

Canonizing Graphs of Bounded Tree Width in Logspace

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Graph canonization is the problem of computing a unique representative, a canon, from the isomorphism class of a given graph. This implies that two graphs are isomorphic exactly if their canons are equal. We show that graphs of bounded tree width can be canonized by logarithmic-space (logspace) algorithms. This implies that the isomorphism problem for graphs of bounded tree width can be decided in logspace. In the light of isomorphism for trees being hard for the complexity class logspace, this makes the ubiquitous class of graphs of bounded tree width one of the few classes of graphs for which the complexity of the isomorphism problem has been exactly determined.

CCS Concepts: • **Theory of computation** → **Problems, reductions and completeness**;

Additional Key Words and Phrases: Algorithmic graph theory, computational complexity, graph canonization, graph isomorphism, logspace algorithms, tree width

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1 INTRODUCTION

The *graph isomorphism problem* (ISOMORPHISM)—deciding whether two given graphs are the same up to renaming vertices—is one of the few fundamental problems in NP for which we neither know that it is polynomial-time solvable nor that it is NP-complete. The fastest known algorithm has quasipolynomial running time (Babai 2016). Since NP-hardness would imply a collapse of the polynomial hierarchy to its second level (Boppana et al. 1987; Schöning 1988), significant effort has been put into better understanding the graph-theoretic requirements on input graphs that make ISOMORPHISM polynomial-time solvable. A classical result of Bodlaender (1990) shows that ISOMORPHISM is solvable in polynomial time for graphs of *bounded tree width*. Polynomial-time algorithms are also known for other graph classes like planar graphs (Hopcroft and Wong 1974; Tarjan 1971) and more general graphs with a crossing-free embedding into a fixed surface (Filotti and Mayer 1980; Grohe 2000; Miller 1980). A deeper complexity-theoretic insight behind the polynomial-time algorithms for embeddable graphs is given by the fact that ISOMORPHISM for graphs embeddable into the plane (Datta et al. 2009a) or a fixed surface (Elberfeld and Kawarabayashi 2014) can be decided by logarithmic-space (*logspace*) algorithms. These algorithms, which are polynomial-time algorithms using at most a logarithmic amount of memory, define the complexity class L.

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So far, it has been an open problem whether for graphs of bounded tree width the isomorphism problem can be solved in logspace. Guided by the goal to determine the exact complexity of the isomorphism problem for these graphs, there has been a sequence of ever stronger partial results. Bodlaender's algorithm (Bodlaender 1990) placing ISOMORPHISM for graphs of bounded tree width in P was first refined to an upper bound in terms of logarithmic-depth circuits with threshold gates (i.e., circuits defining the complexity class TC^1) (Grohe and Verbitsky 2006) and later improved to use semi-unbounded fan-in Boolean gates (i.e., circuits defining the complexity class SAC^1) (Das et al. 2012). Since the chain $L \subseteq SAC^1 \subseteq TC^1 \subseteq P$ is all we know about the inclusion relations of these classes, these works leave the question for a logspace approach that applies to every class of graphs of bounded tree width open. Logspace algorithms are known for small constant bounds on the tree width. Indeed, Lindell's classical approach to testing isomorphism of trees (Lindell 1992) provides us with a logspace algorithm for graphs of tree-width at most 1. This was generalized to graphs of tree width at most 2 (Arvind et al. 2008) and results of Datta et al. (2009b) for graphs without K_5 as a minor apply to graphs of tree width at most 3. Moreover, k -trees, the maximal tree-width- k graphs, admit logspace isomorphism tests (Arvind et al. 2012), as well as graphs with a bounded tree depth (Das et al. 2015). While providing us with ever larger classes of graphs with logspace algorithms for ISOMORPHISM, the general question for bounded tree width graphs remained open.

1.1 Results

Our first main result answers the above question in its most general way by showing that the isomorphism problem for graphs of bounded tree width can be solved by logspace algorithms. Together with a result of Jenner et al. (2003), showing that the isomorphism problem for trees is L-hard, this pinpoints the complexity of ISOMORPHISM for graphs of bounded tree width.

THEOREM 1.1. *For every positive $k \in \mathbb{N}$, there is a logspace algorithm that decides whether two given graphs of tree width at most k are isomorphic. Moreover, this problem is complete for L with respect to first-order reductions.*

For testing whether two graphs are isomorphic, it is in practice often helpful to perform a two-step approach that first computes a canonical representative for each isomorphism class, called the *canon*, and then declares the two graphs to be isomorphic exactly if their canons are equal (rather than isomorphic). To also be able to construct an *isomorphism* between the input graphs (that means, a bijective function between the vertex sets of given graphs that preserves their edge relations), it is helpful to have an additional access to an isomorphism from the input graphs to their canons. Such an isomorphism to the canon is called a *canonical labeling* of a graph. An isomorphism between the input graphs can be constructed by composing canonical labelings.

For most isomorphism algorithms that have been developed so far, it was possible, with varying amounts of extra effort, to turn them into an algorithm that computes canons and canonical labelings. Hence, deciding ISOMORPHISM and computing canons are often known to have the same complexity. However, the current situation for graphs of bounded tree width is different: While the approach from Das et al. (2012) puts the isomorphism problem for graphs of bounded tree width into SAC^1 , this is not done by providing a canonization procedure. In fact, the best known upper bound for canonizing graphs of bounded tree width uses logarithmic-depth circuits with unbounded fan-in Boolean gates (i.e., circuits defining the complexity class AC^1) (Wagner 2011). Between these classes, only the relation $SAC^1 \subseteq AC^1$ is known. Our second main result clarifies this situation by providing a logspace algorithm for canonizing graphs of bounded tree width.

THEOREM 1.2. *For every $k \in \mathbb{N}$, there is a logspace algorithm that, given as input a graph G of tree width at most k , outputs an isomorphism-invariant encoding of G (a canon) and an isomorphism to it (a canonical labeling).*

1.2 Techniques

The known logspace approaches for canonizing certain classes of bounded tree width graphs are based on first computing an isomorphism-invariant tree decomposition for the given input graph and then adjusting Lindell's tree canonization approach to canonize the graph with respect to the decomposition. For example, for k -trees (Arvind et al. 2012) an isomorphism-invariant tree decomposition arises by taking a graph's maximal cliques and their intersections as the bags of the decomposition and connecting two bags based on inclusion. The resulting tree decomposition is both isomorphism-invariant, which is required for a canonization procedure to be correct, and has width k , which enables the application of an extension of Lindell's approach by taking (the constant number of) orderings of the vertices of the bags into account.

Ingredient 1: Isomorphism-invariant tree decomposition into bags without clique separators. In general, for graphs of tree width at most k , there is no isomorphism-invariant tree decomposition of width k . A simple example of graphs demonstrating this are cycles, which have tree width 2, but no isomorphism-invariant tree decomposition of width 2. We could hope to find an isomorphism-invariant tree decomposition by allowing approximate tree decompositions (that means, allowing an increase of the width to some constant k'). Again, cycles show that such tree decompositions do not always exist. To address this issue, we could consider not just one tree decomposition, but an isomorphism-invariant and polynomial-size collection of tree decompositions. However, for all $k' \in \mathbb{N}$, there are graphs of tree width at most 3 for which the smallest isomorphism-invariant collection of tree decompositions of width k' has exponential size. Simple graphs demonstrating this fact are given by forming the disjoint union of n cycles of length n , and adding a vertex that is adjacent to every other vertex.

We work around this problem by considering isomorphism-invariant tree decompositions that may have bags of unbounded size but with bags that are easier from a graph-theoretic and algorithmic perspective than the original graph. An algorithm developed recently (Lokshtanov et al. 2014a) (which refined the time complexity for ISOMORPHISM on graphs of tree width k from Bodlaender's $n^{O(k)}$ bound to $g(k) \cdot n^{O(1)}$ for a function g) applies a technique from Leimer (1993) that turns the input graph into its isomorphism-invariant collection of maximal induced subgraphs without clique separators called *maximal atoms*. (In the example above, the maximal atoms are exactly the enriched cycles.) We adapt this idea as a first step for the proofs of our main results but need to adjust it both with respect to the graph-theoretic concepts as well as the algorithmic ideas involved: While the collection of subgraphs produced by Leimer's approach is isomorphism-invariant, the tree underlying the resulting decomposition highly depends on the order in which subgraphs are considered and, thus, is not isomorphism-invariant. While it is always sufficient to have an isomorphism-invariant set of potential bags capturing a tree decomposition to perform polynomial-time isomorphism tests (see Otachi and Schweitzer (2014)), to apply or work toward logspace techniques it is necessary to have an isomorphism-invariant tree decomposition. Our first main technical contribution develops a tree decomposition of a graph whose bags are maximal atoms that is isomorphism-invariant and logspace-computable for graphs of bounded tree width.

Ingredient 2: Nested tree decomposition and a quasi-complete isomorphism-based ordering. The approach of Lindell (1992) for canonizing trees is based on using a weak order on the class of all trees whose incomparable elements are exactly the isomorphic ones, and showing that the order can be computed in logspace. Das et al. (2012) extended this to also work for graphs with respect to given tree decompositions of bounded width. This is done by adding the idea that, for bounded width, it is possible in logspace to guess partial isomorphisms between bags and recursively check whether they can be extended to isomorphisms between the whole graphs and the tree decompositions. When working with the tree decompositions into maximal atoms described above, it is not

possible to just guess and check partial isomorphisms between bags, since they have an unbounded width.

To handle the width-unbounded bags of the above decomposition, we use the fact that (as shown in Lokshtanov et al. (2014a)), after appropriate preprocessing, the maximal atoms have polynomial-size isomorphism-invariant families of approximate tree decompositions. To compute these families, we combine an approach for constructing separator-based tree decompositions from Elberfeld et al. (2010) to work with the isomorphism-invariant separators from Lokshtanov et al. (2014a). If we choose a bounded width tree decomposition for each atom, and replace each atom by the chosen tree decomposition, then we can turn the width-unbounded decomposition into a width-bounded decomposition for the whole graph. However, since each maximal atom may be associated with several decompositions, we need to consider for each atom a family of decompositions. We call the structure that is obtained a *nested tree decomposition*. To extend the approach that canonizes with respect to width-bounded decompositions of Das et al. (2012) to nested tree decompositions, we incorporate a bag refinement step into the weak ordering. It turns root bags of unbounded width into width-bounded tree decompositions. For each candidate tree decomposition of the root bag, this triggers a modification of the original tree decomposition. However, it turns out that determining whether there is an isomorphism between two graphs that respects two given nested tree decompositions is as hard as the general graph isomorphism problem (see Remark 5.4). Having a polynomial-time algorithm for this, let alone a logspace algorithm, would thus put the general graph isomorphism problem into P. Consequently, we do not generalize the idea of using isomorphism-based orderings with respect to decompositions in a direct way to nested tree decompositions. Instead, we define an approximation of the isomorphism-based ordering. This approximation has the property that it is isomorphism-invariant (i.e., graphs that are isomorphic with respect to given nested decompositions are incomparable) but is only quasi-complete, by which we mean that graphs that are incomparable must be isomorphic but not necessarily via an isomorphism that respects the nested decompositions. Developing the notion of nested tree decompositions along with just the right notion of a quasi-complete isomorphism-based ordering is our second main technical contribution.

Ingredient 3: Recursive logspace algorithm implementing the quasi-complete ordering. Trying all choices of a decomposition on all of the atoms yields exponentially many refined decompositions in total. Avoiding this exponential blowup, our third main technical contribution is a dynamic-programming approach along the tree decomposition that shows how to cycle through candidate decompositions of the maximal atoms while, still, canonizing the graph along the coarser tree decomposition in logspace.

Since recursively cycling through tree decompositions of a bag needs space, we cannot just use the polynomial-size family of tree decompositions that we get from applying the results of Elberfeld et al. (2010) to those of Lokshtanov et al. (2014a) as described above. To implement the recursion in logspace, we compute nested tree decompositions that satisfy a certain additional (quite technical) property, which we call *p-boundedness*. It allows us to maintain a trade-off between the number of candidate tree decompositions chosen for each bag and the size of the subdecomposition sitting below the bag. This makes a recursive algorithm that uses only logarithmic space possible.

1.3 Organization of the Article

Section 2 provides background on standard graph-theoretic notions and logspace. The remaining part of the article is structured along the proofs of the main theorems: In Section 3 we show how to compute isomorphism-invariant tree decompositions into clique-separator-free graphs in logspace, while Section 4 contains the decomposition approach for graphs without clique separators. Section 5 defines the notion of nested tree decompositions and a weak ordering that

is recursively defined along these decompositions, while Section 6 proves that the ordering is logspace-computable for width-bounded and p -bounded decompositions. Section 7 finally proves the article's main theorems. Section 8 concludes with a summary and an outlook.

2 BACKGROUND

We denote the set of natural numbers, which start at 0, by \mathbb{N} , and use shorthands $[n, m] := \{n, \dots, m\}$ and $[m] := [1, m]$ for every $n \in \mathbb{N}$ and $m \in \mathbb{N} \setminus \{0\}$.

2.1 Graphs and Connectivity

For a graph $G = (V, E)$ with vertices V and edges $E \subseteq V \times V$, we define $V(G) := V$ and $E(G) := E$. All graphs considered in the present article are finite, undirected, and simple (neither parallel edges nor loops are present). We denote the class of all finite graphs by \mathcal{G} . To simplify later definitions, we define the *coloring function* $\text{col}_G : V(G) \times V(G) \rightarrow \mathbb{Z}$ of a graph G as follows. $\text{col}_G(u, v)$ equals -1 if $v = u$, 1 if $v \neq u$ and $\{u, v\} \in E(G)$, and 0 if $v \neq u$ and $\{u, v\} \notin E(G)$. If the vertices or the edges of G are *colored*, then we extend the coloring function to return natural number encodings of colors.

Subgraphs and *induced subgraphs* are defined as usual. We write $G[V']$ to denote the subgraph induced by a vertex set $V' \subseteq V(G)$ in a graph G . A *path* is an alternating sequence $v_0 e_0 \dots v_{m-1} e_{m-1} v_m$ of distinct vertices and edges from G with $e_i = \{v_i, v_{i+1}\}$ for every $i \in \{0, \dots, m-1\}$.

A *separation* of a graph G is a pair (A, B) of subsets of $V(G)$ with (1) $A \cup B = V(G)$, and (2) $(E(G) \cap ((A \setminus B) \times (B \setminus A))) = \emptyset$. The intersection $A \cap B$ is the *separator* of (A, B) and it has size $|A \cap B|$. The definition of how a separation (A, B) separates parts of a graph commonly distinguishes between whether the separated parts are vertices or sets of vertices: A separator (A, B) *separates* vertex sets $X, Y \subseteq V(G)$ with $X \subseteq A$ and $Y \subseteq B$. A separator (A, B) *separates* vertices $x, y \in V(G)$ with $x \in A \setminus B$ and $y \in B \setminus A$. The *connectedness* of sets $X, Y \subseteq V(G)$ in a graph is the size of a smallest separator that separates them, it is denoted by $\kappa_G(X, Y)$. The *connectedness* of vertices $x, y \in V(G)$, denoted by $\kappa_G(x, y)$, is defined in the same way, except that we set $\kappa_G(x, y) := \infty$ if x and y are adjacent.

2.2 Graph Isomorphism

An *isomorphism* from a (colored) graph G to a (colored) graph H is a bijective mapping $\varphi : V(G) \rightarrow V(H)$, such that $\text{col}_G(u, v) = \text{col}_H(\varphi(u), \varphi(v))$ holds for every $u, v \in V(G)$. Graphs G and H that admit an isomorphism between them are *isomorphic*. This gives rise to an equivalence relation that partitions \mathcal{G} into *isomorphism classes*. The *graph isomorphism problem* is the language $\text{ISOMORPHISM} := \{(G, H) \in \mathcal{G} \times \mathcal{G} \mid G \text{ and } H \text{ are isomorphic}\}$. Here an encoding of the graphs as strings is assumed. (We can assume that the graphs are given as adjacency matrices, which is, however, irrelevant, since the reasonable encodings are logspace equivalent.)

A *canonization* is a mapping $\text{can} : \mathcal{G} \rightarrow \mathcal{G}$ where G is isomorphic to $\text{can}(G)$, such that for every two graphs G and H , we have $\text{can}(G) = \text{can}(H)$ exactly if G and H are isomorphic. The graph $\text{can}(G)$ is the *canon of G (under can)*, and an isomorphism φ between G and $\text{can}(G)$ is a *canonical labeling of G (under can)*. Comparing the canons of two graphs G and H suffices to test whether they are isomorphic, and canonical labelings of two graphs can be used to construct an isomorphism.

A mapping that associates an object $\text{inv}(G)$ with every graph $G \in \mathcal{G}$, for example a tree decomposition or a family of tree decompositions, is *isomorphism-invariant* if for every isomorphism φ between two graphs the result of applying φ and inv is independent of the order in which they are applied. That means, for every isomorphism φ from a graph G to a graph H , replacing all occurrences of vertices $v \in V(G)$ in $\text{inv}(G)$ by their image $\varphi(v)$ yields $\text{inv}(H)$.

2.3 Tree Decompositions

A (tree) decomposition $D = (T, \mathcal{B})$ of a graph G is a tree T together with a family of bags $\mathcal{B} = (B_n)_{n \in V(T)}$ with $B_n \subseteq V(G)$ for each $n \in V(T)$, such that (*connectedness property*) for each vertex $v \in V(G)$ the induced subtree $T[\{n \in V(T) \mid v \in B_n\}]$ is nonempty and connected, and (*covering property*) for each edge $\{u, v\} \in E(G)$ there is a node $n \in V(T)$ with $\{u, v\} \subseteq B_n$. For every edge $\{n, m\} \in E(T)$, $B_{\{n, m\}} := B_n \cap B_m$ is the *adhesion set* between nodes n and m . The *torso* of a node $n \in V(T)$ is the graph obtained from the induced graph $G[B_n]$ by adding for every neighboring node n' of n the clique on the adhesion set $B_n \cap B_{n'}$. Given a tree decomposition $D = (T, \mathcal{B})$, its *size* is $|D| := |V(T)|$, and its (*tree*) *width* is the maximum over all $|B_n| - 1$ for $n \in V(T)$. The *tree width* of G , denoted by $\text{tw}(G)$, is the minimum width of a tree decomposition for it.

When working with trees underlying *rooted* tree decompositions $D = (T, \mathcal{B})$, which have a distinguished root node $r \in V(T)$, we talk about a *parent*, *ancestor*, *child*, and *descendant* of a node $n \in V(T)$ with respect to the root $r \in V(T)$ in the usual way. Given a rooted tree decomposition $D = (T, \mathcal{B})$, a *subdecomposition* $D' = (T', \mathcal{B}')$ is a decomposition that arises by using a node $n \in V(T)$ and all its ancestor nodes to form a tree decomposition. A *child decomposition* is a subdecomposition that contains a child of the root node, but not the root.

Tree decompositions are commonly studied in both their unrooted and rooted variants. In the context of logspace and the isomorphism problem, we do not need to restrict ourselves to a particular definition (as formalized by the following fact). However, to facilitate a clear presentation, we use rooted tree decompositions.

FACT 2.1. *There is a logspace-computable and isomorphism-invariant mapping that turns an unrooted tree decomposition D for a graph G into a rooted tree decomposition D' for G with the same adhesion sets.*

Fact 2.1 seems to be a folklore and special cases have, for example, been used in Arvind et al. (2012) and Datta et al. (2009a). Its proof takes an unrooted tree decomposition and turns it into a rooted decomposition by declaring the *center of the tree*, the unique node or edge with the maximum distance to the tree's leaves, to be the root. If the center is an edge, then we subdivide it by inserting a new node whose bag is the intersection of the edge's incident bags.

Two graphs G and G' are *isomorphic with respect to tree decompositions* $D = (T, \mathcal{B})$ and $D' = (T', \mathcal{B}')$, respectively, if there exists an isomorphism φ from G to G' and an isomorphism ψ from T to T' satisfying $B'_{\psi(n)} = \{\varphi(v) \mid v \in B_n\}$ for every node $n \in V(T)$. Under these conditions, we say that φ *respects* D and D' . Based on this definition and the way of how it refines the isomorphism equivalence relation among graphs, we also consider *canons of graphs with respect to tree decompositions*.

2.4 Logspace

A deterministic Turing machine whose working space is logarithmically bounded by the input length is called a *logspace DTM*. The functions $f : \{0, 1\}^* \rightarrow \{0, 1\}^*$ computed by such machines are *logspace-computable* (or *in logspace*). The complexity class L , called (*deterministic*) *logspace*, contains all languages $P \subseteq \{0, 1\}^*$ whose characteristic functions are in logspace. Functions in logspace are closed under composition (Stockmeyer and Meyer 1973; Jones 1975) and also under queries to oracles for languages from L (Ladner and Lynch 1976). Reingold (2008) studied the problem $\text{UNDIRECTED-REACHABILITY} := \{(G, s, t) \mid \text{there is a path from } s \in V(G) \text{ to } t \in V(G) \text{ in the undirected graph } G\}$, and showed that it is in L . Furthermore, we can test whether a graph's tree width is bounded by a constant, since $\text{TREE-WIDTH-}k := \{G \mid \text{tw}(G) \leq k\} \in L$ for every $k \in \mathbb{N}$ (Elberfeld et al. 2010). Details about the circuit complexity classes that are mentioned in the introduction are given in the book of Vollmer (1999), but we do not require them in the following.

3 DECOMPOSING GRAPHS INTO PARTS WITHOUT CLIQUE SEPARATORS

A *clique* is a graph with an edge between every two vertices, including the empty graph by definition. A separation (A, B) is a *clique separation* with *clique separator* $A \cap B$ in a graph G if it (1) separates two vertices $x, y \in V(G)$, and (2) $G[A \cap B]$ is a clique.

We construct isomorphism-invariant tree decompositions for graphs of bounded tree width whose bags induce subgraphs without clique separators and whose adhesion sets are cliques (that means, the torsos are exactly the subgraphs induced by the bags). In this context isomorphism-invariance means that every isomorphism from a graph G_1 to G_2 induces a map from the tree decomposition of G_1 to that of G_2 . These tree decompositions serve as an intermediate decomposition step in the proofs of our main theorems.

LEMMA 3.1. *For every $k \in \mathbb{N}$, there is a logspace-computable and isomorphism-invariant mapping that turns a graph G with tree width at most k into a tree decomposition D for G in which*

- (1) *subgraphs induced by the bags do not contain clique separators, and*
- (2) *adhesion sets are cliques.*

The tree decomposition we construct to prove the lemma is a refined version of a decomposition of Leimer (1993) of graphs into their collections of maximal induced subgraphs without clique separators. The crucial point is that we need to adjust his method to not only output the collection of maximal induced subgraphs without clique separators, which suffices for its application in Lokshtanov et al. (2014a), but also an isomorphism-invariant tree decomposition that is based on it. To do that, we replace the approach of Leimer (1993), which is based on finding clique-separator-free parts in a single phase via computing elimination orderings, by several steps. In these steps, we compute graphs that are clique-separator-free with respect to cliques up to a certain size. The size bound grows when going from one step to the next. While Leimer's method runs in polynomial-time and applies to general graphs, our approach needs logarithmic space and applies to graphs of bounded tree width, which suffices for our applications. The decomposition is depicted at an example in Figure 1.

Let us make some remarks on the relation of this decomposition to others. The first step of our decomposition, the decomposition with respect to cliques of size at most 1 is actually the block-cut tree of the graph, that is the decomposition into biconnected components (including bridges) and cut-vertices. However, such an analogy does not continue toward higher connectivity. In fact, consider a cycle, we see that there is no isomorphism-invariant decomposition into 3-connected components. While there are still ways to remedy this issue and obtain useful decompositions into 3-connected components, this is not the case for separators of larger size. The main observation for us is that if we only consider clique separators, invariant decompositions actually do exist.

Section 3.1 presents the definition and logspace-computability of maximal subgraphs without size-bounded clique separators. Section 3.2 presents a transformation of graphs that is used in Section 3.3 to prove Lemma 3.1

3.1 Atoms with Respect to Constant-Size Clique Separators

Let $c \in \mathbb{N}$, which we use as an upper bound on the size of clique separators we consider. A *c-atom* is a graph that does not contain clique separators of size at most c . *Atoms*, which are the graphs Leimer (1993) deals with, are $|V(G)|$ -atoms (or, alternatively, graphs without clique separators). A *maximal c-atom* of a graph G is a maximal induced subgraph $G[A]$ for some $A \subseteq V(G)$ that is a *c-atom* (that means, every extension of it contains a clique separator of size at most c). A

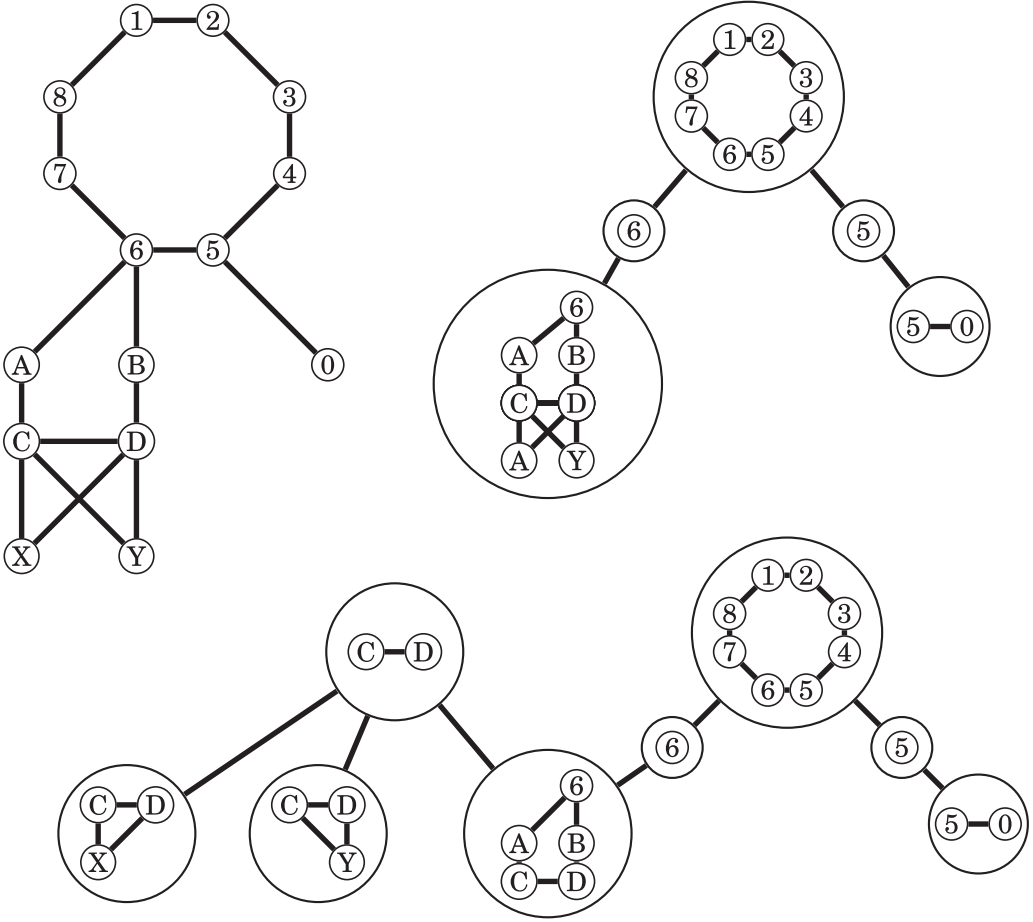


Fig. 1. The figure shows a graph (top left), the invariant decomposition into 1-atoms (top right) as well as the invariant decomposition into 2-atoms (bottom).

maximal atom in a graph G is a maximal $|V(G)|$ -atom. For every $c \in \mathbb{N}$, c -atoms are nonempty and connected.

Two vertices $a_1, a_2 \in V(G)$ are *c-inseparable* (with respect to clique separations in G) if there is no clique separator of size at most c in G separating a_1 and a_2 and *c-separable*, otherwise. A set $A \subseteq V(G)$ is *c-inseparable* if all distinct $a_1, a_2 \in A$ are *c-inseparable*. The set A is *maximal c-inseparable* if no extension of A is *c-inseparable*.

Note that the definition of induced subgraphs $G[A]$ that are c -atoms only considers separations of $G[A]$ while *c-inseparability* of a vertex set A is based on separations in the (ambient) graph G . If we look at maximal c -atoms and maximal *c-inseparable* sets, then these notions coincide.

LEMMA 3.2. *For every $c \in \mathbb{N}$, graph G , and $A \subseteq V(G)$, $G[A]$ is a maximal c -atom of G if and only if A is maximal c -inseparable in G .*

PROOF. (From c -atoms to c -inseparable sets.) Two vertices a_1 and a_2 from a vertex set A that are *c-separable* in G are also *c-separable* in $G[A]$. This implies that, if $G[A]$ is a c -atom (meaning that A is *c-inseparable* in $G[A]$), then A is *c-inseparable* in G .

(From c -inseparable sets to c -atoms.) Let $A \subseteq V(G)$ be maximal c -inseparable. Assume, for the sake of contradiction, that $a_1, a_2 \in A$ are separated by a clique separator C of size at most c in $G[A]$. Since C does not separate a_1 from a_2 in G , there is a path from a_1 to a_2 in G that avoids C . Let P be a shortest path of this kind. Since C separates a_1 from a_2 in $G[A]$, there is an $x \in V(G) \setminus A$ on the path P . Since $x \notin A$, but A is chosen to be maximal c -inseparable, there is a vertex $a' \in A$ and a clique separator C' of size at most c that separates a' from x . Since A is c -inseparable in G , C' cannot separate elements of A in G . Thus, we know that either C' separates x from a_1 or $a_1 \in C'$, and either C' separates x from a_2 or $a_2 \in C'$. That means, the path P , which starts in a_1 , passes through x , and ends in a_2 , must intersect C' in some vertex $p_1 \neq x$ before reaching x , and intersect C' again in some vertex $p_2 \neq x$ after leaving x . Since C' is a clique, we can take a shortcut by directly taking the edge $\{p_1, p_2\} \in E(G)$ without visiting x . This contradicts the fact that P is a shortest path. \square

LEMMA 3.3. *Let G be a graph with c -inseparable set $I \subseteq V(G)$ with $|I| \geq c + 1$ for some $c \in \mathbb{N}$. Then $A := \{a \in V(G) \mid I \cup \{a\} \text{ is } c\text{-inseparable}\}$ is the unique maximal c -inseparable set in G with $I \subseteq A$.*

PROOF. We first argue that A is c -inseparable: Assume, for sake of contradiction, that vertices $a_1, a_2 \in A$ are c -separable in G via a clique separator C of size at most c . Since $|C| \leq c < c + 1 \leq |I|$, there is a vertex $v \in I \setminus C$. Moreover, since C separates a_1 from a_2 , C separates a_1 from v or it separates a_2 from v . This contradicts $a_1, a_2 \in A$, since both $I \cup \{a_1\}$ and $I \cup \{a_2\}$ are c -inseparable in G by construction. To see that A is unique and maximal among the c -inseparable sets containing I , note that every candidate $a \in V(G) \setminus A$ is c -separable from at least one vertex in I . \square

Combining Lemma 3.2 with Lemma 3.3, we are able to compute the family of c -atoms of a given graph for every constant $c \in \mathbb{N}$ in logspace.

LEMMA 3.4. *For every $c \in \mathbb{N}$, the mapping that turns a graph G into its family of maximal c -atoms is logspace-computable and isomorphism-invariant.*

PROOF. We show how to compute the family of all maximal sets A that are c -inseparable in G . By Lemma 3.2, these are exactly the vertex sets of the maximal c -atoms of G . We first check for all sets A of size at most c whether they are maximal c -inseparable in G , and output the sets that pass the test. To find maximal c -inseparable sets with more than c vertices, we consider every c -inseparable set $I \subseteq V(G)$ with $|I| = c + 1$ and output $A := \{a \in V(G) \mid I \cup \{a\} \text{ is inseparable}\}$. The correctness of the algorithm follows from Lemma 3.3. It can be implemented by a logspace DTM, since we only cycle through vertex sets of (constant) size at most c and use oracle calls to UNDIRECTED-REACHABILITY, which is in L (Reingold 2008).

The mapping that turns a graph G for a $c \in \mathbb{N}$ into its family of c -inseparable sets is isomorphism-invariant by definition. \square

3.2 Chordal Completions with Respect to Constant-Size Clique Separators

Instead of working with a given graph G directly, some of the arguments of Leimer (1993) are based on working with its *chordal completion* G^* , which is the graph that arises from G by replacing every maximal atom in the graph with a clique on the atom's vertices. Interestingly, the vertex sets of the maximal atoms are the same for G and G^* . The same property holds when moving from a graph G to the *c -chordal completion* G^c , for every $c \in \mathbb{N}$, that arises from G by replacing every maximal c -atom with a clique on its vertex set. For a formal proof of this fact we first show that the intersection of two c -atoms is a clique.

LEMMA 3.5. *Let G be a graph and A_1 and A_2 be two distinct maximal c -inseparable sets in G . Then $A_1 \cap A_2$ is a clique of size at most c .*

PROOF. Since $I = A_1 \cap A_2$ is c -inseparable, it has size at most c to be contained in two distinct maximal c -inseparable sets by Lemma 3.3. Let $a_1 \in A_1 \setminus A_2$ and $a_2 \in A_2 \setminus A_1$ be c -separable vertices and C be a clique separator of size at most c that separates them. Assume, for the sake of contradiction, that there exists a vertex $a' \in (A_1 \cap A_2) \setminus C$. Since both $a' \in A_1$ and $a' \in A_2$ and both A_1 and A_2 are c -inseparable, we can find a path $P_{a_1, a'}$ between a_1 and a' and a path P_{a', a_2} between a' and a_2 in $G - C$. Thus, there is also a path P between a_1 and a_2 . This contradicts the existence of $a' \in (A_1 \cap A_2) \setminus C$. Thus, the clique C contains all of $A_1 \cap A_2$. In particular, $A_1 \cap A_2$ is a clique. \square

LEMMA 3.6. *Let $c \in \mathbb{N}$, G be a graph, $A \subseteq V(G)$, and $C \subseteq V(G)$ with $|C| \leq c$. Then,*

- (1) *$G[A]$ is a maximal c -atom in G if and only if $G^c[A]$ is a maximal c -atom in G^c , and*
- (2) *C is a clique separator in G if and only if C is a clique separator in G^c .*

PROOF. We start to prove the first property. The arguments of the proof are based on properties of vertex sets of maximal c -atoms and maximal c -inseparable vertex sets. We can freely switch between both points of view, since they are equivalent by Lemma 3.3.

Let $A \subseteq V(G)$ be maximal c -inseparable in G . Since $G^c[A]$ is a clique, A is also c -inseparable in G^c . To prove that A is maximal with this property in G^c , assume, for the sake of contradiction, that A is not maximal with this property in G^c . Then there is a vertex $x \in V(G) \setminus A$, such that $A \cup \{x\}$ is c -inseparable in G^c . Since A is maximal c -inseparable in G , there is a clique separator C of size at most c that separates a vertex $a \in A$ from x in G . By assumption, C does not separate a from x in G^c . Thus, we can find a path P between a and x in $G^c - C$. We show that P can be modified to a walk P' between a and x in $G - C$ by inspecting each of its edges and, if necessary, redirecting it. Let $\{u, v\}$ be an edge from P . If $\{u, v\} \in E(G)$, then we are done. If $\{u, v\} \in E(G^c) \setminus E(G)$, then we know by the construction of G^c that u and v are c -inseparable in G (in particular, they are part of a common maximal c -inseparable set). Thus, we can find a path $P_{\{u, v\}}$ between u and v in $G - C$ and modify P to use $P_{\{u, v\}}$ instead of $\{u, v\}$, which is only present in G^c . Overall, this leads to the construction of a walk P' in $G - C$ between a and x . This contradicts the fact that C separates a and x in G .

We are left to prove the converse direction. Let a_1 and a_2 be two vertices that are c -inseparable in G^c . For the sake of contradiction, assume they are c -separable in G . Let (B_1, B_2) be a clique separation in G with clique separator $C = B_1 \cap B_2$ of size at most c and $a_1 \in B_1 \setminus B_2$ and $a_2 \in B_2 \setminus B_1$. In particular, this means that every vertex $a'_1 \in B_1 \setminus B_2$ is c -separable from every vertex $a'_2 \in B_2 \setminus B_1$. Hence, no edges are constructed between the sets $B_1 \setminus B_2$ and $B_2 \setminus B_1$, and C is also a clique separator of size at most c in G^c . This contradicts the initial choice of a_1 and a_2 as being c -inseparable vertices in G^c . Thus, a_1 and a_2 are c -inseparable in G , too. The arguments above imply that every c -inseparable set of G^c is a c -inseparable set in G . In particular, this holds for the maximal c -inseparable sets and, thus, for the maximal c -atoms.

For the second property, let C be a clique separator of size at most c in G , which does not need to be a minimum clique separator. Then C is also a clique separator in G^c by the arguments from the last paragraph. For the other direction, let C be a clique separator of size at most c in G^c . It also separates two distinct vertices in G , and we are left to prove that it is a clique. Since C is a clique separator of size at most c , it lies in the intersection of two distinct maximal c -inseparable sets A_1 and A_2 in G^c . The sets A_1 and A_2 are also maximal c -inseparable sets of G by the first property of the lemma proved above, and $A_1 \cap A_2$ is a clique by Lemma 3.5. Thus, C is a clique separator in G , too. \square

3.3 Isomorphism-Invariant Tree Decompositions into Atoms

Our goal is to compute an isomorphism-invariant tree decomposition of a graph into its c -atoms.

A *minimum clique separator (with respect to x and y in a graph G)* is an inclusion-wise minimal clique that separates x and y in G . For every $c \in \mathbb{N}$ and graph G , we define the graph $T_c = T_c(G)$

whose node set consists of all c -atoms of G and all minimum clique separators of size at most c . An edge is inserted between every c -atom $G[A]$ and minimum clique separator C with $C \subseteq A$. We define the class of bags $\mathcal{B}_c(G) = (B_n)_{n \in V(T_c)}$ as follows. If $n \in V(T_c)$ is identified with a c -atom $G[A]$, then $B_n := A$, and if $n \in V(T_c)$ is identified with a minimum clique separator C , then $B_n := C$.

PROPOSITION 3.7. *For every $c \in \mathbb{N}$, the mapping defined by $G \mapsto (T_c(G), \mathcal{B}_c(G))$ is isomorphism-invariant.*

The graph $T_c(G)$ is typically not a tree. However, as stated by Lemma 3.8, $T_c(G)$ is a tree and, moreover, $(T_c(G), \mathcal{B}_c(G))$ is a tree decomposition if G is a $(c - 1)$ -atom.

LEMMA 3.8. *For every positive $c \in \mathbb{N}$ and $(c - 1)$ -atom G , $(T_c(G), \mathcal{B}_c(G))$ is a tree decomposition for G . Moreover, $T_c(G)$ has a unique center.*

PROOF. Instead of working with G , we use the graph G^c , whose maximal c -inseparable vertex sets and clique separators of size at most c are exactly the respective ones of G by Lemma 3.6. Thus, we set $G := G^c$ throughout the proof, which does not alter the construction of T_c . To simplify the notations of the proof, we also set $T := T_c(G)$ and $\mathcal{B} := \mathcal{B}_c(G)$.

CLAIM 1. *T is connected.*

PROOF OF CLAIM 1. If T is a single node, then the claim holds. If $V(T) \geq 2$, then we argue as follows.

Since every clique separator is contained in some maximal c -atom, it suffices to show that distinct maximal c -atoms are connected in T . Let A_1 and A_2 be maximal c -atoms. Since A_1 and A_2 are distinct, there is a clique separator C of size at most c separating A_1 from A_2 in G . For an atom A and a clique separator C separating A from a vertex $x \in V(G) \setminus A$, define $\Delta(A, C)$ to be the minimum $\sum_{i=1}^c |P_i|$ among all c -tuples of vertex-disjoint paths P_1, \dots, P_c that start in A and end in C . Such paths exist by Menger's theorem (Diestel 2005) after the preprocessing mentioned above. Note that $\Delta(A, C) = 0$ if and only if $C \subseteq A$. We show that distinct atoms A_1 and A_2 are connected in T by induction on $\Delta(A_1, C) + \Delta(A_2, C)$. If $\Delta(A_1, C) = \Delta(A_2, C) = 0$, then $C \subseteq A_1$ and $C \subseteq A_2$ so A_1 and A_2 are connected in T by definition. Thus, we assume that for all cliques C of size c separating A_1 and A_2 we have $\Delta(A_1, C) + \Delta(A_2, C) > 0$. To continue the proof, we distinguish two cases.

For the first case, assume there exists a clique C of size c separating A_1 and A_2 with $\Delta(A_1, C) > 0$ and $\Delta(A_2, C) > 0$. This implies both $C \not\subseteq A_1$ and $C \not\subseteq A_2$. Since C is c -inseparable, there is a c -atom A' that contains C . Since every path from A_1 to A_2 must intersect C , we have $\Delta(A', C) + \Delta(A_i, C) < \Delta(A_1, C) + \Delta(A_2, C)$ for $i \in \{1, 2\}$. Thus, A' is connected to A_i in T for every $i \in \{1, 2\}$ and, thus, A_1 is connected to A_2 .

For the second case, suppose that for all clique separators C of size c separating A_1 and A_2 , we have $\Delta(A_1, C) = 0$, or $\Delta(A_2, C) = 0$. Let C be such a clique separator. Without loss of generality, we may assume $\Delta(A_1, C) = 0$. Since $\Delta(A_2, C) > 0$, there is an element $v \in C \setminus A_2$. Let C' be a clique that separates v from A_2 . Since $v \in C \setminus A_2$, we conclude that $\Delta(A_2, C') = 0$ and, thus, $C' \subseteq A_2$. If there is an atom A' containing C and C' , then A' is adjacent to A_1 and A_2 and, thus, A_1 and A_2 are connected. Otherwise, there must be a clique separator C'' separating a vertex in C from a vertex in C' . However, this implies $C'' \not\subseteq A_1$ and $C'' \not\subseteq A_2$. This brings us to the previous case with $\Delta(A_1, C'') > 0$ and $\Delta(A_2, C'') > 0$. \square

CLAIM 2. *If $A_1, C_1, A_2, \dots, C_{t-1}, A_t$ is a path in T between two c -atoms, then for every $i \geq 2$, the c -atom A_i does not contain a vertex from $A_1 \setminus A_2$.*

PROOF OF CLAIM 2. Let v_1 be a vertex in $A_1 \setminus A_2$. Such a vertex must exist, since A_1 and A_2 are distinct maximal c -atoms. We show for $2 \leq i \leq t - 1$ that the separator C_i contains a vertex v_i that

is separated from v_1 by C_1 . For $i \geq 2$, we choose v_i as a vertex in $C_i \setminus C_1$. Such a vertex exists, since the C_i are distinct subsets of $V(G)$ of the same size c . To see that C_1 separates v_1 from v_i , observe that by induction C_1 separates v_1 from v_{i-1} , but v_{i-1} and v_i cannot be separated by C_1 , since they lie in the same c -atom A_i . Since $v_{t-1} \in A_t$, C_1 separates all vertices in $A_1 \setminus A_2$ from all vertices in $A_t \setminus A_2$. This proves the claim. \square

To see that T is a tree, assume that $A_1, C_1, A_2, \dots, C_{t-1}, A_t = A_1$ is a cycle. By the second claim, $A_t = A_1$ does not contain a vertex from $A_1 \setminus A_2$, but this contradicts A_1 and A_2 being maximal c -atoms that are distinct. To see that the center of T is a unique node, it suffices to observe that a separator cannot be a leaf of T .

To prove the connectedness property of decompositions, let $A_1, C_1, A_2, \dots, C_{t-1}, A_t$ be a path, such that A_1 and A_t contain a common vertex v that is not contained in A_i for $i \in \{2, \dots, t-1\}$. Since separators are always contained in some adjacent atom, this is the only case that needs to be considered. However, the existence of such a path directly contradicts the second claim above. The covering property of tree decompositions follows from the fact that every edge of G is part of some c -atom. Hence, T is a tree decomposition of G . \square

LEMMA 3.9. *For every $d, c \in \mathbb{N}$, with $d \leq c$, there is a logspace-computable mapping that turns a d -atom G into a tree decomposition $D = (T, \mathcal{B})$ of G such that*

- (1) *the bags are c -atoms,*
- (2) *the adhesion sets are cliques, and*
- (3) *the mapping is isomorphism-invariant.*

PROOF. We show the lemma by induction on $c - d$. If $c - d = 0$, then the c -atom G is the unique bag of the tree decomposition D , which satisfies all requirements of the lemma. If $c - d > 0$, then we construct and prove the correctness of the constructed tree decomposition as follows.

(Construction.) We use Lemma 3.8 to construct a tree decomposition $D' = (T', \mathcal{B}')$ whose bags are the graph's $(d + 1)$ -atoms and minimum clique separators. Applying the induction hypothesis, we compute for each $(d + 1)$ -atom A an isomorphism-invariant tree decomposition $D_A = (T_A, \mathcal{B}_A)$ into its c -atoms. We continue combining D' with the decompositions D_A to construct $D = (T, \mathcal{B})$. We use nodes $V(T) := \{(B, A) \mid B \in V(T_A) \text{ and } A \in V(T')\}$ for T . Two nodes (B_1, A_1) and (B_2, A_2) of T are adjacent if (1) $A_1 = A_2$ and B_1 and B_2 are adjacent in D_{A_1} , or (2) A_1 and A_2 are adjacent in T' and for each $i \in \{1, 2\}$, B_i contains $A_1 \cap A_2$ and is closest to the root with this property in D_{A_i} . To each node of (B, A) of T , \mathcal{B} assigns the bag $B_{(B,A)} = B$.

Since distances in trees are logspace-computable, we can determine the bag closest to the root in the definition above. Thus, T can be constructed in logspace based on constructing T by Lemma 3.8 and T_A by induction.

(Correctness of construction.) The tree T is well defined, since the intersection of two atoms A_1 and A_2 that are adjacent in D' is a clique and every clique must be contained in some bag of a tree decomposition. Moreover, the bags that contain a clique form a connected subtree and the bag closest to the root is well defined.

For the isomorphism invariance of T note first that the construction of D' and for each $(d + 1)$ -atom A the construction of D_A is isomorphism-invariant by Proposition 3.7. This implies that $V(T)$ is isomorphism-invariant. To see that also the edge set $E(T)$ is isomorphism-invariant note that whether (B_1, A_1) and (B_2, A_2) are adjacent is determined by D_{A_1} if $A_1 = A_2$ and thus isomorphism invariant. If $A_1 \neq A_2$, however, then whether (B_1, A_1) and (B_2, A_2) are adjacent depends on A_1 and A_2 being adjacent in T' , which is isomorphism-invariant, and whether for $i \in \{1, 2\}$, we have $B_i \subseteq A_1 \cap A_2$ and that B_i is closest to the root with this property in D_{A_i} . This is a combinatorial property of B_i determined by the structure of the graph that is preserved under isomorphisms.

We argue that T is a tree. It is connected, since D' and each D_A is connected. To argue that it is cycle free, suppose $(B_1, A_1), \dots, (B_t, A_t)$ with $(B_t, A_t) = (B_1, A_1)$ is a cycle. Note that for two atoms A and A' that are adjacent in D' the bag B and B' for which (B, A) is adjacent to (B, A') is unique. This implies that either the walk A_1, \dots, A_t contains a cycle, or there are indices $1 \leq j < k \leq t$, such that $A_j = A_{j+1} = \dots = A_k$ and $(B_j, A_j), (B_{j+1}, A_j), \dots, (B_k, A_j)$ is a closed walk, which implies that B_j, B_{j+1}, \dots, B_t is a cycle. Since both D' and D_A are acyclic, this yields a contradiction.

It remains to show that with this definition T is a tree decomposition whose adhesion sets are cliques. If two vertices (B_1, A_1) and (B_2, A_2) are adjacent, then $B_1 \cap B_2$ is an adhesion set either in $D_{A_1} = D_{A_2}$, or in D . In either case it is a clique. To show the connectivity property of tree decompositions, let $(B_1, A_1), \dots, (B_t, A_t)$ be a path in T such that B_1 and B_t contain a vertex v that does not appear in B_i for $2 \leq i \leq t-1$. This implies, since D' is a tree decomposition, that v is contained in all A_i . In turn, this implies that v is contained in all B_i , since each D_A is a tree decomposition. \square

PROOF OF LEMMA 3.1. Let G be the input graph of tree width at most k . Without loss of generality, we assume that it is connected. Then the graph G is a 0-atom. We apply Lemma 3.9 to G with $d = 0$ and $c = k + 1$. Since G has tree width at most k , the size of a largest clique in G is bounded by $k + 1$. Thus, the subgraphs induced by the bags, which do not contain clique separators of size at most $k + 1$ by their construction, do not contain clique separators (of any size). \square

Let us emphasize that, while we used the c -chordal completion G^c for some structural considerations throughout this section, the decomposition computed by the algorithm of Lemma 3.1 is a decomposition of the unmodified graph G .

4 DECOMPOSING GRAPHS WITHOUT CLIQUE SEPARATORS

The decomposition procedure from the previous section provides us with a tree decomposition whose bags are clique-separator-free. In the present section, we decompose clique-separator-free graphs further into isomorphism-invariant tree decompositions of bounded width (formalized by Lemma 4.1). This needs two additional assumptions that we later meet during the proofs of Theorems 1.1 and 1.2. First, the decomposition is based on two distinguished nonadjacent vertices from the graph. Second, we assume that the given graph is improved as defined next.

Let $\text{impr} : \mathcal{G} \rightarrow \mathcal{G}$ be the mapping that takes a graph G and adds edges between all vertices $u, v \in V(G)$ with $\kappa(u, v) > \text{tw}(G)$. The impr -operator *improves* the graph by adding edges of G based on its tree width. To avoid losing information, we introduce a function $\text{col}_{\text{impr}(G)}$ that colors edges that appear originally in the inputs with a different color than those coming from the improvement. The mapping impr is isomorphism-invariant by definition. Besides this, we use three further properties of the mapping impr . First, the graph we get from applying impr is saturated in the sense that a second application of it does not add new edges. Formally, this means $\text{impr}(G) = \text{impr}(\text{impr}(G))$ for every graph G as proved in Lokshtanov et al. (2014b, Lemma 2.5). Second, the tree decompositions of a graph G are exactly the tree decompositions of $\text{impr}(G)$. This implies $\text{tw}(G) = \text{tw}(\text{impr}(G))$ and is proved in Lokshtanov et al. (2014b, Lemma 2.6). Third, the mapping impr is logspace-computable for graphs of bounded tree width. This follows from Reinhold's algorithm for UNDIRECTED-REACHABILITY, and the fact that the tree width of a graph bounds the size of the separators, we need to consider to compute impr .

LEMMA 4.1. *For every $k \in \mathbb{N}$, there is a $k' \in \mathbb{N}$ and a logspace-computable and isomorphism-invariant mapping that turns every graph G with a distinguished non-edge $\{u, v\} \notin E(G)$, where G*

- (1) *has tree width at most k ,*
- (2) *does not contain clique separators, and*
- (3) *is improved (that means, $G = \text{impr}(G)$),*

into a width- k' tree decomposition $D = (T, \mathcal{B})$ for G .

Note that in this lemma, to achieve isomorphism invariance, having the distinguished edge (or something similar) is crucial for the existence of an isomorphism-invariant decomposition of width at most k (consider for example cycles).

The rest of the section is devoted to the proof of the lemma. The construction of the decomposition is based on recursively splitting the graph into smaller subgraphs using size-bounded and isomorphism-invariant separators. To do this, we adapt in a first step the isomorphism-invariant separators from Lokshtanov et al. (2014a) and show their logspace-computability (this is done in Section 4.1). Then, we combine this with a logspace approach for handling the recursion involved in this approach from Elberfeld et al. (2010) (this is done in Section 4.2).

4.1 Constructing Isomorphism-Invariant Separators

Lokshtanov et al. (2014a) identified an isomorphism-invariant family of separators that can be used as part of a recursive algorithm for constructing isomorphism-invariant tree decompositions. We adapt their approach of constructing separators, which is tailored to find (time-efficient) algorithms proving fixed-parameter tractability, to work in logspace. For this, we need to adjust some terminology.

For a graph G and a set of vertices $V' \subseteq V(G)$, we define the neighborhood of V' in G to be the set $N_G(V') := \{v \in V(G) \setminus V' \mid \text{there exists } w \in V' \text{ with } \{v, w\} \in E(G)\}$. A *graph with interface* is a pair (G, I) consisting of a graph G and an *interface* $I \subseteq V(G)$ where

- (1) $G \setminus I$ is connected, and
- (2) $I = N_G(V(G) \setminus I)$.

We split a graph into several components based on separators for its interface. Let G be a graph and $X, Y \subseteq V(G)$. It is well-known (see, for example, Lokshtanov et al. (2014a)) that there is a unique separator (A, B) for X and Y of minimum size and (inclusion-wise) minimal A . We denote it by (A_X, B_Y) and set $\text{sep}(X, Y) := A_X \cap B_Y$. Exactly the same property holds when considering vertices $x, y \in V(G)$. In this case, we denote the corresponding separation by (A_x, B_y) , and set $\text{sep}(x, y) := A_x \cap B_y$. We use these separations to define the *separator* $\text{sep}_s(G, I)$ of a graph with interface (G, I) with respect to a threshold value $s \in \mathbb{N}$: If $|I| \leq s$, then we set

$$\text{sep}_s(G, I) := I \cup \bigcup_{\substack{x, y \in I, x \neq y, \text{ and} \\ \kappa_G(x, y) \leq \text{tw}(G)}} \text{sep}(x, y), \text{ and}$$

$$\text{sep}_s(G, I) := I \cup \bigcup_{\substack{X, Y \subseteq I, X \cap Y = \emptyset, |X| = |Y| = \text{tw}(G) + 1, \text{ and} \\ \kappa_G(X, Y) \leq \text{tw}(G)}} \text{sep}(X, Y), \text{ otherwise.}$$

The definition of $\text{sep}_s(G, I)$ is isomorphism-invariant with respect to G and I . In fact, the definition of $\text{sep}_s(G, I)$ is tailored as to construct an isomorphism-invariant extension of the interface $\text{sep}_s(G, I) \supseteq I$, typically resulting in the rest of the graph decomposing into several components, while keeping control of the size of the new interfaces to the new components that arise. Intuitively, in case the interface is small ($|I| \leq s$), this is guaranteed by the fact that $|\text{sep}_s(G, I)|$ will not be too large (say only of medium size). In the case, however, that I is not small, then $|\text{sep}_s(G, I)|$ may become large, but a submodularity argument (Lokshtanov et al. 2014b) guarantees that the new interfaces that arise are at most of medium size.

In more detail, the following fact on size bounds for separators and neighborhoods follows from the statements and proofs of Lokshtanov et al. (2014b, Lemmata 3.3 and 3.4).

FACT 4.2. *There are functions $\text{small} \in O(k)$, $\text{medium} \in O(k^3)$, and $\text{large} \in O(2^{k \log k})$ with the following properties: Let (G, I) be a graph with interface, such that $\text{tw}(G) \leq k$, G is improved and an atom such that*

- (1) $G[I]$ is not a clique, and
- (2) $|I| \leq \text{medium}(\text{tw}(G))$.

Moreover, let $S := \text{sep}_{\text{small}(\text{tw}(G))}(G, I)$. Then $S \setminus I \neq \emptyset$, $|S| \leq \text{large}(\text{tw}(G))$, and for every component C_i of $G \setminus S$ with its graph with interface $(G_i, I_i) := (G[V(C_i) \cup N_G(V(C_i))], N_G(V(C_i)))$

- (1) $G[I_i]$ is not a clique, and
- (2) $|I_i| \leq \text{medium}(\text{tw}(G))$.

For our purpose, we also need to ensure computability of $\text{sep}_s(G, I)$ in logspace. The following proposition follows from the definition of $\text{sep}_s(G, I)$, and the constant bound on the tree width of the given graphs. In this case, to compute $\text{sep}_s(G, I)$, we only need to enumerate vertex sets of constant size combined with reachability queries in undirected graphs. Moreover, for $s \in \mathbb{N}$, we know that $(G, I) \mapsto \text{sep}_s(G, I)$ is isomorphism-invariant by definition.

PROPOSITION 4.3. *For every $k \in \mathbb{N}$, there is a logspace DTM that, given a graph with interface (G, I) where $\text{tw}(G) \leq k$ and $s \in \mathbb{N}$, outputs $\text{sep}_s(G, I)$.*

4.2 Constructing Isomorphism-Invariant Tree Decompositions

To construct isomorphism-invariant tree decompositions using the previously defined (isomorphism-invariant) separators for graphs, we encapsulate their recursive computation using the concept of descriptor decompositions from Elberfeld et al. (2010). We slightly adjust the terminology from Elberfeld et al. (2010) by using graphs with interfaces directly instead of using descriptors.

A *descriptor decomposition* for a graph G is a pair (M, \mathcal{R}) consisting of a directed graph M and a collection \mathcal{R} of subgraphs with interfaces $R_n = (H, I)$ for every node $n \in V(M)$, where $(V(G) \setminus (V(H) \setminus I), V(H))$ is a separator in G . Beside this, (M, \mathcal{R}) contains a *root node* r with $R_r = (G, I)$ for some I . Moreover, for every node $n \in V(M)$ with $R_n = (H, I)$ and children n_1, \dots, n_m of n in M with $R_{n_i} = (H_i, I_i)$ for $i \in [m]$ the following properties hold:

- (1) for each (H_i, I_i) , we have $V(H_i) \subseteq V(H)$ and $(V(H_i) \setminus I_i) \subseteq (V(H) \setminus I)$ and at least one inclusion is proper,
- (2) for each (H_i, I_i) , $V(H_i)$ contains at least one vertex of $V(H) \setminus I$,
- (3) for all distinct (H_i, I_i) and (H_j, I_j) , $(V(H_i) \setminus I_i) \cap (V(H_j) \setminus I_j) = \emptyset$, and
- (4) each edge of H is present in $G[I]$ or some H_i .

Descriptor decompositions contain tree decompositions in the following way (Elberfeld et al. 2010, Lemma III.4). Given a descriptor decomposition (M, \mathcal{R}) rooted at $r \in V(M)$, the subgraph of M reachable from r is a tree T that can be turned into a tree decomposition (T, \mathcal{B}) by setting B_n for each $n \in V(T)$ to be the union of the interface I of $R_n = (H, I)$ and all vertices x that are in interfaces of at least two of the I, I_1, \dots, I_m . The *width* of (M, \mathcal{R}) is the width of (T, \mathcal{B}) .

Mapping a descriptor decompositions (M, \mathcal{R}) to its tree decompositions (T, \mathcal{B}) is isomorphism-invariant. Moreover, from Elberfeld et al. (2010, Lemma III.5), we know that turning descriptor decompositions into their tree decompositions is logspace-computable. In the light of these facts, all we need to do to, finally, to prove Lemma 4.1 is to construct an isomorphism-invariant descriptor decomposition $(M, \mathcal{R})_{\text{inv}}$ of a bounded (approximate) width $k' \in \mathbb{N}$.

LEMMA 4.4. *For every $k \in \mathbb{N}$, there is a $k' \in \mathbb{N}$ and a logspace-computable and isomorphism-invariant mapping that turns every graph G with a distinguished non-edge $\{u, v\} \notin E(G)$, where G*

- (1) *has tree width at most k ,*
- (2) *is an atom, and*
- (3) *is improved,*

into a width- k' descriptor decomposition (M, \mathcal{R}) .

PROOF. Let $\text{small}, \text{medium}, \text{large} : \mathbb{N} \rightarrow \mathbb{N}$ be the functions satisfying Fact 4.2. We consider the directed graph M whose nodes correspond to all subgraphs with interfaces (H, I) , where $(V(G) \setminus (V(H) \setminus I), V(H))$ is a separation in G with separator size $|I| \leq \text{medium}(\text{tw}(G))$. We insert an edge from a node n to a node n' if the graph with interface of n' arises (as a component) by applying Fact 4.2 to the one of n . In addition, we insert an edge from n to a node $(G[S], S)$, which represents the corresponding separator S . We declare the graph with interface (G, \emptyset) to be the root r of (M, \mathcal{R}) and, in addition, connect it to all $(G[C \cup \{u, v\}], \{u, v\})$, where $C \subseteq V(G)$ is the vertex set of a component of $G - \{u, v\}$.

For a constant bound on the tree width, constructing (M, \mathcal{R}) can be done by iterating over all candidate subgraphs with interfaces of a given graph G and using Proposition 4.3 to construct the corresponding separator and connecting it with the children.

To show that (M, \mathcal{R}) is a descriptor decomposition, we first observe that it has a root node r where R_r is G with interface $\{u, v\}$. Moreover, we need to check Properties (1) to (4) of descriptor decompositions: Property (1) follows from the fact that each separator covers the interface I and extends it. Properties (2) and (3) follow from the fact that we always consider nonempty components that are disjoint, respectively. Every edge is contained in $G[I]$, in $G[S]$, or in a component. The edges that are only in $G[S]$ are covered by $(G[S], S)$, which ensures Property (4). The bound on the width follows from the definition of S . \square

PROOF OF LEMMA 4.1. To prove the lemma, we first apply Lemma 4.4 to construct a descriptor decomposition (M, \mathcal{R}) that is isomorphism-invariant and has a bounded width. We turn it into a tree decomposition (T, \mathcal{B}) of the same width as discussed above, which is an isomorphism-invariant mapping. Thus, the combined mapping from G with distinguished pair $\{u, v\}$ to the tree decomposition is isomorphism-invariant as well. \square

5 ISOMORPHISM-BASED ORDERING OF NESTED TREE DECOMPOSITIONS

Nested tree decompositions are tree decompositions whose parts are not just bags but where every bag is associated with a family of tree decompositions for the bag's torso. We use polynomial-size nested tree decompositions to represent exponential-size families of width-bounded tree decompositions that arise by replacing bags with tree decompositions from their families. To solve the isomorphism problem with the help of nested tree decompositions, we use a recursively defined weak ordering on pairs of graphs and nested tree decompositions. Incomparable elements in this weak ordering represent isomorphic graphs. (Recall that a weak ordering of a set is a linear ordering of equivalence classes of the set, where equivalence is defined as incomparability.)

In Section 5.1, we define nested tree decompositions. To define the ordering on nested tree decompositions in Section 5.3, we first define concepts related to weak orderings in Section 5.2.

5.1 Definition of Nested Tree Decompositions

A *nested (tree) decomposition* $\bar{D} = (T, \mathcal{B}, \mathcal{D})$ for a graph G consists of a tree decomposition (T, \mathcal{B}) for G , and a family $\mathcal{D} = (\mathcal{D}_n)_{n \in V(T)}$ where every \mathcal{D}_n is a family of tree decompositions $D \in \mathcal{D}_n$ for the torso of n . Normal tree decompositions can be viewed as nested decompositions where

\mathcal{D}_n is empty for every $n \in V(T)$. We adjust some terminology that usually applies to tree decompositions for the use with nested decompositions. Let $\bar{D} = (T, \mathcal{B}, \mathcal{D})$ be a nested decomposition. The definition of the *width* of a bag B_n in a nested decomposition depends on whether \mathcal{D}_n is empty or contains a set of tree decompositions. If $|\mathcal{D}_n| = 0$, then we set $\text{tw}(B_n) := |B_n| - 1$ and $\text{tw}(B_n) := \max\{\text{tw}(D) \mid D \in \mathcal{D}_n\}$, otherwise. The *width* of \bar{D} is $\text{tw}(\bar{D}) := \max\{\text{tw}(B_n) \mid n \in V(T)\}$. The *size* of \bar{D} is $|\bar{D}| := \sum_{n \in V(T)} (1 + \max\{|D| + 1 \mid D \in \mathcal{D}_n\})$, where $|\mathcal{D}_n| = 0$ implies $\max\{|D| + 1 \mid D \in \mathcal{D}_n\} = 0$.

An (*unordered*) *root set* M of a nested decomposition $\bar{D} = (T, \mathcal{B}, \mathcal{D})$ is a subset $M \subseteq B_r$ of the root bag B_r of \bar{D} with (1) $M = B_r$ in case $|\mathcal{D}_r| = 0$, and (2) every $D \in \mathcal{D}_r$ has a bag B with $M \subseteq B$ in case $|\mathcal{D}_r| > 0$. An *ordered root set* σ is an ordering of an unordered root set. A nested decomposition is shown in Figure 2 on the left.

Refining a nested decomposition $\bar{D} = (T, \mathcal{B}, \mathcal{D})$ with respect to a tree decomposition $D \in \mathcal{D}_r$ for the root $r \in V(T)$ and an ordered root set σ is done as follows. First, we decompose $G[B_r]$ using D . Then, for each child bag B_c of B_r in \bar{D} , we find the highest bag in D that contains the adhesion set $B_{\{r,c\}} = B_r \cap B_c$ and make B_c adjacent to it. A bag of this kind exists, since, by definition, D is a tree decomposition of the torso of B_r . We add a new bag containing the elements of σ . This bag is the new root of the obtained decomposition and adjacent to the highest bag in D that contains all elements of σ (in particular, this operation may change which bag of D is highest). The newly constructed nested decomposition is said to be the decomposition obtained by *refining* \bar{D} with respect to $D \in \mathcal{D}_r$ and denoted by $\bar{D}_{D,\sigma}$. Suppose $\bar{D}_{D,\sigma} = (T', \mathcal{B}', \mathcal{D}')$, then we still need to define \mathcal{D}' . The family \mathcal{D}' is defined to agree with \mathcal{D} on all unchanged bags (those in $V(T) \cap V(T')$) and for each newly added bag $n \in V(T') \setminus V(T)$, we let \mathcal{D}'_n be the empty set. The refinement of a nested decomposition is shown in Figure 2 on the right. The size of a nested decomposition decreases when it is refined. The reason for this is that the contribution of the root bag of \bar{D} to $|\bar{D}|$ is larger than $|D| + 1$, the total size of the newly added bags together with the new root bag. This means $|\bar{D}_{D,\sigma}| < |\bar{D}|$ holds. We use this property for proofs by induction.

To be able to distinguish original bags and bags from refining decompositions, we could mark the bags of D , which arise from the refinement step. We circumvent the need to mark the bags by assuming that the bags B_n with empty \mathcal{D}_n are exactly the marked ones. In turn, we require from all nested decompositions \bar{D} we consider that the set of bags B_n with empty \mathcal{D}_n form a connected subtree in \bar{D} that, if nonempty, contains the root.

PROPOSITION 5.1. *The mapping that turns a nested decomposition $\bar{D} = (T, \mathcal{B}, \mathcal{D})$ with decomposition $D \in \mathcal{D}_r$ and an ordered root set σ into $\bar{D}_{D,\sigma}$ is logspace-computable and isomorphism-invariant.*

5.2 Isomorphism-Based Ordering of Graphs with Vertex Sequences

To define the isomorphism-based ordering for nested decompositions, we review notions related to composed orderings and define an ordering of graphs with given vertex sequences.

Let $<$ be a *weak ordering* on a set M and $a \equiv a'$ denote that two elements $a, a' \in M$ are *incomparable* with respect to $<$. That means, neither $a < a'$ nor $a' < a$ holds. We define the *weak ordering on sequences* from $M^* := \cup_{n \in \mathbb{N}} M^n$ with respect to $<$ as follows. We set $a = a_1 \dots a_s < a'_1 \dots a'_t = a'$ for $a, a' \in M^*$ if $s < t$, or $s = t$ and there is an $i \in [s]$ with $a_i < a'_i$, while $a_j \equiv a'_j$ holds for every $j \in [i - 1]$. The *weak ordering on tuples* from $M_1 \times \dots \times M_k$ with respect to weak orderings $<_i$ for sets M_i , respectively, is defined in the same way except that tuples always have the same length. We denote it by $<_{(1,\dots,k)}$. We define a *weak ordering on finite subsets* of M by setting $M_1 < M_2$ for two finite $M_1, M_2 \subseteq M$ based on comparing the sequences we get by sorting their elements to be monotonically increasing with respect to $<$.

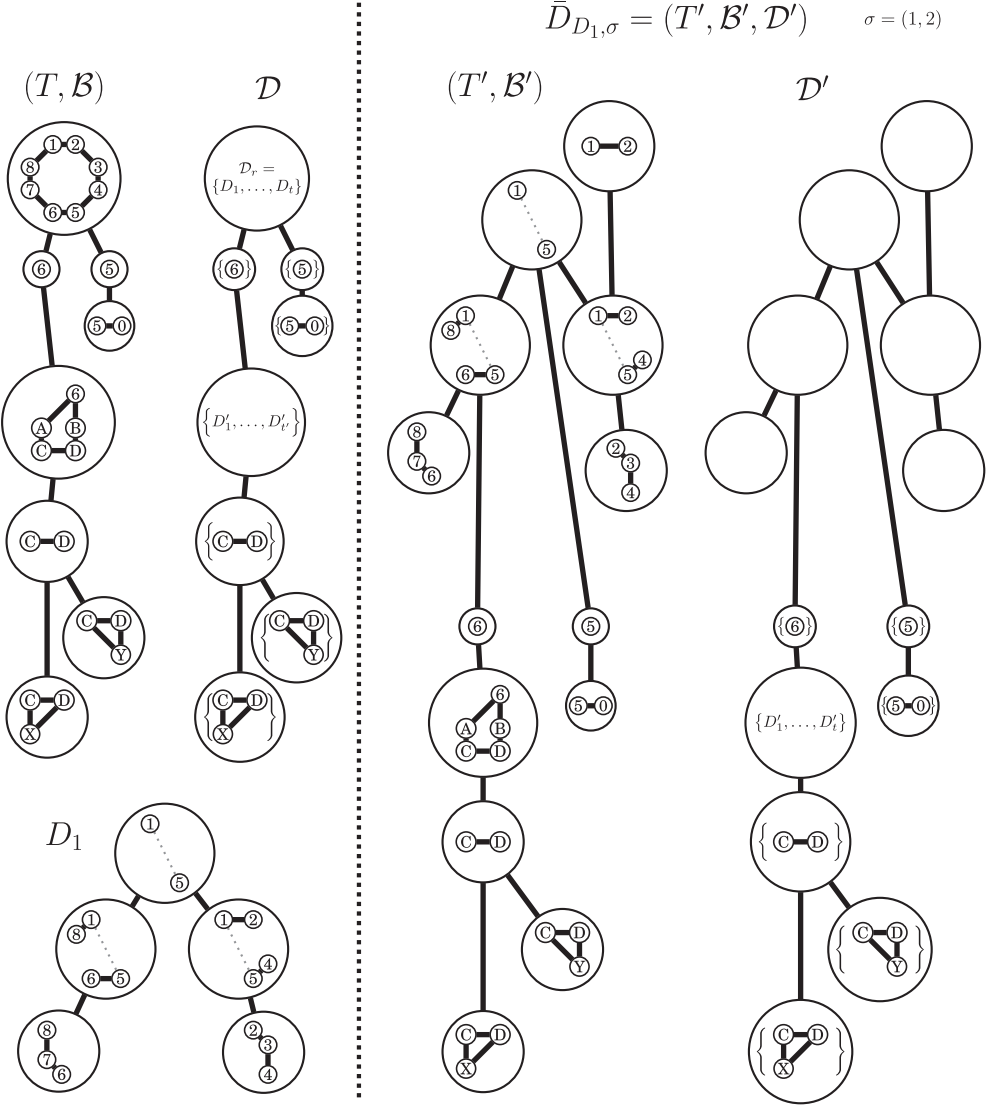


Fig. 2. The figure shows a nested decomposition on the left for the graph from Figure 1. One particular decomposition D_1 for the root bag is shown (bottom left). It corresponds to the distinguished non-edge $\{1, 5\}$ shown dotted. On the right, a refined decomposition is depicted which is obtained with ordered root set $(1, 2)$ and by refining the root bag with D_1 . In \mathcal{D}' , the bags \mathcal{B}'_n with $|\mathcal{D}'_n| = 0$ (upper half) are empty (i.e., marked). The new root bag, determined by σ , is the bag containing $\{1, 2\}$ at the top.

We write the *concatenation* of sequences σ and τ as $\sigma\tau$. Suppose that (G, σ) and (G', σ') are pairs consisting of graphs G and G' with sequences of vertices $\sigma = v_1 \dots v_s$ and $\sigma' = v'_1 \dots v'_t$ from the respective graphs. We set $(G, \sigma) <_{\text{seq}} (G', \sigma')$ if the sequence $\text{col}_G(v_1, v_1) \dots \text{col}_G(v_1, v_s) \text{col}_G(v_2, v_1) \dots \text{col}_G(v_s, v_1) \dots \text{col}_G(v_s, v_s)$ is smaller than the sequence $\text{col}_{G'}(v'_1, v'_1) \dots \text{col}_{G'}(v'_1, v'_t) \text{col}_{G'}(v'_2, v'_1) \dots \text{col}_{G'}(v'_t, v'_1) \dots \text{col}_{G'}(v'_t, v'_t)$ with respect to the (standard) ordering $<$ of \mathbb{N} . We write $(G, \sigma) \equiv_{\text{seq}} (G', \sigma')$ if (G, σ) and (G', σ') are *incomparable* with respect to $<_{\text{seq}}$.

The ordering $<_{\text{seq}}$ is logspace-computable by enumerating all pairs of vertices in lexicographic order of the indices.

Graphs G and G' are *isomorphic with respect to sequences of vertices* $\sigma = v_1 \dots v_s$ and $\sigma' = v'_1 \dots v'_t$ from the respective graphs if $s = t$ and there is an isomorphism φ from G to G' with $\varphi(v_i) = v'_i$ for every $i \in [s]$. We say that φ *respects* σ and σ' in this case. Based on this definition, we also consider *canons of graphs with respect to vertex sequences*.

Due to the following statement, which we immediately get from the definition, we call $<_{\text{seq}}$ an *isomorphism-based ordering of graphs with vertex sequences*.

PROPOSITION 5.2. *Let G and G' be graphs with sequences of vertices $\sigma = v_1 \dots v_s$ and $\sigma' = v'_1 \dots v'_t$ from the respective graphs.*

- (“invariance”-property.) *If G and G' are isomorphic with respect to σ and σ' , then $(G, \sigma) \equiv_{\text{seq}} (G', \sigma')$.*
- (“quasi-completeness”-property.) *If $(G, \sigma) \equiv_{\text{seq}} (G', \sigma')$, then $G[\{v_1, \dots, v_s\}]$ and $G'[\{v'_1, \dots, v'_t\}]$ are isomorphic with respect to σ and σ' .*

5.3 Isomorphism-Based Ordering of Graphs with Nested Tree Decompositions

We define an ordering of graphs with nested decompositions by recursively ordering the child decompositions and combining this with the root bags. If a root bag has no refining tree decompositions, then this is done by trying all possible orderings of the vertices of the bag. If the root bag has refining tree decompositions, then this is done by first refining it before going into recursion.

For each child c of the root node r of a nested decomposition $D = (T, \mathcal{B}, \mathcal{D})$, we define a set $\Pi(c)$ of orderings of a vertex set as follows. If $|\mathcal{D}_c| = 0$, then $\Pi(c)$ contains all orderings of the vertices of B_c . If $|\mathcal{D}_c| > 0$, then $\Pi(c)$ is the set of orderings of the adhesion set $B_{\{r, c\}} = B_r \cap B_c$. We use the sequences from $\Pi(c)$ as ordered root sets for the child decomposition of \bar{D} rooted at c .

For all tuples (G, \bar{D}, σ) and (G', \bar{D}', σ') of graphs with nested decompositions and ordered root sets, we define whether $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$ holds based on the following case distinction:

– (“size”-comparison.) If $|\bar{D}| < |\bar{D}'|$, or $|\bar{D}| = |\bar{D}'|$ and $|\mathcal{D}_r| < |\mathcal{D}'_r|$, then set $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$.

– (“bag”-comparison.) If $|\bar{D}| = |\bar{D}'| = 1$ (which implies $|\mathcal{D}_r| = |\mathcal{D}'_r| = 0$), then set $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$ if $(G, \sigma) <_{\text{seq}} (G', \sigma')$.

– (“recursive”-comparison.) If $|\bar{D}| = |\bar{D}'| > 1$, and $|\mathcal{D}_r| = |\mathcal{D}'_r| = 0$, then we compare the decompositions recursively. Let c_1, \dots, c_s be the children of r in \bar{D} with respective child decompositions $\bar{D}_1, \dots, \bar{D}_s$ and subgraphs G_1, \dots, G_s . Let c'_1, \dots, c'_t be the children of r' in \bar{D}' with respective child decompositions $\bar{D}'_1, \dots, \bar{D}'_t$ and subgraphs G'_1, \dots, G'_t . Set $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$ if the following relation holds, which compares sets of sets that contain tuples to which $<_{(\text{dec}, \text{seq})}$ applies directly:

$$\begin{aligned} & \{ \{ (G_i, \bar{D}_i, \tau), (G, \sigma\tau) \} \mid \tau \in \Pi(c_i) \} \mid i \in [s] \} \\ & <_{(\text{dec}, \text{seq})} \{ \{ (G'_i, \bar{D}'_i, \tau'), (G', \sigma'\tau') \} \mid \tau' \in \Pi(c'_i) \} \mid i \in [t] \}. \end{aligned}$$

– (“refinement”-comparison.) If $|\bar{D}| = |\bar{D}'| > 1$, and $|\mathcal{D}_r| = |\mathcal{D}'_r| > 0$, then set $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$ if $\{ (G, \bar{D}_{D, \sigma}, \sigma) \mid D \in \mathcal{D}_r \} <_{\text{dec}} \{ (G', \bar{D}'_{D', \sigma'}, \sigma') \mid D' \in \mathcal{D}'_r \}$ holds.

Graphs G and G' are *isomorphic with respect to nested decompositions* $\bar{D} = (T, \mathcal{B}, \mathcal{D})$ and $\bar{D}' = (T', \mathcal{B}', \mathcal{D}')$ as well as ordered root sets σ and σ' , respectively, if there exists an isomorphism φ from G to G' that

- (1) respects the (normal) tree decompositions (T, \mathcal{B}) and (T', \mathcal{B}') ,
- (2) respects the sequences σ and σ' , and

- (3) for every $n \in V(T)$ there is a bijection π_n from \mathcal{D}_n to $\mathcal{D}_{n'}$, such that φ restricted to B_n respects D and $\pi(D)$ for all $D \in \mathcal{D}_n$.

Based on how this definition refines the isomorphism equivalence relation among graphs, we consider *canons of graphs with respect to nested decompositions*.

We call $<_{\text{dec}}$ an *isomorphism-based ordering of graphs with nested decompositions*, which is justified by the following lemma.

LEMMA 5.3. *Let (G, \bar{D}, σ) and (G', \bar{D}', σ') be tuples consisting of graphs with respective nested decompositions and ordered root sets.*

- (“invariance”-property.) *If G and G' are isomorphic with respect to \bar{D} and \bar{D}' as well as σ and σ' , then $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$.*
- (“quasi-completeness”-property.) *If $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$, then G and G' are isomorphic with respect to σ and σ' .*

PROOF. We prove each property by induction on the sizes of $\bar{D} = (T, \mathcal{B})$ and $\bar{D}' = (T', \mathcal{B}')$.

(Proof of the “invariance”-property.) Let φ be an isomorphism from G to G' respecting \bar{D} and \bar{D}' as well as σ and σ' . From the above definition, we know that φ respects the normal tree decompositions (T, \mathcal{B}) and (T', \mathcal{B}') for G and G' , respectively, via some isomorphism $\psi: V(T) \rightarrow V(T')$.

First, this implies $|\bar{D}| = |\bar{D}'|$ as well as $|\mathcal{D}_r| = |\mathcal{D}'_{r'}|$ and, hence, the “size”-comparison does not distinguish (G, \bar{D}, σ) and (G', \bar{D}', σ') . If $|\bar{D}| = |\bar{D}'| = 1$, then we deal with the “bag”-comparison. Since φ respects σ and σ' , Proposition 5.2 implies $(G, \sigma) \equiv_{\text{seq}} (G', \sigma')$. In turn, this implies $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$.

If $|\bar{D}| = |\bar{D}'| > 1$ and $|\mathcal{D}_r| = |\mathcal{D}'_{r'}| = 0$, then we are dealing with the “recursive”-case. Let c_i be a child of r and consider the (isomorphic) child $c'_j = \psi(c_i)$ of r' . Moreover, consider an ordering $\tau \in \Pi(c_i)$ and the (isomorphic) ordering $\tau' = \varphi(\tau) \in \Pi(c'_j)$. By construction of τ and τ' , we know $(G, \sigma\tau) \equiv_{\text{seq}} (G', \sigma'\tau')$, and by applying the induction hypothesis, we also know $(G_i, \bar{D}_i, \tau) \equiv_{\text{dec}} (G'_j, \bar{D}'_j, \tau')$. Since this observation holds for all children c_i of r and all $\tau \in \Pi(c_i)$, we have $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$.

If $|\bar{D}| = |\bar{D}'| > 1$ and $|\mathcal{D}_r| = |\mathcal{D}'_{r'}| > 0$, then we deal with the “refinement”-case. We know that there exists a bijection $\pi = \pi_r$ from \mathcal{D}_r to $\mathcal{D}'_{r'}$, such that φ restricted to the vertices from B_r and $B'_{r'}$ respects every pair of tree decompositions $D \in \mathcal{D}_r$ and $\pi(D) \in \mathcal{D}'_{r'}$ via some isomorphism ψ_D . We claim that G and G' are also isomorphic with respect to each pair of refined nested decompositions $\bar{D}_{D, \sigma}$ and $\bar{D}_{\pi(D), \sigma'}$ as well as ordered root sets σ and σ' . Since the size of nested decompositions decreases when refining them, this claim implies $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$ by induction. To prove it, we start to use the above isomorphism φ from G to G' . We construct an isomorphism ρ from the tree $T_{D, \sigma}$ underlying $\bar{D}_{D, \sigma}$ to the tree $T'_{\pi(D), \sigma'}$ underlying $\bar{D}'_{\pi(D), \sigma'}$ as follows. The newly established root node of $T_{D, \sigma}$, whose bag consists of the vertices of σ , is mapped to the newly established root node of $T'_{\pi(D), \sigma'}$, whose bag consists of the vertices of σ' . Every other node is mapped according to either ψ or ψ_D depending on whether it is a tree node of either \bar{D} or D , respectively. The only property we need to show is that ρ preserves the newly established edges, which lie between nodes from D and \bar{D} as well as nodes from $\pi(D)$ and \bar{D}' . Let m be a node of D that gets connected to a node c of \bar{D} during the refinement process. Then m is the highest bag that contains all vertices of $B_r \cap B_c$. Due to the definition of ψ and ψ_D , we know that $\psi_D(m)$ is a bag in $\pi(D)$ that contains all vertices of $B_{\psi(r)} \cap B_{\psi(c)}$. Thus, there is also an edge from $\psi_D(m)$ to $\psi(c)$ in $\bar{D}'_{\pi(D), \sigma'}$. This proves the claim.

(Proof of the “quasi-completeness”-property.) For this direction, assume $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$ holds. This implies $|\bar{D}| = |\bar{D}'|$ and $|\mathcal{D}_r| = |\mathcal{D}'_r|$. If, in addition, we have $|\bar{D}| = |\bar{D}'| = 1$, then the statement follows from Proposition 5.2.

If $|\bar{D}| = |\bar{D}'| > 1$ and $|\mathcal{D}_r| = |\mathcal{D}'_r| = 0$, then we are in the “recursive”-comparison. Thus, we can choose a bijection π from $[s]$ to $[t]$ satisfying for each $i \in [s]$:

$$\{((G_i, \bar{D}_i, \tau), (G, \sigma\tau)) \mid \tau \in \Pi(c_i)\} \equiv_{(\text{dec}, \text{seq})} \{((G_{\pi(i)}, \bar{D}_{\pi(i)}, \tau'), (G', \sigma\tau')) \mid \tau' \in \Pi(c_{\pi(i)})\}.$$

By induction, we can choose for each $i \in [s]$ orderings $\tau_i \in \Pi(c_i)$ and $\tau'_{\pi(i)} \in \Pi(c_{\pi(i)})$, such that there is an isomorphism from the graph G_i decomposed by \bar{D}_i to the graph $G'_{\pi(i)}$ decomposed by $\bar{D}'_{\pi(i)}$ that respects τ_i and $\tau'_{\pi(i)}$. If subgraphs G_i and G_j of G that correspond to child decompositions \bar{D}_i and \bar{D}_j , respectively, contain a common vertex, then this vertex appears in the root bag B_r (due to the connectedness property of tree decompositions) and thus in σ , since $|\mathcal{D}_r| = 0$ and thus the ordered root set σ contains all vertices of B_r . Moreover, the same property holds for the same kind of subgraphs of G and σ' . Since $(G, \sigma\tau_i) \equiv_{\text{seq}} (G', \sigma'\tau'_{\pi(i)})$, the mapping of common vertices agrees with the isomorphisms chosen for G_i and G_j . Thus, we can find a common extension to map G to G' that respects σ and σ' .

If $|\bar{D}| = |\bar{D}'| > 1$ and $|\mathcal{D}_r| = |\mathcal{D}'_r| > 0$, then we are in the “refinement”-comparison and know that $\{(G, \bar{D}_{D, \sigma}, \sigma) \mid D \in \mathcal{D}_r\} \equiv_{\text{dec}} \{(G', \bar{D}'_{D', \sigma'}, \sigma') \mid D' \in \mathcal{D}'_r\}$ holds. Since the size of nested decompositions decreases when refining them, we know by induction that G and G' are isomorphic with respect to σ and σ' . \square

Remark 5.4. The ordering $<_{\text{dec}}$ is defined to satisfy the “quasi-completeness”-property stated in Lemma 5.3, but not a “completeness”-property saying that $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$ implies that G and G' are isomorphic with respect to σ and σ' as well as \bar{D} and \bar{D}' , too. The reason behind this lies in the fact that deciding an ordering of this kind for nested decompositions of a bounded width is as hard as the (general) graph isomorphism problem. (In fact, this even holds in the case of, more restrictive, p -bounded decompositions as defined in Section 6.) This can be seen by the following reduction. Take two graphs G and H for which we want to know whether they are isomorphic. Consider now two (empty) graphs G' and H' with $V(G') = V(G)$, $V(H') = V(H)$, and $E(G') = E(H') = \emptyset$. For G' , construct a nested decomposition by starting with a single bag $B = V(G')$. Then, for each edge $\{v, w\} \in E(G)$, construct a refining decomposition whose tree is a star graph where the center bag equals $\{v, w\}$ and the adjacent bags each contain a single vertex of G' . The nested decomposition for H' is constructed in the same way. Since the refining decompositions exactly encode the edges of the respective graphs, G and H are isomorphic exactly if G' and H' are isomorphic with respect to their nested decompositions. Thus, if \equiv_{dec} were defined as to be in exact correspondence with isomorphisms respecting nested decompositions, then deciding whether $G' \equiv_{\text{dec}} H'$ would be graph isomorphism complete even on graphs without edges.

6 COMPUTING THE ORDERING FOR NESTED TREE DECOMPOSITIONS

We now investigate methods to space-efficiently evaluate the isomorphism based ordering described in the previous section. The nested decompositions we are working with always have a bounded width. This makes it possible to implement the “recursive”-comparison of the isomorphism-based ordering space-efficiently. If the child decompositions are small enough (more precisely, they are smaller by a constant fraction in comparison to their parent), then it is possible to store a constant amount of information, and in particular to store orderings of the size-bounded root bag, before descending into recursion, without exceeding a desired logarithmic space bound. If there is a large child decomposition, of which there can be only one, then we can use Lindell’s

classic technique of precomputing the recursive information before storing anything at all. However, for the “refinement”-comparison, a space-efficient approach turns out to be more challenging. In this case, the ordering asks us to compare various refinements of the root bag. Cycling through these refinements as part of a recursive approach requires too much space, even if the number of decompositions is bounded by a polynomial in the size of the root bag. While it is not clear how to remedy this difficulty in general, the nested decompositions we construct in the proofs of our main theorems satisfy an additional technical condition, called p -boundedness below. This makes it possible to find a trade-off between the recursive space requirement and the space required for cycling through the refinements.

Let \bar{D} be a nested decomposition. Consider a bag n with $|\mathcal{D}_n| > 1$. Let c_1, \dots, c_t be the children of n sorted by monotonically decreasing size of the respecting subdecompositions D_1, \dots, D_t . If it exists, then let $j \in [t]$ be maximal such that $G[A_n]$ with

$$A_n := (B_n \cap B_{c_1}) \cup \dots \cup (B_n \cap B_{c_j})$$

is a clique, and $|D_j| > |D_{j+1}|$ holds or $j = t$ holds. Otherwise, set $j := 0$ and $A_n := \emptyset$. We call the children c_1, \dots, c_j of n the *special children* and A_n is the *attachment clique of the special children*. A nested decomposition \bar{D} is p -bounded for a polynomial $p : \mathbb{N} \rightarrow \mathbb{N}$ if for every $n \in V(T)$ and non-special child c of n we have $|\mathcal{D}_n| \leq p(|\bar{D}|/|\bar{D}_c|)$. For non-special nodes, we use the p -boundedness condition to trade the number of candidate refining decompositions against the size of subdecompositions. This enables an overall space-efficient recursion.

LEMMA 6.1. *For every $k \in \mathbb{N}$ and polynomial $p : \mathbb{N} \rightarrow \mathbb{N}$, there is a logspace DTM that, given as input graphs G and G' along with respective nested decompositions \bar{D} and \bar{D}' and ordered root sets σ and σ' where \bar{D} and \bar{D}'*

- (1) *have width at most k , and*
- (2) *are p -bounded,*

decides $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$.

In Section 6.1, we review a technique of Lindell (1992) used to compute (composed) weak orderings on sets space-efficiently. Section 6.2 contains the proof of Lemma 6.1.

6.1 Comparing Sets via Cross Comparing Elements

We repeatedly apply a technique of Lindell (1992) to compare two sets A and A' with respect to a weak ordering $<$ defined for their elements. Comparing A and A' is performed by repeatedly comparing a single element of A with a single element of A' . Such a comparison is called a *cross comparison*. To apply the technique subsequently, we state it as an abstract fact as follows. Let $<$ be a weak ordering for elements of a set M and $A, A' \subseteq M$ finite subsets of M . The *cross comparison matrix* of $A = \{a_1, \dots, a_s\}$ and $A' = \{a'_1, \dots, a'_t\}$ with respect to $<$ is the binary matrix $C_{A,A'} \in \{0, 1\}^{s \times t}$ with $C_{A,A'}[i, j] = 1$ exactly if $a_i < a'_j$.

FACT 6.2. *Let $<$ be a weak ordering of elements of a set M . There is a logspace DTM that, given the cross comparison matrix $C_{A,A'}$ for sets $A, A' \subseteq M$ (which are not part of the input), decides $A < A'$.*

To apply Fact 6.2, it is important to observe that inputs to its DTM consist only of the cross comparison matrix without encodings of the sets A and A' . Thus, the space used by the machine is in $O(\log(|A| \cdot |A'|))$. The fact can be used to build an algorithm for comparing sets on top of an algorithm for comparing individual elements.

PROPOSITION 6.3. *Let $<$ be a weak ordering of elements of a set M that can be decided by a DTM in space at most $s(a, a')$ for every $a, a' \in M$. There is a DTM that, given sets $A, A' \subseteq M$, decides $A < A'$ in space $O(\log(|A| \cdot |A'|) + \max\{s(a, a') \mid a \in A \text{ and } a' \in A'\})$.*

We apply the proposition in a scenario where the weak ordering $<$ is partially known: A weak ordering $<'$ is *coarser* than a weak ordering $<$ if $a_1 <' a_2$ implies $a_1 < a_2$. In our application, the weak ordering $<'$ compares nested subdecompositions based on their sizes, which can be done during a logspace reduction before the more challenging recursive computation starts.

6.2 Proof of Lemma 6.1

A child decomposition \bar{D}_i of a nested decomposition \bar{D} is *large* if $|\bar{D}_i| > |\bar{D}|/2$; the size bound implies that every nested decomposition has at most one large child.

PROOF OF LEMMA 6.1 The logspace procedure $\text{COMPARISON}(\cdot, \cdot)$ we design implements the recursive definition of $<_{\text{dec}}$. To streamline the recursion, it is more convenient to perform a computational task that is slightly more general than the one required by the lemma. Given input graphs G and G' with nested tree decompositions \bar{D} and \bar{D}' and unordered root sets M and M' (not ordered root sets σ and σ' as described by the lemma), the output of $\text{COMPARISON}((G, \bar{D}, M), (G', \bar{D}', M'))$ is the cross comparison matrix of the sets $\{(G, \bar{D}, \sigma) \mid \sigma \text{ is an ordering of } M\}$ and $\{(G', \bar{D}', \sigma') \mid \sigma' \text{ is an ordering of } M'\}$ with respect to $<_{\text{dec}}$.

The recursive procedure starts at the root bags of both decompositions and descends into them to compute the recursively defined ordering. To have access to the current positions in the decompositions, we do not use a stack, but maintain *node pointers* to the current node and the previous node of the recursive process in each decomposition. These pointers direct us to nodes from the coarser, nested decompositions as well as to nodes from refining decompositions. They only require a logarithmic amount of space. To be able to reconstruct vertex sequences as well as already refined parts of the decomposition, we store sequences of vertices in a bag relative to the bag using the pointers. Thus, to store an ordering of a bag of bounded size, we only require a constant amount of space, provided we have the pointer to the bag at hand.

To analyze the space requirement apart from these pointers, we use two separate tapes. A *decomposition* tape is used to store data related to the “recursive”-comparison and a *refinement* tape is used to store data related to the “refinement”-comparison. We prove separately for each tape that the used space is bounded by $O(\log(|\bar{D}| + |\bar{D}'|))$.

The procedure closely follows the definition of the isomorphism-based ordering and, in particular, distinguishes the same cases.

(“size”-comparison.) By counting refining decompositions and nodes in decompositions, we can determine whether $|\bar{D}| \neq |\bar{D}'|$ or $|\mathcal{D}_r| \neq |\mathcal{D}'_r|$ and in either case directly decide $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$ for all orderings σ and σ' of M and M' , respectively.

(“bag”-comparison.) If $|\bar{D}| = |\bar{D}'| = 1$ (and, thus, $|\mathcal{D}_r| = |\mathcal{D}'_r| = 0$), then we decide $(G, \bar{D}, \sigma) <_{\text{dec}} (G', \bar{D}', \sigma')$ for all orderings σ and σ' of M and M' , respectively, based on the definition of $<_{\text{seq}}$ in logspace.

(“recursive”-comparison.) If $|\bar{D}| = |\bar{D}'| > 1$ and $|\mathcal{D}_r| = |\mathcal{D}'_r| = 0$, then we handle large and non-large children differently.

If \bar{D} has a large child decomposition \bar{D}_i and \bar{D}' has a large child decomposition \bar{D}_j , then we first compute the comparison matrix of the sets $\{(G_i, \bar{D}_i, \sigma) \mid \sigma \in \Pi(c_i)\}$ and $\{(G'_j, \bar{D}'_j, \sigma') \mid \sigma' \in \Pi(c'_j)\}$. The descent into the large children can be performed without storing data on the decomposition tape by just updating the node pointers. The comparison matrix of (constant) size $s_{\text{large}} \in O((k!)^2)$ is stored on the decomposition tape.

After handling large children, we continue to compare $\{((G_i, \bar{D}_i, \tau), \sigma\tau) \mid \tau \in \Pi(c_i)\} \mid i \in [s]\}$ and $\{((G'_j, \bar{D}'_j, \tau'), \sigma'\tau') \mid \tau' \in \Pi(c'_j)\} \mid j \in [t]\}$ with respect to $<_{(\text{dec}, \text{seq})}$. To apply Proposition 6.3 with a smaller space requirement, we first define a weak order $<_{(\text{dec}, \text{seq})}'$ that is coarser than $<_{(\text{dec}, \text{seq})}$ by setting $\{((G_i, \bar{D}_i, \tau), \sigma\tau) \mid \tau \in \Pi(c_i)\} <_{(\text{dec}, \text{seq})}' \{((G'_j, \bar{D}'_j, \tau'), \sigma'\tau') \mid \tau' \in \Pi(c'_j)\}$ if $|\bar{D}_i| < |\bar{D}'_j|$. Ordering the child decompositions with respect to $<$ can now be done by first ordering with respect to $<'$ with higher priority and, then, applying the cross comparison idea behind Proposition 6.3 to pairs whose child decompositions are of the same size. Moreover, we do not need to recurse on large children. Note that to compare $\{((G_i, \bar{D}_i, \tau), \sigma\tau) \mid \tau \in \Pi(c_i)\}$ and $\{((G'_j, \bar{D}'_j, \tau'), \sigma'\tau') \mid \tau' \in \Pi(c'_j)\}$ with respect to $<_{(\text{dec}, \text{seq})}$, it suffices to know the result of the call $\text{COMPARISON}((G_i, \bar{D}_i, N), (G'_j, \bar{D}'_j, N'))$, where N and N' denote the unordered sets of the ordered root sets in $\Pi(c_i)$ and $\Pi(c'_j)$, respectively.

We investigate the space requirement of the method. We first observe that there are at most $\ell \leq |\bar{D}|/m$ child decompositions of size m . Moreover $m \leq |\bar{D}|/2$ for non-large children and, thus, $m \leq \min\{|\bar{D}|/2, |\bar{D}|/\ell\}$. By Fact 6.2, we can compare sets of size at most ℓ using $O(\log(\ell^2))$ space, plus the space required for the recursion, which is at most $O(\min\{\log(|\bar{D}|/2), \log(|\bar{D}|/\ell)\})$ by induction. The current orderings σ and σ' are stored using space $s_{\text{stack}} \in O((k!)^2)$ on the decomposition tape. (Recall that sequences are stored relative to pointers to nodes of the decomposition as described above.) Defining $s_{\text{dec}}(n)$ to be the space requirement for the recursive comparison of decompositions of size n where $s_{\text{dec}}(1)$ depends on the constant k , we have

$$s_{\text{dec}}(n) \leq s_{\text{large}} + \max_{2 \leq \ell \leq n} \{s_{\text{stack}} + O(\log(\ell^2)) + s_{\text{dec}}(\lfloor n/\ell \rfloor)\} \in O(\log n).$$

(“refinement”-comparison.) In case $|\bar{D}| = |\bar{D}'| > 1$ and $|\mathcal{D}_r| = |\mathcal{D}'_r| > 0$, we need to describe how our procedure handles the recursive refinement of the root bag. In the simplest case, if $|\mathcal{D}_r| = |\mathcal{D}'_r| = 1$ holds, we choose the unique refinement and recurse without using any space on the refinement tape. If there are multiple refining decompositions for the roots, then we first handle the special children before taking refining decompositions into account.

For the special children, we compute and store all information about comparisons between them that might ever be required subsequently as follows. Let C and C' be the attachment cliques of the special child decompositions of \bar{D} and \bar{D}' , respectively. Consider arbitrary $P \subseteq \text{pow}(C)$ and $P' \subseteq \text{pow}(C')$, where $\text{pow}(\cdot)$ is the power set operator. Let τ be an ordering of C and τ' be an ordering of C' . Given τ and τ' , we want to compare the special children whose adhesion set is in P or P' . Let $Z_P = (G_P, \bar{D}_P, C)$ be defined as follows. G_P is the subgraph of G induced by the vertices in C and all vertices contained in a special child of \bar{D} with an adhesion set in P and \bar{D}_P is the decomposition of G_P obtained by replacing the root of \bar{D} with the set C and out of the child decompositions of the root only maintaining the special child decompositions with adhesion sets in P . Note that Z_P is a graph with a nested decomposition and unordered root set. We define $Z_{P'}$ similarly for G' . With this definition, we compute and store all cross comparison matrices $\text{COMPARISON}(Z_P, Z_{P'})$ for all choices of $P \subseteq \text{pow}(C)$ and $P' \subseteq \text{pow}(C')$.

Since the sets have only bounded size, the entire outcome of the computation can be stored using (constant) space. More precisely, there are at most 2^{2^k} choices for P and at most equally many for P' and each comparison matrix has size at most $O((k!)^2)$, so overall we require at most $s_{\text{clique}} \in O(2^{2^{2k}} (k!)^2)$ space. The information will be stored on the refinement tape. To store information vertices in C and C' , we only ever store relative indices. For example, to store the i th vertex of C with respect to input ordering of vertices, we store the relative index i . Since we can determine in logspace whether a vertex is contained in C by iterating over all vertices, we can recover the vertex. All information about C and C' is stored in such relative indices.

We argue that all of this information can be computed recursively without exceeding the logarithmic space bound. Indeed, there is at most one large child decomposition \bar{D}_L of \bar{D} with graph G_L and at most one large child decomposition \bar{D}'_L of \bar{D}' with graph G'_L . Before using any space, we first compute the recursive call $\text{COMPARISON}((G_L, \bar{D}_L, M_L), (G'_L, \bar{D}'_L, M'_L))$, where M_L and M'_L are the unordered root sets of \bar{D}_L and \bar{D}'_L , respectively. The result is stored using a constant amount of space on the refinement tape. We then compute for all choices of $P \subseteq \text{pow}(C)$ and $P' \subseteq \text{pow}(C')$ the result of $\text{COMPARISON}(Z_P, Z'_{P'})$. Recall that s_{clique} denotes the amount of refinement space required to store the entire outcome. Since the size of every non-large child decompositions of Z_P is at most $|\bar{D}|/2$, we obtain a recursion for the space satisfying

$$s_{\text{refine}}(n) \leq s_{\text{clique}} + s_{\text{refine}}(\lfloor n/2 \rfloor) \in O(\log n).$$

Having computed $\text{COMPARISON}(Z_P, Z'_{P'})$ for all choices of $P \subseteq \text{pow}(C)$ and $P' \subseteq \text{pow}(C')$, our goal is now to compare the sets $A_\sigma = \{(G, \bar{D}_{D,\sigma}, \sigma) \mid D \in \mathcal{D}_r\}$ and $A'_{\sigma'} = \{(G', \bar{D}'_{D',\sigma'}, \sigma') \mid D' \in \mathcal{D}'_{r'}\}$ for all orderings σ and σ' of M and M' , respectively. Using Fact 6.2, we can compare these sets using $O(\log(|\mathcal{D}_r| \cdot |\mathcal{D}'_{r'}|))$ space, which we will write on the refinement tape, in addition to the recursive space required to compare two elements $(G, \bar{D}_{D,\sigma}, \sigma)$ and $(G, \bar{D}'_{D',\sigma'}, \sigma')$. For two such elements, whenever we would go into recursion, if a recursive result is contained in the precomputed information for special children, we will not go into recursion and rather use the precomputed information. Note that this means that we will never have to recursively descend into a special child of \bar{D} or \bar{D}' again. This observation uses the fact that in \bar{D} the set of bags n with $|\mathcal{D}_n| > 0$ (and similarly in \bar{D}') forms a connected subtree containing the root.

For the refinement space consumption, we now note the following. Since comparisons of special children are precomputed, due to the p -boundedness for every subsequent recursive call, the size of the decompositions of the recursive call is at most $\max\{|\bar{D}|/|\mathcal{D}_r|, |\bar{D}'|/|\mathcal{D}'_{r'}|\}$. Thus, we obtain a recursion for the space bound of

$$s_{\text{refine}}(n) \leq s_{\text{clique}} + \max_{2 \leq \ell \leq n} \{O(\log(\ell^2) + s_{\text{refine}}(\lfloor n/\ell \rfloor))\} \in O(\log n).$$

Thus, the total space requirement of our procedure is logarithmic. \square

7 TESTING ISOMORPHISM AND CANONIZING BOUNDED TREE WIDTH GRAPHS

We show how to compute isomorphism-invariant width-bounded and p -bounded nested decompositions for graphs of bounded tree width and, then, apply this to prove Theorems 1.1 and 1.2.

To compute nested decompositions, we combine the decomposition into atoms described in Section 3 with the decomposition of atoms into width-bounded tree decompositions described in Section 4.

LEMMA 7.1. *For every $k \in \mathbb{N}$, there is a $k' \in \mathbb{N}$, a polynomial $p : \mathbb{N} \rightarrow \mathbb{N}$, and a logspace-computable and isomorphism-invariant mapping that turns every graph G of tree width at most k into a nested decomposition \bar{D} for G that*

- (1) *has width at most k' , and*
- (2) *is p -bounded.*

PROOF. Instead of the original input graph G , we work with its improved version, which we can compute in logspace, since the tree width of G is bounded by k . Mapping the input graph to its improved version is isomorphism-invariant and the improved version has exactly the same tree decompositions. In the following, we denote the improved version of the input graph by G .

Let $D = (T, \mathcal{B})$ be the isomorphism-invariant tree decomposition we get from G by applying Lemma 3.1. Since the lemma guarantees that in D the adhesion sets are cliques, the torso of each

bag is equal to the bag itself. To turn D into a nested decomposition it thus suffices to find a family of tree decompositions of width at most k for each bag. We will apply Lemma 4.1 to find such a family. Since D decomposes an improved graph and the adhesion sets are cliques, every $G[B_n]$ for $n \in V(T)$ is also improved.

Thus, based on D , we construct a nested decomposition \bar{D} by considering every node n of D and defining an isomorphism-invariant family \mathcal{D}_n of tree decompositions of the bag B_n . If B_n has size at most $k + 1$, then we let the family \mathcal{D}_n consist of a single tree decomposition that is just B_n . Note that by this choice, the bag B_n satisfies both the width bounded and the p -boundedness restriction (for every polynomial p with $p(i) \geq 1$ for all $i \in \mathbb{N}$). If the size of B_n exceeds $k + 1$, then we would like to apply Lemma 4.1 to further decompose B_n . However, for the lemma, we need a pair $\{u, v\} \notin E(G)$ in B_n to serve as the root of the decomposition. We cannot simply iterate over all $\{u, v\} \notin E(G)$ in B_n , since the result may violate the p -boundedness condition. We proceed as follows. Let c_1, \dots, c_t be the children of n sorted by decreasing size of the respecting child decompositions D_{c_1}, \dots, D_{c_t} . If it exists, then let $j \in [t]$ be the maximum, such that $G[A_n]$ with

$$A_n := (B_n \cap B_{c_1}) \cup \dots \cup (B_n \cap B_{c_j})$$

is a clique, and $|D_{c_j}| > |D_{c_{j+1}}|$ holds or $j = t$ holds. Otherwise, set $j := 0$ and $A_n := \emptyset$. Thus, A_n is the attachment clique of the special children as defined above. We construct a collection of tree decompositions \mathcal{D}_n for B_n based on whether we have $j < t$ or $j = t$. If $j < t$, then let $m \geq 1$ be the largest integer with $|D_{c_{j+1}}| = |D_{c_{j+m}}|$. By construction, we can find at least one and at most $((k + 1)(m + 1))^2$ pairs of nonadjacent vertices $\{u, v\}$ in $G[A'_n]$ for

$$A'_n := A_n \cup (B_n \cap B_{c_{j+1}}) \cup \dots \cup (B_n \cap B_{c_{j+m}}).$$

This follows from the fact that A'_n is not a clique and A'_n has size at most $(k + 1)(m + 1)$, since all the sets $A_n, B_n \cap B_{c_{j+1}}, \dots, B_n \cap B_{c_{j+m}}$ have size at most $(k + 1)$. We define \mathcal{D}_n to be the collection of tree decompositions we obtained by applying Lemma 4.1 to $G[B_n]$ with pairs $\{u, v\}$ of nonadjacent vertices in $G[A'_n]$. We have $|\mathcal{D}_n| \leq ((k + 1)(m + 1))^2$. This set of decompositions satisfies the p -boundedness restriction with the polynomial $p(m) = ((k + 1)(m + 1))^2$. If $j = t$, then we consider every pair of nonadjacent vertices $\{u, v\}$ in B_n . Again, for every such $\{u, v\}$, we construct a decomposition for $G[B]$ using Lemma 4.1. We have $1 \leq |\mathcal{D}_n| \leq |B_n|^2$ in this case, satisfying the p -boundedness condition, since B_n only has special children. Since the construction of the collections \mathcal{D}_n is isomorphism-invariant, the entire construction is isomorphism-invariant. \square

We have assembled all the required tools to prove our main theorems showing that isomorphism of graphs of bounded tree width and canonization of graphs of bounded tree width can be performed in logarithmic space.

PROOF OF THEOREM 1.1. Given two graphs G and G' , by Lemma 7.1, we can compute in logarithmic space isomorphism-invariant p -bounded nested decompositions \bar{D} and \bar{D}' . By Lemma 5.3, the graphs are isomorphic if and only if there exist ordered root sets σ and σ' with $(G, \bar{D}, \sigma) \equiv_{\text{dec}} (G', \bar{D}', \sigma')$. By Lemma 6.1, this can be checked in logarithmic space by iterating over all suitable choices of σ and σ' .

The L-hardness for every positive $k \in \mathbb{N}$ follows from the L-hardness of the isomorphism problem for trees (connected graphs of tree width at most 1) proved by Jenner et al. (2003). \square

For our canonization procedure, we would like to recursively order the vertices according to $<_{\text{dec}}$. However, due to the fact that there is no exact correspondence between $<_{\text{dec}}$ and isomorphism for graphs with nested decompositions and ordered root sets (recall Remark 5.4 and that we only have a “quasi-completeness”-property not a “completeness”-property), we need to ensure that the

process is canonical. However, as the following proof shows, to ensure canonicity it is sufficient to work with an isomorphism-invariant decomposition.

PROOF OF THEOREM 1.2. We use the isomorphism-invariant mapping from Lemma 7.1 to turn G into a width-bounded and p -bounded nested decomposition $\bar{D} = (T, \mathcal{B}, \mathcal{D})$. The canonical sequence of G 's vertices is based on (G, \bar{D}, σ) , where σ is the empty vertex sequence. To compute a canonical sequence with respect to $<_{\text{dec}}$, we repeatedly apply Lemma 6.1.

If $|\mathcal{D}_r| = 0$, then let $\bar{D}_1, \dots, \bar{D}_s$ be the child decompositions of G containing at least one vertex that is not in σ . We obtain an order on them by defining $\bar{D}_i < \bar{D}_j$ if

$$\{((G_i, \bar{D}_i, \tau), (G, \sigma\tau)) \mid \tau \in \Pi(c_i)\} <_{(\text{dec}, \text{seq})} \{((G_j, \bar{D}_j, \tau), (G, \sigma\tau)) \mid \tau \in \Pi(c_j)\}.$$

Ties are broken arbitrarily, for example, by considering the smallest vertex in the child according to the input ordering. For each child \bar{D}_i , we compute an ordering $\tau_i \in \Pi(c_i)$ that minimizes (G_i, \bar{D}_i, τ_i) . We recursively create a canonical sequence outputting the canonical sequence of (G_i, \bar{D}_i, τ_i) for each child in the order of children just defined.

If $|\mathcal{D}_r| > 0$, then we iterate over all decompositions in \mathcal{D}_r and choose a tuple from $\{(G, \bar{D}_{D, \sigma}, \sigma) \mid D \in \mathcal{D}_B\}$ that is minimal with respect to $<_{\text{dec}}$. Ties are, again, broken based on the input ordering. For computing the canonical sequence, we continue recursively on a minimal $(G, \bar{D}_{D, \sigma}, \sigma)$ only. To obtain a canonical sequence, we alter the nested decomposition slightly whenever we go into the recursion using colored edges. More specifically, Lemma 7.1 constructs D based on two vertices u and v that form a distinguished non-edge. We insert an edge between u and v and color it with a color that does not appear in G (for example, we use -2). In other words, we set $\text{col}_G(u, v) = -2$. This modification is isomorphism-invariant based on the choice of D . The new edge is covered by a bag of D by construction. Inserting the edge only depends on D and, thus, it is stored recursively in an implicit way. The modification has the consequence that distinguished edges are preserved under isomorphism.

The logspace-computability of the sequence follows from Lemma 6.1. Thus, we are left to prove that the sequence is canonical. For this, we need to show that whenever a tie is broken arbitrarily between two options, then the two options are equivalent. There are two situations when a tie can occur.

For the first one, suppose $\{(G_i, \bar{D}_i, \tau), (G, \sigma\tau) \mid \tau \in \Pi(c_i)\} \equiv_{(\text{dec}, \text{seq})} \{((G_j, \bar{D}_j, \tau), (G, \sigma\tau)) \mid \tau' \in \Pi(c_j)\}$ for two child decompositions both containing a vertex not in σ . By Lemma 5.3, there is an isomorphism from the graph induced by the vertices in \bar{D}_i to the graph induced by the vertices in \bar{D}_j fixing σ . This extends to an automorphism of G by fixing all vertices neither in \bar{D}_i nor \bar{D}_j . Since \bar{D} is isomorphism-invariant, this automorphism respects \bar{D} , therefore mapping \bar{D}_i to \bar{D}_j .

For the other case, where a tie can occur, suppose $(G_i, (\bar{D}_i)_{D, \sigma}, \sigma) \equiv_{(\text{dec}, \text{seq})} ((G_j), (\bar{D}_j)_{D', \sigma}, \sigma)$. By Lemma 5.3, there is an isomorphism from G_i to G_j . This isomorphism preserves the distinguished edge. This isomorphism extends to an automorphism of G that fixes all vertices that neither appear in $(\bar{D}_i)_{D, \sigma}$ nor in $(\bar{D}_j)_{D', \sigma}$. Since \bar{D} is isomorphism-invariant, this automorphism of G respects \bar{D} , and since the distinguished edge is preserved it maps $(\bar{D}_i)_{D, \sigma}$ to $(\bar{D}_j)_{D', \sigma}$.

This shows that the computed sequence is canonical. \square

8 CONCLUSION

Summary. We showed how to canonize and compute canonical labelings for graphs of bounded tree width in logspace, and this implies that deciding isomorphism graphs and computing isomorphisms can be done in logspace for graphs of bounded tree width. For the proof, we first developed a tree decomposition into clique-separator-free subgraphs that is isomorphism-invariant and logspace-computable. Then, we showed how to compute, for each bag, an isomorphism-invariant

family of width-bounded tree decompositions in logspace. Finally, we combined both decomposition approaches to construct nested tree decompositions and developed a recursive canonization procedure that works on nested tree decompositions.

Outlook. Testing ISOMORPHISM for graphs that are embeddable into the plane (Datta et al. 2009a), as well as any fixed surface (Elberfeld and Kawarabayashi 2014), can be done in logspace. These graph classes can be described in terms of forbidding fixed minors, which also holds for classes of graphs whose tree width is bounded by constants. This opens up the question of whether these logspace results generalize to any class of graphs excluding fixed minors. For these classes, polynomial-time ISOMORPHISM procedures are known (Ponomarenko 1991). Partial results are known for graphs that exclude the minors K_5 or $K_{3,3}$ (Datta et al. 2009b), but the available techniques are tailored to the respective graph classes. Looking at algorithmic proofs related to the structure of graphs excluding fixed minors (Grohe 2012; Grohe et al. 2013), it seems promising to combine the earlier logspace approach for embeddable graphs (Elberfeld and Kawarabayashi 2014) with our logspace approach for bounded tree width graphs.

Of course, the basic question of determining the exact complexity of ISOMORPHISM on general graphs remains open, and in particular whether the problem is polynomial time solvable. With respect to this question, our work might help to clarify the difference between graphs to which the best known complexity-theoretic lower bounds for ISOMORPHISM (Torán 2004) apply, which are given in terms of classes defined via nondeterministic logarithmic-space-bounded Turing machines and graphs for which (deterministic) logspace algorithms are possible.

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