tuple sets, Discover builds a join tree, with conditions placed according to the edges of the CN. In our example, the edge between  $V\{k_1, k_2\}$  and T translates into the foreign-key join between  $\sigma_{k1} \wedge_{k2} \wedge \neg_{k3} V$  and T.

Note that executing the tree in Figure 5(a) may yield joined tuples that are not minimal. A tuple  $v \in V$  must have keywords  $k_1,k_2$  to pass the corresponding selections. If there is a tuple  $t \in T$  containing  $k_3$  Discover may output v-t-u, because it does not impose any restriction on keywords in T. However, v-t-u is not a minimal result; node u is redundant, since v-t already includes all keywords. Moreover, the tree in Figure 5(a) may produce MTJNT that do not map to the corresponding CN (bottom of Figure 4) and thus lead to duplicates. Assume tuples v'-t'-u' joined by the operator tree. If t' contains keyword  $k_1$ , the resulting MTJNT v'-t'-u' maps to a different CN ( $V\{k_1,k_2\}$ ,  $T\{k_1\}$ ,  $U\{k_3\}$ ), whose operator tree produces a duplicate copy. Finally, complex operator trees may generate results that are not even JNT, let alone MTJNT. Consider the tree shown in Figure 5(b), translated from the topmost CN in Figure 4. Since node  $T\{$ } is involved twice, it is possible for a result to contain the same tuple  $t \in T$  twice, for example, u-t-s-t-v, violating the requirement that JNT be acyclic.

In addition, the CN generation of *Discover* suffers from duplicates. Its algorithm for creating CN maintains a queue Q of CN fragments. Initially, Q contains only one fragment, consisting of a single node  $T\{k_1\}$ , where T is an arbitrary relation, and  $k_1$  is the first keyword in q. At each step, *Discover* extracts a CN fragment from Q, and adds a new node, forming a larger fragment. A complete CN is generated whenever a fragment contains all keywords. Duplicate candidate networks result from the undirected addition of nodes. Assume that an initial fragment  $(T\{k_1\})$  is expanded by adding  $U\{k_2\}$  and  $V\{k_3\}$  (among others). Consequently, two new CN-fragments are inserted into the heap. When  $(T\{k_1\}, U\{k_2\})$  is later dequeued, it is expanded by adding  $V\{k_3\}$ , generating the  $CN(T\{k_1\}, U\{k_2\}, V\{k_3\})$ . Similarly,  $(T\{k_1\}, V\{k_3\})$  is expanded by adding  $U\{k_2\}$ , generating the CN ( $T\{k_1\}$ ,  $V\{k_3\}$ , $U\{k_2\}$ ). However, these two CN are identical. Duplicate CN translate into identical operator trees, and perform redundant computations, exacerbating the problem of duplicate MTJNT. Besides Discover, there are several other OB implementations, including Mragvati Sarda and Jain 2001], DBXplorer [Agrawal et al. 2002], and SPARK [Luo et al. 2007]. Unfortunately, they do not describe CN generation, nor address the previous issues. Markowetz et al. [2009] eliminate unproductive CN through reachability indexing, in order to accelerate query processing.

## 2.3 Additional Work on Relational Keyword Search

Databases were first modeled as a graph by Dar et al. [1998]. Besides GB and OB, there is a Materialization-Based (MB) approach to R-KWS, followed by Ekso [Su and Widom 2005] and Ease [Li et al. 2008]. Relying on extensive precomputations, MB enumerates all subgraphs in G of a certain diameter during preprocessing. For each such subgraph  $G_S$ , it constructs a virtual document with the attribute values of all nodes in  $G_S$ . These documents are materialized on disk and organized in an inverted index. Upon query arrival, the system uses this index to determine the subgraphs containing all keywords, from which it removes unnecessary nodes to restore minimality, and eliminates duplicate results. Such an index grows quickly and soon exceeds the size of the database.

In addition, MB is not applicable to dynamic scenarios involving data streams, since the data are not available in advance, rendering precomputations impossible.

R-KWS has been extended in various directions. Hristidis et al. [2003] relax the notion of minimality of *Discover* by requiring each terminal node in a result to contain a keyword, but not necessarily a unique one. Other systems, such as Liu et al. [2006], support phrases (e.g., the title of an article) and lists of synonyms (e.g., "car OR automobile OR vehicle"). Several authors propose ranking schemes for top-k processing, based on IR ranking functions [Hristidis et al. 2003; Balmin et al. 2004; Chaudhuri et al. 2004; Golenberg et al. 2008; Li et al. 2008; Liu et al. 2006; Luo et al. 2007]. Additionally, Golenberg et al. [2008] penalize results for overlap (redundancy). KWS has also been extended to XML databases, using various semantics [Hristidis et al. 2003; Guo et al. 2003; Cohen et al. 2005; Xu and Papakonstantinou 2005; Liu and Chen 2007]. XML elements form nodes in a data graph; edges connect elements that are related through containment or by reference. Query processing depends on the database's physical layout. Ease [Li et al. 2008] supports uniform keyword search over structured, unstructured, and semistructured data. Furthermore, R-KWS has been applied to distributed systems containing multiple, possibly heterogeneous, data sources [Sayyadian et al. 2007]. The methods of Yu et al. [2007] and Vu et al. [2008] determine the most promising source for an R-KWS query among a set of distributed databases, using their summaries. Finally, Wu et al. [2007] propose keyword-driven OLAP, and De Felipe et al. [2008] investigate R-KWS on spatial databases.

## 2.4 Data Streams

There is an extensive body of literature on relational data streams. Under this paradigm, data elements (relational tuples) from various sources are collected at a *Data Stream Managing System* (DSMS), where users register continuous queries. When a new tuple arrives, all relevant queries are re-evaluated. Query processing is usually performed by routing tuples through trees of operators, resembling their traditional counterparts such as selections or joins. Influential DSMS systems include: (i) *Aurora* [Abadi et al. 2003], targeting mainly sensor data, (ii) *TelegraphCQ* [Chandrasekaran et al. 2003], focusing on the novel Eddy operator [Avnur and Hellerstein 2000], (iii) *Stream* [Arasu et al. 2006], designed as a general-purpose DSMS, and (iv) *Pipes* [Krämer and Seeger 2004]. In-depth surveys on DSMS can be found in Babock et al. [2002] and Golab and Öszu [2003].

Depending on the application characteristics, DSMS follow different models regarding the validity of tuples. One popular approach assumes a *sliding window* of a given time frame w, that is, a tuple s expires w time units after its arrival. In this case, all arrivals in the system correspond to insertions; deletions are implicit. Another common model assumes *positive-negative* tuples, that is, the DSMS receives a negative tuple -s that takes the same route through the operator tree as s, and erases all occurrences of its positive counterpart. In both cases, the *lifespan* of a tuple s is the interval  $[s.t_{start}, s.t_{end})$ 

between its arrival s.t<sub>start</sub> and the (implicit or explicit) deletion s.t<sub>end</sub>. Two tuples can be joined while their lifespans overlap. An abstract join operator jcontains two input buffers j.left-buffer and j.right-buffer, storing alive tuples from the left and right input. When a new tuple s arrives from the left input, it is inserted in *j.left-buffer* and subsequently compared against all tuples t in *j.right-buffer*. For every pair that fulfills the join condition, a new composite tuple c is created and passed onto higher operators. The lifespan of c is commonly defined as the intersection of participating tuples' lifespans; for example,  $c.t_{start} = max(s.t_{start} + t.t_{start})$  and  $c.t_{end} = min(s.t_{end} + t.t_{end})$ . For sliding windows, c's lifespan is known at its creation time, and buffers are purged periodically of expired tuples. In the case of positive-negative tuples, the expiration time is not known. Instead, when a negative tuple -s arrives in the left input, s is removed from j.left-buffer and subsequently probed against j.right-buffer. For any join result c, a negative output -c is passed on to higher operators, informing them about c's expiration. Tuples from the right input are handled symmetrically. Various implementations differ mainly in the organization of buffers, such as, lists or hash tables.

KWS has also been applied to streaming documents (e.g., continuously arriving news articles). With few exceptions [Yan and Garcia-Molina 1999; Fabret et al. 2001; Irmak et al. 2006], most related work is proprietary. The main difference with respect to our work is that documents do not have to be joined, but are evaluated individually (as in traditional KWS). In a poster, Hristidis et al. [2006] propose KWS over multiple textual streams. Similarly to our work, results are constructed by combining units of information (emails, news articles) from several streams. The authors, however, do not follow a relational model, leading to several key differences with our problem setting. First, tuples in Hristidis et al. [2006] have only one attribute: their text. Second, only tuples that contain keywords can contribute to a result. Third, and most significantly, combinations (joins) of several tuples are not evaluated upon their (textual) attribute, but tuples can always be joined, as long as the data streams from which they origin are sufficiently correlated. The correlation between streams is continuously updated, and stored in a stream schema. Unfortunately, the poster does not provide a formal definition of semantics, or details about algorithms and experiments.

In Markowetz et al. [2007], we introduce keyword search over relational streams, focusing exclusively on OB processing. This article extends our previous work on the following aspects: (i) We cover both static tables and streams, (ii) we present homogenized R-KWS semantics and query processing techniques for conventional databases, and remedy several common problems encountered by previous systems, (iii) we include GB processing, and (iii) we develop a novel optimization for OB processing.

## 3. RELATIONAL KEYWORD SEARCH ON TABLES

This section discusses methods for graph- and operator-based processing that avoid the shortcomings of prior systems and significantly improve performance of R-KWS in conventional databases. In our discussion, we assume an

```
GB (DataGraph G, InvertedIndex I, Query q) // Performs Graph Based R-KWS

1. Initialize the result set RS to Ø

2. For each sn \in I-invertedList(k_I) // for every node sn containing a selected keyword k_I of q

3. List RS_{sn} = GSearch(G, q, sn)

4. For each MTJNT r \in RS_{sn}

5. If (r includes a node n containing k_I) AND (n.nid < sn.nid)

6. Discard r // this MTJNT has already been created by GSearch(G, q, n)

7. Else Insert r into RS

8. Return RS
```

Fig. 6. Algorithm GB.

undirected data graph G that contains neither directional information nor backedges. Two tuples are connected by an edge iff they satisfy the join condition (symmetric or asymmetric), as specified by the application. In practice, many important join conditions, such as the similarity of textual attributes [Chaudhuri et al. 1995; Gravano et al. 2003], are symmetric. Undirected edges also avoid the de facto duplicates of Banks. If an application requires results to contain directed edges, it can fetch this information from the schema during post-processing at negligible cost.

Similar to Discover, we answer R-KWS queries with the set of MTJNT in G that contain up to  $T_{max}$  nodes. Recall from Section 2.2 that the concepts related to MTJNT do not involve edge directions, and thus directly extend to the undirected model. The parameter  $T_{max}$  allows adjusting runtime cost without restricting R-KWS semantically. Depending on the application, it can be set to an arbitrarily high number, including infinity. We follow the definitions of Discover because they are well understood, popular, and extend easily to data streams. However, we do not champion any particular R-KWS semantics. Similarly, we do not emphasize the ranking of results, since output from streams naturally follows a temporal order. The next two sections describe GB and OB query processing on static data. Both frameworks extend to alternative semantics and support arbitrary monotonous ranking functions.

### 3.1 Graph-Based Processing

Our method loosely follows the general paradigm of previous systems. Specifically, given an inverted index I (on disk), it traverses an undirected data graph G (in memory), searching for results. We reduce the problem of finding all results in the graph to retrieving those containing a particular node. By definition, every MTJNT includes a node sn containing an arbitrary (but fixed) query keyword, say  $k_1$ . It is hence sufficient to identify tuples/nodes containing this term (using I), and then traverse G locally, searching for MTJNT. Figure 6 depicts the pseudocode for this query processing method, referred to as GB. Specifically, GB initiates a graph traversal around each node  $sn \in I$ .invertedList( $k_1$ ), using the GSearch procedure (line 3). Note that it is possible for an MTJNT r to contain multiple nodes with  $k_1$ . In this case, r could wrongly be discovered by multiple calls to GSearch, each starting with a different seed node. To avoid such duplicates, GB assigns a distinct ID n.nid to each node  $n \in G$ , and reports a result r only if the current seed node sn of GSearch has the smallest ID from all nodes in r that contain  $k_1$  (lines 5 through 7). The concrete values for node

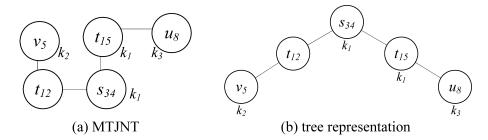


Fig. 7. An MTJNT and its unique representation as a tree.

```
GSearch (DataGraph G, Query q, Node sn)
                                                                      // Finds all MTJNT that contain a given node sn
    Initialize result set RS to empty
                                        // stores intermediate trees
     Initialize queue Q to empty
     Insert a tree into Q consisting of a single node sn (the root)
3.
4.
     While (Q is not empty)
          De-queue the first tree t_{old} in Q
          For each node nl on the rightmost root-to-leaf path in t_{old}
6.
              For each neighbor n_{new} of nl in G, where n_{new} \notin t_{old}
                                                                                 // n_{new} must not already exist in t_{old}
8.
                  If (n_{new}.nid > nc.nid, for every child nc of nl)
                                                                                // the nid must exceed the siblings
                      Create a new tree t_{new} by adding n_{new} as the rightmost child of nl
10.
                      If (t_{new} is an MTJNT)
11.
                          Insert t_{new} into RS;
                      Else
12.
13.
                          If (t_{new} \text{ contains less than } T_{max} \text{ nodes})
14.
                          AND (each terminal node, except those on the rightmost path, contains a unique keyword in q)
                              Append t_{new} to Q;
                                                                      // t_{new} still has the potential of forming an MTJNT
15.
16.
                          Else, discard t_{new}
17. Return RS
```

Fig. 8. Algorithm GSearch.

IDs do not influence correctness as long as no two nodes share the same ID. For ease of presentation, in the following we suppose that node IDs follow a lexicographic order on relation and tuple-ID; for example, for  $s_1, s_2 \in S$  and  $t_1 \in T$ , we have  $s_1.nid < s_2.nid < t_1.nid$ .

It remains to clarify the data graph traversal. In general, starting from the seed node sn, an MTJNT can be discovered in many ways. Any efficient algorithm, however, should enumerate each MTJNT precisely once. Additionally, its traversal should not stretch over the entire data graph, but must remain constrained to sn's vicinity. At the same time, it has to be extensive enough to ensure that all MTJNT containing sn are discovered. In order for GSearch to meet these criteria, we model each MTJNT as a unique tree, whose root consists of the seed node sn, while child nodes are ordered left-to-right by increasing nid. Given  $q = \{k_1, k_2, k_3\}$  and assuming  $sn = s_{34}$ , Figure 7 depicts an MTJNT and its tree representation. Node  $s_{34}$  serves as the root and its children ( $t_{12}$  and  $t_{15}$ ) are ordered left-to-right.

The *GSearch* algorithm, illustrated in Figure 8, enumerates all possible trees in G rooted at sn, in the lexicographic order of their preorder traversal. Whenever the algorithm encounters a tree that corresponds to an MTJNT, it reports a result. Specifically, *GSearch* maintains a queue Q of trees, each constituting a fraction of a potential MTJNT. At every step, one such tree  $t_{old}$  is de-queued

(line 5) and expanded by adding one new node  $n_{new}$ , resulting in a new tree  $t_{new}$ . To ensure that the preorder traversal of  $t_{new}$  is lexicographically greater than that of  $t_{old}$ , we enforce the following two requirements. First,  $n_{new}$  can only become child of a node nl on the rightmost root-to-leaf path of  $t_{old}$  (line 6). Second, the sid of  $n_{new}$  must exceed that of its siblings' (line 8). For example, a new node  $n_{new}$  can only be added to nodes  $s_{34}$ ,  $t_{15}$  or  $u_8$  in the tree of Figure 7(b). If  $n_{new}$  is added to under  $s_{34}$ , the nid of  $n_{new}$  has to be larger than that of nodes  $t_{12}$  and  $t_{15}$ . The new tree  $t_{new}$  falls into one of three categories: (i) It forms an MTJNT, and is included in the result set (lines 10 and 11); (ii) it has the potential to become an MTJNT, and is inserted in Q to be expanded later (lines 13 through 15); (iii) none of the previous and the tree can be safely discarded (line 16). In order to form an MTJNT at a later time (the second category), a tree must contain less than  $T_{max}$  nodes (line 13), and all terminal nodes, except on the rightmost path, must have a unique keyword (line 14), that is, a keyword that does not exist in any other node in  $t_{new}$ . The algorithm terminates when Q becomes empty.

GSearch can accommodate a looser definition of minimality [Hristidis et al. 2003] by simply relaxing the condition of line 14. Furthermore, it can capture phrases and synonyms [Liu et al. 2006] by treating them as single keywords. In particular, a node is considered to contain a synonym list l (respectively, phrase), if and only if the corresponding tuple contains some (respectively, all) keywords in l. Similar modifications for phrases and synonyms also apply to the rest of the proposed methods. The following two lemmas establish the correctness of GSearch and GB, respectively.

Lemma 3.1. GSearch computes the set of MTJNT containing node sn correctly, completely, and without duplicates.

PROOF. Correctness is trivial since GSearch explicitly verifies that every output constitutes an MTJNT. Completeness is established because: (i) without pruning, GSearch enumerates every possible tree in the data graph containing sn, including those corresponding to MTJNT, and (ii) the pruning conditions eliminate exclusively trees that are guaranteed not to lead to MTJNT. Finally, GSearch does not generate duplicates, because it discovers trees at most once, in increasing order of their preorder traversal.  $\Box$ 

Lemma 3.2. GB answers an R-KWS query q correctly, completely, and without duplicates.

PROOF. The correctness of GB follows directly from the corresponding property of GSearch. Its results are complete, because each MTJNT must include at least one node containing  $k_1$  and GSearch is called for every such node. Finally, since GSearch is free of duplicates, GB could only produce duplicates if the same MTJNT was discovered by separate calls of GSearch. Such an MTJNT would have to include several nodes containing  $k_1$  (potential root nodes). However, lines 5 and 6 of GB ensure that MTJNT are recorded only when GSearch has been called for the node with the smallest nid. Thus, GB is free of duplicates.  $\Box$ 

Finally, note that the pseudocode of Figure 8 is *iterative* and maintains state (subgraphs that have not yet been expanded) by storing trees in a queue. In addition, we perform the duplicate check (lines 5 and 6 of GB) as part of GSearch; that is, if  $n_{new}$  contains keyword  $k_1$ , its nid must be larger than that of sn, in order to be added to  $t_{new}$ . Although this design avoids duplicates earlier, we present this check outside GSearch in order to separate the functionality of GB and GSearch. GB can be adapted to produce results according to a monotone ranking function f, as follows. Instead of calling GSearch for each seed node individually, we initialize Q with the set of all seed nodes, and generate results through a single instance of the algorithm. Internally, Q is organized as a priority queue according to an estimate of the upper bound of f. At each step, GSearch retrieves the most promising partial result, and processes it. The highest ranking results are thus generated first.

## 3.2 Operator-Based Processing

The concepts of operator-based systems are oblivious to edge direction, and extend to our semantics more readily than their graph-based counterparts. However, as discussed in Section 2.2, Discover incurs duplicate CN, duplicate MTJNT, and invalid results. In the following, we remedy this behavior and simultaneously adapt operator-based query processing to our semantics. In particular, we introduce three modifications: (i) UniqueJoin, (ii) a redefinition of  $S\{\}$ , and (iii) the CNGen algorithm for generating CN without duplicates. Compared to the former two, the latter is rather complicated, but inherits many characteristics from GSearch, simplifying the presentation.

Recall from Section 2.2 that a join operator consumes composite tuples from its left, and single records from its right input. Furthermore, the only valid output of each operator tree (i.e., MTJNT) is *acyclic*. This gives rise to the following invariant: All intermediate results, generated by join operators at lower levels, must also be free of cycles. To enforce this property, we replace all join operators with their *UniqueJoin* counterpart. *UniqueJoin* verifies that the compound tuple from its left input does not already contain the tuple from the right, in which case the tuples would form a cycle. Assume, for instance, a traditional operator  $j_{trad}$  that joins a compound tuple  $t_{left}$  and a basic tuple  $t_{right}$ , iff  $j_{trad}$ .condition( $t_{left}$ ,  $t_{right}$ ) = true. The corresponding *UniqueJoin* evaluates that: (i)  $j_{trad}$ .condition( $t_{left}$ ,  $t_{right}$ ) = true, and (ii)  $t_{right} \notin t_{left}$ . For the remainder of the article, we assume all join operators to use *UniqueJoin*.

Next, we redefine  $S\{\}$  to denote the set of tuples in relation S that do not contain any keyword; that is,  $S\{\} = \{s \in S | \forall k \in q, s \text{ does } not \text{ contain } k\}$ . The definition of candidate networks remains the same, but leaf operators are adapted to the redefined selections. For instance, given  $q = \{k_1, k_2, k_3\}$ ,  $S\{\}$  is translated into  $\sigma \neg_{k1} \wedge \neg_{k2} \wedge \neg_{k3} S$ . For the following discussion, we exclusively use the new definition of  $S\{\}$ . The combination of  $S\{\}$  allows the following statements.

Lemma 3.3. An operator tree op-tree translated from a CN only generates correct output.

PROOF. *Op-tree* outputs tuples that can be joined and hence form a subgraph of G. The *UniqueJoin* operator ensures that this subgraph is *acyclic*, namely a JNT. The selection operators ensure that: (i) the JNT contains all keywords, that is, is total, and (ii) has unique keywords in all external nodes, that is, is minimal. Tuples contain *only specified* keywords, and no others, and thus cannot impair minimality. Hence, all output constitutes MTJNT.  $\Box$ 

Lemma 3.4. If each CN is translated into one operator tree, then every MTJNT is retrieved exactly once.

PROOF. Assume an MTJNT r generated by two different operator trees  $op\text{-}tree_1$  and  $op\text{-}tree_2$ . Let  $cn_1$  and  $cn_2$  be their corresponding candidate networks. For every node  $s \in S$  of r containing a set of keywords K, each candidate network contains a node  $S\{K\}$ . Furthermore, both CN share the same structure as r. With identical nodes arranged in an identical structure, we deduce that  $cn_1 = cn_2$ . Since every CN was translated into a single operator tree, we thus find  $op\text{-}tree_1 = op\text{-}tree_2$ , violating our initial assumption. Hence, we conclude that every MTJNT is retrieved precisely once.  $\square$ 

The preceding modifications avoid erroneous and duplicate results. OB can generate output according to a monotone ranking function f, by applying the method of Hristidis et al. [2003]. In particular, OB stores all CN in a heap, with the most promising one ranking first. Then, it keeps executing the operator tree of the first CN, until it becomes less promising than the second-topmost, at which time it switches to the latter. Next we demonstrate how to compute the set of CN. Our CNGen algorithm constructs candidate networks by traversing the expanded schema, a process resembling GSearch. Both algorithms face similar challenges and lend themselves to comparable solutions. Akin to MTJNT, we model CN as unique trees, generated according to their unique preorder traversal. CN share the same structure as MTJNT, and must be total and minimal. In contrast to MTJNT, they can contain multiple identical nodes; for example, the top CN in Figure 4 contains two instances of T{}. In the following, we focus on this distinction and its implications for duplicate avoidance.

Similar to *GSearch*, *CNGen* requires a total ordering on nodes of the expanded schema. For our presentation, we assume that each relation (e.g., S) has an ID (S.rid). Nodes in the expanded schema have a keyword bitmap (kbit) according to the contained keywords, such as,  $S\{k_2\}.kbit = 010_b = 4$  and  $S\{k_1,k_3\}.kbit = 101_b = 5$ . The node order nid is a lexicographic combination of rid and kbit, for instance,  $S\{k_1,k_3\}.nid < T\{k_2\}.nid$  and  $S\{k_2\}.nid < S\{k_1,k_3\}.nid$ . Having defined this order, CN can be represented as unique trees. The node containing keyword  $k_1$  serves as root  $n_{root}$ . If several such nodes exist, the smallest nid breaks the tie. Child nodes are ordered left-to-right by nid. We observe two special cases, caused by the inclusion of multiple copies of the same node. In the first, a CN contains multiple instances of the root node. Here, the lexicographically smaller preorder traversal determines the CN's unique tree representation. Figures 9(a) and 9(b) depict the tree representation of a CN containing two instances of  $T\{k_1\}$ . Only the tree in Figure 9(a) represents the

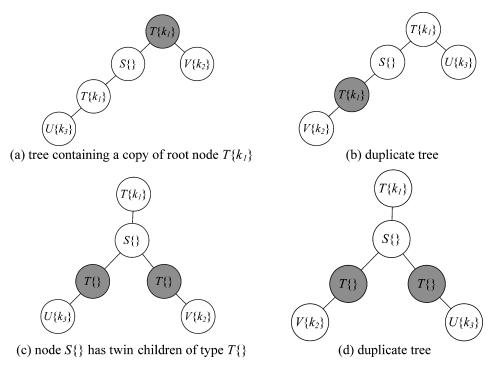


Fig. 9. Candidate networks with multiple identical nodes.

CN correctly because its preorder traversal  $(T\{k_1\}, S\{\}, T\{k_1\}, U\{k_3\}, V\{k_2\})$  is lexicographically smaller than that of Figure 9(b)  $(T\{k_1\}, S\{\}, T\{k_1\}, V\{k_2\}, U\{k_3\})$ . In the second special case, a node has twin children. To establish a unique tree representation, we demand that the subtree rooted at the left twin is lexicographically smaller than the one rooted at its right sibling. Consider  $S\{\}$  in Figure 9(c) having two identical children of type  $T\{\}$ . The traversal of the subtree rooted at the left twin  $(T\{\}, U\{k_3\})$  is smaller than that rooted at the right sibling  $(T\{\}, V\{k_2\})$ . This tree hence represents the CN correctly, whereas the tree of Figure 9(d) is incorrect.

Figure 10 provides pseudocode for CNGen, generating all CN whose tree representation is rooted at a node  $n_{root}$  of the expanded schema. Similar to GSearch, it starts with a tree  $t_{first}$  containing only  $n_{root}$ . At each step, the algorithm dequeues one tree  $t_{old}$  from Q, and spawns new trees by adding a node to the rightmost root-to-leaf path (lines 6 to 9). In contrast to GSearch, it allows multiple instances of the same nodes to be added (line 7). Again, we distinguish three cases: (i) If  $t_{new}$  constitutes a CN, it is included in the result set (lines 10 and 11); (ii) if the tree is not a CN, but still has the potential of becoming one, it is inserted in Q (line 24); and (iii) in any other case, the tree is discarded (line 26). In order to have the potential for forming a CN in the future, the current tree must have less than  $T_{max}$  nodes, and every terminal node (except those on the rightmost path) must contain a unique keyword (lines 14 and 15).

```
InitCNGen (Expanded Schema E)
     Initialize result set RS to empty
     For each node n_{root} containing k_1
          Call CNGen(E, n_{root}) and add all its returned results to RS
          Remove n_{root} from E
     Return RS
5.
CNGen (Expanded Schema E, Node n<sub>root</sub>)
                                                                       // Generates all CN rooted at a given node n_{root}
     Initialize result set RS to empty
     Initialize queue Q to empty
                                        // stores intermediate trees
     Insert a tree into Q consisting of a single node n_{root} (the root)
3.
     While (O is not empty)
          De-queue the first tree t_{old} in Q
6.
          For each node nl on the rightmost root-to-leaf path in t_{old}
              For each neighbor nn of nl in G // Unlike GSearch, there is no check if t_{old} already contains n_{now}
7.
8.
                  If (n_{new}.nid \ge nc.nid, for every child nc of nl) // the nid must be no smaller than the siblings'
                      Create a new tree t_{new} by adding nn as the rightmost child of nl
                      If (t_{new} is a CN)
10.
                          Insert t_{new} into RS;
11.
12.
                          Boolean has potential = false
13.
                          If (t_{new} \text{ contains less than } T_{max} \text{ nodes})
14
15.
                          AND (each terminal node, except those the rightmost path, contains a unique keyword in q)
16.
17.
                          If (t_{new} contains a second node n_{copy} that is identical to n_{root})
                                                                                                      // anywhere in the tree
18.
                              Create a copy t_{copy} of t_{new} and rotate it, so that n_{copy} serves as root
                              If (t_{copy}.lexiorder < t_{new}.lexiorder)
19.
20.
                                  has_potential = false
21.
                          If (\exists \text{ node } n_{right} \text{ on the rightmost root-leaf path, with an identical sibling } n_{left})
                              If (tree-rooted-at-n_{left}.lexiorder \geq tree-rooted-at-n_{right}.lexiorder)
22.
23.
                                  has_potential = false
                                                                     // t_{new} still has the potential of forming an CN
24
                          If (has potential), append t_{row} to O:
25.
                          Else, discard t_{new}
26. Return RS
```

Fig. 10. The iterative CNGen algorithm.

Similar to GB, a looser definition of minimality akin to Hristidis et al. [2003] can be implemented by relaxing the conditions in line 15. Note that these checks resemble lines 13 and 14 of GSearch. However, CNGen imposes two additional conditions in order to avoid the aforesaid duplicates. First, line 17 evaluates whether  $t_{new}$  contains a second instance  $(n_{copy})$  of its root node  $n_{root}$  (not necessarily  $n_{new}$ , but any older node in  $t_{new}$ ). In this case, CNGen compares the preorder traversal of two trees: (i)  $t_{new}$  rooted at  $n_{root}$  and (ii) a version  $t_{copy}$  of  $t_{new}$  that has been rearranged so that  $n_{copy}$  serves as root. If the former is lexicographically larger,  $t_{new}$  constitutes a duplicate of  $t_{copy}$  and can be discarded. The latter tree is generated from a different seedling in Q and does not have to be explored at this point. Second, line 21 checks all nodes along the rightmost root-to-leaf path in  $t_{new}$ . If any such node has an identical sibling, CNGen lexicographically compares the preorder traversal of the trees rooted at both nodes. If the left is larger,  $t_{new}$  can be discarded; the correct representation of the CN will be generated from another tree in Q.

The outer function InitCNGen calls CNGen for all nodes containing  $k_1$ , for example,  $S\{k_1\}$ ,  $S\{k_1,k_2\}$ ,  $T\{k_1\}$ . These nodes then serve as roots of CN-trees. Since every CN must include at least one node containing  $k_1$ , the set of expansions initiated by InitCNGen eventually produces all CN for the given query

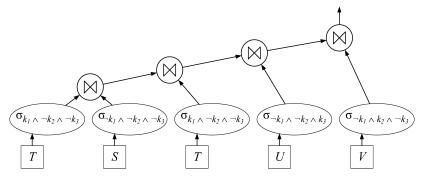


Fig. 11. Operator tree for the CN of Figure 9(a).

and schema. To ensure that individual calls of CNGen do not yield duplicates, CN are generated according to their unique tree representation. Furthermore, separate calls to CNGen must not generate identical CN. This could only happen for CN with multiple nodes containing  $k_1$ . To avoid such behavior, InitCNGen removes  $n_{root}$  from the expanded schema after  $CNGen(E, n_{root})$  terminates (line 4). Note that this does not cause any loss of results, since all CN containing  $n_{root}$  have been generated by  $CNGen(E, n_{root})$ .

Lemma 3.5. InitCNGen computes the set of CN (i) correctly, (ii) completely and (iii) without duplicates.

The proof follows from the previous discussion.

Finally, we address the translation of CN into operator trees. Similar to Discover, we map CN to left-deep trees, where leaf nodes are source operators that perform selections, and interior operators are UniqueJoins. Sources are ordered left-to-right in the order of their addition during CNGen: the leftmost source corresponds to the node  $(n_{root})$  of the expanded schema from which the CN was discovered; the rightmost node was added last. Figure 11 shows the operator tree for the CN in Figure 9(a). Join conditions correspond to parent-child relationships in the CN tree.

In our description of CNGen, we chose the lexicographic order of rid and kbit for simplicity. Since all operator trees are left-deep, it would be more efficient to arrange sources with the highest selectivity to the left. We should thus add such nodes as early as possible during CNGen. For instance, instead of  $k_1$ , we could choose the rarest keyword to determine  $n_{root}$ . Furthermore, without any specific knowledge about table size and distribution of keywords, it is reasonable to assume that the selectivity increases with the number of keywords; that is, nodes with a large number of keywords should be visited first, and hence receive a small nid. In summary, the preceding algorithms adapt operator-based R-KWS to our semantics of general join conditions, eliminating duplicate and invalid results.

## 4. RELATIONAL KEYWORD SEARCH ON STREAMS

This section focuses on continuous R-KWS queries on relational streams. Section 4.1 presents formal semantics. Sections 4.2 and 4.3 propose GB and OB query processing methodologies, further optimized later in the article.

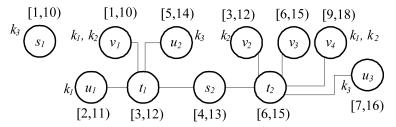


Fig. 12. Instantaneous data graph at  $\tau = 9$ .

#### 4.1 Semantics

We assume multiple relational data streams, whose tuples arrive in increasing t<sub>start</sub> and may be deleted explicitly (through a negative tuple) or implicitly (according to the sliding window model). A Streaming Relation (SR) is the union of several streams with a common structure and meaning. For example, all cash id, price, time> that can be wrapped into a single SR. We suppose all streams to be bundled into SR, and hence use both terms interchangeably. A streaming schema denotes which streams can be joined and on what attributes. Nodes in this graph represent SR, and are connected by an edge iff a common join attribute has been defined. In our examples, we assume four SR: S, T, U, and V, arranged in a schema identical to Figure 1(a) that is, the only joins permitted are between tuples in T and those in S, U, or V. The schema would usually be provided by the system operator, but may also be altered by individual users (e.g., by excluding SR that are not relevant to a query). If a new data stream is instantiated, it can be integrated by either: (i) merging it with an existing SR (if it adheres to the same format) or (ii) introducing a new SR.

We assume a continuous keyword query of the form  $q := \{k_1, \ldots, k_m\}$ , and define its semantics by identifying results on instantaneous views (snapshots) of the system. At every time instant  $\tau$ , the instantaneous data graph  $G(\tau)$  contains a node for each tuple s that is alive at  $\tau$ . Tuples are connected by an edge iff they can be joined. Figure 12 shows  $G(\tau=9)$  for the example schema, including lifespans of tuples. Note that in case of positive-negative tuples, the end of a lifespan is not known in advance. In our example, we assume a query with three keywords  $k_1, k_2, k_3$  whose appearance are denoted next to the tuples. Results are defined using the concept of MTJNT. Similar to keyword search over static tables, we impose a limit  $T_{max}$  of tuples per MTJNT. Let  $R(\tau)$  be the set of MTJNT in  $G(\tau)$  that do not exceed  $T_{max}$  nodes. The result R of a continuous R-KWS query is the union of  $R(\tau)$ , for all  $\tau$ . In Figure 12,  $(v_1, t_1, u_2), (v_1, t_1, s_2, t_2, u_3)$ , and  $(u_1, t_1, s_2, t_2, v_2, u_3)$  are results in  $R(\tau=9)$ . At time  $\tau=10, v_1$  expires and so do the former two MTJNT, while  $(u_1, t_1, s_2, t_2, v_2, u_3)$  continues as an element of  $R(\tau=10)$ . We require results to be produced in ascending  $t_{start}$  order

The following lemma allows for an efficient generation and compact representation of R.

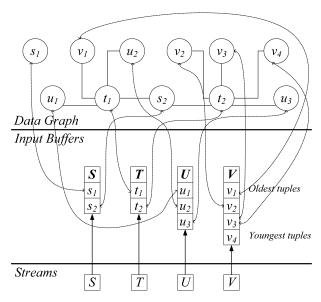


Fig. 13. A graph-based architecture for continuous R-KWS.

LEMMA 4.1. Let  $r \in R(\tau)$  be an MTJNT on  $G(\tau)$ . If every node n in r is alive at  $\tau + 1$ , then r is an MTJNT in  $G(\tau + 1)$ ; that is,  $r \in R(\tau + 1)$ .

PROOF. Since attributes do not change values over time, any tuple that contains keyword k at time  $\tau$  also contains k at time  $\tau+1$ . Similarly, edges do not change: If r is connected at  $\tau$ , it is also connected at  $\tau+1$ . Therefore, if r is total and minimal at  $\tau$ , it is also total and minimal at  $\tau+1$ .  $\square$ 

According to this lemma, an MTJNT r is not affected by insertions or deletions of external nodes. Consequently, every MTJNT r needs to be constructed and reported only once (at  $r.t_{start}$ ), rather than at every instant of its lifespan. The termination of a result r depends on the stream model. Using a sliding window of duration w, we can compute the lifespan of r directly upon its creation, as:  $r.t_{start} = max(n.t_{start})$  and  $r.t_{end} = min(n.t_{start}+w)$ , where n are the component tuples of r. For example,  $(v_1, t_1, u_2)$  in Figure 12 is first discovered at  $\tau = 5$ , in combination with its lifespan [5, 10). In the positive-negative model (where  $n.t_{end}$  is not known in advance), r is terminated when any of its constituent tuples are deleted. At this point, the user must receive a negative result tuple -r. Finally, the proposed semantics also capture conventional relational tables (e.g., containing "static" information about product characteristics) by modeling them as streams of everlasting tuples (i.e.,  $t_{start} = 0$  and  $t_{end} = \infty$ ).

## 4.2 Graph-Based Processing

This approach extends the paradigm of graph-based query processing to data streams. Continuous GB maintains the instantaneous data graph  $G(\tau)$  in memory and retrieves MTJNT by means of graph traversal. Figure 13 depicts the general architecture, using the data of Figure 12. The two main components are:

(i) a set of input buffers, one for each stream, and (ii) a data graph. The buffers store tuples that are currently alive; for example, T.buffer collects tuples  $t_1$  and  $t_2$  from data stream T. Arrivals and expirations of tuples necessitate updates of the data graph, and trigger a search for (expired) MTJNT. A new tuple is first inserted in the appropriate buffer. A corresponding node is created in the data graph, where it must be connected to neighboring nodes. The latter are determined by joining the new tuple against the input buffers of neighboring relations.

For example,  $t_2$  is probed against S.buffer, U.buffer, and V.buffer, thereby found to join with  $s_2$  and  $v_2$  (among others), and consequently connected to these nodes via edges. After each insertion, the resulting data graph is traversed for new MTJNT. Assume tuple  $s \in S$ , inserted at instant  $\tau$ . A call to  $GSearch(G(\tau), q, s)$ , retrieves the MTJNT in  $G(\tau)$  that contain s. These results are returned to the user. Calling GSearch for every new tuple answers the continuous R-KWS query. Tuple expirations are handled according to the stream model. Given a negative tuple -s at instant  $\tau$ , the user must receive a negative MTJNT -r for every MTJNT r that contains s. These MTJNT are identified by calling  $GSearch(G(\tau-1), q, s)$  on the current data graph. After this operation has completed, s is removed from the buffer and data graph, completing the update of  $G(\tau)$ . In the case of the sliding window model, there is no need to inform the user about expired MTJNT; instead it is sufficient to periodically scan the input buffers for expired tuples, and update the data graph accordingly.

Lemma 4.2. GB produces results (i) correctly, (ii) completely, (iii) without duplicates, and (iv) in the proper temporal order.

PROOF. According to Lemma 3.1, results generated by GSearch are correct, complete, and free of duplicates. The correctness GB follows directly from this property. The algorithm calls GSearch for every new tuple and hence finds all MTJNT. It avoids duplicates because every MTJNT is found only once, from its youngest constituting tuple. Finally, results are generated directly upon the arrival of their youngest tuple, and hence in correct temporal order.  $\Box$ 

The GB architecture easily adapts to changes in the schema. When a new data stream sends tuples for the first time, the system creates the corresponding buffer, and commences inserting tuples in the data graph. When a data stream is removed from the schema, the corresponding buffer is discarded. Relational tables are integrated as (static) input buffers, whose tuples straightforwardly participate in the data graph. If possible, these buffers should be stored in memory to avoid disk access for every newly arriving tuple of neighboring streams. Section 5 optimizes GB further through keyword labeling.

## 4.3 Operator-Based Processing

This approach uses relational stream operators to process continuous R-KWS queries. Figure 14 illustrates the system architecture. A query triggers the generation of candidate networks, subsequently translated into operator trees. These trees resemble their counterparts in conventional databases (e.g., Figure 11), receive streaming data from the bottom, and produce results

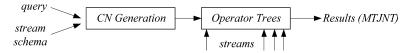


Fig. 14. Operator-based query processing for continuous R-KWS.

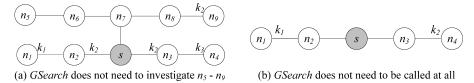


Fig. 15. Optimizing GSearch.

(MTJNT) at the top. Join operators follow the description of Section 2.4, and additionally implement the *UniqueJoin* pattern (Section 3.2), ensuring that no basic tuple contributes more than once to a single result.

Lemma 4.3. OB produces results that are (i) correct, (ii) complete, (iii) without duplicates, and (iv) in the proper temporal order.

Proof. Correctness, completeness, and the absence of duplicates follow Lemma 3.4. MTJNT are produced directly upon arrival of their youngest tuple and hence in correct temporal order.  $\Box$ 

Operator-based systems do not adapt as easily to changes in the schema as their graph-based counterparts. When a new data stream connects to the system, an entire range of operator trees must be generated before any data can be processed. Section 6.6 addresses the efficient creation of operator trees during this time-critical operation. In comparison, removing a data stream is easy. All operator trees that consume its tuples are guaranteed not to generate output, and can be removed whenever convenient. Again, we treat relational tables as streams of everlasting tuples which are scanned once and "stream" into to operator trees at the beginning of query processing. Section 6 introduces several effective optimizations for OB.

## 5. OPTIMIZATIONS FOR CONTINUOUS GB

Graph-based R-KWS over data streams can be greatly accelerated by restricting graph traversals. Assume  $q = \{k_1, k_2, k_3\}$  and the graph of Figure 15(a) where the current state (tree  $t_{new}$ ) of GSearch(G, q, s) consists of nodes s and  $n_2$ . The algorithm visits nodes  $n_5$  through  $n_9$ , among others, although none of these nodes contains (or leads to) the missing keywords  $k_1$  or  $k_3$ . If GSearch had been informed accordingly, the algorithm could have avoided traversing this branch. Similarly, in Figure 15(b), none of the tuples contains  $k_3$ . If this was known to the algorithm, graph traversal from s could be omitted altogether. Since keywords appear highly infrequently, both cases are common in practice. In this section we introduce keyword labeling, a simple and effective method to summarize reachable keywords for a given node. It improves performance

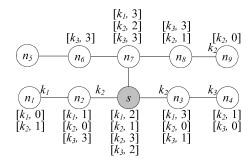


Fig. 16. A min-complete labeled graph for  $T_{max} = 4$ .

by avoiding unnecessary calls to *GSearch* and constraining graph traversals. For ease of presentation, we avoid stream-specific notation (e.g., timestamps) whenever possible.

A keyword label (KL) of format  $[k_i, h]$ , stored at node n, indicates a path of hedges in the data graph, connecting n to an occurrence of keyword  $k_i$ . We use  $n:[k_i, h]$  to indicate that the label  $[k_i, h]$  is located at node n. There is no need to consider paths exceeding  $T_{max} - 1$  edges, since MTJNT are limited to  $T_{max}$ nodes. Figure 16 depicts labels for the data of Figure 15(a), assuming  $T_{max} =$ 4. Node s stores four KL:  $[k_1, 2]$ ,  $[k_2, 1]$ ,  $[k_2, 3]$ , and  $[k_3, 2]$ . For example,  $s:[k_1, 2]$ 2] corresponds to the path  $(s-n_2-n_1)$ , connecting s to an occurrence of  $k_1$ , via two edges. We require keyword labels to be *min-complete*, that is, each node must store a KL for every reachable keyword, indicating the *shortest* path. The labeling in Figure 16 is min-complete. Note that node s has two labels for  $k_2$ , namely  $[k_2, 1]$  and  $[k_2, 3]$ , corresponding to paths of different lengths. The KL for the longer path (i.e.,  $[k_2, 3]$ ) is not required. On the other hand, removing the KL for the shortest path ( $[k_2, 1]$ ) violates min-completeness. The benefits of a min-complete labeling during query processing are twofold. First, GSearch(G, G)q, s) only needs to be called if s node can reach all query terms, that is, if the node stores a KL for every  $k \in q$ . In any other case, s is guaranteed not to participate in an MTJNT. Second, whenever a call to GSearch cannot be avoided, the labeling allows pruning the graph traversals. Recall that the original GSearch (Figure 8) uses two conditions (lines 13 and 14) to verify whether a new tree  $t_{new}$  has the *potential* of becoming an MTJNT. Keyword labels enable stricter requirements that eliminate more trees. Since GSearch expands  $t_{new}$  from nodes in the rightmost root-to-leaf path, these must be able to reach the missing keywords, without exceeding a total of  $T_{max}$  nodes. Otherwise,  $t_{new}$  cannot become total and is discarded.

Assume the query  $q = \{k_1, \ldots, k_5\}$ , and  $T_{max} = 9$ . The tree in Figure 17(a) lacks keyword  $k_2$ , and new nodes can only be added to  $n_1$ ,  $n_2$ , or  $n_3$  (for simplicity, we omit KL that are stored on the remaining nodes, or regard other keywords). Node  $n_1$  can reach  $k_2$  in four hops, the shortest path to any occurrence of  $k_2$ . However, the tree already contains six nodes, and would thus exceed  $T_{max}$ . It can hence be discarded directly without further expansions. If more than one keyword is missing, we have to take into account that paths may overlap.

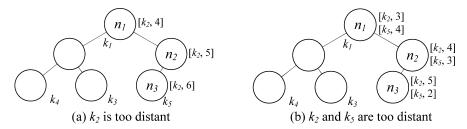


Fig. 17. Intermediate trees abandoned by *KL-aware GSearch*.  $(T_{max} = 9)$ .

Consider the tree in Figure 17(b), lacking keywords  $k_2$  and  $k_5$ . The KL indicate paths of three nodes to  $k_2$  (via  $n_1$ ), and two nodes to  $k_5$  (via  $n_3$ ), and the tree could become total by adding five nodes. However, this is not the minimal extension, since paths potentially overlap. The KL at  $n_2$  indicate paths to  $k_2$  and  $k_5$ , of four and three edges, respectively. If  $k_5$  is situated along the path to  $k_2$ , four nodes suffice to reach both keywords. In our example, this marks the smallest number of nodes that could possibly be added. Since the tree already contains six nodes, it is bound to exceed  $T_{max} = 9$  and can hence be discarded.

In general, a KL-aware GSearch only inserts  $t_{new}$  into Q iff there exists a set NL of labels, situated at nodes on the rightmost root-to-leaf path of  $t_{new}$ , that meets the following criteria.

- (i) The KL in NL can reach all missing keywords; that is, NL. $keywords \cup t_{new}$ .keywords = q.
- (ii)  $|t_{new}| + \Sigma_n(max(h|n:[k,h] \in NL) \leq T_{max}$ .

The second criterion takes the potential overlap between paths into account, by considering the longest path leaving a node n rather than their sum. Assuming  $T_{max} = 10$  in Figure 17(b), the set  $NL = \{n_1: [k_2, 3], n_1: [k_5:4]\}$  satisfies both conditions. Specifically, (i) the KL in NL contain all missing keywords  $(k_2 \& k_5)$ , and (ii) it is possible for the only concerned node  $n_1$  to reach both keywords in four edges, while  $|t_{new}| = 6$ ; hence,  $|t_{new}| + 4 \le T_{max}$ . The following lemma states the correctness of the resulting KL-aware GSearch.

### Lemma 5.1. KL-aware GSearch yields the same results as GSearch.

PROOF. KL-aware GSearch does not produce results that are not generated by the original algorithm because the keyword labels only filter unpromising trees. On the other hand, KL-aware GSearch does not miss any correct result because min-completeness ensures that discarded trees cannot reach the remaining keywords within  $T_{max}$  nodes.  $\square$ 

In order to remain min-complete, keyword labels must be updated whenever tuples arrive or expire. In the following, we present two implementations. The first is applicable to both stream models, sliding windows as well as explicit deletions; the second is more efficient, but restricted to sliding windows.

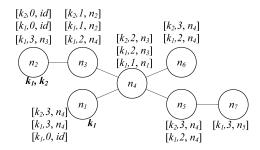


Fig. 18. A min-complete labeling with predecessor-KL.

#### 5.1 Predecessor-KL

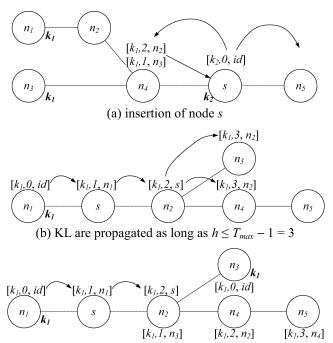
A predecessor-KL is a triplet of the form [k, h, p]. Stored at node n, it indicates a path of length h, connecting n to an occurrence of keyword k. In this path, p is n's predecessor. We write n:[k, 0, id] if n already contains term k. Figure 18 depicts a graph with occurrences of keywords  $k_1$  and  $k_2$ . Node  $n_4$  reaches  $k_2$  via the path  $n_2$ - $n_3$ - $n_4$ , and stores  $[k_2, 2, n_3]$ , indicating that its predecessor in this path is  $n_3$ . The same path connects  $k_1$  to  $n_4$ , and the node also stores  $[k_1, 2,$  $n_3$ ]. The fact that  $n_4$  can also reach  $k_1$  in its neighbor  $n_1$  is indicated by  $n_4$ :[ $k_1$ ,  $1, n_1$ ]. To guarantee min-completeness, we maintain the following invariant: Every node n must contain a predecessor-KL [h, h, p], for the shortest path leading from n through p to the occurrence of k. For example, in Figure 18,  $n_4$ must keep both KL  $[k_1, 2, n_3]$  and  $[k_1, 1, n_1]$ , since they represent the shortest path via predecessors  $n_3$  and  $n_1$  respectively. Node  $n_6$  reaches  $k_1$  contained in  $n_2$  and  $n_1$  through three and two steps, respectively. However, since both paths (to  $n_2$  and  $n_1$ ) share the same predecessor  $n_4$ , it suffices to keep only  $[k_1, 2, n_4]$ . Assuming  $T_{max} = 4$ , each path contains at most  $T_{max} - 1 = 3$  edges and there is no need to store  $n_7$ :[ $k_2$ , 4,  $n_5$ ]. This invariant yields a min-complete labeling, and allows a correct *KL-aware GSearch*.

An arriving tuple s can itself contain a keyword, or create new paths between keywords and nodes. Both cases require KL insertions and updates. Changes are not limited to s and its immediate neighbors, but are propagated in a radius of  $T_{max}-1$ . Figure 19 describes the underlying algorithms InitSendKL and SendKL. The former initializes labels at node s; the latter propagates changes. InitSendKL works in two stages. First, it constructs KL at s, for all keywords contained in the node (lines 2 and 3), and transmits this information to all neighboring nodes (lines 4 and 5), using SendKL. Second, it propagates older KL from neighboring nodes to s (lines 6 and 7), again by using SendKL. Consider inserting s in the graph of Figure 20(a). First, s receives label  $[k_2, 0, id]$ , since it contains  $k_2$ . This label is then passed to the neighboring nodes  $n_4$  and  $n_5$ . Second, older labels  $[k_1, 1, n_3]$  and  $[k_1, 2, n_2]$  from  $n_4$  are sent to s, and propagated further. As discussed shortly, Send-KL is recursive, and pushes labels deep into the graph. For example, the new KL  $[k_2, 0, id]$  is passed from s to  $n_4$ , and from there to  $n_3$ ,  $n_2$  and finally  $n_1$ .

*SendKL* propagates KL in a store-and-forward fashion. A node receiving a KL: (i) stores this locally (line 4), and (ii) forwards it to all neighbors, except for

```
InitSendKL (Node s) // Adjusts Predecessor-KL to the arrival of a new node s
// s: the newly inserted node
     Insert s in the data graph
2.
     For every keyword k, occurring in s
                                                        // Create KL at new node
         Add [k, 0, s] to s.KLlist
3.
         For every neighbor n of s in the data graph // Pass KL to neighbors – and further
             SendKL(s, n, [k, 1, s])
     For every neighbor n of s in the data graph
                                                        // Pass KL from neighbors to new node – and further
6.
             For every KL [k, h, pr] \in n.KLlist where pr \neq s
                 SendKL(n, s, [k, h+1, s])
SendKL (Node pn, Node rn, KL newKL)
                                                        // Propagates a KL to a neighboring node
// pn: the node passing newKL
// rn: the neighboring node receiving newKL
// Assume that newKL = [k, h, pn]
    If h > T_{max} - 1, Return
If (\exists [k, h', pn] \in rn.KLlist \mid h' \le h)
2.
                                                        // an existing KL indicates a shorter route from rn through pn
         Return
//
     Lines 4-6: store the new KL, and propagate it to neighboring nodes
4.
     Add newKL to rn.KLlist
     For each neighbor nn of rn \mid nn \neq pn
                                                        // do not send the KL back to the sender
         SendKL(rn, nn, [k, h+1, rn])
6.
```

Fig. 19. SendKL and InitSendKL.



(c) KL are propagated as long as they announce shorter routes

Fig. 20. Examples of InitSendKL and SendKL ( $T_{max} = 4$ ).

```
InitRevokeKL (Node s) // Adjusts KL-labeling to the expiration of node s
     Remove s from the data graph
     For all former neighbors n of s
3.
           For all keywords k
              RevokeKL(n, k, s, \infty)
4.
RevokeKL (Node pn, Node rn, Keyword k, int new-dist)
                                                                                 // informs of increased path length
// k: keyword of concern
// pn: node whose distance to k has increased
// new-dist: the new distance between pn and k
// rn: a neighbor of nn, receiving the message
          Remove [k, old-dist, pn] from rn.KLlist
          KL kl_1 = [k, h_1, n_1] \in rn.KLlist, such that h_1 is minimal.
2.
                                                                                 // kl_1 indicates the shortest path to k
          KL \ kl_2 = [k, h_2, n_2] \in rn.KLlist \setminus kl_1, such that h_2 is minimal. // kl_2 indicates the second shortest path to k
3.
          If (new-dist \le T_{max} - 1)
              Add [k, new-dist, pn] to rn.KLlist
// Case 1: if the best path to k used to lead through pn; all neighbors are informed
          If (old\text{-}dist < h_1)
7.
              For all neighbors nn of rn \mid nn \neq pn
                  bdist = \min(h_{best} \mid [k, h_{best}, n_{best}] \in rn.KLlist \mid n_{best} \neq nn) \text{ or } \infty \text{ if no such KL exists}
8.
                  RevokeKL(nn, rn, k, bdist + 1)
10.
          Else
// Case 2: If the second best path to k used to lead through pn, only node n_l supplying the shortest route is informed
12.
                  bdist = \min(h_{best} \mid [k, h_{best}, n_{nest}] \in rn.KLlist \mid n_{best} \neq n_I) or \infty if no such KL exists
                  RevokeKL(n_1, rn, k, bdist + 1)
13.
```

Fig. 21. RevokeKL and InitRevokeKL.

the transmitter (lines 5 and 6). In Figure 20(b), the new node s receives a KL from its left neighbor  $n_1$  and forwards it to  $n_2$ . This node stores the KL (as  $[k_1, 2, s]$ ) and propagates it to  $n_3$  and  $n_4$ . There are two terminating conditions. First, a label is only propagated as long as its hop-count h does not exceed  $T_{max}-1$  (line 1). For instance, assuming  $T_{max}=4$ , the KL in Figure 20(b) is sent to  $n_4$ , but not to  $n_5$ . Second, KL are propagated only as long as they announce shorter paths (lines 2 and 3). Consider Figure 20(c). In contrast to Figure 20(b), a second tuple  $(n_3)$  already contains  $k_1$ . The KL from  $n_1$  is forwarded to s and further to  $n_2$ . Because  $n_2$  can already reach  $k_1$  via a shorter path  $(n_2-n_3)$ , there is no need to send any new label to  $n_3$  and  $n_4$ .

Similarly, expiring tuples can remove keyword occurrences and destroy paths that connect nodes with keywords. Both cases lead to KL updates and deletions. Figure 21 describes the functionality of InitRevokeKL and RevokeKL. The former algorithm removes a deleted tuple s, and calls the latter to propagate this news (within a radius of  $T_{max}$ ). Using RevokeKL, a node pn instructs its neighbor rn that its shortest path to keyword k ceased to exist, and that the new (longer) path from rn to k via pn takes new-dist hops. If pn cannot reach k anymore at all, it sets new-dist  $= \infty$ . Upon receiving such a message, node rn replaces its KL [k, old-dist, pn] with [k, new-dist, pn] (lines 1, 4, and 5) and informs its neighbors (lines 6 through 13). Consider the disappearance of s in Figure 22(a). The node notifies its neighbor  $n_2$ , by calling  $RevokeKL(s, n_2, k_1, \infty)$ . This node removes the old KL  $n_2$ :[ $k_1$ , 2, s], and in turn informs its neighbors (Figure 22(b)), for instance, by calling  $RevokeKL(n_2, n_3, k_1, 4)$ . The receiver (e.g.,  $n_3$ ) then replaces its label [ $k_1$ , 3,  $k_2$ ] with [ $k_1$ , 4,  $k_2$ ], and continues propagating the message.

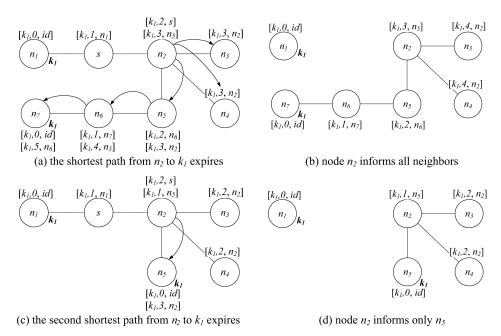
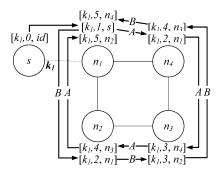
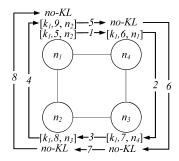


Fig. 22. RemoveKL and InitRemoveKL.

When a node rn receives a call of RevokeKL, there are three cases. In the first, the expiring path (via pn) has been the shortest path between rn and keyword k (lines 6 through 9). When this happens, all neighbors of rn must be notified. Consider node  $n_2$  in Figure 22(a), whose shortest route to  $k_1$  leads through node s. If  $n_2$  receives  $RevokeKL(s, n_2, k_1, \infty)$ , it informs all neighbors  $n_3$ ,  $n_4$ , and  $n_5$ . These nodes update their KL accordingly (Figure 22(b)) and subsequently notify their own neighbors. In the second case, the expiring path (via sn) has been the second shortest path between rn and k (lines 7 through 11). Most neighbors of rndo not need to update their KL, because these indicate already the shortest path via rn. The only node that requires an update is the neighbor of rn that supplies the shortest path (lines 11 through 13). Assume the expiration of s in Figure 22(c). When  $n_2$  receives  $RevokeKL(s, k_1, n_2, \infty)$ , it updates its KL but does not inform  $n_3$  and  $n_4$  because its shortest path to  $k_1$  (via  $n_5$ ) did not change. The only label that has to be updated (by calling  $RevokeKL(n_2, k_1, n_5, \infty)$ ) resides at the node  $n_5$ , which provides the shortest path between  $n_2$  and  $k_1$ . Figure 22(d) depicts the resulting keyword labels. In the last case, where the expiring path (via pn) is neither the shortest nor the second shortest path between rn and k, RevokeKL terminates directly after updating the KL at rn, and does not inform neighboring nodes.

In applications involving graphs, updates along cycles commonly pose a challenge. Even though RevokeKL does not address cycles explicitly, it handles them effectively. In Figure 23(a), tuple s contains keyword  $k_1$ , and nodes  $n_1$  through  $n_4$  form a circle. Accordingly, KL have been propagated in two directions: clockwise (A) and counterclockwise (B). Now, assume that s expires. This is the only occurrence of  $k_1$ , and all KL must be removed. Figure 23(b) investigates the





- (a) KL on a data graph containing a cycle
- (b) KL are removed when  $h = T_{max}$

Fig. 23. Revoking KL on cycles ( $T_{max} = 10$ ).

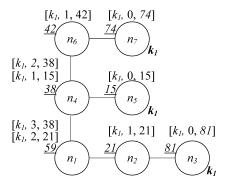


Fig. 24. KL with timestamps.

updates on the clockwise-oriented KL. Upon expiration, s calls RevokeKL on  $n_1$ . This node removes  $[k_1, 1, s]$ , and then informs its neighbors. Due to  $n_1$ : $[k_1, 5, n_2]$ , it assumes that it is able to reach  $k_1$  via  $n_2$  and hence updates  $n_4$  by calling  $RevokeKL(n_4, n_1, k_1, 6)$ . This node replaces  $[k_1, 2, n_1]$  with  $[k_1, 6, n_1]$  and informs  $n_5$ . Update operations thus complete a circle, return to node  $n_1$ , and set out on a second tour. So far, only the hop-count h has increased. Yet, this process stops when h reaches  $T_{max}-1$ . Assume  $T_{max}=10$ . The fourth call to RevokeKL causes  $n_1$ : $[k_1, 5, n_2]$  to be replaced with  $n_1$ : $[k_1, 9, n_2]$ , and triggers updates on  $n_4$ , by calling  $RevokeKL(n_4, n_1, k_1, 10)$ . This node removes  $n_4$ : $[k_1, 6, n_1]$ , does not create a new KL because  $h=T_{max}$ , and calls  $RevokeKL(n_3, k_1, n_4, 11)$ . The KL at  $n_3$  is removed accordingly and the update request is propagated further. After the circle has been completed, all KL have been removed correctly. In practice,  $T_{max}$  is small, and RevokeKL converges rapidly.

## 5.2 Time-KL

For sliding windows, we propose a more efficient labeling that does not require explicit removal. Specifically, a time-KL is a triplet  $[k, h, t_{end}]$  indicating a path of length h to an occurrence of keyword k, which exists until  $t_{end}$  (when its earliest tuple expires). Figure 24 depicts a data graph with time-KL. Node  $n_4$ 

```
InitSendTimeKL (Node s)
                                                                // Updates TimeKL for the arrival of a new node s
     Insert s in the data graph
     For every keyword k, occurring in s
                                                               // Create KL at new node
          Add [k, 0, s.t_{end}] to s.KLlist
          For every neighbor n of s in the data graph
                                                               // Pass KL to neighbors - and further
4.
              SendTimeKL(s, n [k, 1, s.t_{end}])
     For every neighbor n of s in the data graph
                                                               // Pass KL from neighbors to new node - and further
          For every KL [k, h, t_{end}] \in n.KLlist
7.
8.
              SendTimeKL(n, s, [k, h + 1, t_{end}])
                                                               // Propagates a TimeKL to a neighboring node
SendTimeKL (Node pn, Node rn, KL newKL)
// pn: the node passing the KL
// rn: the neighboring node receiving a KL
// newKL = [k, h, t_{end}] the keyword label
     If (h \ge T_{max} - 1), Return
     If (\exists [k, h_{old}, t_{end-old}] \in rn.KLlist \mid h_{old} \leq h \& t_{end-old} \geq t_{end})
                                                                         // if there exists a dominating entry, abort
3.
          Return
                                                                         // remove dominated KL
     For all [k, h_{old}, t_{end-old}] \in rn.KLlist
          If (h_{old} \ge h) AND (t_{end-old} \le t_{end})
              Remove [k, h_{old}, t_{end-old}] from rn.KLlist
     Add [k, h, min(newKL.t_{end}, rn.t_{end})] to rn.KLlist
                                                                         // Add the new KL
                                                                         // Propagate to neighbors
     For each neighbor nn of rn
          SendTimeKL(rn, nn, [k, h + 1, min(newKLt_{end}, rn.t_{end})]
```

Fig. 25. SendTimeKL and InitSendTimeKL.

is connected to  $k_1$  in  $n_7$  via two hops. Among the path's nodes,  $n_7$  expires at 74,  $n_6$  at 42, and  $n_4$  at 38. The latter marks the minimum, and  $n_4$  stores  $[k_1, 2, 38]$ . Similarly, there is a path of length one from  $n_4$  to  $k_1$  (in  $n_5$ ). This node expires at time 15, and  $n_4$  stores the KL  $[k_1, 1, 15]$ . A third path connects  $n_4$  to  $k_1$  (at  $n_3$ ). The path has a length of three, and the earliest node  $(n_2)$  expires at 21. Recall that the min-completeness property requires the labels to indicate the shortest path between a node and a keyword at any time. Both KL  $[k_1, 2, 38]$  and  $[k_1, 1, 15]$  must be stored, since each indicates the shortest path for some period of time. On the other hand, the third path does not need to be recorded because it is longer and expires sooner than the other two. We say that KL  $[k, h_1, t_{end-1}]$  dominates another  $[k, h_2, t_{end-2}]$  iff  $h_1 \leq h_2$  and  $t_{end-1} \geq t_{end-2}$ . The mincomplete property is met iff the graph contains all KL that are not dominated by others (i.e., those in the skyline). The labeling in Figure 24 satisfies this invariant.

A new tuple s invokes new time-KL, thereafter propagated in a store-and-forward fashion. Figure 25 shows the pseudocode for InitSentTimeKL and Send-TimeKL. The former algorithm: (i) creates KL for keywords in the new node s, thereafter broadcasted through the data graph (lines 2 through 5), and (ii) propagates KL from neighboring nodes to s and onward (lines 6 through 8). The latter algorithm transfers a label newKL from one node pn to a neighbor rn. There are two terminating conditions. First, if the hop-count h exceeds  $T_{max}$ , newKL is disregarded. Second, if newKL is dominated by an older label at rn, it is also ignored (lines 2 and 3). In any other case, newKL indicates the shortest path from rn to k, at least for some period, and is stored (line 8) and propagated further to neighboring nodes (lines 9 and 10). Older labels that are now dominated by newKL become obsolete, and are removed (lines 5 through 7).

Outdated time-KL are ignored by *GSearch* and removed whenever convenient, such as, when their node expires.

#### 6. OPTIMIZATIONS FOR CONTINUOUS OB

As illustrated in Sections 2.2 and 3.2, an R-KWS query on static tables commonly involves an immense number of operator trees. For continuous queries over data streams, this obstacle becomes even more pronounced. If a selection on a table (e.g.,  $T\{k_1\}$ ) returns no tuples, all operator trees using this input (e.g., Figure 11) can be discarded immediately. For data streams, this is not permissible. Even though the selection  $T\{k_1\}$  does not currently produce tuples, it may do so in the future, and all operator trees must thus be maintained. This section proposes optimizations that enable efficient OB R-KWS over data streams. In particular, Section 6.1 integrates individual operator trees into a single mesh, sharing common subexpressions. A first approach creates a Full-Mesh (FM) during preprocessing, allowing to dedicate runtime resources exclusively to tuple processing. Sections 6.2 and 6.3 describe two optimizations for FM, namely demand-driven operator execution and temporary operator disconnection. Section 6.4 presents a Partial-Mesh (PM) that does not require preprocessing, but grows and shrinks dynamically. Section 6.5 addresses the purging of dead tuples. Finally, Section 6.6 addresses changes in the schema.

#### 6.1 Operator Mesh

We integrate all operator trees into an operator mesh, reducing CPU cost (for evaluating joins) as well as memory overhead (for intermediate results). The mesh has  $|SR| \cdot 2^{|K-1|}$  clusters, where |SR| is the number of streaming relations and |K| the number of query keywords. Each cluster contains the operator trees for all CN discovered from a certain  $n_{root}$ . The trees in a cluster overlap on their left because they include at least the same  $\underline{n_{root}}$ , but usually they share larger parts. The entire operator mesh has  $|SR| \cdot 2^{|K|}$  leafs/sources, one for each node of the extended schema. The maximum depth of the mesh is  $T_{max}+1$ . The number of edges depends on the schema complexity. Output is produced at all levels, since operator trees vary in height. Different clusters are interconnected only through their source operators; joins from different clusters do not connect directly. In addition, we introduce a central output operator that collects results from all topmost operators (those producing MTJNT). Figure 26 shows the shared execution of four operator trees. Their corresponding CN all have been created by CNGen for  $n_{root} = S\{k_1\}$ . The join  $j_1(S\{k_1\} \bowtie T\{\})$  is shared by  $(S\{k_1\}, T\{\}, V\{k_2\}, T\{k_1\}, U\{k_3\}), (S\{k_1\}, T\{\}, U\{k_2, k_3\}), \text{ and } (S\{k_1\}, T\{\}, V\{k_2\}, K_3\})$  $V(k_2,k_3)$ ). Note that the figure depicts only a small subset of the particular cluster.

Mesh creation is performed in parallel to CN generation. Specifically, the first node in a cluster corresponds to the root node  $n_{root}$ , from which CNGen starts. Whenever the algorithm generates a new tree  $t_{new}$  from  $t_{old}$  (by adding a new child  $n_{new}$  to a parent  $n_{old}$ ), a join  $t_{new}.op$  is added to the mesh. The left child of  $t_{new}.op$  is  $t_{old}.op$  (the operator that was inserted when  $t_{old}$  was created), and the

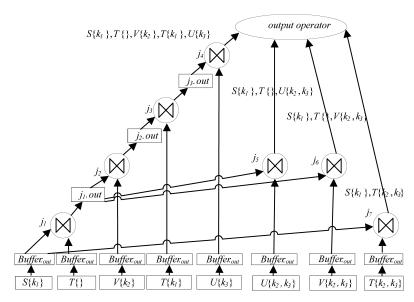


Fig. 26. Meshed trees for four CN in the same cluster.

right child is the source of  $n_{new}$ . For each tree t in CNGen, we maintain a pointer to the corresponding operator t.op, in order to decide where to place subsequent joins when t is expanded. The algorithm is initialized with  $t_{first}.op$  pointing to the source of  $n_{root}$ . For instance, the mesh of Figure 26 at first contains only  $S\{k_1\}$ . When CNGen visits  $T\{\}$ , we add  $j_1$  and connect it to  $S\{k_1\}$  on its left and  $T\{\}$  on its right. Subsequent insertions of  $V\{k_2\}$ ,  $T\{k_1\}$ ,  $U\{k_3\}$  in the CN cause the addition of  $j_2$ ,  $j_3$  and  $j_4$ . Similarly, when at a later point CNGen inserts  $U\{k_2, k_3\}$  to the tree containing  $S\{k_1\}$  and  $T\{\}$ ,  $j_5$  is added to the mesh and connected to  $j_1$  (representing  $S\{k_1\} \bowtie T\{\}$ ) and source  $U\{k_2, k_3\}$ .

We further compact the mesh by sharing buffers. In traditional DSMS, a join operator j has two individual input buffers, j.left-buffer and j.right-buffer. In our system, these buffers are replaced by the output buffers of the child operators, for example, in Figure 26,  $j_1.out$  replaces  $j_5.left$ -buffer and  $j_6.left$ -buffer. Because a single operator may have thousands of parents, this concept of state sharing greatly reduces memory consumption. Note that tuples in the buffers are naturally ordered by  $t_{start}$  (the instant at which they were produced). More complex indexing schemes are not required, since buffers in the mesh commonly contain few tuples, if any. R-KWS meshes are larger and more densely connected than any other operator graph for relational data. However, they also show beneficial characteristics. In particular: (i) they have a distinct structure, that is, clustered left-deep trees, and (ii) their join and selection operators are rather selective. In the following, we exploit these characteristics for further optimization.

#### 6.2 Demand-Driven Operator Execution

Our first solution generates a Full-Mesh (FM) of operators, prior to the actual query processing. This mesh is maintained in main memory throughout the

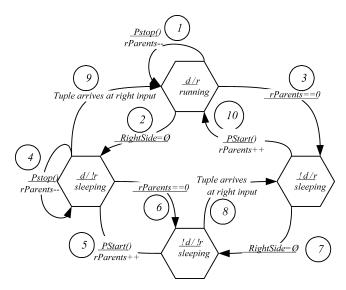


Fig. 27. States and transitions for join operators.

lifespan of the query. It allows demand-driven operator execution, an interoperator messaging system that eliminates ineffective join operations. We observe that join operators commonly execute without any prospect of contributing to an actual result, because joins at higher levels lack input from their right child. In Figure 26, assume tuples from  $S\{k_1\}$  and  $T\{\}$ , while  $V\{k_2\}$ ,  $U\{k_2, k_3\}$ , and  $V\{k_2, k_3\}$  are empty. None of the joins  $j_2$ ,  $j_5$ , or  $j_6$  requires the output of  $j_1$  because they do not receive right input. In the worst case,  $j_1$ 's results expire before the arrival of any tuples from  $V\{k_2\}$ ,  $U\{k_2, k_3\}$ , or  $V\{k_2, k_3\}$ . The join has wasted CPU and memory, without any contribution to the query. Even if  $V\{k_2\}$  had tuples available and  $j_2$  consumed input from  $j_1$ , the execution of both operators could still be wasted, for instance, if  $j_4$  happens to lack right input.

Under demand-driven operator execution, a join is considered to be either running or sleeping. Running operators process input; sleeping ones ignore it. A join operator is sent to sleep if: (i) it has no input from the right child (a source), or (ii) all its parents are sleeping. Sending operators to sleep does not affect the result's correctness or completeness because either: (i) the operator cannot produce output, or (ii) its output would not be consumed. Figure 27 shows the state diagram for a join operator. States are characterized by two binary flags: d indicating that at least one parent operator is running, and r specifying that the operator's right input is not empty. An operator only runs in the topmost state, (d/r). When it leaves this state (transition 2 or 3) it goes to sleep (or halts), to wake up (or restart) later (transitions 9 and 10). Operators exchange messages regarding their state, in order to ensure that all d and r flags are up-to-date. Particularly, a join operator communicates changes (running/sleeping) to its left child that adjusts its d flag accordingly. Likewise, sources inform their parents (i.e., joins for which they constitute the right child), whenever their buffer runs

empty, or when a new tuple arrives to a previously empty buffer, so that these joins maintain correct r flags.

Assume the operators in Figure 26, where all sources produce tuples and consequently all join operators are running. When  $U\{k_2, k_3\}$  dries up, it informs its parent  $j_5$ , which turns off its r flag, goes to sleep (transition 2), and informs its left child  $(j_1)$ , by calling  $j_1.Pstop$ . Upon receiving this notification,  $j_1$  decreases its counter of running parents (transition 1), but takes no further action because it still has other running parents  $(j_2$  and  $j_6)$ . When  $V\{k_2, k_3\}$  stops producing output,  $j_6$  halts, and  $j_1$  is left with a single running parent  $(j_2)$ . If now  $T\{k_1\}$  also dries up,  $j_3$  adjusts its r flag, goes to sleep, and informs  $j_2$ . This operator decreases its counter (rParents = 0), halts (transition 3), and calls  $j_1.Pstop$ . This join also finds all its parents sleeping, and likewise halts.

Before going to sleep, an operator sets a local timestamp stopTime = now. When it later wakes up, it processes all tuples from its left and right input that are: (i) alive and (ii) arrived after stopTime. To ensure the correct temporal order of results and to avoid duplicates, tuples are processed according to increasing order of  $t_{start}$ , and joined against those of the opposite input that have a smaller  $t_{start}$ . Before processing tuples, the newly awaking join has to ensure that its left input buffer is up-to-date. After all, the left child may also be sleeping, causing its output buffer to be incomplete. Thus, the operator calls leftChild.Pstart, asking its left input to wake up and update its output buffer.

Continuing the example of Figure 26, consider that the only sources with output are  $S\{k_1\}$ ,  $T\{\}$ ,  $V\{k_2\}$ ,  $U\{k_3\}$ , and  $T\{k_2, k_3\}$ , and the only running join operators are  $j_4$  and  $j_7$ . (The output operator is always running.) Join  $j_4$  does not generate results, due to lack of left input  $(j_3)$  is sleeping). When  $T\{k_1\}$  begins producing output, it causes  $j_3$  to adjust its r flag, wake up (transition 9), and call  $j_2$ . Pstart. This operator  $(j_2)$  restarts and informs  $j_1$ . Consequently, all joins except  $j_5$  and  $j_6$ , are running again. This concludes our discussion of demanddriven operator execution. Note that this method is not restricted to keyword search; it can equally benefit other data stream applications.

#### 6.3 Temporary Operator Disconnection

While the previous optimization targets individual join operations, a significant portion of computational expenses is inherent to the structural properties and complexity of the operator mesh. Consider the selection  $T\{\}$  in Figure 26 that supplies  $j_1$  with right input. For every tuple  $t \in T\{\}$ , the system: (i) checks if  $j_1$  has demand, and (ii) verifies whether  $j_1.left$ -buffer contains any tuples. If both conditions are met, it: (iii) compares t with all tuples in  $j_1.left$ -buffer to identify join partners. Compared to the first two steps, the latter only takes place rarely, since: (i) demand-driven operator execution is highly effective, and the majority of operators are dormant, and (ii) due to high join selectivity and keyword scarceness most operators on higher levels have empty left input buffers. The actual comparison of join attributes: (iii) is thus performed too infrequently to incur significant cost. In contrast, steps (i) and (ii) have a substantial impact. At first sight, both operations only involve a simple lookup. Yet, their tremendous cost is due to the immense frequency at which they are executed. First, for a

single tuple t, they are performed on several thousands of join operators that have  $T\{\}$  as their right input (the number of joins that have  $T\{\}$  as left input is negligible). Second, this large number of lookups has to be performed *for every arriving tuple*.

We minimize this effect through temporary operator disconnection. Whenever a join operator lacks demand or left input, it temporarily disconnects from its right child. In Figure 26, the operator  $j_1$  disconnects from  $T\{\}$ , whenever  $S\{k_1\}$  dries up, or when all parents  $(j_2, j_3, j_4)$  cease to request output. None of the tuples arriving later at  $T\{\}$  causes access to  $j_1$  or  $j_1.left$ -buffer, and the CPU cycles for steps (i) and (ii) are saved. If a disconnected join operator (e.g.,  $j_1$ ) encounters both demand and left input, it reconnects to its right child (e.g.,  $T\{\}$ ). The temporary disconnection does not compromise the completeness of results, since during this period new tuples from the right child are guaranteed not to find join partners. Yet, the simple measure significantly reduces CPU consumption.

As shown in the experimental evaluation, FM combined with demand-driven operator execution and the temporary operator disconnection is highly efficient. However, data processing has to be delayed until the mesh is complete. For certain applications, this is not acceptable. Furthermore, the size of the mesh can exceed the available main memory, especially if multiple queries are active in parallel. Our next approach avoids initialization and reduces memory consumption by adapting the mesh dynamically.

#### 6.4 Partial-Mesh

A *Partial-Mesh* (PM) is built at runtime and breaks the distinction between operator initialization and tuple processing. The method maintains relatively few *active* operators in memory, that is, only those with input. Specifically, it is each operator's responsibility to create its parents before it can produce output. Conversely, it destroys its parents (and other operators up the tree) if it cannot supply them with input. Especially in large meshes, operators are commonly idle; some never execute throughout the query lifespan. Their absence does not affect result's completeness, but dramatically reduces memory consumption. In the following we describe how to grow and shrink the operator mesh.

Initially, the partial mesh contains only the  $|SR| \cdot 2^{|K|}$  sources. Join operators are created later, as tuples travel upwards. For our left-deep operator trees, we demand that a join must be part of the mesh iff it has left input. Recall that the operator mesh is composed of  $|SR| \cdot 2^{|K-1|}$  clusters, one for each source containing  $k_1$ . Figure 28 illustrates the generation of a part of the cluster in Figure 26. When the leftmost source  $S\{k_1\}$  first produces output, it creates its direct parents  $j_1$  and  $j_7$ , along with others that are not depicted (Figure 28(a)). Joins that produce MTJNT (e.g.,  $j_7$ ) connect to the (permanent) output operator. In contrast, when  $j_1$  generates results, it creates its own parents, for example,  $j_2$ ,  $j_5$  and  $j_6$  (Figure 28(b)). These directly process their input; for instance, when  $j_1$  outputs a first tuple t and instantiates  $j_2$ , this operator immediately probes t against  $T\{\}$ . Again, first results from  $j_2$  trigger the addition of new join operators.

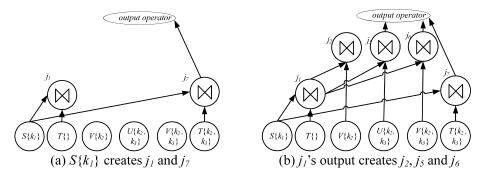


Fig. 28. Growing a cluster of operators from  $S\{k_1\}$ .

#### **TreeGen** (Operator $t_{new}$ .op)

- 1. if  $t_{new}.op$  is a selection
- 2. Tree  $t_{new} = a$  tree with a single node  $n_{root}$
- 3. else //  $t_{new}.op$  is a join
  - Tree  $t_{old}$  = TreeGen(left child of  $t_{new}$ .op)
- 5. Let  $n_{new}$  be the node corresponding to the right child of  $t_{new}.op$
- 6. Let  $n_{old}$  be the node joined with  $n_{new}$  in  $t_{new}.op$
- 7. Tree  $t_{new}$  = add  $n_{new}$  as the rightmost child of  $n_{old}$  in  $t_{old}$
- 8. return  $t_{no}$

Fig. 29. Algorithm TreeGen.

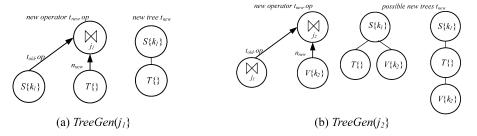


Fig. 30. Examples of TreeGen.

It remains to show how an operator at an arbitrary level in the mesh determines its direct parents. Recall from Sections 3.2 and 6.1 that whenever CNGen creates a new tree  $t_{new}$  (by adding a node  $n_{new}$  to a previous tree  $t_{old}$ ), a join  $t_{new}.op$  is inserted into the operator mesh. The left input of  $t_{new}.op$  is  $t_{old}.op$  and the right one is the source of  $n_{new}$ . In PM the problem is reversed: We have an operator  $t_{new}.op$ , but we need the corresponding tree  $t_{new}$  in order to decide which parents to create. Figure 29 illustrates TreeGen, an algorithm for reconstructing a tree  $t_{new}$ , given its last added operator  $t_{new}.op$ .

In essence, the algorithm checks the join condition of  $t_{new}.op$ : If  $n_{old}$  is the source joined with  $n_{new}$ , then  $t_{new}$  is generated by adding  $n_{new}$  as the rightmost child of  $n_{old}$  in  $t_{old}$ . Tree  $t_{old}$  is reconstructed recursively in the same manner. Figure 30 explains TreeGen by retracing the steps of Figure 28. When  $S\{k_1\}$  produces its first output,  $TreeGen(S\{k_1\})$  returns a tree  $t_0$  that contains a single node  $S\{k_1\}$ . The parents of  $S\{k_1\}$  in the mesh are computed by simulating

one loop of  $CNGen(S\{k_1\})$ , that is, adding nodes to  $t_0$  according to the rules of Section 3.2. Each parent (e.g.,  $j_1, j_7$ ) is inserted in the mesh and connected to its left and right inputs. Similarly, when  $j_1 = S\{k_1\} \bowtie T\{\}$  starts generating results, it has to create the layer of its parents. The call  $TreeGen(j_1)$  returns the tree  $t_1$  of Figure 30(a), derived by adding a child  $T\{\}$  to the only node  $S\{k_1\}$  of  $t_0$ . The expansion of  $t_1$  reveals the parents of  $j_1$  (e.g.,  $j_2$ ,  $j_5$ ,  $j_6$ ). Continuing the example, when  $j_2$  starts producing results, it has to create its own parents.  $TreeGen(j_2)$  checks which component of  $j_1$  joins with  $V\{k_2\}$  in  $j_2$ . If  $V\{k_2\}$  is joined with  $S\{k_1\}$ ,  $t_2$  is derived by adding  $V\{k_2\}$  as the rightmost child of  $S\{k_1\}$  in  $t_1$  (left tree in Figure 30(b)). Otherwise ( $V\{k_2\}$  is joined with  $T\{\}$ ),  $t_2$  is derived by adding  $V\{k_2\}$  to  $T\{\}$  (right tree in Figure 30(b)). Note that during the computation of  $t_2$ , we must also reconstruct  $t_1$ , since previous trees have been discarded. Keeping intermediate trees would require a large amount of memory, defeating the purpose of PM.

Conversely to generating parents, any operator without output destroys its parents, thereby freeing memory. In Figure 28(b), if  $j_1$  stops producing output, its buffer eventually runs dry. Consequently, the parents  $j_2$ ,  $j_5$ , and  $j_6$  are removed, leading back to the partial mesh of Figure 28(a). Join operators that have been removed from main memory are regenerated whenever necessary, for instance, fresh output by  $j_1$  at a later time leads to the anew creation of  $j_2$ ,  $j_5$ , and  $j_6$ . The destruction of parent operators recursively travels up the operator mesh, for example, if  $S\{k_1\}$  dries up, the entire cluster in Figure 28(b) is reduced to its sources.

### 6.5 Purging Expired Tuples

When a source tuple s is deleted, all intermediate results that include s must be removed from the system. Under the positive-negative stream model, purging is part of query processing; that is, a negative tuple -s travels up the mesh, expunging all composite tuples containing s. The sliding window model allows different variants for removing tuples. Under this model, source buffers are ordered by  $s.t_{end}$  (since  $s.t_{end} = s.t_{start} + w$ ), and can be purged by simply inspecting the topmost tuples. In contrast, the output buffers of joins are not sorted on  $t_{end}$  (since join results do not expire according to their creation order), and deletions involve complete buffer scans. Thus, in the sequel we assume that source buffers are immediately purged, and propose two algorithms, eager and lazy, for removing tuples from the output buffers of join operators.

The eager approach, illustrated in Figure 31, mimics the bottom-up method used under the positive-negative model. Specifically, whenever a source tuple expires, the corresponding leaf operator removes the tuple from its output buffer and informs its parents. Any join operator receiving such a note checks its own output buffer, and (should it find expired tuples) informs its parents. The approach is memory-optimal, since deleted tuples are removed immediately from all affected operators. However, it is CPU-intensive, due to the recursive call for all parents (potentially thousands) in lines 7 and 8.

In contrast, the lazy approach reduces CPU consumption by removing expired tuples only when these are encountered during join execution. Assume,

```
      Eager (Operator op)

      1. boolean tell_parents = false

      2. For all tuples s in op.out

      3. If s expires

      4. tell_parents = true

      5. Remove s from op.out

      6. If (tell_parents)

      7. For all parent operators p of op

      8. Eager(p)
```

Fig. 31. Eager purging.

for instance, that in Figure 26  $S\{k_1\}$  and  $T\{\}$  have tuples from which  $j_1$  produces output. Whenever a tuple in  $V\{k_2\}$ ,  $U\{k_2, k_3\}$ , or  $V\{k_2, k_3\}$  arrives, it is probed against  $j_1.out$ . The probe loops over the buffer and inspects each tuple for matching join attributes. During the loop, all dead tuples in  $j_1.out$  are removed, essentially for free. Lazy thereby incurs minimal CPU overhead, but provides no guarantee regarding when a dead tuple is removed. If  $V\{k_2\}$ ,  $U\{k_2, k_3\}$ , or  $V\{k_2, k_3\}$  dry up,  $j_1.out$  will not be purged and continue to waste memory.

For FM, we combine lazy with demand-driven operator execution, in order to limit the time that expired tuples remain in the system. Recall that the troublesome case involves an operator  $(j_1)$  with output, whose parents  $(j_2, j_5, j_6)$  have no right input. Using demand-driven operator execution,  $j_1$  must be sleeping, since all its parents are also dormant. The problem of deleting expired tuples is hence reduced to purging the output buffers of sleeping operators. When an operator op halts, its output buffer may still contain live tuples that cannot be expunged since op may wake up soon. However, after op sleeps for w seconds, its entire output has expired, and its buffer can be discarded. On the other hand, if op restarts before w, the expired tuples will be removed by join processing. Even if a tuple in the output buffer expires before op halted, it cannot remain in the system for more than 2w after its expiration.

In order to monitor outdated buffers, lazy maintains a doubly linked list Q of sleeping operators. If an operator op halts, an entry  $e = \langle op, stopTime \rangle$  is appended to Q. Additionally, op keeps a pointer to e. A continuous process watches Q's head. When the topmost operator  $op_{top}$  (the first to halt in Q) has been sleeping for w ticks  $(op_{top}.stopTime + w = now)$ , it is dequeued and its buffer is cleared of all content. Should an operator wake up before it is dequeued, it removes its entry from Q by following the corresponding pointer. Since the removal of outdated buffers relies on demand-driven operator execution, this optimization is only applicable to FM. In contrast, lazy purging for PM does not provide any guarantees regarding when an expired tuple is deleted.

#### 6.6 Changes in the Schema

Schema changes may be caused by the appearance or disappearance of either a source (SR), or an edge indicating which SR can be joined. In the following, we focus on changes due to SR; those incurred by edges are handled similarly. First, we address appearances. A new SR  $S_{new}$  at time  $t_{new}$ , introduces  $2^{|K|}$  new nodes in the expanded schema and produces an equal number of source operators. Let  $M_{old}$  ( $M_{new}$ ) be the operator mesh before (after)  $t_{new}$ . Directly

switching from  $M_{old}$  to an empty  $M_{new}$  is not permissible, since older tuples (and intermediate results) that are still alive at  $t_{new}$  would be lost. Instead,  $M_{new}$  is generated on top of  $M_{old}$ , so that all operators of  $M_{old}$  (and their intermediate results) become part of  $M_{new}$ . Specifically, we apply CNGen using the same nid as  $M_{old}$  for old nodes, and assign to each new node a nid that is larger than that of all older sources. Consequently, every operator cluster in  $M_{old}$  becomes part of a cluster in  $M_{new}$ . Additionally,  $M_{new}$  contains  $2^{|K-1|}$  additional clusters, rooted at sources of  $S_{new}$ . In order not to suspend query processing, the migration from  $M_{old}$  to  $M_{new}$  occurs successively. During the transition, tuples are routed up the mesh as usual. Each new join operator that receives tuples from both children processes them directly, ensuring that tuples which arrived after  $t_{new}$  are properly joined with older ones, and no results are lost during mesh migration.

The disappearance of an SR causes the removal of  $2^{|K-1|}$  sources from the mesh. All direct parents of these sources are also purged. The removal of parents travels recursively up the mesh. This process may cause some other operators to remain without parents. Such operators must also be deleted from the mesh. To achieve this, for every direct parent p of a deleted source, we insert the left child into a list  $l_{rem}$ , and delete p. After this stage terminates, each operator in  $l_{rem}$  that has no parents is removed and its left child is appended to  $l_{rem}$ . The process terminates when  $l_{rem}$  is empty. SR disappearances require no immediate attention and can be performed whenever the system has resources to spare. The previous discussion applies to FM as well as PM. The only difference is that in PM new operators are only created as high as there is data. This concludes the algorithmic part of the article; next, we evaluate the proposed methods experimentally.

## 7. EXPERIMENTAL EVALUATION

Section 7.1 compares OB and GB on snapshot queries using the TPC-H benchmark. Section 7.2 evaluates the two methodologies and the impact of various optimizations on continuous queries over relational streams. Section 7.3 concludes with guidelines regarding the most suitable method depending on the problem characteristics. All algorithms are implemented in C++ and experiments are performed on a 3.2 GHz Dual-Pentium IV with 2GB of RAM.

### 7.1 Snapshot R-KWS Queries over Tables

We compare GB and OB implemented as described in Section 3 (without the optimizations of later sections). We loaded the TPC-H dataset into a relational database powered by MySQL 5.1, using the MyISAM storage engine and its internal full-text index. For our experiments we focus on the six tables of Figure 32: Part~(0.2M~entries),~Supplier~(10K),~PartSupp~(0.8M),~Customer~(150K),~Orders~(1.5M),~and~LineItem~(6M). Two tables can join if and only if there is a foreign-key to primary-key between them, shown as arrows in Figure 32. We restrict the length of join sequences to  $T_{max}$ , which ranges between 4 and 6.

We designed seven sets of R-KWS queries  $QS_1$ - $QS_7$ , listed in Table I. We use the following types of keywords in the queries: (i) people's or

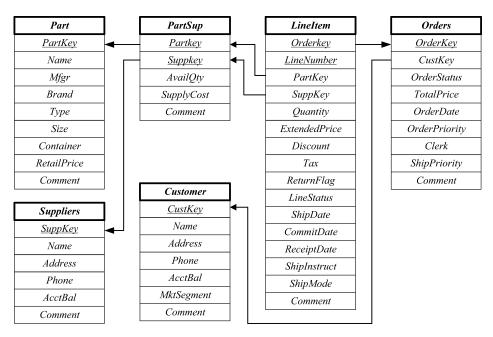


Fig. 32. TCP-H tables used in the experiments.

Table I. Query Set

| Query Set | Template   |
|-----------|--|
| $QS_1$    | $PeopleName_1, PeopleName_2$   |
| $QS_2$    | $PeopleName_1, PeopleName_2, PeopleName_3$                                       |
| $QS_3$    | $PartName_1, PartName_2$   |
| $QS_4$    | $PartName_1, PartName_2, PartName_3$   |
| $QS_5$    | $PeopleName_1, PeopleName_2, Year_1$   |
| $QS_6$    | $PartName_1, PartName_2, Year_1$   |
| $QS_7$    | $PartBrand_1$ , $PeopleName_1$ , $PartMfgr_1$ , $PartType_1$ , $PartContainer_1$ |

companies' names (denoted as PeopleName), which appear in the columns Customer.Name, Supplier.Name, and Orders.Clerk; (ii) terms from the name of a part, for example, "ivory", from the Part.Name attribute; (iii) years, which are present in LineItem.ShipDate, LineItem.CommitDate, LineItem.ReceiptDate, Orders.OrderDate; and (iv) terms from Part.Brand, Part.Mfgr, Part.Size, and Part.Container. We manually selected all queries such that they produce at least one result at the minimum  $T_{max}$  value (i.e., 4). This simulates the fact that users usually have background knowledge on the data. Furthermore, every class of queries mimics a practical search task. Specifically, queries in: (i)  $QS_1$  and  $QS_2$  retrieve connections between multiple people, (ii)  $QS_3/QS_4$  find co-occurrences of different parts, (iii)  $QS_5$  (respectively,  $QS_6$ ) are similar to  $QS_1$  (respectively,  $QS_3$ ), with an additional keyword specifying the year that these relationships occur; (iv)  $QS_7$  consist of attributes of one particular part.

Figure 33 depicts the total runtime (y-axis) of GB and OB, as well as the result set cardinality |R| (below the x-axis) for the seven query sets. For each query set, we generated 5 queries, and report the median values after setting

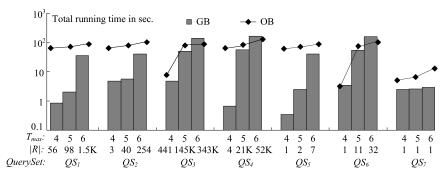


Fig. 33. Query processing time for various  $T_{max}$ .

 $T_{max}$  to 4, 5, and 6. Queries in  $QS_3$  and  $QS_4$  yield numerous results because terms from part names have low selectivity (the TPC-H generator uses a small dictionary to generate part names). In contrast, people's names are much more selective, leading to fewer results in the corresponding queries. Naturally, |R| and the query cost increases with  $T_{max}$ . For fixed  $T_{max}$ , all queries in the same set have similar cost because they search similar parts of the data graph (in GB), and create identical operator trees (in OB).

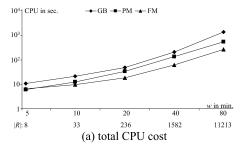
GB usually outperforms OB, sometimes by orders of magnitude because it utilizes the data graph precomputed at an initialization phase; consequently, during query processing it only accesses the database to read from the inverted index. The precomputation cost (around 220 seconds) is not included in the diagram because the same data graph is used by all queries. However, the graph consumes 0.737GB of RAM, which is a constant overhead of GB given that it must reside in main memory. The parameter  $T_{max}$  generally has a significant impact on the performance of GB because a higher  $T_{max}$  increases the portion of the data graph traversed for each query. This effect depends on the density of the subgraph where the search is performed. For  $QS_7$ , queries often comprise of keywords extracted from various attributes of a single record. Consequently, nodes far from the seeds may not reach new keywords, leading to early pruning of the partial result by GSearch, independently of the  $T_{max}$  value. The performance of OB is dominated by join operations. The cost is relatively low for when the joins involve small tables ( $QS_3$  at  $T_{max} = 4$ ,  $QS_6$  at  $T_{max} = 4$ , and  $QS_7$ ). In all other cases, the overhead of OB is high due to the size of the tables (e.g., *LineItem* has over 6M records).  $T_{max}$  generally does not have a significant impact on OB (except for  $QS_3$  and  $QS_6$ , where a small  $T_{max}$  avoids expensive joins) since a long join sequence often terminates after not obtaining results from first few tables due to high selectivity of the keywords.

#### 7.2 Continuous R-KWS Queries over Streams

Our stream implementations follow the *Pipes* data stream framework [Krämer and Seeger 2004]. Due to lack of real datasets, we resort to synthetic streams. In particular, we construct a schema containing |SR| streaming relations, connected in the shape of a ternary tree: Each SR can be joined with up to four other SR (one parent and three children). An SR has one attribute for each

Parameter Range & Default W 5, 10, **20**, 40, 80 minutes KWF 0.003, 0.007, **0.01**, 0.013, 0.016 |SR|5, 10, **15**, 20, 25 1/sel 1/500, 1/750, 1/1000, 1/1250, 1/1500 |K|2, 3, 4, 5  $T_{ma}$ 2, 3, 4, 5, 6, 7, 8 |Q|1, 2, 3, 4, 8, 16, 32, 64

Table II. Parameters under Investigation



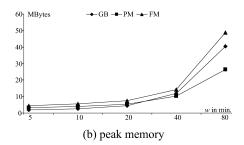


Fig. 34. Window size w.

edge, used to evaluate equijoins with the corresponding neighbor. Note that while the schema forms a tree, the data graph can take arbitrary shapes, and it may contain cycles. Additionally, the schema is larger and more densely connected than that of TPC-H (Figure 32). Each SR generates one tuple per second. Attribute values are randomly and independently chosen in the range [1, sel]. Two tuples of neighboring SR can thus be joined with probability 1/sel, the join selectivity. A tuple contains several different keywords, each with an independent probability KWF. We assume a sliding window of w minutes, and answer a continuous R-KWS query with |K| keywords over five hours. Table II illustrates the ranges and the default values (in boldface) of the experimental parameters.

Initially, we investigate the three core methodologies for continuous R-KWS queries: graph-based (GB), a full mesh of operators (FM), and a partial mesh of operators (PM). Subsequently, we study the effect of individual optimizations. The GB implementation uses keyword labels including temporal information (time-KL). FM includes lazy purging, demand-driven operator execution, and temporary operator disconnection. Recall that these optimizations are not applicable to PM. We investigate peak memory and total CPU as a function of w, KWF, |SR|, sel, |K|, and  $T_{max}$ . In each experiment, we vary one parameter and set the remaining ones to their default. Additionally, we report the number of generated results |R| (shown under the x-axis of the CPU chart). If affected by the parameter under investigation, we also state the duration (in seconds) of FM mesh-initialization (below the x-axis of the memory chart), which indicates the size and complexity of the operator mesh.

Figures 34(a) and 34(b) illustrate the impact of the window size w on the total CPU time (in seconds) and the peak memory (in Mbytes). While the number of live tuples grows linearly with w, the number of combinations in which they



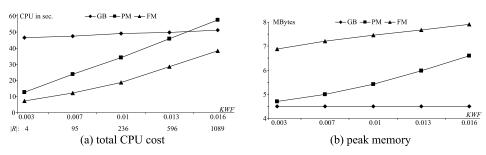


Fig. 35. Keyword frequency KWF.

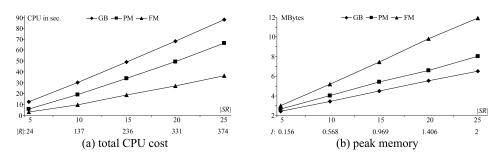


Fig. 36. Number of stream relations |SR|.

can be joined (i.e., edges in the data graph) grows quadratically, as reflected in the number of results |R|. Consequently, GB consumes more space and CPU, to store the data graph and construct edges, respectively. Both OB approaches behave similarly, as tuples are more likely to travel up the mesh, requiring CPU for construction and space for storage of intermediate results. As expected, FM incurs the least computational overhead, while PM excels in terms of space.

Figure 35 investigates the impact of the keyword frequency KWF. The relative performance of FM and PM remains similar to Figure 34. Resource consumption of both OB systems grows with KFW, as tuples are more likely to pass selection operators and climb the operator mesh. The increasing number of intermediate results is reflected in the cardinality of the result set |R|. In comparison, GB requires less memory, but uses significant amounts of CPU. This approach remains almost insensitive to varying values of KWF, because the dominant factor, the maintenance of the materialized data graph, is independent of keyword appearances. The slight increase in the CPU and memory consumption of GB are attributed to: (i) a growing number of graph traversals, and (ii) storage and maintenance of additional keyword labels. Later in this section, we investigate the impact of KL individually.

Figure 36 evaluates the effect of the number of streaming relations |SR|. Because each SR is limited to four neighbors, an increasing |SR| causes a linearly more complex streaming schema and data graph. The resource consumption of all methods grows proportionally. In the case of GB, the enlarged data graph requires more memory (for storage) and CPU (for construction). For OB systems, the greater streaming schema causes the operator mesh to grow, as can be seen from the prolonged initialization phase (I) of FM (see Figure 36(b)). The

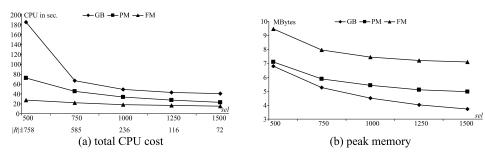


Fig. 37. Join selectivity 1/sel.

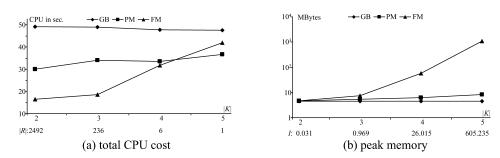
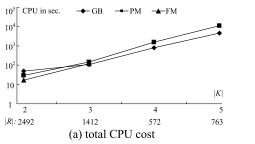


Fig. 38. Number of keywords |K|.

same is true for the number of tuples reaching intermediate operators rises, as reflected in the number of results |R|. The relative performance of all three systems is similar to the diagrams of Figure 35.

Figure 37 investigates the impact of the join selectivity. An increase in sel (drop in 1/sel) causes a quadratic decrease in the number of edges in the data graph, and the number of results |R| shrinks accordingly. Consequently, GB incurs a quadratically smaller overhead for creating and storing the data graph. Likewise, KL have to be propagated over a reduced number of links. FM and PM display an analogous behavior, as tuples are less likely to reach higher levels of the operator mesh. PM benefits most from a high sel, as fewer operators have to be instantiated and executed. Again, FM is most efficient in terms of CPU, and GB in terms of space.

Figure 38 depicts the effect of the query keywords |K|. Since this parameter has no impact on the number of tuples or the way they can be joined, GB remains almost unaffected. For OB, however, more query terms cause an exponential growth in the size and complexity of the operator mesh, as observed from the initialization time of FM (parameter I in Figure 38(b)). Three keywords require only 1 second of initialization, whereas five keywords take about 10 minutes. Since most operators in this mesh are commonly idle, they are never created by PM, hence the increasing advantage of this approach in terms of memory. Whereas in Figure 38  $T_{max}$  is fixed to 4 (default value), Figure 39 repeats the experiment after setting  $T_{max}$ :=|K|+2. The comparison with Figure 38 reveals that in this case the number of results increases fast with |K| and  $T_{max}$ , which leads to an analogous growth of processing costs. Notably, for



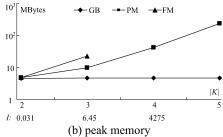


Fig. 39. Number of keywords |K|, with  $T_{max} := |K| + 2$ .

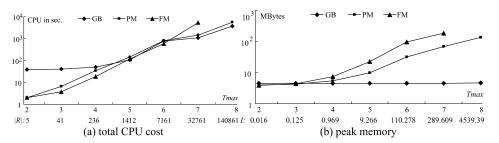


Fig. 40. Limit  $T_{max}$  of nodes per MTJNT.

|K| > 3, the memory requirement of FM exceeds the amount of main memory in our system (i.e., 2GBytes); at |K| = 5, the CN generation module alone takes longer than 5 hours. GB is most scalable with respect to |K| and  $T_{max}$ , as its memory consumption remains almost constant in all settings.

Figure 40 investigates the effect of  $T_{max}$ , which has a similar impact to |K| because it does not influence the number of tuples, or the way that they can be joined. GB's memory consumption thus remains constant. However, for high values of  $T_{max}$ , MTJNT can become larger and more complex. Thus, keyword labels must be propagated in a greater radius, and every call to GSearch explores a larger fraction of the data graph. Consequently, GB's CPU overhead grows with  $T_{max}$ . For OB systems, an increase in  $T_{max}$  leads to an exponential growth of the operator mesh. In case of FM, operator initialization for  $T_{max}=6$  exceeds two minutes, compared to less than one second for  $T_{max}=4$ . Akin to the mesh size, CPU and memory grow fast, since: (i) the mesh requires more space, (ii) the number of intermediate results increases, and (iii) their generation incurs additional CPU. The number of intermediate tuples is reflected in the cardinality of the result set |R|. For  $T_{max}=8$ , FM was too slow to process tuples at the given arrival rate and PM is the only viable OB solution.

Whereas the previous experiments assume a single query, Figure 41 depicts the resources consumed by |Q| parallel queries. For queries consisting of random keywords drawn from a large dictionary, the chance that two queries share any common term is negligible; hence, we assume all |Q| queries to be disjoint. Consequently, both PM and FM execute |Q| independent queries, and their costs increase linearly with |Q|. In contrast, a large portion of GB's overhead is due to the storage and maintenance of the data graph, which is shared among

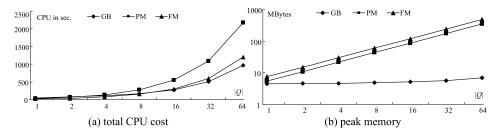


Fig. 41. Number of concurrent queries |Q|.

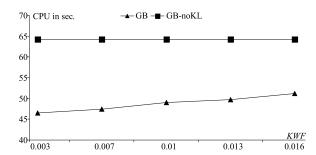


Fig. 42. Effect of keyword labels on GB vs. keyword frequency KWF.

all queries. Therefore, GB scales better than OB approaches with the number of parallel queries.

Next, we evaluate specific optimizations. Figure 42 evaluates the effect of  $Keyword\ Labeling\ (KL)$  on GB, using  $KeyWord\ Frequency\ (KWF)$  as an exemplary parameter. While peak memory for both approaches remains identical and almost constant, KL reduces the overall CPU consumption. The unoptimized approach explores the data graph in a diameter of  $T_{max}$  around every new tuple. Consequently, the workload stays unaffected by KWF. In contrast, the optimized system only traverses tuples near a keyword occurrence. As KWF increases, this case becomes progressively more frequent, rendering KL less beneficial. Eventually, for a very high KWF, the workload of both approaches converges. On the other end of the spectrum, a smaller KWF promises even greater savings from KL.

Figure 43 illustrates the importance of duplicate elimination in GSearch, by comparing the number of results returned by our duplicate-free GSearch with a straightforward breath-first graph traversal. For w=80, GSearch finds 11,213 unique results, whereas the naïve approach outputs 18,844, including duplicates. In addition to the overhead for retrieving these duplicates, the naïve method entails the cost of eliminating them during a post-processing step.

The next experiment evaluates the benefits of the operator mesh for OB systems. As an exemplary parameter, we chose the query cardinality |K|. Figure 44 depicts the resource utilization of two systems: (i) FM, using an operator mesh, and (ii) a forest of independent operator trees. The operator mesh reduces both CPU and memory requirements by orders of magnitude. Indeed, the forest approach crashes for any complex scenario, for instance, an increased  $T_{max}$ .

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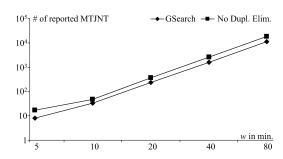


Fig. 43. Effect of duplicate elimination in GSearch vs. window size w.

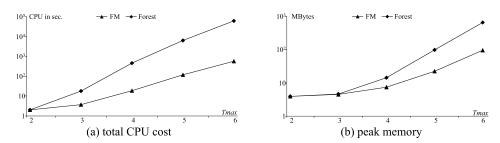


Fig. 44. Effect of mesh for OB vs. the number of keywords |K|.

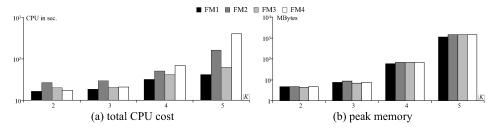


Fig. 45. Effect of FM optimizations vs. the number of keywords |K|.

Next, we evaluate four different variations of FM. The first (FM1) is fully optimized, and uses demand driven-operator execution, lazy purging, and temporary operator disconnection (this is the default implementation used in the previous experiments). The second variation (FM2) does *not* use demand-driven operator execution; instead it pushes intermediate tuples up the operator mesh, even if they are not needed. In contrast, FM3 is fully optimized, but applies eager purging, as described in Section 6.5. Finally, FM4 uses demand-driven operator execution and lazy purging, but omits temporary operator disconnection, that is, its join operators remain connected to their right child at all times. Since the benefits of the various optimizations do not become apparent under all settings, we evaluate the four implementations on two exemplary parameters, the query cardinality |K|, and the join selectivity 1/sel. Figure 45(a) illustrates the impact of |K| on CPU consumption. As suggested by the high cost

<sup>&</sup>lt;sup>2</sup>FM4 served as the default FM implementation in Markowetz et al. [2007].

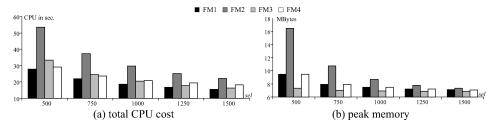


Fig. 46. Effect of FM optimizations vs. the join selectivity 1/sel.

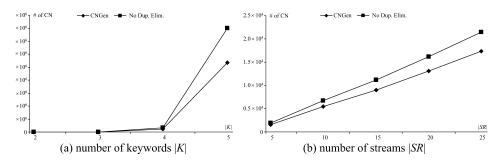


Fig. 47. Duplicates avoided by CNGen.

of FM2, demand-driven operator execution reduces computational expenses at all settings. Similarly, the poor performance of FM4 illustrates the benefits of temporary operator disconnection on large operator meshes (|K|=4 and |K|=5). Finally, lazy (FM1) consistently outperforms eager pruning (FM3). For this particular parameter, all four implementations consume roughly the same amount of memory (Figure 45(b)).

Figure 46 studies the effect of the join selectivity. Small values of sel lead to more intermediate results, reflected in an increasing consumption of CPU as well as memory. A comparison of FM1 and FM2 reveals that demand-driven operator execution achieves the most significant gains, especially for low sel. The relatively good performance of FM4 indicates that temporary operator disconnection is not as beneficial under this setting as in Figure 45. As predicted in Section 6.5, lazy purging (FM1) outperforms eager (FM3) in terms of CPU, but consumes more peak memory.

Finally, we evaluate the importance of duplicate elimination during CN generation. Figure 47 illustrates the number of CN generated by two algorithms. The first is CNGen, which avoids duplicates by observing a unique preorder traversal. The second removes all duplicate-specific checks from CNGen (i.e., line 8 and lines 17 through 23 in Figure 10), resulting in a CN generation module similar to that proposed in Discover.<sup>3</sup> For the default parameters (|K| = 3, |SR| = 15, and  $T_{max} = 4$ ), the naïve algorithm generates roughly 3,000

<sup>&</sup>lt;sup>3</sup>For static databases, *Discover* executes a partial CN as soon as it is generated, and prunes it if the join yields an empty result set. Depending on the data, it may produce fewer CN than in Figure 47. However, in a stream setting, all CN must be created before their execution, and this optimization is no longer applicable.

duplicates, in addition to about 9,000 unique CN. For more complex settings, this number increases rapidly. Duplicate elimination is a key feature of *CNGen* since duplicate CN are expensive to create, detect, and remove.

# 7.3 Summary of Experimental Evaluation

For conventional tables, GB is more efficient than OB, often by a wide margin. Moreover, since the construction of the data graph is relatively efficient and the graph can be dynamically maintained, GB is preferable for datasets with frequent updates. However, GB consumes a considerable amount of main memory to store the data graph. In contrast, OB utilizes the functionality provided by a DBMS, and, thus, can answer R-KWS queries using much less memory than GB. Therefore, the choice between the two methods depends on the application scenario. On servers dedicated for R-KWS queries, GB is the best choice due to its high performance. On servers running multiple applications and only answering R-KWS queries infrequently, OB might be preferable due to its low memory footprint. Finally, OB is the only feasible solution if the data graph is too large to fit into main memory.

For continuous R-KWS queries over data streams, FM is usually the most CPU-efficient method for a single query, except for queries involving numerous keywords and/or a large value of  $T_{max}$ . On the other hand, GB and PM are more economical in terms of memory consumption. In particular, GB is the best choice for long queries and large values of  $T_{max}$ . Moreover, GB scales better with the number of registered queries, and it is the clear choice for systems involving numerous simultaneous queries. Compared to previous GB methods, GSearch avoids duplicate results, which reduces the total cost (to retrieve and then eliminate duplicates). Furthermore, the proposed keyword labeling schemes limit the scope of graph traversals, further enhancing performance. The optimizations of OB also achieve significant benefits with respect to the basic methods.

#### 8. CONCLUSION

R-KWS has several advantages over conventional query languages; most notably, it handles broad query tasks whose complexity does not permit hand-coded structured queries. At the same time, it presents considerable algorithmic challenges because query processing has to explore a vast search space. We face these challenges through a series of contributions. First, we provide R-KWS semantics that are well defined and easily extensible to streaming environments. Then, we develop GB and OB processing techniques that match these semantics and remedy problems encountered in previous systems. Subsequently, we adapt our framework to relational streams, and propose a wide range of optimizations. Finally, we support our claims through an extensive set of experiments. In the future, we plan to further improve R-KWS performance by means of indexing. In parallel, we intend to integrate ranking into continuous R-KWS query processing. For example, if there are a sudden burst of results, it may be desirable to report only the top-k answers for the affected period.

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