Trapezoid Graphs and Generalizations, Geometry and Algorithms

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Abstract.

Trapezoid graphs are a class of cocomparability graphs containing interval graphs and permutation graphs as subclasses. They were introduced by Dagan, Golumbic and Pinter [DGP]. They propose an $O(n^2)$ algorithm for chromatic number and a less efficient algorithm for maximum clique on trapezoid graphs. Based on a geometric representation of trapezoid graphs by boxes in the plane we design optimal, i.e., $O(n \log n)$, algorithms for chromatic number, weighted independent set, clique cover and maximum weighted clique on such graphs. We also propose generalizations of trapezoid graphs called k-trapezoid graphs. The ideas behind the clique cover and weighted independent set algorithms for trapezoid graphs carry over to higher dimensions. This leads to $O(n \log^{k-1} n)$ algorithms for k-trapezoid graphs. We also propose a new class of graphs called *circle trapezoid graphs*. This class contains trapezoid graphs, circle graphs and circular-arc graphs as subclasses. We show that clique and independent set problems for circle trapezoid graphs are efficiently solvable. The algorithms solving these two problems require algorithms for trapezoid graphs as subclasses.

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1. Introduction

Trapezoid graphs were introduced by Dagan, Golumbic, and Pinter [DGP]. Consider a channel, i.e., a pair of two horizontal lines. A *trapezoid* between these lines is defined by two points on the top and two points on the bottom line. A graph is a *trapezoid graph* if there exists a set of trapezoids corresponding to the vertices of the graph such that two vertices are joined by an edge iff the corresponding trapezoids intersect (see Figure 1). Dagan, Golumbic, and Pinter propose an algorithm computing the minimum number of colors in a proper coloring of such a graph in time $O(n^2)$ and a less efficient backtracking algorithm finding a maximum clique in such a graph (throughout the paper we assume that n is the number of vertices of the graph or order in question).

The problem of finding maximum cliques or minimum colorings for trapezoid graphs arises in connection with channel routing problems in VLSI design. Given some labeled terminals on the upper and lower side of a two sided channel, terminals with the same label will be connected in a common net. Such a net may be modeled by a trapezoid connecting the rightmost resp. leftmost terminals with the same label. Nets then may be routed without intersection iff the

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corresponding trapezoids do not intersect, i.e., iff they are independent. The number of colors needed to color the trapezoid graph is the number of layers needed to route the nets without intersection.



Figure 1. A trapezoid graph G, the order P and two representations.

For our algorithms we will make use of another equivalent characterization of trapezoid graphs. To give this alternative characterization it is convenient to fix some terminology. If $x = (x_1, \ldots, x_k)$ and $y = (y_1, \ldots, y_k)$ are points in \mathbb{R}^k , then x is said to be dominated by y, denoted x < y, if x_i is less than y_i for all $i = 1, \ldots, k$. The order thus given between points in \mathbb{R}^k is also called a dominance order. This order can be extended to boxes, i.e., sets of the form $\{(x_1, \ldots, x_k) \in \mathbb{R}^k : l_i \le x_i \le u_i, 1 \le i \le k\}$ where (l_1, \ldots, l_k) is the lower corner and (u_1, \ldots, u_k) is the upper corner of the box. A box b dominates a box b' if the lower corner of b dominates the upper corner of b'. Note that points may be understood as boxes where the lower and upper corner coincides. If one of the two boxes dominates the other we say that they are comparable. Otherwise they are incomparable. Now the vertices of trapezoid graph may be represented by boxes with two boxes incomparable iff the corresponding vertices are joined by an edge.

The connection between the box representation and the trapezoid representation of a trapezoid graph is the following. Interpret the points on the lower of the two lines of the channel as lying on the x-axis and that of the upper line as lying on the y-axis of the Euclidean plane. Each trapezoid then corresponds to an axis-parallel box in the plane whose projection on the

x- and y-axis coincides with the lower and upper side of the trapezoid (see Figure 1). It is easily seen that two trapezoids are disjoint exactly if the corresponding boxes are comparable.

What makes the box representation useful is the additional dominance order on boxes that may be exploited by sweep line algorithms. All computation is done in a single sweep leading to $O(n \log n)$ algorithms for clique, independent set and cover problems on trapezoid graphs. Hence, these graphs are another class of graphs where very efficient algorithms for such problems can be given. There exists a lower bound for the number of comparisons needed to compute maximum increasing subsequences in permutations, Fredman [Fre]. Permutations correspond to permutation graphs in such a way that increasing sequences correspond to either cliques or independent sets. As permutation graphs are trapezoid graphs, Fredman's bound shows that our algorithms are optimal in the same sense.

Algorithms for trapezoid graphs should be compared with algorithms for general cocomparability graphs. For these graphs the maximum independent set and the minimum clique cover problem can be solved in $O(m \log n)$, see [McSp], here *m* is the number of edges in the graph, i.e., *m* is in $O(n^2)$. The bottleneck of the computation is the complexity of transitive orientation. The maximum clique and chromatic number problems on cocomparability graphs seem to be harder. To the best of our knowledge the complexity is dominated by finding a maximum matching in a bipartite graph. The time needed to solve this problem is almost $O(n^{2.5})$ (see [ABMP]), and $O(n^3)$ in the weighted case (see [PaSt]).

In Section 2 we give some definitions and replace graph terminology by order terminology that proves to be more convenient in designing our algorithms. We assume the vertices of the trapezoid graph to have some weights. To compute maximum weighted cliques or independent sets turns out to impose no additional difficulty. In Section 3 we present an algorithm computing a maximum weighted independent set and a minimum clique cover at the same time (or in order terminology, a maximum weighted chain and a minimum antichain partition). We also show how to extend this algorithm from boxes in the plane to boxes in \mathbb{R}^k . Section 4 shows how to compute a minimum coloring (or a minimum chain partition). Unfortunately, this algorithm cannot be turned into an efficient one finding a maximum weighted clique (maximum weighted antichain). Hence, a different approach is proposed in Section 5 giving an efficient algorithm for the last problem.

In Section 6 we discuss a new class of graphs, called circle trapezoid graphs. A *circle trapezoid* is the region between two non-crossing chords of a circle. Alternatively, it is the convex hull of two disjoint arcs on the circle. *Circle trapezoid graphs*, *CT-graphs* for short, are the intersection graphs of families of circle trapezoids on a fixed circle. It is easily seen, that CT-graphs are a common generalization of trapezoid graphs, circle graphs and circular-arc graphs. We show, that in this large class of graphs the maximum clique and maximum independent set problems can still be solved efficiently.

2. Trapezoid graphs and trapezoid orders

The k-dimensional box representation (V, l, u) of a graph $\mathbf{G} = (V, E)$ consists of mappings $l: V \to \mathbb{R}^k$ and $u: V \to \mathbb{R}^k$ such that l(v) is the lower and u(v) the upper corner of a box box(v) where two vertices of the graph are joined by an edge iff their corresponding boxes are incomparable. If a graph has such a representation it is a k-trapezoid graph. If we additionally have a weight $w: V \to \mathbb{R}$ on the vertices of \mathbf{G} then the k-trapezoid graph is weighted. The weight of a clique, i.e., a set of mutually joined vertices in the graph, is the sum of the weights of its elements. Similarly, the weight of an independent set, i.e., a set of vertices with no two of them joined by an edge, is the sum of the weights of its elements. We are mainly interested in the case k = 2 where we simply deal with trapezoid graphs.

As already mentioned in Section 1, we switch to the richer structure given by the dominance order on the boxes of a box representation. Let the boxes of a box representation of a trapezoid graph together with the dominance order be the corresponding *trapezoid order*. A set of mutually comparable elements of an order is a *chain* and a set of mutually incomparable elements is an *antichain*. Recall that two boxes are incomparable iff the corresponding vertices of the trapezoid graph are joined. Let **G** be a trapezoid graph and **P** be a corresponding trapezoid order. Then it is easily verified that

- A minimum clique cover of G is a minimum antichain partition of P.
- A maximum weighted independent set in G is a maximum weighted chain in P.
- A minimum coloring of G is a minimum chain partition of P.
- A maximum weighted clique in G is a maximum weighted antichain in P.

A maximal element of a dominance order is one with no element dominating it. Each chain has exactly one maximal element. In contrast to the weight w(v) of a box v in a trapezoid order we will often attribute a *chain weight* W(v) to v which is the maximum weight of a chain with v as its maximal element.

Note that in the limiting case the box representation (V, l, u) of a trapezoid graph (V, E) may consist of points, i.e., l(v) = u(v), for all $v \in V$. Such graphs are known as *permutation graphs* and the points with the dominance order in the plane as 2-dimensional order. We denote such an order by (V, p) with p(v) = l(v) = u(v). Before giving the actual algorithms for the trapezoid orders we will sometimes recall algorithms for 2-dimensional orders since they are easier to grasp while showing important features extendible to the general case. The class of k-trapezoid orders is known as the class of orders of intervall dimension at most k, consult the book of Trotter [Tro] for further information on dimension and interval dimension of orders.

We will often have to maintain a finite set of real numbers such that values may be inserted or deleted from it and the predecessor or successor of a given query value can be found. Using balanced trees (e.g., red-black trees described in Cormen, Leiserson, Rivest [CLR]) all these operations can be done in $O(\log n)$ time and linear space. If we further assume the benefits of a random access machine and assume that the values are taken from a finite range U then the above operations take only $O(\log \log n)$ time and linear space when implemented on a data structure of van Emde Boas [vEB]. Hence, under these assumptions, the $\log n$ factor in the running time of the algorithms for 2-dimensional trapezoid orders may be replaced by a $\log \log n$ factor.

Throughout the paper we assume that the points l(v) and u(v) of a box representation have mutually different x- and y-coordinates. Otherwise, we may obtain a box representation of the same order fulfilling this requirement by perturbing the corner points with two line sweeps in the following way. Points with the same x-coordinates are perturbed slightly such that points which are lower corners have smaller x-coordinates than such which are upper corners. A similar perturbation is done for the y-coordinate. The x- and y-coordinate of a point $p \in \mathbb{R}^2$ will be denoted by p_x and p_y , resp. We will always use a vertical sweep line L going from left to right, i.e., from lower to higher x-coordinates.

3. Minimum antichain partition and maximum chain for k-trapezoid orders

We first give a brief description of an algorithm solving the maximum chain problem for a 2-dimensional order (V, p) in the weighted case. Let the weights be given by $w: V \to \mathbb{R}^+$. First, the points are sorted, so that we can access them by increasing *x*-coordinate, i.e., from left to right. Secondly, we compute a function $W: V \to \mathbb{R}$, where W(v) is the chain weight of v, i.e., the weight of a maximum weighted chain having v as its maximal element.

W(v) is computed with the aid of a sweep line L moving from left to right and halting at every point p(v). We maintain a set M of weighted markers on L so that the weight W(m)for $m \in M$ is just the weight of a maximum weighted chain on the set of points dominated by m, i.e., on $\{v \in V : p(v) < m\}$. For each $m \in M$, origin(m) is the maximal element of the maximum weighted chain dominated by m. When reaching a point p(v) we find the first marker m below p(v) on L, set W(v) = W(m) + w(v) and establish a link from v to origin(m). To update L we position a new marker m' with W(m') = W(v) and origin(m') = v at the y-coordinate of p(v). Then we remove those markers above m' that have smaller weight. Note that although the number of markers on L cannot exceed 2n. Finally, starting from a point vwith maximum chain weight W(v) we use the links to construct a heaviest chain.

Now we mimic this algorithm for the case where the box representation (V, l, u) of a trapezoid order **P** is given. Essentially, the idea is to separate the action taken by the algorithms for 2-dimensional orders whenever the sweep line reaches a new element into two parts. The first part of the action, located at l(v), is to compute the chain weight W of the new element v. This is done by finding the element v' of maximum chain weight among the elements with u(v') < l(v) and link v to v'. Note that the maximum weight of v' implies that v' was the maximum element of its chain. The second part of the action, located at u(v), is to make the chain weight of v available for further elements. The main difference to the permutation graph algorithm is that before inserting the information corresponding to v into the structure M we have to check whether the information is still relevant when released. The reason is that there might be an element v' with W(v') > W(v) whose box is completely dominated by the upper corner of v's box. Again, the weight of marker $m \in M$ will be equal to the weight of a maximum weighted chain on the boxes dominated by m, i.e., on the elements $v \in V$ with u(v) < m, in particular the weights on M are increasing with increasing y-coordinate.

The algorithm for computing a maximum weighted chain in a box representation is given next. For convenience, we initialize the sweep line with a dummy point d with W(d) = 0 and origin(d) =nil, such that d is below all points that will ever be inserted into L.

```
MAXCHAIN(V, l, u, w)
```

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for each p from left to right do
     m \leftarrow first marker below p on L
     if p = l(v) for some v \in V then
           W(v) \leftarrow W(m) + w(v)
           link(v) \leftarrow origin(m)
     if p = u(v) for some v \in V then
           if W(v) > W(m) then
                 insert a new m_v at p_y in L
                 W(m_v) \leftarrow W(v)
                 \operatorname{origin}(m_v) \leftarrow v
                 remove all m' that are higher and lighter than m_v from L
x \leftarrow \text{origin}(\text{uppermost marker on } L)
C \leftarrow \{x\}
while link(x) \neq nil do
     x \leftarrow \text{link}(x)
     C \leftarrow C \cup \{x\}
return C
```

LEMMA **3.1**. At the end of the main loop in MAXCHAIN the following invariant holds true. If y is an arbitrary point on L and m the next marker below y, then a maximum weighted chain dominated by point y has weight W(m).

Proof. If L has stopped at some point p = l(v) no new box has become available to increase any maximum weighted chain and no weight of any marker has been changed. But note that the weight of a maximum chain with maximal element v has weight W(m) + w(v), for m the marker below l(v), by the invariance assumption.

On the other hand, suppose L has stopped at p = u(v). If $y < u_y(v)$ or if no new marker is inserted in the sweep line, there can neither be a new maximum weighted chain nor a new marker below y. Hence we assume $y \ge u_y(v)$ and a new marker m_v has been inserted at height $u_y(v)$, i.e., there is a new chain with weight $W(v) = W(m_v)$ available for points above $u_y(v)$. Let m' and m be the markers immediately below y before and after the insertion of m_v . If m' = m then $W(m') > W(m_v)$ (otherwise, m' would have been removed) and W(m) = W(m') remains optimal among all chain weights. If $m' \neq m$ then m can only be m_v and $W(m_v) > W(m')$ either by the condition for removing markers or by the condition for the insertion of m_v . Since W(m') was optimal among all chain weights save the new one ending in $v, W(m_v)$ surely is an optimal weight for y.

Of course, Lemma 3.1 implies that Algorithm MAXCHAIN computes a maximum weighted chain, since all boxes are dominated by the uppermost point on L after the sweep was completed.

As already noted the sweep line can be implemented so that find, insert and delete operations require $O(\log n)$ time. It is easily seen, that 3n is an upper bound for the number of these operations. This proves an $O(n \log n)$ time bound.

The unweighted case can be simulated by unit weights. As the weights of all markers are different the number of markers on L in the unweighted case cannot exceed the length of a maximum chain in \mathbf{P} . If ω is the size of a longest chain in \mathbf{P} then all steps can be carried out in $O(n \log \omega)$. If each element of \mathbf{P} has unit weight, then no two elements with the same chain weight are comparable. Hence, collecting the elements of chain weight i in a set A_i yields a partition A_1, \ldots, A_ω of \mathbf{P} into antichains. It is easily seen that the maximum weighted chain must contain one element of A_i , $i = 1, \ldots, \omega$. This proves this antichain partition to be minimal since a partition into fewer antichains would force at least two elements of the chain into one antichain, which is impossible. Hence, a minimal antichain partition is a byproduct of algorithm MAXCHAIN. We summarize these remarks in

THEOREM 3.2. A maximum weighted chain and a minimum antichain partition of a trapezoid order on n points, given its box representation, can be computed in $O(n \log n)$ time and linear space.

Assume a k-dimensional box representation \mathbf{P} is given, for some higher dimension k > 2. We use dynamic multi-dimensional range trees (see, e.g., Smid [Smi] or Mehlhorn [Meh]) for the construction of a maximum chain for a point set in k dimensions. We need a data structure for a point set P in (k - 1)-dimensional space that, for a given query point q, allows to find some $p \in P$ with maximum chain weight W(p) among all points of P that are dominated by q. We also want to insert new points with some given chain weight. Given such a data structure, it is easy to compute a maximum weighted chain for a point set P in k dimensions. A sweep plane visits all points ordered by increasing last coordinate. For each point q on the sweep plane a point p is found in the range tree that has maximum chain weight W(p) among all points dominated by q in the first k - 1 coordinates. But since all points with smaller values than q in their last coordinate have been swept and have already been inserted in the range tree, p also has maximum weight among points dominated by q in all k dimensions. Hence, we may insert q in the range tree with chain weight W(q) = W(p) + w(q). Along with W(q) we may also store a link to point p. After insertion of all points a maximum weighted chain for point set Pis easily found. At first, a point p_m with the highest weight ever computed during the sweep is searched. Then, beginning with p_m , the chain is extracted by following the corresponding chain of links. Again, if all points have unit weight then W(p) is the size of a chain with maximal element p and two points with the same chain weight cannot be comparable. Hence, a minimum antichain partition is found as byproduct.

With the following changes the above approach is easily adapted to compute a maximum weighted chain for a box order. If point q on the sweep plane corresponds to a lower point l(v) of some box v we calculate W(v) = W(p) + w(v) as above but do not yet insert q in the tree. If q = u(v) for some box v we insert q in the range tree with chain weight W(q) = W(v) that has already been calculated before.

For convenience, let us briefly recall how such a (k-1)-dimensional range tree T works. We set d = k - 1. Let the d coordinates be denoted by x_1, \ldots, x_d . The point set ordered by x_d -coordinate is represented by the leaves of a binary tree T. If d = 1 each node t of T records some point p with chain weight W(p) maximal among all weights in the leaves below t, i.e., in the leaves of the subtree rooted at t. If d > 1 each node t of the main range tree points to a (d-1)-dimensional secondary range tree constructed recursively with respect to the first d-1 coordinates for the points in the subtree of t.

Suppose we want to find, for a given query point q, the point with maximal weight among all points of P with all coordinates smaller than that of q. In a first step the point of P with smallest x_d -coordinate greater than that of q is searched in the main tree. Let the search path be S_q . Let L_q be the set of left children of nodes in S_q that are not in S_q . It is easily seen that each point with x_d -coordinate smaller than or equal to that of q has a leaf below some node in L_q . Hence, to find a point p with W(p) maximal among all points dominated by q we proceed as follows. If d = 1 we check all leaves pointed to by the nodes in L_q and return the leaf with maximum weight. If d > 2 the range trees in nodes $t \in L_q$ allow to find points p_t with maximum weight among leaves below t and dominated by q in the first d - 1 coordinates. In this case the point searched for is that with maximum weight among points p_t , for $t \in L_q$. On the other hand, if we want to insert point q into tree T this may be done by first inserting it in the main tree and then inserting it in all secondary range trees at nodes along the insertion path, if d > 1. If d = 1, pointers along the insertion path are set to the leaf belonging to q if its weight is the new maximum in the corresponding subtree.

It is easily seen that a query takes time $O(\log^d n)$ if all trees are balanced. If some trees become unbalanced during an insertion they must be rebalanced and it can be shown that this takes amortized time $O(\log^d n)$. For d = 1 we need linear space. And since a point is contained in at most $\log n$ secondary trees if d > 1, the total amount of space is $O(n \log^{d-1} n)$, by induction. We leave it to the reader to supply the omitted details that give a complete proof of the following statement.

THEOREM 3.3. If an order $\mathbf{P} = (V, P)$ is given by a box representation in \mathbb{R}^k , then a minimum antichain partition and a maximum chain of \mathbf{P} can both be obtained in $O(n \log^{k-1} n)$ time and $O(n \log^{k-2} n)$ space.

4. Chain partitions of trapezoid orders

In this section we show how to partition a trapezoid order \mathbf{P} into chains such that the number of chains used is minimal. Of course, this only makes sense if we assume unit weights on the elements of \mathbf{P} . Again, we begin with a short description of a similar algorithm for 2-dimensional orders which we then adapt for the case of a given box representation of \mathbf{P} .

An optimal chain partition for a point set can be obtained by a sweep of a line L from left to right in the following way. Assume the set of points to the left of the current position of L to be already optimally partitioned into chains. On L the maximal elements of the chains of this partition are maintained ordered by y-coordinates. When reaching a new point p we search for the point q on L which has maximal y-coordinate among all points on L that are below p. If q exists then p is appended as new maximal element to the chain of q, otherwise, p does not dominate any chain of the actual partition and we initialize a new chain consisting of p only. Finally, L is updated by inserting p and removing q.

Now suppose, that **P** is given by a box representation (V, l, u). We have to separate the action that has to be taken when the sweep line reaches a new element into two parts. The first part of the action, located at l(v), is to find the chain of the already existing partition that will be extended by v. The second part, located at u(v), is to make the chain with maximum v available for further elements. A chain C with maximum element v will be called *closed* as long as u(v) has not been visited by L, otherwise C is *open*.

The algorithm for computing a minimum chain partition in a box representation is given as follows. We initialize the sweep line with a dummy point d such that d is below all points that will ever be inserted into L.

MINCHAINPARTITION(V, l, u)

```
for each p from left to right do

q \leftarrow \text{first element below } p \text{ on } L

if p = l(v) for some v \in V then

if q = u(w) for some w \in V then

chain(v) \leftarrow chain(w) \cup \{v\}

remove q from L

else (q = d)

chain(v) \leftarrow \{v\}

if p = u(v) for some v \in V then

insert p at p_y in L

return \{ chain(v) : u(v) \in L \}
```

The time consuming operations in this algorithm are the search, insert and remove operations for points on the sweep line L. With the use of a balanced search tree the running time of the algorithm is in $O(n \log n)$. If we assume the points to be presorted, the running time is in $O(n \log \alpha)$ where α is the number of chains in the partition.

To prove that the chain partition found by this algorithm is minimum we show how to extract an antichain from **P** that contains an element from each chain in the partition. Let $\mathcal{C} = \{C_1, \ldots, C_{\alpha}\}$ be the chain partition found. Let v be the last element that opened a new chain, say C_{α} . Note that $l_x(v)$ is larger than $l_x(v')$ if v' is the minimal element of a chain C_i with $i \neq \alpha$. Let U be the set of elements v' whose lower corner is dominated by the lower corner l(v) of v and whose upper corner was not yet swept when l(v) was processed, i.e., $l_x(v) < u_x(v')$. It is clear that $U \cup \{v\}$ is an antichain, hence, all these elements belong to

different chains. Let \mathcal{C}_1 be the corresponding set of chains. All chains of \mathcal{C}_1 where closed after the sweep passed l(v). The remaining set of chains, i.e., $\mathcal{C} \setminus (\mathcal{C}_1 \cup \{C_\alpha\})$, is called \mathcal{C}_2 . Let X(v)be the set of elements $v' \in V$, such that either l(v') or u(v') is contained in the quarter-plane $\{(x, y) : x \leq l_x(v) \text{ and } y \geq l_y(v)\}$. It is easily seen that every chain $C \in \mathcal{C}_2$ contains an element of X(v). Let C^* be the subchain of C induced by the elements in X(v) and \mathcal{C}_2^* be the set of these subchains. The next lemma states the crucial property of \mathcal{C}_2^* .

LEMMA 4.1. The chain partition C_2^* of the order induced by X(v) is exactly the chain partition generated by MINCHAINPARTITION, when the input consists of the boxes of elements in X(v) only.

Proof. Let L be the sweep line for input (V, l, u) and L^* be the sweep line for the restricted input, i.e, X(v) replaces V. The lemma is an easy consequence of the following invariant: Considered at the same x-coordinate, $x \leq l_x(v)$, the restriction of L to the half line above $l_y(v)$ and L^* are identical. This is certainly true at the beginning when both lines are empty. Now suppose they are equal and L meets point p. We distinguish four cases.

First consider the situation p = l(v') and $v' \notin X(v)$. Since $v' \notin X(v)$ we have $l_y(v') < l_y(v)$. There may be a removal below $l_y(v')$ in L, but it cannot affect the half line above $l_y(v)$. Now let p = l(v') and $v' \in X(v)$. Suppose, that there is an element $q \neq d$ below p in L^* and let wbe the element with u(w) = q. In this case v' joins the chain of the w and q is removed from L^* . Obviously, the same action takes place on L. If there is only the dummy element below p on L^* , then v' opens a new chain for the restricted input. On L there may be an element below p. Nevertheless, the y-coordinate of this element has to be smaller than $l_y(v)$ and the changes on L will not affect the half line above $l_y(v)$.

If p = u(v') and $v' \notin X(v)$, then $u_x(v') < l_x(v)$ and $v' \notin X(v)$ imply that $u_y(v') < l_y(v)$. Therefore p is inserted in the half line of L below $l_y(v)$. Finally, let p = u(v') and $v' \in X(v)$. We then have $u_y(v') > l_y(v)$ and p is inserted in both, L^* and L.

By induction on the number of boxes in the input we may now assume that the chain partition \mathcal{C}_2^* is optimal for X(v). Choose an antichain B of the order induced on X(v), such that B contains an element from each chain $C^* \in \mathcal{C}_2^*$. Since every element in the antichain $U \cup \{v\}$ is incomparable to every element in X(v), we conclude, that $A = B \cup U \cup \{v\}$ is an antichain. The antichain A consists of a member of every chain of the chain partition \mathcal{C} , i.e., $|A| = |\mathcal{C}|$. Since $|A| \leq |\mathcal{C}|$ for every antichain A and every chain partition \mathcal{C} , equality can only hold if A is maximum and \mathcal{C} minimum. This proves

THEOREM 4.2. A minimum chain partition of a trapezoid order on n points, given its box representation can be computed in time $O(n \log n)$ and linear space.

5. Maximum antichain for trapezoid orders

We first describe the geometry of antichains in a box representation. Our algorithm for maximum weighted antichains of trapezoid orders will be based on this geometric structure rather than on duality as the algorithms presented so far. First, we need some definitions.

Define the shadow of a point p as the set of points in the plane dominating p, shadow(p) = $\{q : p < q\}$. The shadow of a set of points is the union of the shadows of the elements. A downwards staircase is a sequence of horizontal and vertical line segments that may be obtained as the topological boundary of the shadow of a set of points. The staircase of a set of points then is the boundary of the shadow of the set. Note that any two different points on a staircase are incomparable. If S is a staircase and $l, u: V \to \mathbb{R}^2$ a set of boxes we denote the set of vertices whose box intersect S by A(S), i.e., $A(S) = \{v \in V : box(v) \cap S \neq \emptyset\}$.

LEMMA 5.1. Let **P** be an order given by a box representation. If S is a staircase then A(S) is an antichain. Moreover, if A is an antichain of **P** then there exists a staircase S such that $A \subseteq A(S)$.

Proof. Assume that A(S) is not an antichain. Then there are $v, v' \in A(S)$ with v < v'. Consequently, for two different points $p \in box(v) \cap S$ and $p' \in box(v') \cap S$ on the staircase we have p < p'. But this is impossible, as noted above.

If A is an antichain of **P**, let $u(A) = \{u(v) : v \in A\}$. Let staircase S be the boundary of the shadow of u(A). Now suppose that there is an element $v \in A$, such that $box(v) \cap S = \emptyset$. Then u(v) must lie in the shadow of u(A) and it follows that l(v) is contained in the shadow of u(a), for some $a \in A$. By definition, u(a) < l(v) and hence a < v in **P**, a contradiction. \Box

Given a weighted order **P** with a box representation we define the weight of a staircase S as the sum of weights of all boxes intersecting S. If S is a staircase and $p \in S$, then we refer to the part of S above and to the left of p as staircase ending in p and again its weight is the sum of weights of intersecting boxes.

The following algorithm computes an antichain of maximum weight. It uses two different data structures. The sweep line L halts at every point l(v) and u(v), for $v \in V$. Roughly, it contains a list of weighted markers, so that the weight W(m) of marker m is the weight of a heaviest staircase ending in m. Moreover, a heaviest staircase ending in an arbitrary point y on L can be composed by joining the vertical line segment from y to the next marker m above y with a heaviest staircase ending in m. Structure L is initialized with a dummy point d of weight 0, such that d is above all points that will ever be inserted into L. The second structure Δ contains a list of all *open* boxes, i.e., boxes which have their left sides already swept but not their right ones. The total weight of all open boxes the upper sides of which lie between points y_1 and y_2 on L with $y_1 \geq y_2$ is denoted by $\Delta(y_1, y_2)$.

MAXANTICHAIN(\mathbf{P}, l, u, w)

```
for each p from left to right do
     m \leftarrow first marker above p on L
     if p = l(v) for some v \in V then
           add w(v) to all markers in interval [l_u(v), u_u(v)]
           insert a new item in \Delta at height u_{y}(v) with weight w(v)
           m^* \leftarrow \text{next marker below } p \text{ on } L
           while W(m) + \Delta(m, m^*) > W(m^*) do
                 remove m^* from L
                 m^* \leftarrow \text{next marker below } p \text{ on } L
     if p = u(v) for some v \in V then
           insert a new marker m_v at p_u in L
           W(m_v) \leftarrow W(m) + \Delta(m, m_v)
           \operatorname{list}(m_v) \leftarrow \operatorname{list}(m) \cup \{p\}
           remove item at u_u(v) from \Delta
T \leftarrow \text{staircase of points in } \text{list}(\text{lowest}(L))
for each v \in V do
     if v intersects T then A \leftarrow A \cup \{v\}
return A
```

LEMMA 5.2. At the end of the main loop in Algorithm MAXANTICHAIN we have the following invariant. If y is an arbitrary point on L and m the next marker above y, then a maximum weighted staircase that ends in y on L has weight $W(m) + \Delta(m, y)$.

Proof. Let W' denote the sweep line structure and Δ' denote the open box structure before a halt of the sweep line L and let W and Δ be the pair of structures after the halt. Let m be the first marker above y on L. Let S_y denote part of the staircase of $list(m) \cup \{y\}$ that ends in point y. We show that S_y has weight $W(m) + \Delta(m, y)$ and that this weight is maximal among weights of staircases ending in y.

At first, suppose the sweep line L halts at some point l(v) for $v \in V$. If $y > u_y(v)$ there is no change to any staircase ending in y. Otherwise, we have $y < u_y(v)$. If furthermore $u_y(v) < m$ then S_y intersects the new box v and has weight $W(m) + \Delta(m, y) = W'(m) + (\Delta'(m, y) + w(v))$. By the invariance assumption, this weight is maximal among all staircases ending in y. If $l_y(v) < m < u_y(v)$ then S_y has weight $W(m) + \Delta(m, y) = (W'(m) + w(v)) + \Delta'(m, y)$ which again is maximal. If $m < l_y(v)$ then the weight of S_y is $W(m) + \Delta(m, y) = W'(m) + \Delta'(m, y)$ which is the maximum weight of a staircase that avoids v (see Figure 2).



Figure 2. Sweep line L with open boxes and staircase S_y ending in y.

On the other hand, if a staircase S'_y ending in y intersects v then there is a $y' > l_y(v)$, such that S'_y is composed of a staircase ending in y' and a vertical segment from y to y'. Let m' be the first marker above y'. Since the case $m' > y' > l_y(v)$ has already been considered, by the definition of Δ , the weight of S'_y is at most

$$W(m') + \Delta(m', y') + \Delta(y', m) + \Delta(m, y) = W(m') + \Delta(m', m) + \Delta(m, y).$$

By the condition on the removal of markers in the algorithm, we know that $W(m) > W(m') + \Delta(m', m)$. Consequently, the weight of S'_y is less than $W(m) + \Delta(m, y)$, the weight of S_y . Note that the weight of markers m' between $l_y(v)$ and $u_y(v)$ is increased by w(v). But the weight of the associated staircases is increased by the same amount since they all intersect the box of v now.

Now suppose L halts at some point u(v). Since v is no longer open we have to remove $u_y(v)$ from Δ' . On the other hand, we have to maintain the invariant. Thus, a new marker m_v is

inserted in L with weight $W(m_v) = W'(m') + \Delta'(m', m_v)$, where m' is the next marker above $u_y(v)$. If m, the next marker above y, is different from m_v there again is no change in the weight of staircases ending in y. On the other hand, if m_v is the next marker above y, then the weight of S_y is

$$W(m_v) + \Delta(m_v, y) = W'(m) + \Delta'(m, m_v) + \Delta'(m_v, y) = W'(m) + \Delta'(m, y),$$

which is maximal by the invariance assumption, since no new box has to be considered. Note that the associated staircase of m_v is constructed by enlarging that of m in such a way that it additionally intersects all open boxes between m and m_v with total weight $\Delta'(m, m_v)$. This means that it has weight $W(m_v)$ which is maximal.

THEOREM 5.3. Let $\mathbf{P} = (V, P)$ be a trapezoid graph given by a box representation and $w: V \rightarrow \mathbb{R}$ be a weighting of \mathbf{P} . MAXANTICHAIN computes a maximally weighted antichain of \mathbf{P} .

Proof. After all boxes have been swept, structure Δ is empty (i.e., there is no box left open). Hence, the theorem follows from the invariant of Lemma 5.2.

L may be implemented by a balanced binary tree. One has to be careful only about adding some weight w to W(m') for markers m' lying in an interval [l, u]. We give a sketch of a possible implementation and invite the reader to supply the details. Suppose that each marker corresponds to a leaf of the tree and that the leaves are sorted by increasing y-coordinate of the markers. Let h be the height of such a tree. Let each node of the tree have some extra field holding the increment in the weight for all leaves in its subtree. The weight of a marker is then easily computed in h steps by summing up all weights along the search path to the corresponding leaf and finally adding this sum to the weight stored in the leaf. On the other hand, note that the set of all leaves corresponding to an interval [l, u] is the disjoint union of the leaves of at most 2h subtrees which can be found along the search paths of l and u. Consequently, if the weight of all leaves in interval [l, u] have to be increased, it suffices to update the weight increment fields of the at most 2h roots of these subtrees.

The main operation in a rebalancing of a balanced tree, e.g., of a red-black tree (see [CLR]), is a rotation at some internal node t. In the case of a left rotation we propagate the increment from right(t) to each of its children and from t to right(t) before performing the rotation. Right rotations are handled symmetrically. This ensures that the sum of increments on a path to a leaf are equal before and after the rotation. The computing time of the whole algorithm is increased by a constant factor only. Consequently, the addition of some weight to an interval as well as insertion, deletion, predecessor and successor queries, and the computation of the weight of some element can all still be done in time $h = O(\log n)$.

 Δ may be implemented by any one dimensional range tree where insertion, deletion, and query again takes $O(\log n)$ time. The main loop is executed n times and each step therein takes logarithmic time save the while loop. But in total the while loop is executed at most n times since each removed point must have been inserted before. Of course, the test for intersection of a box v with jump line T can be done in time $O(\log n)$. The space requirement of both data structures is linear. In summary, we obtain

THEOREM 5.4. A maximum weighted antichain of a trapezoid order on n points, given its box representation, can be computed in time $O(n \log n)$ and linear space.

Note that one can do without the Δ structure if one uses subtraction in the W structure. But the above algorithm is easier to understand and it can be adapted to the case where no subtraction is allowed (e.g., in semigroups).

We conclude this section with an open problem. We have given optimal algorithms for the classical chain and antichain problems for trapezoid orders. Also, for k-trapezoid orders $k \geq 3$ we have obtained a fast algorithm for maximum weighted chain. Is there an algorithm for maximum antichain for k-trapezoid orders $k \geq 3$ whose running time improves over the complexity of bipartite matching and hence the complexity of the algorithm for general orders?

6. Algorithms for circle trapezoid graphs

A *circle trapezoid* is the region in a circle that lies between two non-crossing chords and CT*graphs* are the intersection graphs of families of circle trapezoids on a common circle. Figure 3 gives an example. In this section we develop polynomial algorithms for the maximum weighted clique and maximum weighted independent set problems on CT-graphs.

6.1. Crossing graphs and independent sets of CT-graphs

Let $\mathbf{G} = (V, E)$ be a CT-graph. Of course, we will assume that a representation of \mathbf{G} is given. Let p be an arbitrary point on the circle and let C_p be the set of vertices of \mathbf{G} whose circle trapezoid contains p. Note that C_p induces a clique of \mathbf{G} , therefore, an independent set of \mathbf{G} can contain at most one element from C_p . Using p as the 'origin' of the circle and fixing an orientation (clockwise) of the circle we define a unique representation for a circle trapezoid. The representation consists of a 5-tuple $(t_1, t_2, t_3, t_4, \sigma)$. The first four components are the corners of the circle trapezoid in clockwise order starting from p. The fifth component σ is a sign, + or -, where + indicates that p is contained in one of the arcs of the circle trapezoid.



Figure 3. A circle trapezoid graph G with a representation

Define a *double interval* as a pair (I_1, I_2) of intervals on the real line, where I_2 is a subinterval of I_1 , i.e., $I_2 \subset I_1$. Let $I = (I_1, I_2)$ and $J = (J_1, J_2)$ be double intervals. We say I contains J if $J_1 \subset I_2$ and call them *disjoint* if $I_1 \cap J_1 = \emptyset$. Two double intervals are called *crossing* if they are not disjoint and neither of them is contained in the other. Call a graph $\mathbf{G} = (V, E)$ a

crossing graph if its vertices can be put in one to one correspondence to a collection of double intervals such that two vertices of \mathbf{G} are adjacent if and only if their corresponding double intervals cross. It is not hard to see that the class of crossing graphs contains both, trapezoid graphs and overlap graphs (recall that a graph is an overlap graph if and only if it is a circle graph). Our next lemma relates CT-graphs and crossing graphs.

LEMMA 6.1. Let $\mathbf{G} = (V, E)$ be a CT-graph given by a representation and C_p be the set of all vertices of \mathbf{G} whose circular trapezoid share a specified point p on the circle. Then for any subset W of $V \setminus C_p$, the subgraph of \mathbf{G} induced by W is a crossing graph.

Proof. Given the circular trapezoid representation we associate to a vertex $v \in W$ with corresponding circular trapezoid $(t_1, t_2, t_3, t_4, -)$ two arcs along the circle. Let A_1 be the arc from t_1 to t_4 and let A_2 be the arc from t_2 to t_3 , in both cases we choose the arc which does not contain p. Obviously, $A_2 \subset A_1$. Cutting the circle at p we obtain a line with a collection of double intervals representing the subgraph of **G** induced by W.

Let us first turn to the case where we find a point p on the circle such that C_p is empty, i.e., there is no $v \in V$ containing p. By the above lemma, the CT-graph **G** is also a crossing graph. Thus, let us first figure out how to compute maximum weighted independent sets in crossing graphs. Our algorithm for this problem is very much alike the algorithm given by Gavril [Gav] (see also Golumbic [Gol]) for the case of overlap graphs.

For a pair of double intervals we have defined the relations containment, disjointness, and crossing. By definition, every pair of double intervals is in exactly one of these relations. Containment is an antisymmetric and transitive relation, i.e., an order relation. Disjointness is defined using only the first interval of each double interval, therefore, we can transitively orient disjoint pairs by the relation 'lies entirely to the left', which gives an interval order.

To compute the maximum independent set of a crossing graph $\mathbf{G} = (V, E)$ given by a family \mathcal{I} of double intervals we proceed as follows. First, the *containment order* $\mathbf{P}_C = (V, P_C)$ and the *interval order* $\mathbf{P}_I = (V, P_I)$ corresponding to \mathcal{I} are extracted and a linear extension $L_C = v_1, \ldots, v_n$ of \mathbf{P}_C is computed, i.e., $I_1(v_j) \subseteq I_2(v_i)$ implies j < i. We artificially extend \mathbf{P}_C and L_C by an element v_{n+1} of weight 0, such that $v_{n+1} > v_i$ for all $i = 1, \ldots, n$. This preprocessing can be accomplished in time $O(n^2)$. Next, the following algorithm is called.

```
\begin{aligned} \text{MAXINDSET-CROSS}(\mathbf{P}_{C}, \mathbf{P}_{I}, L_{C}) \\ \text{for} \quad i = 1 \text{ to } n + 1 \text{ do} \\ & U_{i} \leftarrow \{v_{j} : v_{j} < v_{i} \text{ in } P_{C}\} \\ & C \leftarrow \text{maximum } W \text{-weighted } \mathbf{P}_{I} \text{-chain of elements of } U_{i} \\ & W(v_{i}) \leftarrow w(v_{i}) + \sum_{v \in C} W(v) \\ & I(v_{i}) \leftarrow \{v_{i}\} \cup \bigcup_{v \in C} I(v) \\ & \text{return } I(v_{n+1}) \end{aligned}
```

It is important to note, that U_i contains only elements v_j with j < i. Hence, the weights $W(v_j)$ of all elements in U_i have already been computed before the *i*th round. The following invariance at the end of each round of the algorithm is easily proved. For all $j \leq i$, the weight $W(v_j)$ is the weight of a maximum independent set $I(v_j)$ containing only elements $v \in U_j \cup \{v_j\}$, *i.e.*, elements with $v \leq v_j$ in \mathbf{P}_C . This invariant implies that $I(v_{n+1})$ is a maximum independent set for \mathbf{G} .

Clearly, every instruction but the second in the loop can be executed in O(n) time. The second instruction itself is a maximum weighted chain computation in an interval order. This

problem can be solved in linear time when the endpoints of the intervals are available in increasing order. For completeness we sketch an algorithm for this problem.

Visit the endpoints from left to right and maintain the weight ρ of the longest chain among intervals whose right endpoint has already been seen. When reaching the left endpoint of an interval, say the interval of v, we know that the maximum weighted chain having v as maximal element has weight $W(v) = \rho + w(v)$. At the right endpoint of v's interval we update ρ by the rule $\rho = \max\{\rho, W(v)\}$. Note that this algorithm can be seen as a one dimensional version of algorithm MAXCHAIN in Section 2, i.e., instead of a sweep line we use a sweep point and thus need no search for the relevant marker. In summary, we obtain

LEMMA 6.2. Algorithm MAXINDSET-CROSS solves the maximum weighted independent set problem for crossing graphs in time $O(n^2)$.

If we cannot find a point p on the circle with no $v \in V$ intersecting it then the CT-graph **G** is not necessarily a crossing graph. Nevertheless, we may reduce the detection of a maximum weighted independent set of a CT-graph to one application of MAXINDSET-CROSS and at most n maximum weighted chain computations on interval orders in the following way.

For $v \in V$ let N[v] denote the set of neighbors of v in \mathbf{G} together with v itself and let $\mathbf{G}(v)$ be the subgraph of \mathbf{G} induced by $V \setminus N[v]$. Also, let \mathbf{G}_p denote the subgraph induced by $V \setminus C_p$. As remarked above, the vertices of C_p form a clique in \mathbf{G} . Therefore a maximum independent set I of \mathbf{G} is either a maximum independent set in \mathbf{G}_p or there is a $v \in C_p$, such that $I = I' \cup \{v\}$ where I' is a maximum independent set of $\mathbf{G}(v)$. Since $C_p \subseteq N[v]$ for all $v \in C_p$ Lemma 6.1 shows that each of the above graphs $\mathbf{G}(v)$, as well as \mathbf{G}_p are crossing graphs.

Consequently, the solution for the maximum weighted independent set problem for CTgraphs is either a maximum weighted independent set in \mathbf{G}_p or one of the sets $I = I' \cup \{v\}$ where $v \in C_p$ and I' is a maximum weighted independent set in $\mathbf{G}(v)$.

THEOREM 6.3. The maximum weighted independent set problem for CT-graphs can be solved in time $O(n^2)$.

Proof. The crucial observation is that having applied algorithm MAXINDSET-CROSS to \mathbf{G}_p the problem for each of the graphs $\mathbf{G}(v), v \in C_p$, can be solved by a single maximum chain computation in an interval order, i.e., in O(n) time. Let $v \in C_p$ and let the circular trapezoid of v be given by $(s_1, s_2, s_3, s_4, +)$. The double intervals corresponding to vertices of $\mathbf{G}(v)$ are exactly those with $I_1 \subset (s_1, s_2)$ or $I_1 \subset (s_3, s_4)$. Let v_i be an element of $\mathbf{G}(v)$ and recall that the set U_i is the set of elements whose double interval is contained in the double interval of v_i . It follows that U_i is contained in $\mathbf{G}(v)$ and hence that sets $I(v_i)$ and weights $W(v_i)$ computed by MAXINDSET-CROSS with input \mathbf{G}_p and with input $\mathbf{G}(v)$ are equal. To solve the problem for $\mathbf{G}(v)$ it thus suffices to select the intervals contained in (s_1, s_2) or (s_3, s_4) and compute a maximum weighted chain of this set of intervals.

6.2. Cliques of CT-graphs

Let $\mathbf{G} = (V, E)$ be a CT-graph, given by a circular trapezoid representation. In this section we show how to apply Algorithm MAXANTICHAIN of Section 5, which computes a maximum weighted clique of a trapezoid graph, to compute such a clique of \mathbf{G} .

LEMMA 6.4. Let C be a maximum weighted clique of a CT-graph $\mathbf{G} = (V, E)$ given by a representation. Then there exists a chord of a circular trapezoid v in C such that the set of all

circular trapezoids intersecting contains all circular trapezoids of elements of C. Furthermore, their intersection graph is a trapezoid graph.

Proof. To avoid any unnecessary confusion we may assume that all endpoints of the trapezoids on the circle are different. Choose an arbitrary chord c_1 of an arbitrary circular trapezoid v_1 of C. Either all trapezoids of C intersect this chord or there is a circular trapezoid $v_2 \in C$ intersecting v_1 but not its chord c_1 . Of the two chords of v_2 let c_2 be the one which is nearer to c_1 . Now either all trapezoids of C intersect c_2 or there is a circular trapezoid $v_3 \in C$ intersecting v_2 but not c_2 and again c_3 may be the chord of v_3 nearer to c_2 . Note that c_3 and c_1 lie on opposite sides of c_2 . By repeating the above arguments we finally find a sequence v_1, v_2, \ldots of circular trapezoids of C with nonintersecting chords c_1, c_2, \ldots such that c_i lies between c_{i-1} and c_{i+1} , for $i \geq 2$. Consequently, all chords of the sequence are pairwise different. Thus, the sequence is finite and all trapezoids of C intersect the chord c of the last trapezoid v of the sequence.

To obtain a trapezoid representation for the graph induced by all the circular trapezoids intersecting chord c we cut the circle at the endpoints of c and use the two parts of the circle on one and the other side of c as the two lines of the trapezoid representation. Circular trapezoids with no chord intersecting c (i.e., circular trapezoids containing c) together with v are mapped to trapezoids which intersect all other trapezoids of the representation. All circular trapezoids with both chords intersecting c are mapped to the corresponding trapezoids in the representation without any change. If a circular trapezoid has only one chord intersecting c, too, by giving it a new endpoint. We only have to choose the new endpoint near the appropriate endpoint of c so that the intersected circular trapezoids (among those intersecting c) remain the same. After this change the circular trapezoid is mapped as above to the corresponding trapezoid in the representation.

With the above lemma it is now easy to find a maximum weighted clique in a CT-graph. We consider each chord of each circular trapezoid of \mathbf{G} , find a trapezoid representation of all circular trapezoids intersecting this chord in linear time as in the proof of the above lemma, and compute a maximum weighted clique for this representation by Algorithm MAXANTICHAIN. Lemma 6.4 guarantees that in this manner we finally find the maximum weighted clique of \mathbf{G} . By Theorem 5.3, we have

THEOREM 6.5. A maximum weighted clique for a circular trapezoid graph can be computed in time $O(n^2 \log n)$, given its representation.

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