Exploiting Database Management Systems and Treewidth for Counting

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Abstract
Bounded treewidth is one of the most cited combinatorial invariants in the literature. It was also applied for solving several counting problems efficiently. A canonical counting problem is \#SAT, which asks to count the satisfying assignments of a Boolean formula. Recent work shows that benchmarking instances for \#SAT often have reasonably small treewidth. This paper deals with counting problems for instances of small treewidth. We introduce a general framework to solve counting questions based on state-of-the-art database management systems (DBMSs). Our framework takes explicitly advantage of small treewidth by solving instances using dynamic programming (DP) on tree decompositions (TD). Therefore, we implement the concept of DP into a DBMS (PostgreSQL), since DP algorithms are already often given in terms of table manipulations in theory. This allows for elegant specifications of DP algorithms and the use of SQL to manipulate records and tables, which gives us a natural approach to bring DP algorithms into practice. To the best of our knowledge, we present the first approach to employ a DBMS for algorithms on TDs. A key advantage of our approach is that DBMSs naturally allow for dealing with huge tables with a limited amount of main memory (RAM).

KEYWORDS: Dynamic Programming, Parameterized Algorithmics, Bounded Treewidth, Database Systems, SQL, Relational Algebra, Counting

1 Introduction
Counting solutions is a well-known task in mathematics, computer science, and other areas (Chakraborty et al. 2016; Domshlak and Hoffmann 2007; Gomes et al. 2009; Sang et al. 2005).

* This is an extended version of a paper (Fichte et al. 2020) that appeared in the Proceedings of the 22nd International Symposium on Practical Aspects of Declarative Languages (PADL-20).
In combinatorics, for instance, one characterizes the number of solutions to problems by means of mathematical expressions, e.g., generating functions (Doubilet et al. 1972). One particular counting problem, namely \textit{model counting} (\#SAT) asks to output the number of solutions of a given Boolean formula. While we stay in the realm of model counting, the findings of this work are also relevant for answer set programming. This is particularly true for tight programs (using, e.g., Clark’s completion (Clark 1977)), but also interesting for applications of quantitative reasoning, solved by programs that are compiled to SAT with the help of tools like lp2sat (Janhunen 2006) or lp2acyc (Bomanson et al. 2016).

Model counting and variants thereof have already been applied for solving a variety of real-world applications and questions in modern society related to reasoning (Chakraborty et al. 2016; Choi et al. 2015; Dueñas-Osorio et al. 2017; Xue et al. 2012). Such problems are typically considered very hard, since \#SAT is complete for the class \#P (Bacchus et al. 2003; Roth 1996), i.e., one can simulate any problem of the polynomial hierarchy with polynomially many calls (Toda 1991) to a \#SAT solver. Taming this high complexity is possible with techniques from parameterized complexity (Cygan et al. 2015). In fact, many of the publicly available \#SAT instances show good structural properties after using regular preprocessors like pmc (Lagniez and Marquis 2014), see (Fichte et al. 2018b; Fichte et al. 2019). By good structural properties, we mean that graph representations of these instances have reasonably small \textit{treewidth}. The measure treewidth is a structural parameter of graphs which models the closeness of the graph of being a tree. Treewidth is one of the most cited combinatorial invariants studied in parameterized complexity (Cygan et al. 2015) and was subject to algorithmics competitions (Dell et al. 2018).

The observation, stated above, that various recent problem instances for \#SAT have small treewidth, leads to the question whether a general framework that leverages treewidth is possible for counting problems. The general idea to develop such frameworks is indeed not new, since there are (a) specialized solvers such as dynQBF, gpuSAT, and fvs-pace (Charwat and Woltran 2019; Fichte et al. 2019; Kiljan and Pilipczuk 2018) as well as (b) general systems that exploit treewidth like D-FLAT (Bliem et al. 2016), Jatatosk (Bannach and Berndt 2019), and sequoia (Langer et al. 2012). Some of these systems explicitly use \textit{dynamic programming} (DP) to directly exploit treewidth by means of so-called \textit{tree decompositions} (TDs), whereas others provide some kind of declarative layer to model the problem (and perform decomposition and DP internally). In this work, we solve (counting) problems by means of explicitly specified DP algorithms, where essential parts of the DP algorithm are specified in form of SQL \texttt{SELECT} queries. The actual run of the DP algorithm is then delegated to our system \texttt{dpdb}, which employs database management systems (DBMSs) (Ullman 1989). This has not only the advantage of naturally describing and manipulating the tables that are obtained during DP, but also allows \texttt{dpdb} to benefit from decades of database technology in form of the capability to deal with huge tables using a limited amount of main memory (RAM), dedicated database joins, query optimization, and data-dependent execution plans. Compared to other generic DP systems like D-FLAT (Bliem et al. 2016), our system \texttt{dpdb} uses relational algebra (SQL) for specifying DP algorithms, which is even competitive with specialized systems for model counting, and therefore applicable beyond rapid prototyping.

\textbf{Contribution.} We implement a system \texttt{dpdb} for solving counting problems based on dynamic programming on tree decompositions, and present the following contributions. (i) Our system \texttt{dpdb} uses database management systems to handle table operations needed for performing dynamic programming efficiently. The system \texttt{dpdb} is written in Python and employs PostgreSQL as DBMS, but can work with other DBMSs easily. (ii) The architecture of \texttt{dpdb} allows to solve
general problems of bounded treewidth that can be solved by means of table operations (in form of relational algebra and SQL) on tree decompositions. As a result, dpdb is a generalized framework for dynamic programming on tree decompositions, where one only needs to specify the essential and problem-specific parts of dynamic programming in order to solve (counting) problems. (iii) Finally, we show how to solve the canonical problem \#SAT with the help of dpdb, where it seems that the architecture of dpdb is particularly well-suited. In more detail, we compare the runtime of our system with state-of-the-art model counters. We observe a competitive behavior and promising indications for future work.

Prior Work This is an extended version of a paper (Fichte et al. 2020) that appeared at the 22nd International Symposium on Practical Aspects of Declarative Languages. The new material includes improved and extended examples as well as a detailed description of our DP algorithms and how these algorithms can be implemented for the system dpdb. Further, we added two new DP algorithms for the additional problems MAXSAT and MINIDS, to demonstrate how to use dpdb: The problem MAXSAT is similar to SAT, but consists of hard clauses that need to be satisfied as well as soft clauses. The goal of MAXSAT is to compute the maximum number of soft clauses that can be satisfied using only assignments that also satisfy all the hard clauses. Further, problem MINIDS is a popular graph problem that aims for computing for a given graph, a set of vertices (called independent dominating set) such that there is no edge between these vertices and all the other vertices of the graph have an edge to at least one of the vertices in this set. Both problems can be easily extended to counting, where we require to compute the number of witnessing solutions. Finally, we added new experimental results, where we used the most recent version 12 of PostgreSQL as the underlying database management system, which operated on a ramdisk drive.

2 Preliminaries

We assume familiarity with the terminology of graphs and trees. For details, we refer to the literature and standard textbooks (Diestel 2012).

2.1 Boolean Satisfiability

We define Boolean formulas and their evaluation in the usual way, cf., (Kleine B"uning and Lettman 1999). A literal is a Boolean variable \( x \) or its negation \( \neg x \). A CNF formula \( \varphi \) is a set of clauses interpreted as conjunction. A clause is a set of literals interpreted as disjunction. For a formula or clause \( X \), we abbreviate by \( \text{var}(X) \) the variables that occur in \( X \). An assignment of \( \varphi \) is a mapping \( I : \text{var}(\varphi) \to \{0,1\} \). The formula \( \varphi(I) \) under assignment \( I \) is obtained by removing every clause \( c \) from \( \varphi \) that contains a literal set to 1 by \( I \), and removing from every remaining clause of \( \varphi \) all literals set to 0 by \( I \). An assignment \( I \) is satisfying if \( \varphi(I) = 0 \). Problem \#SAT asks to output the number of satisfying assignments of a formula.

We also allow equality formulas, which are Boolean formulas, where variables are expressions using equality. In more detail: Let \( d \) be a fixed constant over domain \( \text{dom}(v) \), where we call \( d \) term constant. Let \( v \) and \( v' \) be variables over some domain \( \text{dom}(v) \) and \( \text{dom}(v') \), where we call \( v \) and \( v' \) term variables. Then, an equality formula \( \beta \) is an expression of the form \( v = d \) or \( v = v' \). A term assignment \( J \) of equality formula \( \beta \) over term variables \( \text{tvar}(\beta) \) assigns each domain variable \( v \in \text{tvar}(\beta) \) a value over domain \( \text{dom}(v) \). The Boolean formula \( \beta(J) \) under term
assignment $J$ is obtained as follows. First, we replace all expressions $v = d$ in $\beta$ by 1 if $J(v) = d$, all expressions $v = v'$ by 1 if $J(v) = J(v')$, and by 0 otherwise. Second, we remove from the resulting clauses in $\beta(J)$ each clause $c$ that contains a literal set to 1. Finally, we remove from every remaining clause in $\beta(J)$ every literal that is set to 0. We say a term assignment $J$ is satisfying if $\beta(J) \neq 0$.

### 2.2 Tree Decomposition and Treewidth

Treewidth is widely used for fine-grained complexity analyzes and to establish algorithms that provide tractability when bounding the treewidth. While it is only defined for graphs and hence widely used in graph theory (Bodlaender and Koster 2008), one can define graph representations of input instances for a variety of problems. Dedicated techniques then allow to solve problems from many domains such as propositional satisfiability (Samer and Szeider 2010), artificial intelligence (Gottlob and Szeider 2007), knowledge representation (Gottlob et al. 2006), argumentation (Fichte et al. 2019), non-monotonic reasoning (Fichte et al. 2018), abduction in Datalog (Gottlob et al. 2007), and databases (Grohe 2007), probabilistic inference (Dechter 1999) (under the name bucket elimination) including constraint satisfaction, Fourier and Gaussian elimination for solving linear equalities and inequalities, and combinatorial optimization. While theoretical conditional lower bound results seem to discourage using algorithms that exploit bounded treewidth (Pan and Vardi 2006; Fichte et al. 2020), dynamic programming along tree decompositions or related decompositions has recently been used to establish practical algorithms (Fichte et al. 2019; Dudek et al. 2020). An algorithm that exploits small treewidth takes a tree decomposition, which is an arrangement of a graph into a tree, and evaluates the problem in parts, via dynamic programming (DP) on the tree decomposition. Informally speaking, the tree decomposition provides an evaluation ordering, which one employs by a problem specific algorithm where the runtime of combinatorial hard part is bounded by a (potentially exponential) function of the treewidth $w$.

Then, the underlying idea of treewidth is that it provides a measure for the closeness of a potential evaluation ordering which is simply a tree. Below, we provide the formal definitions of the notions tree decomposition and treewidth.

A tree decomposition (TD) of a given graph $G$ is a pair $\mathcal{T} = (T, \chi)$ where $T$ is a rooted tree and $\chi$ is a mapping which assigns to each node $t \in V(T)$ a set $\chi(t) \subseteq V(G)$, called bag, such that (i) $V(G) = \bigcup_{t \in V(T)} \chi(t)$ and $E(G) \subseteq \{ \{u, v\} \mid t \in V(T), \{u, v\} \subseteq \chi(t) \}$; and (ii) for each $r,s,t \in V(T)$, such that $s$ lies on the path from $r$ to $t$, we have $\chi(r) \cap \chi(t) \subseteq \chi(s)$. We let $\text{width}(\mathcal{T}) := \max_{t \in V(T)} |\chi(t)| - 1$. The treewidth $\text{tw}(G)$ of $G$ is the minimum width $\text{width}(\mathcal{T})$ over all TDs $\mathcal{T}$ of $G$. For $k \in \mathbb{N}$, we can compute a tree decomposition of width $k$ or output that none exists in time $2^{O(k)} \cdot |V|$ (Bodlaender and Koster 2008).

**Example 1**

Figure 1 depicts a graph $G$ (left) and a TD $\mathcal{T}$ of $G$ (right) of width 2. The treewidth of $G$ is also 2 since $G$ contains a complete graph with 3 vertices (Kloks 1994).
Next, we give the definition of a standard restriction for TDs, so called nice TDs. A nice TD ensures that one needs to consider only a small number of case distinctions in a DP algorithm later. For a node \( t \in V(T) \), we say that type\((t)\) is leaf if \( t \) has no children and \( \chi(t) = \emptyset \); join if \( t \) has children \( t' \) and \( t'' \) with \( t' \neq t'' \) and \( \chi(t) = \chi(t') = \chi(t'') \); intro ("introduce") if \( t \) has a single child \( t' \), \( \chi(t') \subseteq \chi(t) \) and \( |\chi(t)| = |\chi(t')| + 1 \); rem ("removal") if \( t \) has a single child \( t' \), \( \chi(t') \supseteq \chi(t) \) and \( |\chi(t')| = |\chi(t)| + 1 \). If for every node \( t \in V(T) \), type\((t)\) \( \in \{ \text{leaf, join, intro, rem} \} \), then the TD is called nice. The conditions allow us to focus on each of the cases of our algorithms individually.

3 Dynamic Programming on Tree Decompositions

In the preliminaries, we gave definitions for tree decompositions and treewidth and stated a variety of application areas. We mentioned that treewidth is widely used as a structural measure of the DP works the same for most problems. Hence, we focus on table algorithms and their provide the data structures and the table algorithm for the specific problem as the general outline along a given TD of a graph representation are supposed to be merged.

In practice, one can design a so called dynamic programming (DP) algorithm, which works as follows. Sub-problems are evaluated along the tree decomposition. At each node of the tree, information is gathered in tables. A table contains tuples of a fixed form that are designed to ensure certain properties. Then, a table algorithm maintains these tables during a post-order traversal. Thereby, it handles different cases according to the node contents of the TD and it ensures that properties required to solve the problem in parts sustain. The size of a table depends on the number of items in the bag, but is allowed to be exponential in the size of a bag. Hence, the overall technique works in linear time in the size of the problem and exponential in the bag size. Intuitively, the tree decomposition fixes an order in which we evaluate our problem. As a result, evaluating a problem along a tree decomposition allows for solving the problem at interest in parts, where the tree decomposition provides these parts and directs how solutions to the parts are supposed to be merged.

More formally, a solver based on dynamic programming (DP) evaluates the input \( \cal I \) in parts along a given TD of a graph representation \( G \) of the input. Thereby, for each node \( t \) of the TD, intermediate results are stored in a table \( \tau_t \). This is achieved by running a so-called table algorithm \( A \), which is designed for a certain graph representation, and stores in \( \tau_t \) results of problem parts of \( \cal I \), thereby considering tables \( \tau_t' \) for child nodes \( t' \) of \( t \). DP works for many problems \( \cal P \) as follows.

1. Construct a graph representation \( G \) of the given input instance \( \cal I \).
2. Heuristically compute a tree decomposition \( \cal T = (T, \chi) \) of \( G \).
3. Traverse the nodes in \( V(T) \) in post-order, i.e., perform a bottom-up traversal of \( T \). At every node \( t \) during post-order traversal, execute a table algorithm \( A \) that takes as input \( t \), bag \( \chi(t) \), a local instance \( \cal P(t, \cal I) = \cal I_t \) depending on \( \cal P \), as well as previously computed child tables of \( t \) and stores the result in \( \tau_t \).
4. Interpret table \( \tau_n \) for the root \( n \) of \( T \) in order to output the solution of \( \cal I \).

When specifying a DP algorithm for a specific problem such as \#SAT, it is often sufficient to provide the data structures and the table algorithm for the specific problem as the general outline of the DP works the same for most problems. Hence, we focus on table algorithms and their description in the following.

Next, we state the graph representation and table algorithm that we need to solve the problem \( \cal P = \#SAT \) (Samer and Szeider 2010). First, we need the following graph representation. The
Algorithm 1: Table algorithm \( \text{Sat}(t, \chi(t), \varphi, \langle \tau_1, \ldots, \tau_r \rangle) \) for solving \#SAT (Samer and Szeider 2010).

**In:** Node \( t \), bag \( \chi(t) \), clauses \( \varphi \), and a sequence \( \langle \tau_1, \ldots, \tau_r \rangle \) of child tables.

**Out:** Table \( \tau_c \).

1. if type\((t) = \text{leaf}\) then \( \tau_c := \{ (\emptyset, 1) \} \)
2. else if type\((t) = \text{intr.}, \) and \( a \in \chi(t) \) is introduced then
   \[ \tau_c := \{(I \cup \{a \rightarrow 0\}, c) \mid (I, c) \in \tau_c, \varphi(I \cup \{a \rightarrow 0\}) = \emptyset\} \]
   \[ \cup \{(I \cup \{a \rightarrow 1\}, c) \mid (I, c) \in \tau_c, \varphi(I \cup \{a \rightarrow 1\}) = \emptyset\} \]
3. else if type\((t) = \text{rem.}, \) and \( a \notin \chi(t) \) is removed then
   \[ \tau_c := \{(I \setminus \{a \rightarrow 0, a \rightarrow 1\}, \sum_{c \in C(I)} c) \mid (I, c) \in \tau_c\} \]
   \[ / * C(I) \text{ is the set that contains all counters for which assignments } J \text{ are the same as } I \text{ after we remove } a \text{ from the assignment } I */ \]
   \[ C(I) := \{c \mid (J, c) \in \tau, J \setminus \{a \rightarrow 0, a \rightarrow 1\} = I \setminus \{a \rightarrow 0, a \rightarrow 1\}\} \]
4. else if type\((t) = \text{join} \) then
   \[ \tau_c := \{ (I_c_1, c_2) \mid (I, c_1) \in \tau_c, (I, c_2) \in \tau_2\} \]
5. return \( \tau_c \)

The primal graph \( G_{\varphi} \) of a formula \( \varphi \) has as vertices its variables, where two variables are joined by an edge if they occur together in a clause of \( \varphi \). Given formula \( \varphi \), a TD \( T, \chi \) of \( G_{\varphi} \) and a node \( t \) of \( T \). Sometimes, we refer to the treewidth of the primal graph of a given formula by the treewidth of the formula. Then, we let local instance \#SAT\((t, \varphi) = \varphi_t \) be \( \varphi_t := \{c \mid c \in \varphi, \text{var}(c) \subseteq \chi(t)\} \), which are the clauses entirely covered by \( \chi(t) \).

Table algorithm Sat as presented in Algorithm 1 shows all the cases that are needed to solve \#SAT by means of DP over nice TDs. Each table \( \tau_c \) consist of rows of the form \( (I, c) \), where \( I \) is an assignment of \( \varphi_t \), and \( c \) is a counter. Nodes \( t \) with type\((t) = \text{leaf}\) consist of the empty assignment and counter 1, cf., Line 1. For a node \( t \) with introduced variable \( a \in \chi(t) \), we guess in Line 3 for each assignment \( \beta \) of the child table, whether \( a \) is set to true or to false, and ensure that \( \varphi_t \) is satisfied. When an atom \( a \) is removed in node \( t \), we project assignments of child tables to \( \chi(t) \), cf., Line 5, and sum up counters of the same assignments. For join nodes, counters of common assignments are multiplied, cf., Line 7.

Example 2
Consider formula \( \varphi := \{\neg a, b, c\}, \{a, \neg b, \neg c\}, \{a, d\}, \{a, \neg d\}\}. Satisfying assignments of formula \( \varphi \) are, e.g., \{a \rightarrow 1, b \rightarrow 1, c \rightarrow 0, d \rightarrow 0\}, \{a \rightarrow 1, b \rightarrow 0, c \rightarrow 1, d \rightarrow 0\} or \{a \rightarrow 1, b \rightarrow 1, c \rightarrow 1, d \rightarrow 1\}. In total, there are 6 satisfying assignments of \( \varphi \). Observe that graph \( G_t \) of Figure 1 depicts the primal graph \( G_{\varphi} \) of \( \varphi \). Intuitively, \( T \) of Figure 1 allows to evaluate formula \( \varphi \) in parts. Figure 2 illustrates a nice TD \( T = (\cdot, \chi) \) of the primal graph \( G_{\varphi} \) and tables \( \tau_1, \ldots, \tau_{12} \) that are obtained during the execution of Sat on nodes \( t_1, \ldots, t_{12} \). We assume that each row in a table \( \tau_c \) is identified by a number, i.e., row \( i \) corresponds to \( \omega_{ij} = (I_{ij}, c_{ij}) \).

Table \( \tau_1 = \{ (\emptyset, 1) \} \) has type\((t_1) = \text{leaf} \). Since type\((t_2) = \text{intr} \), we construct table \( \tau_2 \) from \( \tau_1 \) by taking \( I_{12} \cup \{a \rightarrow 0\} \) and \( I_{12} \cup \{a \rightarrow 1\} \) for each \( (I_{12}, \chi_{12}) \in \tau_1 \). Then, \( t_3 \) introduces \( c \) and \( t_4 \) introduces \( b \). \( \varphi_1 = \varphi_2 = \varphi_3 = \emptyset \), but since \( \chi(t_4) \subseteq \text{var}(c_1) \) we have \( \varphi_4 = \{c_1, c_2\} \) for \( t_4 \). In consequence, for each \( I_{4i} \) of table \( \tau_3 \), we have \( \{c_1, c_2\}(I_{4, i}) = \emptyset \) since Sat enforces satisfiability of \( \varphi_1 \) in node \( t \). Since type\((t_5) = \text{rem} \), we remove variable \( c \) from all elements in \( \tau_1 \) and sum up counters accordingly to construct \( \tau_5 \). Note that we have already seen all rules where \( c \) occurs and hence \( c \) can no longer affect interpretations during the remaining traversal. We similarly create \( \tau_6 = \{\{a \rightarrow 0\}, 3\}, \{\{a \rightarrow 1\}, 3\}\} \) and \( \tau_{10} = \{\{a \rightarrow 1\}, 2\}\}. Since type\((t_{11}) = \text{join} \), we
build table $\tau_{11}$ by taking the intersection of $\tau_6$ and $\tau_{10}$. Intuitively, this combines assignments agreeing on $a$, where counters are multiplied accordingly. By definition (primal graph and TDs), for every $c \in \varphi$, variables $\var{c}$ occur together in at least one common bag. Hence, since $\tau_{12} = \{\emptyset, 6\}$, we can reconstruct for example model $\{a \mapsto 1, b \mapsto 1, c \mapsto 0, d \mapsto 1\} = I_{11,1} \cup I_{5,4} \cup I_{9,2}$ of $\varphi$ using highlighted (yellow) rows in Figure 2. On the other hand, if $\varphi$ was unsatisfiable, $\tau_{12}$ would contain no values, i.e., $\tau_{12} = \emptyset$.

### 4 Dynamic Programming on Tree Decompositions Expressed in Relational Algebra

While algorithms that run dynamic programming on bounded treewidth can be quite useful for efficient problem solving in practice, implementations turn out to be tedious already for problems such as the propositional satisfiability problem. In the following of the paper, we aim for rapid prototyping with dynamic programming by a declarative approach that ideally uses existing systems, gets parallel execution for free, and remains fairly efficient.

In the previous section, we explained that the traversal of the tree decomposition and the overall methodology of the procedure stays the same. But the core of dynamic programming on tree decompositions for various problems is mostly the specification of the table algorithm that modifies a table based on previously computed tables. Hence, one can often focus on the table algorithms and their descriptions. When recalling basics from databases 101 from undergraduate studies (Elmasri and Navathe 2016) and taking a closer look on Algorithm 1 above, we can immediately spot that we are effectively describing a query on existing data that produces a new table by Algorithm 1. This motivates our idea to use a database management system to execute the query and specify the query in SQL. Before we can proceed with our idea to use databases for executing DP algorithms, we take a step back and recall that the theory of SQL queries is based on relational algebra.

Relational algebra allows us to describe our algorithms and later use SQL encodings for specifying the table algorithm. The intermediate step of stating the algorithm in a relation algebra description is twofold. First, we can immediately see the connection between the algorithms given in the literature, which allows us to use the existing algorithms without reproving all properties. Second, we obtain a compact mathematical description, which is not just a lengthy and technical SQL query that might be hard to understand to researchers from the community who are usually not very familiar with practical databases and the usage of query languages.
Before we start with details on our approach, we briefly recall basics in relational algebra. The classical relational algebra was introduced by Codd (1970) as a mathematical framework for manipulating relations (tables). Since then, relational algebra serves as the formal background and theoretical basis in relational databases and their standard language SQL (Structured Query Language) for querying tables (Ullman 1989). In fact, in the following, we need extended constructs, which have not been defined in the original framework by Codd, but are standard notions in databases nowadays (Elmasri and Navathe 2016). For the understanding later, we would like to mention that the SQL table model and relational algebra model slightly differ. The SQL table model is a bag (multiset) model, rather than a set (Garcia-Molina et al. 2009, Chapter 5). Below we also use extended projection and aggregation by grouping. Sometimes these are defined on bags. We avoid this in the definitions in order to keep the algorithms close to the formal set based notation. Finally, we would like to emphasize that we are not using relation algebra here as developed by Alfred Tarski for the field of abstract algebra, but really relational algebra as used in database applications and theory.

A column a is of a certain finite domain dom(a). Then, a row r over set cols(r) of columns is a set of pairs of the form (a, v) with a ∈ cols(r), v ∈ dom(a) such that for each a ∈ cols(r), there is exactly one v ∈ dom(a) with (a, v) ∈ r. In order to access the value v of an attribute a in a row r, we sometimes write r.a, which returns the unique value v with (a, v) ∈ r. A table τ is a finite set of rows r over set cols(τ) := cols(r) of columns, using domain dom(τ) := ∪a∈cols(τ) dom(a). We define renaming of τ, given a set A of columns and a bijective mapping m : cols(τ) → A with dom(a) = dom(m(a)) for a ∈ cols(τ), by ρm(τ) := {(m(a), v) | (a, v) ∈ τ}. In SQL, renaming can be achieved by means of the AS keyword.

Selection of rows in τ according to a given equality formula φ over term variables cols(τ) is defined[1] by σφ(τ) := {r | r ∈ τ, φ(ass(r)) = 0}, where function ass provides the corresponding term assignment of a given row r ∈ τ. Selection in SQL is specified using keyword WHERE. Given a relation τ′ with cols(τ′) ∩ cols(τ) = ∅. Then, we refer to the cross-join by τ × τ′ := {r ∪ τ′′ | r ∈ τ, τ′′ ∈ τ′}. Further, a θ-join (according to φ) corresponds to τ = φ τ := σφ(τ × τ′). Interestingly, in SQL a θ-join can be achieved by specifying the two tables (cross-join) and adding the selection according to φ by means of WHERE.

Assume in the following a set A ⊆ cols(τ) of columns. Then, we let table τ projected to A be given by ΠA(τ) := {ra | r ∈ τ}, where ra := {(a, v) | (a, v) ∈ r, a ∈ A}. This concept of projection can be lifted to extended projection ΠA,S, where we assume in addition to A, a set S of expressions of the form a ← f, such that a ∈ cols(τ) \ A. f is an arithmetic function that takes a row r ∈ τ, and there is at most one such expression for each a ∈ cols(τ) \ A in S. Formally, we define ΠA,S(τ) := {ra ∪ rS | r ∈ τ} with rS := {(a, f(r)) | a ∈ cols(r), (a ← f) ∈ S}. SQL allows to specify (extended) projection directly after initiating an SQL query with the keyword SELECT.

Later, we use aggregation by grouping AG(a,g), where we assume a ∈ cols(τ) \ A and a so-called aggregate function g : 2A → dom(a), which intuitively takes a table of (grouped) rows. Therefore, we let AG(a,g)(τ) := {r ∪ {{a,g(τ[r])}} | r ∈ ΠA(τ)}, where τ[r] := {r′ | r′ ∈ τ, r ⊆ r′}. For this purpose, we use for a set S ⊆ S of integers, the functions SUM for summing up values in S, MIN for providing the smallest integer in S, as well as MAX for obtaining the largest integer in S, which are often used for aggregation in this context. The SQL standard uses projection (SELECT)
to specify \( A \) as well as the aggregate function \( g \), such that these two parts are distinguished by means of the keyword \texttt{GROUP BY}.

\textbf{Example 3}

Assume a table \( \tau_1 := \{r_1, r_2, r_3\} \) of 2 columns \( a, b \) over Boolean domain \( \text{dom}(a) = \text{dom}(b) = \{0, 1\} \), where \( r_1 := \{(a, 1), (b, 1)\} \), \( r_2 := \{(a, 0), (b, 0)\} \), \( r_3 := \{(a, 0), (b, 1)\} \).

\[
\begin{array}{c|cc}
\tau_1 & a & b \\
--- & --- & --- \\
r_1 & 1 & 1 \\
r_2 & 0 & 0 \\
r_3 & 0 & 1 \\
\end{array}
\quad
\begin{array}{c|cc}
\tau_2 & b & a \\
--- & --- & --- \\
r_1 & 1 & 1 \\
r_2 & 0 & 0 \\
r_3 & 1 & 0 \\
\end{array}
\]

Then, \( r_3.a = 0 \) and \( r_3.b = 1 \). Rows can be swapped by renaming and we let \( \tau_2 := \rho_{\{a \leftrightarrow b, b = a\}}(\tau_1) \). Observe that, e.g., \( \rho_{\{a \leftrightarrow b, b = a\}}(\{r_3\}) \) corresponds to \( \{(a, 1), (b, 0)\} \), i.e., considering \( r_3 \) and swapping \( a \) and \( b \). We select rows by using the selection \( \sigma \). For example, if we want to select rows where \( b = 1 \) (colored in blue) we can use \( \sigma_{b = 1}(\tau_1) \).

\[
\begin{array}{c|cc}
\tau_1 & a & b \\
--- & --- & --- \\
r_1 & 1 & 1 \\
r_2 & 0 & 0 \\
r_3 & 0 & 1 \\
\end{array}
\]

Hence, applying \( \sigma_{b = 1}(\tau_1) \) results in \( \{r_1, r_3\} \). Table \( \tau_1 \) can be \( \theta \)-joined with \( \tau_2 \), but before, we need to have disjoint columns, which we obtain by renaming each column \( c \) to a fresh column \( c' \) as below by \( \rho_{\{a \leftrightarrow a', b = b'\}}(\tau_2) \). Then, \( \tau_3 := \tau_1 \bowtie_{a = a', b = b'}(\rho_{\{a = a', b = b'\}}(\tau_2)) \).

\[
\begin{array}{c|cc}
\tau_1 & a & b \\
--- & --- & --- \\
r_1 & 1 & 1 \\
r_2 & 0 & 0 \\
r_3 & 0 & 1 \\
\end{array}
\quad
\begin{array}{c|cc|cc}
\rho_{a = a', b = b'}(\tau_2) & b' & a' \\
--- & --- & --- \\
r_1 & 1 & 1 \\
r_2 & 0 & 0 \\
r_3 & 1 & 0 \\
\end{array}
\quad
\begin{array}{c|ccc}
\tau_3 & a & b & b' & a' \\
--- & --- & --- & --- & --- \\
r_1 & 1 & 1 & 1 & 1 \\
r_2 & 0 & 0 & 0 & 0 \\
\end{array}
\]

Consequently, we have \( \tau_3 = \{(a, 0), (a', 0), (b, 0), (b', 0)\}, \{(a, 1), (a', 1), (b, 1), (b', 1)\}\}. Extended projection allows not only to filter certain columns, but also to add additional columns. As a result, if we only select column \( a \) of each row of \( \tau_1 \), but add a fresh column \( c \) holding the sum of the values for \( a \) and \( b \), then \( \Pi_{\{a, \{c \leftarrow a+b\}\}}(\tau_1) \) corresponds to \( \{(a, 1), (c, 2)\}, \{(a, 0), (c, 0)\}, \{(a, 0), (c, 1)\}\).

\[
\begin{array}{c|cc}
\Pi_{\{a, \{c \leftarrow a+b\}\}}(\tau_1) & a & c \\
--- & --- & --- \\
r_1 & 1 & 2 \\
r_2 & 0 & 0 \\
r_3 & 0 & 1 \\
\end{array}
\]

Grouping \( \tau_1 \) according to the value of column \( a \), where we aggregate each group by summing
up the values of columns $b$ in a fresh column $d$, results in \( \{a\}G_d \leftarrow \tau \mapsto \text{SUM}(\{r.b| r \in \tau\})(\tau_1) \), which simplifies to \( \{(a, 1), (d, 1)\}, \{(a, 0), (d, 1)\} \) as illustrated below.

\[
\begin{array}{c|cc}
\tau_1 & a & b \\
\hline
r_1 & 1 & 1 \\
r_2 & 0 & 0 \\
r_3 & 0 & 1 \\
\end{array}
\]

\[
\begin{array}{c|cc}
\{a\}G_d \leftarrow \tau \mapsto \text{SUM}(\{r.b| r \in \tau\})(\tau_1) & a & d \\
\hline
r_1 & 1 & 1 \\
r_3 & 0 & 1 \\
\end{array}
\]

### 4.2 Table Algorithms in Relational Algebra

Now, we are in the position to use relational algebra instead of set theory based notions to describe how tables are obtained during dynamic programming. The step from set notation to relational algebra is driven by the observation that in these table algorithms one can identify recurring patterns and one mainly has to adjust problem-specific parts of it. We continue the description with our problem \#SAT. We picture tables $\tau$ for each TD node $t$ as relations, where $\tau$ distinguishes a unique column $x$ for each $x \in \chi(t)$. In addition, we require a column $cnt$ for counting in \#SAT, or a column for modeling costs or weights in case of optimization problems.

Algorithm 2 presents a table algorithm for problem \#SAT that is equivalent to Algorithm 1 but relies on relational algebra for computing tables. Since our description in relation algebra yields the same results as the set based-notation above, we omit formal correctness proofs. Nonetheless, we briefly explain below why both notations are identical. We highlight the crucial parts by coloring Algorithm 1. In particular, one typically derives for nodes $t$ with $\text{type}(t) = \text{leaf}$, a fresh initial table $\tau$, cf., Line 1 of Algorithm 2. Then, whenever a variable $a$ is introduced, such algorithms often use $\theta$-joins with a fresh initial table for the introduced variable $a$. Hence, the new column represents the potential values for variable $a$. In Line 5 the selection of the $\theta$-join is performed according to $\varphi$, i.e., corresponding to the local instance of \#SAT. Further, for nodes $t$ with $\text{type}(t) = \text{rem}$, these table algorithms typically need projection. In case of Algorithm 2 Line 5 also needs grouping to sum up the counters for those rows of $\tau_1$ that concur in $\tau$. Thereby, rows are grouped according to values of columns $\chi(t)$ and we keep only one row per group in table $\tau$, where the fresh counter $cnt$ is the sum among all counters in $\tau$. Finally, in Line 7 for a node $t$ with $\text{type}(t) = \text{join}$, we use extended projection and $\theta$-joins, where we join on the same truth assignments. This allows us later to leverage database technology for a usually expensive

**Algorithm 2:** Table algorithm Sat\(^t\)(\(t, \chi(t), \varphi, (\tau_1, \ldots, \tau_1)\)) for solving \#SAT.

| In: Node $t$, bag $\chi(t)$, local formula $\varphi$, sequence $(\tau_1, \ldots, \tau_1)$ of child tables. |
| Out: Table $\tau_1$. |
| 1 if $\text{type}(t) = \text{leaf}$ then $\tau_1 := \{(\text{cnt}, 1)\}$ |
| 2 else if $\text{type}(t) = \text{intr}$, and $a \in \chi(t)$ is introduced then |
| 3 | $\tau_1 := \tau_1 \bowtie \varphi \{(a, 0)\}, \{(a, 1)\}$ |
| 4 else if $\text{type}(t) = \text{rem}$, and $a \notin \chi(t)$ is removed then |
| 5 | $\tau_1 := \chi(t)G_{\text{cnt}} \leftarrow \tau \mapsto \text{SUM}(\{r.\text{cnt}| r \in \tau\})(\pi_{\text{cols}(\tau_1)}(\tau_1)$ |
| 6 else if $\text{type}(t) = \text{join}$ then |
| 7 | $\tau_1 := \pi_{\chi(t), \{\text{cnt} \bowtie \text{cnt} \text{cnt}\}, \chi(t)}(\tau_1 \bowtie_{\varphi_{\text{icol}}(t_1)} w = w' \{w \rightarrow w'\} \tau_2)$ |
| 8 return $\tau_1$. |
operation. Extended projection is needed for multiplying the counters of the two rows containing the same assignment.

4.3 Table algorithms for selected problems

Dynamic programming algorithms are known for a variety of problems. Standard texts in the area of parameterized algorithms and introductory lectures provide various specifications. For formal properties and detailed algorithm descriptions, we refer to other works ([Bodlaender 1988], [Cygan et al. 2015, Chapter 7], [Dechter 1999], [Bannach and Berndt 2019]). Below, we present the table algorithms for a selection of combinatorial problems in relational algebra notation. In order to simplify the presentation, we assume that the instance is given by \( \mathcal{I} \) and that the used tree decomposition is nice and given by \( \mathcal{I} = (T, \chi) \). If the problem is a graph problem \( \mathcal{I} \) is a TD of \( \mathcal{I} \), otherwise we implicitly assume that \( \mathcal{I} \) is a TD of the primal graph of instance \( \mathcal{I} \). For graph problems \( \mathcal{I} \) and a given node \( t \) of \( T \), we refer to the local instance of \( \mathcal{I} = G = (V, E) \) by local graph \( G_t \) and define it by \( G_t := (V \cap \chi(t), E \cap [\chi(t) \times \chi(t)]) \).

**Problem \#o-COL**

| Algorithm 3: Table algorithm Col\((t, \chi(t), G_t, \{\tau_1, \ldots, \tau_i\})\) for solving \#o-COL. |
|---|
| **In:** Node \( t \), bag \( \chi(t) \), local graph \( G_t \), and a sequence \( \{\tau_1, \ldots, \tau_i\} \) of child tables. |
| **Out:** Table \( \tau_i \). |
| 1 if type\((t) = \text{leaf} \) then \( \tau_i := \{(\text{cnt}, 1)\} \) |
| 2 else if type\((t) = \text{intr}, \) and \( a \in \chi(t) \) is introduced then |
| 3 \( \tau_i := \tau_1 \bowtie \Delta_{\chi(a) \in \chi(t), \uin E, \uin \notin \chi(t)} \{(\{a, 0\}), \{(a, 1)\}, \ldots, \{(a, o)\}\} \) |
| 4 else if type\((t) = \text{rem}, \) and \( a \notin \chi(t) \) is removed then |
| 5 \( \tau_i := \chi(t)\bigcup_{\text{cnt} \in \chi(t)} (\Pi_{\text{viol}}(\tau_i), (a) \tau_i) \) |
| 6 else if type\((t) = \text{join} \) then |
| 7 \( \tau_i := \Pi_{\chi(t)} (\chi(t) \bowtie \text{cnt} \text{ cnt}) \bigcup_{\text{cnt} \in \text{cnt} \text{ cnt}} (\tau_1 \bowtie \chi(a) \in \chi(t), \uin E, w = a) \bigcup_{\text{cnt} \in \text{cnt} \text{ cnt}} (w \neq a\tau_2) \) |
| 8 return \( \tau_i \) |

For a given graph instance \( \mathcal{I} = G = (V, E) \), an o-coloring is a mapping \( t : V \to \{1, \ldots, o\} \) such that for each edge \( \{u, v\} \in E \), we have \( t(u) \neq t(v) \). Then, the problem \#o-COL asks to count the number of o-colorings of \( G \), whose local instance \#o-COL\((t, G)\) is the local graph \( G_t \). The table algorithm for this problem \#o-COL is given in Algorithm 3. Similarly to Algorithm 2 for (empty) leaf nodes, the counter \text{cnt} is set to 1 in Line 1. Whenever a vertex \( a \) is introduced, in Line 3, one of the \( o \) many color values for \( a \) are guessed and \( \theta \)-joined with the table \( \tau_1 \) for the child node of \( t \) such that only colorings with different values for two adjacent vertices are kept. Similarly to Algorithm 2, whenever a vertex \( a \) is removed, Line 5 ensures that the column for \( a \) is removed and that counters \text{cnt} are summed up for rows that concur due to the removal of column \( a \). Then, the case for join nodes in Line 7 is again analogous to Algorithm 2 where only rows with the same colorings in both child tables are kept and counters \text{cnt} are multiplied accordingly.

**Problem MINVC**

Given a graph instance \( \mathcal{I} = G = (V, E) \), a vertex cover is a set of vertices \( C \subseteq V \) of \( G \) such that for each edge \( \{u, v\} \in E \), we have \( \{u, v\} \cap C \neq \emptyset \). Then, MINVC asks to find the minimum
Algorithm 4: Table algorithm $\text{VC}(t, \chi(t), G_t, \langle \tau_1, \ldots, \tau_l \rangle)$ for solving M\text{INVC}.

**In:** Node $t$, bag $\chi(t)$, local graph $G_t$, and a sequence $\langle \tau_1, \ldots, \tau_l \rangle$ of child tables.

**Out:** Table $\tau_t$.

1. if type$(t) = \text{leaf}$ then $\tau_t := \{ (\text{card}, 0) \}$
2. else if type$(t) = \text{intr}$, and $a \in \chi(t)$ is introduced then
   3. $\tau_t := \tau_t \bowtie_{\chi(a) \in G_t, u' \in t} \{ (a, 0), (a, 1) \}$
3. else if type$(t) = \text{rem}$, and $a \notin \chi(t)$ is removed then
   5. $\tau_t := \chi(t) G_{\text{card} \leftarrow t \rightarrow \text{MIN} \{ (\text{card} + r, a) \in t \}} (\Pi_{\text{cols}(\tau_t)} \setminus \{a\} \tau_t)\$
6. else if type$(t) = \text{join}$ then
   7. $\tau_t := \Pi^\chi(t), \{ \text{card} = \text{card}' \} (\tau_t \bowtie_{\chi(a) \in G_t, u' \in t} \rho \setminus \{m \rightarrow u'\} \tau_t)\$
8. return $\tau_t$.

Cardinality $|C|$ among all vertex covers $C$, i.e., $C$ is such that there is no vertex cover $C'$ with $|C'| < |C|$. Local instance M\text{INVC}$(t, G) := G_t$, where the local graph $G_t$ is defined above. We use an additional column card for storing cardinalities. The table algorithm for solving M\text{INVC} is provided in Algorithm 4, where, for leaf nodes the cardinality is 0, cf., Line 1. Then, when introducing vertex $a$, we guess in Line 3 whether $a$ shall be in the vertex cover or not, and enforce that for each edge of the local instance at least one of the two endpoint vertices has to be in the vertex cover. Note that the additional cardinality column only takes removed vertices into account. More precisely, when a vertex $a$ is removed, we group in Line 5 according to the bag columns $\chi(t)$, where the fresh cardinality value is the minimum cardinality (plus 1 for $a$ if $a$ shall be in the vertex cover), among those rows that concure due to the removal of $a$. The join node is similar to before, but in Line 7 we additionally need to sum up the cardinalities of two adjoining child table rows.

**Problem MAXSAT**

Algorithm 5: Table algorithm $\text{MSat}(t, \chi(t), \mathcal{F}_t, \langle \tau_1, \ldots, \tau_l \rangle)$ for solving MAXSAT.

**In:** Node $t$, bag $\chi(t)$, local instance $\mathcal{F}_t = (\emptyset, \psi_t)$, and a sequence $\langle \tau_1, \ldots, \tau_l \rangle$ of child tables.

**Out:** Table $\tau_t$.

1. if type$(t) = \text{leaf}$ then $\tau_t := \{ (\emptyset, 0) \}$
2. else if type$(t) = \text{intr}$, and $a \in \chi(t)$ is introduced then
   3. $\tau_t := \tau_t \bowtie_{\chi(a) \in \emptyset, \psi_t, \{a\}} \{ (a, 0), (a, 1) \}$
3. else if type$(t) = \text{rem}$, $a \notin \chi(t)$ is removed, and $\psi' \text{ are removed local soft-clauses then}$
   5. $\tau_t := \chi(t) G_{\text{card} \leftarrow t \rightarrow \text{MAX} \{ (\text{card} + \Sigma_{\psi' \in \text{soft-clauses}} u', a) \in t \}} (\Pi_{\text{cols}(\tau_t)} \setminus \{a\} \tau_t)\$
6. else if type$(t) = \text{join}$ then
   7. $\tau_t := \Pi^\chi(t), \{ \text{card} = \text{card}' \} (\tau_t \bowtie_{\chi(a) \in \emptyset, \psi_t, \{a\}} \rho \setminus \{m \rightarrow u'\} \tau_t)\$
8. return $\tau_t$.

Given Boolean formulas $\varphi$ and $\psi$, an instance of problem MAXSAT is of the form $\mathcal{F} = (\varphi, \psi)$ and we assume that $\mathcal{F}$ is a TD of primal graph $G_{\varphi, \psi}$. A solution to MAXSAT is a satisfying assignment $I$ of hard-clauses $\varphi$ such that $|\{c \mid c \in \psi, c(I) = \emptyset\}|$ is maximized, i.e., $I$ is a satisfying assignment of $\varphi$ that satisfies the maximum number of soft-clauses $\psi$ among all satisfying assignments of $\varphi$. We define the local instance $\mathcal{F}_t := (\varphi, \psi_t)$ consisting of local formula $\varphi$, referred to by local hard-clauses and local formula $\psi_t$, called local soft-clauses.

The table algorithm for problem MAXSAT is given in Algorithm 5, where we use column card...
for holding satisfied soft-clauses. Leaf tables only hold a cardinality value of 0 as in Line 4. Then, similar to the table algorithm Sat’ (cf., Algorithm 2), when introducing a variable a, we guess the truth value and keep those rows, where local formula $\varphi_t$ is satisfied. Whenever a variable a is removed in a node t, we remove column a and group rows that have common values over columns $\chi(t)$. Thereby, the new cardinality $\text{card}$ for each group is the maximum among the values of $\text{card}$ including the number of satisfied local soft-clauses $\psi'$ of the child node of t that are removed in $\psi_t$ (due to removal of a). Finally, similar to Algorithm 4, a join node sums up cardinalities of two child rows containing the same assignment.

**Problem MINIDS**

Algorithm 6: Table algorithm $\text{IDS}(t, \chi(t), G_t, \langle \tau_1, \ldots, \tau_l \rangle)$ for solving MINIDS.

In: Node $t$, bag $\chi(t)$, local graph $G_t$, and a sequence $\langle \tau_1, \ldots, \tau_l \rangle$ of child tables.

Out: Table $\tau_t$.

1. **if** type$(t)$ = leaf **then** $\tau_t := \{\langle \text{card}, 0 \rangle\}$
2. **else if** type$(t)$ = intr, and $a \in \chi(t)$ is introduced **then**
   
   $\tau_t := \Pi_{\chi(t)}(\cup_{u \in G_t}(d_u \cup \{1 \text{ or } 0 \text{ if } a \in \chi(u) \}) \cup \text{card} \cup \text{card}^d) \{t_1 \Rightarrow (a, 0, 0), \{(a, 1), (d_u, 1)\}\}$

3. **else if** type$(t)$ = rem, and $a \notin \chi(t)$ is removed **then**
   
   $\tau_t := \chi(t) \cup \text{card} \cup \text{card}^d \cup \text{card} \cup \text{card}^d \{t_1 \Rightarrow (a, 0, 0), \{(a, 1), (d_u, 1)\}\}$

4. **else if** type$(t) = \text{join then}$
   
   $\tau_t := \Pi_{\chi(t)}(\cup_{u \in G_t}(d_u \cup \text{card} \cup \text{card}^d) \{t_1 \Rightarrow (a, 0, 0), \{(a, 1), (d_u, 1)\}\}$

5. **return** $\tau_t$

Given a graph instance $G = (V, E)$, a dominating set of $G$ is a set of vertices $D \subseteq V$ of $G$, where each vertex $v \in V$ is either in $D$ or is adjacent some vertex in $D$, i.e., there is a vertex $d \in D$ with $\{d, v\} \in E$. A dominating set $D$ is an independent dominating set of $G$, if there is no edge in $E$ between vertices of $D$. Then, the problem MINIDS asks to find the minimum cardinality $|D|$ among all independent dominating sets $D$ of $G$ (if exists). We define local instance by $\text{MINIDS}(t, G_t) := G_t$.

The table algorithm for solving MINIDS is given in Algorithm 6 where a table $\tau_t$ of a node $t$ uses column $\text{card}$ for cardinalities of potential dominating sets, and an additional Boolean column $d_u$ per bag vertex $u \in \chi(t)$. Intuitively, $d_u$ indicates whether vertex $u$ is already “dominated”, i.e., either $u$ is in the dominating set or $u$ has an adjacent vertex to the dominating set. Similar to before, leaf nodes set cardinality $\text{card}$ to 0, cf., Line 4. For a node $t$ with an introduced vertex $a$, we guess in Line 6 whether $a$ shall be in the dominating set or not (and set $d_a$ to 0 or 1, respectively).

Then, we only keep rows that are independent, i.e., a can not be in the dominating set and adjacent to $u$ in edges $E(G_t)$ of local graph $G_t$ at the same time. Finally, values $d_u$ (dominance status) for $a$ and for neighbors of $a$ are updated accordingly. When a vertex $a$ is removed in a node $t$, Line 5 only keeps rows, where $d_a$ is true, i.e., $a$ is indeed dominated, and removes columns $a, d_u$. Further, we group rows according to their values to $\chi(t) \cup \{d_u \mid u \in \chi(t)\}$ and for each group we set the cardinality to the minimum among the cardinalities of the group rows (including $a$ if $a$ is in the set). For join nodes $t$, Line 7 sums up cardinalities of rows holding the same dominating set and treats a vertex $u \in \chi(t)$ as dominated if it is dominated in at least one of the two rows.

Similar to MINVC and #o-Col one can model several other (graph) problems. One could also think of counting the number of solutions to problem MINVC, where both a column for
cardinalities and one for counting is used. There, in addition to grouping, we additionally could use conditions over groups where only rows are kept whose column values for \( \text{card} \) form the minimum within the group.

### 4.4 Generalizing the Patterns of Table Algorithms

In the previous sections, we presented the table algorithms for solving a selection of combinatorial problems, namely, \texttt{SAT}, \texttt{o-COL}, \texttt{MINVC}, \texttt{MAXSAT}, and \texttt{MINIDS}, by dynamic programming. As mentioned in Section 2.2, there are a variety of application areas where such algorithms allow for solving problems efficiently. When specifying most algorithms, we focus on the table algorithm \( A \), which is executed for each node \( t \) of \( T \) of the considered tree decomposition \( \mathcal{T} = (T, \chi) \) and computes a new table depending on the previously computed tables at the children of \( t \). From the descriptions above, it is easy to see that the algorithms effectively follow standard patterns. Therefore, we present a general template in Algorithm 7, where parts of table algorithms for problems that are typically problem-specific are replaced by colored placeholders of the form \#placeHolder\#, cf., Algorithm 2. The general template of table algorithms works for many problems, including decision problems, counting problems as well as optimization problems.

The intuition behind these placeholders is as follows: For leaf nodes, the initial table (typically empty) can be specified using \#leafTab\#. For introduce nodes, the potential cases for the introduced vertex \( a \) are given with the help of \#intrTab\#. Then, according to the local instance, we only keep those rows that satisfy \#intrFilter\#. The placeholder \#intrAddCols\# allows to add additional columns, which we often need when solving problems that involve counting or optimizing a value. In other words, placeholder \#intrAddCols\# in Line 3 of Algorithm 7 uses extended projection, which is needed for problems requiring changes on vertex introduction. Nodes, where an atom \( a \) is removed sometimes require to filter rows, which do not lead to a solution using \#remFilter\#, and to remove columns concerning \( a \) by \#remCols\#. Further, one oftentimes needs to aggregate rows according to the values of the columns of the bag and additional columns (given by \#remGroupCols\#), where the aggregation is specified by \#remAggr\#. Finally, for join nodes, one can specify an additional filter \#joinAddFilter\# that goes beyond checking equivalence of row values in the \( \theta \)-join operation. Further, depending on the problem one might need to add and update the values of additional columns by using extended projection in form of placeholder \#joinAddCols\#.

Note that while the algorithms presented here assume for simplicity nice tree decompositions, the whole architecture does not depend on certain restrictions of TDs, or whether it is nice or not. Instead, a table algorithm of any TD is simply specified by handling \textit{problem-specific}
Figure 3: Architecture of Dynamic Programming with Databases. Steps highlighted in red are provided by the system depending on the specification of yellow and blue parts, which is given by the user for specific problems $P$. The yellow “E”s represent events that can be intercepted and handled by the user. The blue part concentrates on table algorithm $A$, where the user specifies how SQL code is generated in a modular way.

Implementations of the placeholders of Algorithm 7, where the system following this architecture is responsible for interleaving and overlapping these cases within a node $t$. In fact, we discuss an implementation of a system according to this architecture next, where it is crucial to implement non-nice TDs to obtain higher efficiency.

5 System dpdb: Dynamic Programming with Databases & SQL

In this section, we present a general architecture to model table algorithms by means of database management systems. We move from relational algebra definitions to specifications of the table algorithms in terms of SQL queries. The overall architecture is follows the DP approach as presented in Section 3. It works as depicted in Figure 3, where the steps highlighted in yellow and blue need to be specified depending on the problem $P$. Steps outside Step 3 are mainly setup tasks, the yellow “E”s indicate events that might be needed to solve more complex problems on the polynomial hierarchy. For example, one could create and drop auxiliary sub-tables for each node during Step 3 within such events. Observe that after the generation of a TD $T = (T, \chi)$, Step 2b automatically creates tables $\tau_t$ for each node $t$ of $T$, where the corresponding table columns of $\tau_t$ are specified in the blue part, i.e., within $A$. The default columns of such a table $\tau_t$ that are assumed in this section foresee one column for each element of the bag $\chi(t)$, where additional columns that are needed for solving the problem can be added. This includes additional auxiliary columns, which can be also counters or costs for counting or optimization, respectively. Besides the definition of table schemes, the blue part concerns the specification of the table algorithm by means of a procedural generator template that describes how to obtain SQL code for each node $t$, thereby depending on $\chi(t)$ and on the tables for child nodes of $t$. This generated SQL code is then used internally for manipulation of tables $\tau_t$ during the tree decomposition traversal in Step 3 of dynamic programming.

We implemented the proposed architecture of the previous section in the prototypical system dpdb. The system is open-source written in Python 3, and uses PostgreSQL as DBMS. We are certain that one can easily replace PostgreSQL by any other state-of-the-art relational database that uses SQL. In the following, we discuss implementation specifics that are crucial for a performant system that is still extendable and flexible.

Our system dpdb is available under GPL3 license at [github.com/hmarkus/dp_on_dbs](http://github.com/hmarkus/dp_on_dbs).
Computing TDs. TDs are computed mainly with the library htd version 1.2 with default settings (Abseher et al. 2017), which finds TDs extremely quick also for interesting instances (Fichte et al. 2019) due to heuristics. Note that dpdb directly supports the TD format of recent competitions (Dell et al. 2018), i.e., one could easily replace the TD library. It is important not to enforce htd to compute nice TDs, as this would cause a lot of overhead later in dpdb for copying tables. However, in order to benefit from the implementation of \( \theta \)-joins, query optimization, and state-of-the-art database technology in general, we observed that it is crucial to limit the number of child nodes of every TD node. In result, when huge tables are involved, \( \theta \)-joins among child node tables cover at most a limited number of child node tables. Hence, the query optimizer of the database system can still come up with meaningful execution plans depending on the contents of the table. Nonetheless we prefer \( \theta \)-joins with more than just two tables, since binary \( \theta \)-joins already fix in which order these tables shall be combined, which already limits the query optimizer. Apart from this trade-off, we tried to outsource the task of joining tables to the DBMS, since the performance of database systems highly depends on query optimization. The actual limit, which is a restriction from experience and practice only, highly depends on the DBMS that is used. For PostgreSQL, we set a limit of at most 5 child nodes for each node of the TD, i.e., each \( \theta \)-join covers at most 5 child tables.

Towards non-nice TDs. Although this paper presents the algorithms for nice TDs (mainly due to simplicity), the system dpdb interleaves these cases as presented in Algorithm 7. More precisely, the system executes one query per table \( \tau \) for each node \( t \) during the traversal of TD \( T \). This query consists of several parts and we briefly explain its parts from outside to inside in accordance with Algorithm 7. First of all, the inner-most part concerns the row candidates for \( \tau \) consisting of the \( \theta \)-join among all child tables of \( \tau \) as in Line 7 of Algorithm 7. If there is no child node of \( t \), table \#leafTab\# of Line 1 is used instead. Next, the result is cross-joined with \#intrTab\# for each introduced variable as in Line 3, but without using the filter \#intrFilter\# yet. Then, the result is projected by using extended projection involving \( \chi(t) \) as well as both \#joinAddCols\# and \#intrAddCols\#. Actually, there are different configurations of how dpdb can deal with the resulting row candidates. For debugging (see below) one could (1) actually materialize the result in a table, whereas for performance reasons, one should use (2) common table expressions (CTEs or \WITH\-queries) or (3) sub-queries (nested queries), which both result in one nested SQL query per table \( \tau \). On top of these row candidates, selection according to \#intrFilter\#, cf., Line 3, is executed. Finally, the resulting table is plugged as table \( \tau \) into Line 5 where in particular the result is grouped by using both \( \chi(t) \) and \#remGroupCols\# and each group is aggregated by \#remAggr\# accordingly. It turns out that PostgreSQL can do better with sub-queries than CTEs, since we observed that the query optimizer oftentimes pushes (parts of) outer selections and projections into the sub-query if needed, which is not the case for CTEs, as discussed in the PostgreSQL manual (PostgreSQL Global Development Group 2020, Sec. 7.8.1). On different DBMSs or other vendors, e.g., Oracle, it might be better to use CTEs instead.

Example 4
Consider again Example 2 and Figure 1 and let us use table algorithm Sat’ with dpdb on formula \( \phi \) of TD \( T \) and Option (3): sub-queries, where the row candidates are expressed via a sub-queries. Then, for each node \( t_i \) of \( T \), dpdb generates a view \( vi \) as well as a table \( \tau_i \) containing in the end

5 Actually, dpdb keeps only columns relevant for the table of the parent node of \( t \).
the content of \( v_i \). Observe that each view only has one column \( a \) for each variable \( a \) of \( \phi \) since the truth assignments of the other variables are not needed later. This keeps the tables compact, only \( \tau_1 \) has two rows, \( \tau_2 \), and \( \tau_3 \) have only one row. We obtain the following views.

CREATE VIEW v1 AS SELECT a, sum(cnt) AS cnt FROM
(WITH intrTab AS (SELECT 0 AS val UNION ALL SELECT 1)
SELECT i1.val AS a, i2.val AS b, i3.val AS c, 1 AS cnt
FROM intrTab i1, intrTab i2, intrTab i3)
WHERE (NOT a OR b OR c) AND (a OR NOT b OR NOT c) GROUP BY a

CREATE VIEW v2 AS SELECT a, sum(cnt) AS cnt FROM
(WITH intrTab AS (SELECT 0 AS val UNION ALL SELECT 1)
SELECT i1.val AS a, i2.val AS d, 1 AS cnt FROM intrTab i1, intrTab i2)
WHERE (a OR d) AND (a OR NOT d) GROUP BY a

CREATE VIEW v3 AS SELECT a, sum(cnt) AS cnt FROM
(SELECT \( \tau_1 \).a, \( \tau_1 \).cnt * \( \tau_2 \).cnt AS cnt FROM \( \tau_1 \), \( \tau_2 \) WHERE \( \tau_1 \).a = \( \tau_2 \).a)
GROUP BY a

**Parallelization.** A further reason to not over-restrict the number of child nodes within the TD, lies in parallelization. In \( dpdb \), we compute tables in parallel along the TD, where multiple tables can be computed at the same time, as long as the child tables are computed. Therefore, we tried to keep the number of child nodes in the TD as high as possible. In our system \( dpdb \), we currently allow for at most 24 worker threads for table computations and 24 database connections at the same time (both pooled and configurable). On top of that we have 2 additional threads and database connections for job assignments to workers, as well as one dedicated watcher thread for clean-up and connection termination, respectively.

**Logging, Debugging and Extensions.** Currently, we have two versions of the \( dpdb \) system implemented. One version aims for performance and the other one tries to achieve comprehensive logging and easy debugging of problem (instances), thereby increasing explainability. The former does neither keep intermediate results nor create database tables in advance (Step 2b), as depicted in Figure 3, but creates tables according to an SQL SELECT statement. In the latter, we keep all intermediate results, we record database timestamps before and after certain nodes, provide statistics as, e.g., width and number of rows. Further, since for each table \( \tau_i \), exactly one SQL statement is executed for filling this table, we also have a dedicated view of the SQL SELECT statement, whose result is then inserted in \( \tau_i \). Together with the power and flexibility of SQL queries, we observed that this helps in finding errors in the table algorithm specifications.

Besides convenient debugging, system \( dpdb \) immediately contains an extension for **approximation**. There, we restrict the table contents to a maximum number of rows. This allows for certain approximations on counting problems or optimization problems, where it is infeasible to compute the full tables. Further, \( dpdb \) foresees a dedicated **randomization** on these restricted number of rows such that we obtain different approximate results on different random seeds.

Note that \( dpdb \) can be easily extended. Each problem can overwrite existing default behavior and \( dpdb \) also supports problem-specific argument parsers for each problem individually. Out-of-the-box, we support the formats DIMACS SAT and DIMACS graph \( (Liu et al. 2006) \) as well as the common format for TDs \( (Dell et al. 2018) \).
Implementing table algorithms with dpdb for selected problems

The system dpdb allows for easy prototyping of DP algorithms on TDs. In the following, we present the relevant parts of table algorithm implementations according to the template in Algorithm 7 for our selection of problems below. More precisely, we give the SQL implementations of the table algorithms of the previous section in form of specifying the corresponding placeholders as given by the template algorithm $A$. Thereby, we only specify the placeholders needed for solving the problems, i.e., placeholders of template algorithm $A$ that are not used (empty) are left out. To this end, we assume in this section for each problem a not necessarily nice TD $\mathcal{F} = (T, \chi)$ of the corresponding graph representation of our given instance $\mathcal{I}$, as well as any node $t$ of $T$ and its child nodes $t_1, \ldots, t_\ell$.

**Problem #Sat.** Given instance formula $\mathcal{F} = \varphi$. Then, the specific placeholders of the template for #Sat for a node $t$ with $\varphi = \{l_1, \ldots, l_{k_1}\}, \ldots, \{l_n, \ldots, l_{k_n}\}$ that are required for dpdb to solve the problem are as follows.

- $\#\text{leafTab}#$: $\text{SELECT } 1 \text{ AS cnt}$
- $\#\text{intrTab}#$: $\text{SELECT } 0 \text{ AS val UNION ALL SELECT } 1$
- $\#\text{intrFilter}#$: $(l_{1,1} \lor \ldots \lor l_{1,k_1}) \land \ldots \land (l_{n,1} \lor \ldots \lor l_{n,k_n})$
- $\#\text{remAggr}#$: $\text{SUM}(\text{cnt}) \text{ AS cnt}$
- $\#\text{joinAddCols}#$: $\tau_1.\text{cnt} \land \ldots \land \tau_\ell.\text{cnt} \text{ AS cnt}$

If one compares this specification to Algorithm 2, one sees that conceptually the same idea is given above. However, for efficiency dpdb does not rely on nice TDs. Observe that for the plain decision problem SAT, where the goal is to decide only the existence of a satisfying assignment for given formula $\varphi$, placeholder $\#\text{leafTab}#$ would need to return the empty table and parts $\#\text{remAggr}#$ and $\#\text{joinAddCols}#$ are just empty since no counter cnt is needed.

**Problem #o-Col.** Recall the problem #o-Col and Algorithm 3. Let $\mathcal{F} = G = (V,E)$ be a given input graph. Then, specific implementation parts for #o-Col for a node $t$ with $E(G_t) = \{\{u_1, v_1\}, \ldots, \{u_n, v_n\}\}$ is given as follows.

- $\#\text{leafTab}#$: $\text{SELECT } 1 \text{ AS cnt}$
- $\#\text{intrTab}#$: $\text{SELECT } 0 \text{ AS val UNION ALL SELECT } 1$
- $\#\text{intrFilter}#$: $(u_1 \lor \ldots \lor u_n) \land \ldots \land (v_1 \lor \ldots \lor v_n)$
- $\#\text{remAggr}#$: $\text{SUM}(\text{cnt}) \text{ AS cnt}$
- $\#\text{joinAddCols}#$: $\tau_1.\text{cnt} \land \ldots \land \tau_\ell.\text{cnt} \text{ AS cnt}$

**Problem MinVC.** Given any input graph $\mathcal{F} = G = (V,E)$ of MinVC. Then, problem MinVC for a node $t$ with $E(G_t) = \{\{u_1, v_1\}, \ldots, \{u_n, v_n\}\}$ and removed vertices $\chi(t_1) \cup \ldots \cup \chi(t_\ell) = \{r_1, \ldots, r_m\}$ is specified by the following placeholders (cf., Algorithm 4).

- $\#\text{leafTab}#$: $\text{SELECT } 0 \text{ AS card}$
- $\#\text{intrTab}#$: $\text{SELECT } 0 \text{ AS val UNION ALL SELECT } 1$
- $\#\text{intrFilter}#$: $(u_1 \lor v_1) \land \ldots \land (u_n \lor v_n)$
- $\#\text{remAggr}#$: $\text{MIN}(\text{card} + r_1 + \ldots + r_m) \text{ AS card}$
- $\#\text{joinAddCols}#$: $\tau_1.\text{card} \land \ldots \land \tau_\ell.\text{card} \text{ AS card}$

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4 Prototypical implementations for problems #Sat as well as MinVC are readily available in dpdb.
Problem MAXSAT. Given an instance $\mathcal{I} = (\varphi, \psi)$ of problem MAXSAT. Then, the problem for a node $t$ with local hard clauses $\varphi_t = \{\{l_{1,1} \ldots l_{1,k_1}\}, \ldots, \{l_{n,1} \ldots l_{n,k_n}\}\}$ and local soft clauses $\psi_t = \{\{l'_{1,1} \ldots l'_{1,k'_1}\}, \ldots, \{l'_{p,1} \ldots l'_{p,k'_p}\}\}$ is specified by the following placeholders (cf., Algorithm 5).

- **#leafTab#:** SELECT 0 AS card
- **#intrTab#:** SELECT 0 AS val UNION ALL SELECT 1
- **#intrFilter#:** $(l_{1,1} OR ... OR l_{1,k_1}) \text{ AND ... AND } (l_{n,1} OR ... OR l_{n,k_n})$
- **#remAggr#:** MIN(card + $(l'_{1,1} OR ... OR l'_{1,k'_1}) + ...$
- **#joinAddCols#:** $\tau_1$.card + ... + $\tau_i$.card AS card

Problem MINIDS. Recall an instance $\mathcal{I} = G = (V,E)$ of problem MINIDS and table algorithm IDS as given in Algorithm 6. The implementation of table algorithm IDS for MINIDS for a node $t$ assumes that $E(G_t) = \{\{u_1, v_1\}, \ldots, \{u_n, v_n\}\}$. Further, we let bag $\chi(t) = \{a_1, \ldots, a_k\}$, removed vertices $\chi(t) \setminus (\chi(t_1) \cup \ldots \cup \chi(t_l)) = \{r_1, \ldots, r_m\}$, and we let the $w_i$ many neighbors $N_i$ of each vertex $a_i$ in $G$ (with $1 \leq i \leq k$) be given by $N_i = \{e_{a_i,1}, \ldots, e_{a_i,w_i}\}$. Then, the SQL implementation can be specified as follows.

- **#leafTab#:** SELECT 0 AS card
- **#intrTab#:** SELECT 0 AS val, 0 AS d UNION ALL SELECT 1, 1
- **#intrFilter#:** $(\text{NOT } u_1 \text{ OR NOT } v_1) \text{ AND ... AND } (\text{NOT } u_n \text{ OR NOT } v_n)$
- **#intrAddCols#:** card, $d_{a_1}$ OR $e_{a_1,1}$ OR ... OR $e_{a_1,w_1}$ AS $d_{a_1}$, ...
- **#remFilter#:** $d_{a_1}$ AND ... AND $d_{a_k}$
- **#remAggr#:** MIN(card + $r_1 + ... + r_m$) AS card
- **#remGroupCols#:** $d_{a_1}, \ldots, d_{a_k}$
- **#joinAddCols#:** $\tau_1$.card + ... + $\tau_i$.card AS card,
- **#leafTab#:** $\tau_1$.card + ... + $\tau_i$.card AS card

Note that implementations could generate and apply parts of this specification, as for example in #intrFilter# only edges that have not been checked so far in any descending node, need to be checked.

Similar to MINVC, #\(\alpha\)-COL, and MINIDS one can model several related (graph) problems. One could also think of counting the number of solutions to problem MAXSAT, where both, a column for cardinals and one for counting is used. There, in addition to grouping with GROUP BY in \(dpdb\), we additionally use the HAVING construct of SQL, where only rows are kept, whose column card is minimal.

6 Experiments

We conducted a series of experiments using publicly available benchmark sets for #SAT. Our tested benchmarks (Fichte et al. 2018a) are publicly available and our results are also on github at [github.com/hmarkus/dp_on_dbs/tplp].
Measure & Resources. We mainly compare wall clock time and number of timeouts. In the time we include preprocessing time as well as decomposition time for computing a TD with a fixed random seed. For parallel solvers we allowed access to 24 physical cores on machines. We set a timeout of 900 seconds and limited available RAM to 14 GB per instance and solver. However, since our solver dpdb is a solver using multiple threads, we restricted the results of dpdb to a total of 900 seconds of CPU time. While allowing for all the other (parallel) solvers more than 900 seconds of CPU time. For dpdb, we only allowed 900 seconds of CPU time, but at the same time restricted to 900 seconds wall clock time.

Benchmark Instances. We considered a selection of overall 1494 instances from various publicly available benchmark sets #SAT consisting of fre/meel benchmarks (1480 instances), and c2d benchmarks (14 instances). We preprocessed the instances by the #SAT preprocessor pmc (Lagniez and Marquis 2014), similar to results of recent work on #SAT (Fichte et al. 2019), where it was also shown that more than 80% of the #SAT instances have primal treewidth below 19 after preprocessing. For preprocessing with pmc we used the recommended options -vivification -eliminateLit -litImplied -iterate=10 -equiv -orGate -affine, which ensures that model counts are preserved.

Benchmarked system dpdb. We used PostgreSQL 12 for our system dpdb on a tmpfs-ramdisk (/tmp) that could grow up to at most 1 GB per run. To ensure comparability with previous results (Fichte et al. 2020), where we had employed PostgreSQL 9.5 for our system dpdb, we also considered the configuration dpdb pg9 that uses the preinstalled database system PostgreSQL 9.5 without any ramdisk at all (plain hard disk). However, we observed major performance increases of dpdb compared to dpdb pg9. We allow parallel execution for the database management system PostgreSQL with access to all cores of the benchmarking system. However, we restrict the total CPU time to ensure that we do not bias system resources towards dpdb. 

Other benchmarked systems. In our experimental work, we present results for the most recent versions of publicly available #SAT solvers, namely, c2d 2.20 (Darwiche 2004), d4 1.0 (Lagniez and Marquis 2017), DSHARP 1.0 (Muse et al. 2012), miniC2D 1.0.0 (Oztok and Darwiche 2015), cnf2eadt 1.0 (Koriche et al. 2013), bdd_minisat 1.0.2 (Toda and Soh 2015), and sdd 2.0 (Darwiche 2011), which are all based on knowledge compilation techniques. We also considered rather recent approximate solvers ApproxMC2, ApproxMC3 (Chakraborty et al. 2014), ganak (Sharma et al. 2019), and srs 1.0 (Ermon et al. 2012), as well as CDCL-based solvers Cachet 1.21 (Sang et al. 2004), sharpCDCL 7 and sharpSAT 13.02 (Thurley 2006). Finally, we also included multi-core solvers gpusat 1.0 and gpusat 2.0 (Fichte et al. 2018b; Fichte et al. 2019), which both are based on dynamic programming, as well as countAntom 1.0 (Burchard et al. 2015) on 12 physical CPU cores, which performed better than on 24 cores. Experiments were conducted with default solver options. Note that we excluded distributed solvers such as dCountAntom (Burchard et al. 2016) and DMC (Lagniez et al. 2018) from our experimental setup. Both solvers require a cluster with access to the Open-MPI framework (Gabriel et al. 2004) and fast physical interconnections.
Figure 4: Runtime for the top 15 solvers over all #SAT instances. The x-axis refers to the number of instances and the y-axis depicts the runtime sorted in ascending order for each solver individually.

Unfortunately, we do not have access to OpenMPI on our cluster. Nonetheless, our focus are shared-memory systems and since dpdb might well be used in a distributed setting, it leaves an experimental comparison between distributed solvers that also include dpdb as subsolver to future work.

Benchmark Hardware. Almost all solvers were executed on a cluster of 12 nodes. Each node is equipped with two Intel Xeon E5-2650 CPUs consisting of 12 physical cores each at 2.2 GHz clock speed, 256 GB RAM and 1 TB hard disc drives (not an SSD) Seagate ST1000NM0033. The results were gathered on Ubuntu 16.04.1 LTS machines with disabled hyperthreading on kernel 4.4.0-139. As we also took into account solvers using a GPU, for gpusat1 and gpusat2 we used a machine equipped with a consumer GPU: Intel Core i3-3245 CPU operating at 3.4 GHz, 16 GB RAM, and one Sapphire Pulse ITX Radeon RX 570 GPU running at 1.24 GHz with 32 compute units, 2048 shader units, and 4GB VRAM using driver amdgpu-pro-18.30-641594 and OpenCL 1.2. The system operated on Ubuntu 18.04.1 LTS with kernel 4.15.0-34.

6.2 Results

Figure 4 illustrates the top 15 solvers, where instances are preprocessed by pmc, in a cactus-like plot, which provides an overview over all the benchmarked #SAT instances. The x-axis of these plots refers to the number of instances and the y-axis depicts the runtime sorted in ascending order for each solver individually. Overall, dpdb seems to be quite competitive and beats most of the solvers, as for example d4, countAntom, c2d, ganak, sharpSAT, dsharp, and approxmc. Interestingly, dpdb solves also instances, whose treewidth upper bounds are larger than 41. Surprisingly, dpdb pg9 shows a different runtime behavior than the other solvers. We believe that the reason lies in an initial overhead caused by the creation of the tables that seems to depend on the number of nodes of the used TD. There, I/O operations of writing from main memory to hard disk seem to kick in. This disadvantage is resolved if benchmarking dpdb on
Table 1: Number of solved \#SAT instances, preprocessed by pmc and grouped by intervals of upper bounds of the treewidth. time[h] is the cumulated wall clock time in hours, where unsolved instances are counted as 900 seconds.

| solver     | 0-20 | 21-30 | 31-40 | 41-50 | 51-60 | >60 | best | unique | ∑ | time[h] |
|------------|------|-------|-------|-------|-------|-----|------|--------|---|--------|
| miniC2D   | 1193 | 29    |       | 2     | 1     | 7   | 11   | 0      | 1242 | 68.77  |
| dpdb      | 1193 | 31    | 7     | 2     | 0     | 0   | 2    | 1      | 1233 | 70.44  |
| gpusat2   | 1196 | 32    | 1     | 0     | 0     | 154 | 1    | 0      | 1229 | 71.27  |
| d1        | 1163 | 20    | 10    | 2     | 4     | 28  | 33   | 1      | 1227 | 76.86  |
| countAntom| 1141 | 18    | 10    | 5     | 4     | 13  | 102  | 0      | 1191 | 84.39  |
| dpdb pg9  | 1159 | 19    | 5     | 2     | 0     | 0   | 0    | 0      | 1184 | 100.99 |
| c2l       | 1124 | 31    | 10    | 3     | 3     | 10  | 24   | 0      | 1181 | 84.41  |
| ganak     | 1031 | 16    | 10    | 2     | 4     | 29  | 633  | 0      | 1092 | 107.25 |
| sharpSAT  | 1029 | 16    | 10    | 2     | 4     | 30  | 80   | 0      | 1091 | 106.88 |
| sdd       | 1014 | 4     | 7     | 1     | 0     | 2   | 0    | 0      | 1028 | 124.23 |
| sts       | 927  | 4     | 8     | 7     | 5     | 52  | 35   | 21     | 1003 | 128.43 |
| dsharp    | 853  | 3     | 7     | 2     | 0     | 0   | 29   | 0      | 865  | 157.87 |
| cnf2eadt  | 799  | 3     | 7     | 2     | 0     | 7   | 157  | 0      | 818  | 170.17 |
| approxmc3 | 794  | 3     | 7     | 2     | 0     | 6   | 1    | 0      | 812  | 173.35 |
| bdd_minisat| 791 | 4     | 1     | 0     | 0     | 46  | 0    | 0      | 796  | 175.09 |

Table 7 Final Discussion & Conclusions

We presented a generic system dpdb for explicitly exploiting treewidth by means of dynamic programming on databases. The idea of dpdb is to use database management systems (DBMSs) for table manipulation, which makes it (i) easy and elegant to perform rapid prototyping for problems with DP algorithms and (ii) allows to leverage decades of database theory and database system tuning. It turned out that all the cases that occur in dynamic programming can be handled quite elegantly with plain SQL queries. Our system dpdb can be used for both decision and counting problems, thereby also considering optimization. We see our system particularly well-suited for counting problems, especially, since it was shown that for model counting (#SAT) instances of practical relevance typically have small treewidth (Fichte et al. 2019). In consequence, we carried out preliminary experiments on publicly available instances for #SAT, where we see competitive behavior compared to most recent solvers.

Future Work

Our results give rise to several research questions. We want to push towards other database systems and vendors. For example, we expect major improvements in commercial database management systems due to the availability of efficient enterprise features. In particular, we expect in the

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These upper bounds were obtained via decomposer htd in at most two seconds.
DBMS Oracle that the behavior when we use different strategies on how to write and evaluate our SQL queries, e.g., sub-queries compared to common table expressions. Currently, we do not create or use any indices, as preliminary tests showed that meaningful B*tree indices are hard to create and creation is oftentimes too expensive. Further, the exploration of bitmap indices, as available in Oracle enterprise DBMS would be worth trying in our case (and for #SAT), since one can efficiently combine database columns by using extremely efficient bit operations. It would also be interesting to investigate whether operating system features to handle memory access can be helpful (Fichte et al. 2020). In addition, one might consider dpdb in the setting of distributed algorithms such as dCountAntom (Burchard et al. 2016) and DMC (Lagniez et al. 2018).

It might be worth to rigorously test and explore our extensions on limiting the number of rows per table for approximating #SAT or other counting problems, cf., (Chakraborty et al. 2016; Dueñas-Osorio et al. 2017; Sharma et al. 2019) and compare to the recent winners of the newly established model counting competition (Fichte et al. 2020). Recent results (Hecher et al. 2020) indicate that by using hybrid solving and abstractions our results can also be extended to projected model counting (Fichte et al. 2018).

Another interesting research direction is to study whether efficient data representation techniques on DBMSs can be combined with dynamic programming in order to lift our solver to quantified Boolean formulas.

It would also be interested to consider other measures such as (fractional) hypertree width (Fichte et al. 2020; Dzulfikar et al. 2019) and investigate whether tree decompositions with additional properties (Jégou and Terrioux 2014) or other heuristics to compute tree decompositions improve solving (Strasser 2017). Furthermore, interesting directions for future research would be to implement counting various problems in our framework, such as in constraint satisfaction (Durand and Mengel 2015; Khamis et al. 2016), constraint networks (Jégou and Terrioux 2014), argumentation (Fichte et al. 2019), description logics (Fichte et al. 2021), or epistemic logic programming (Hecher et al. 2020).

**System and License**

Our system dpdb is available under GPL3 license at [github.com/hmarkus/dp_on_dbs](https://github.com/hmarkus/dp_on_dbs)

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