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A NOTE ON FINDING A MAXIMUM EMPTY RECTANGLE

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Abstract

Given a rectangle A and a set S of n points in A , the *maximum empty rectangle* problem is that of finding a largest-area rectangle which is contained in A , has its sides parallel to those of A , and does not contain any of the points in S . This note describes an efficient algorithm for solving this problem.

1. Introduction

Given a rectangle A and a set S of n points in A , a *valid rectangle* is one which is contained in A , has its sides parallel to those of A , and does not contain any of the points in S . The *maximum empty rectangle* (MER) problem is that of finding the largest-area valid rectangle. This problem was first posed by Naamad, Lee and Hsu [NHS], who gave an $O(\min(n^2, s \log n))$ time algorithm for solving it. The parameter s is defined as the number of *restricted rectangles* of the problem, where a restricted rectangle (RR for short) is a valid rectangle such that each of its four edges either contains a point of S or coincides with an edge of A . Naamad et. al. prove that $s = O(n^2)$ and give an example in which $s = \Theta(n^2)$. They also show that when the points are drawn from a uniform distribution, the expected value of s is $O(n \log n)$, so that the average time

complexity of their algorithm is $O(n \log^2 n)$. More recently, an elegant algorithm of time complexity $O(n \log^3 n)$ has been given by Chazelle et. al. [CDL]. This is an improvement over [NLH] for large s , but is actually inferior in the average case. In this paper we describe an algorithm of time complexity $O(s+n \log^2 n)$. Thus its average time complexity matches that of [NLH], and its worst-case complexity improves on that of [NLH] in the range $n \log n < s < n^2$, and on that of [CDL] in the range $n \leq s \leq n \log^3 n$.

2. Preliminaries

We use p_1, \dots, p_n to denote the n input points, and $X(p_i), Y(p_i)$ to denote the coordinates of p_i . To simplify the exposition, we assume that no two points have same x -coordinate, and similarly for y -coordinates. Throughout, s denotes the total number of RR 's.

Let Q be an RR . Q is *bounded from above (resp. below, left, right)* by p_i iff its top (resp. bottom, left, right) edge contains p_i . Similarly, Q is *bounded from above (resp. below, left, right)* by A iff its top (resp. bottom, left, right) edge coincides with the top (resp. bottom, left, right) edge of A . For example, the RR shown dotted in Figure 1 is bounded from above by A , from the left by p_3 , from below by p_2 and from the right by p_5 .

For reasons which will become clear later, we choose to use two distinct sets of coordinate axes x_1, y_1 and x_2, y_2 as shown in Figure 1. Note that the origin of x_1, y_1 is at the southwest corner of A , while that of x_2, y_2 is at its southeast corner. We use $X_1(p_i), Y_1(p_i)$ to denote the coordinates of p_i with respect to x_1, y_1 , while $X_2(p_i), Y_2(p_i)$ denote its coordinates with respect to x_2, y_2 . Note that $Y_1(p_i) = Y_2(p_i)$, and that $X_2(p_i) = L - X_1(p_i)$ where L is the horizontal dimension of A (see Figure 1).

A point p_i is said to *dominate* p_j with respect to the coordinates x_1, y_1 iff $X_1(p_i) > X_1(p_j)$ and $Y_1(p_i) > Y_1(p_j)$. We use $DOM_1(p_i)$ to denote the set of points in S that are dominated by point p_i w.r.t. x_1, y_1 . $DOM_2(p_i)$ is similarly defined w.r.t. x_2, y_2 . In other words, $DOM_1(p_i)$ (resp. $DOM_2(p_i)$) are the points of S that are below and to the left (resp. right) of p_i . Of course

$DOM_1(p_i) \cup DOM_2(p_i)$ is simply the set of points that are below p_i . For example, in Figure 1, $DOM_1(p_3) = \{p_1\}$ and $DOM_2(p_3) = \{p_2, p_4\}$.

If H is a set of points, then a point of H is a *maximum* w.r.t. x_1, y_1 if no other point of H dominates it w.r.t. x_1, y_1 . From now on we use $MAX_1(H)$ to denote the set of maxima of H w.r.t. x_1, y_1 . $MAX_2(H)$ is similarly defined w.r.t. x_2, y_2 . For example, for the set S of points shown in Figure 1, $MAX_1(S) = \{p_4, p_5\}$ and $MAX_2(S) = \{p_1, p_3, p_5\}$. Observe that listing the elements of $MAX_1(H)$ by increasing x_1 -components is equivalent to listing them by decreasing y_1 -components, so that we can unambiguously talk about 'sorting $MAX_1(H)$ ' without specifying with respect to which coordinate. A similar remark holds for $MAX_2(H)$.

3. The Algorithm

The algorithm enumerates all the RR 's, and chooses the largest-area one. View the RR 's as being partitioned into two classes: Those that are bounded from above by p_i 's, and those that are bounded from above by A . There are $2n+1$ RR 's that are bounded from above by A , and it is not hard to enumerate them in time $O(n \log n)$ (see [NLH] for details). However it is more difficult to enumerate all of the RR 's that are bounded from above by p_i 's in time $O(s+n \log^2 n)$. From now on, we use SR_i to denote the set of RR 's that are bounded from above by p_i , and $BEST(i)$ to denote the largest-area RR in SR_i . The algorithm *MER* (below) computes the largest-area RR .

Algorithm MER

Step 1: Enumerate the RR 's that are bounded from above by A , and let $BEST(0)$ be the largest-area one. As pointed out above, this can be done in time $O(n \log n)$.

Step 2: For $i=1, \dots, n$ do the following: Compute SR_i and let $BEST(i)$ be the largest-area RR in SR_i .

Note: Step 2 is done using algorithm *MER1* which is described later in this Section, and whose

running time is $O(s + n \log^2 n)$.

Step 3: Return the largest-area *RR* among $BEST(0), \dots, BEST(n)$. This takes time $O(n)$.

End of Algorithm MER

Correctness of the algorithm follows from the fact that every *RR* either belongs to one of the SR_i 's or is bounded from above by A . The crucial issue is that of implementing Step 2 in time $O(s + n \log^2 n)$. The rest of this Section deals with this problem.

Before describing the algorithm *MERI* for computing the $BEST(i)$'s, we make the following observation.

Observation 3.1 Suppose the elements of $MAX_1(DOM_1(p_i))$ are given in sorted order, and also those of $MAX_2(DOM_2(p_i))$. (In Figure 2, $MAX_1(DOM_1(p_2)) = \{p_4, p_3, p_9\}$, $MAX_2(DOM_2(p_2)) = \{p_7, p_1\}$, and the rectangles in SR_i are shown dotted). Then it is possible to compute SR_i (and hence $BEST(i)$) in time $O(|SR_i|)$.

Proof: First we observe that

$$|SR_i| = O(|MAX_1(DOM_1(p_i))| + |MAX_2(DOM_2(p_i))|).$$

Thus it suffices to show that SR_i can be computed in time $O(|MAX_1(DOM_1(p_i))| + |MAX_2(DOM_2(p_i))|)$. This can indeed be done by simultaneously scanning the elements of $MAX_1(DOM_1(p_i))$ by increasing x_1 -coordinate, and those of $MAX_2(DOM_2(p_i))$ by increasing x_2 -coordinate. This 'scan' is reminiscent of the way two sorted sequences are merged, and we leave its detailed specification to the reader. \square

Before describing algorithm *MERI*, we recall the following result of Overmars and Van Leeuwen: There exists a data structure for dynamically maintaining the maxima of a set of points in the plane w.r.t. a set of coordinate axes x, y , such that insertions and deletions take time $O(\log^2 n)$. Such an *augmented tree structure* (as it is called in [OV]) takes $O(n)$ storage space, and can initially be created in time $O(n \log n)$. At any time, the maxima are available at the root, in sorted order. If the points are stored in the augmented tree structure according to their x -

coordinate, then a *split* operation about any vertical line $x=x_0$ can be implemented in time $O(\log^2 n)$. Such a *split* operation results in two augmented tree structures: One for the points to the left of the vertical line, and one for those to its right. A *concatenate* operation also takes $O(\log^2 n)$ time and has the reverse effect of a *split*.

Now we have all the ingredients for describing algorithm *MER1*.

Algorithm MER1

Input: p_1, \dots, p_n, A

Output: $BEST(1), \dots, BEST(n)$

Step 1: Create two augmented tree structures $T1$ and $T2$, each of which initially contains all of the p_i 's. $T1$ (resp. $T2$) is for the maxima w.r.t. x_1, y_1 (resp. x_2, y_2), and the p_i 's are stored in it according to their x_1 (resp. x_2) coordinate. As previously mentioned, both $T1$ and $T2$ can be created in time $O(n \log n)$.

Step 2: Sweep a horizontal line from the top edge of A down to the bottom edge of A , and whenever the line encounters a p_i do the following (i)-(v):

- (i) Delete p_i from both $T1$ and $T2$. This takes $O(\log^2 n)$ time.
- (ii) Split $T1$ about $X_1(p_i)$, resulting in $T1_left$ and $T1_right$. Note that the elements of $MAX_1(DOM_1(p_i))$ are available in sorted order at the root of $T1_left$ (note that this is true only because all the points above p_i have already been encountered by the downward-sweeping line and hence deleted). This takes time $O(\log^2 n)$.
- (iii) Similarly to (ii), split $T2$ about $X_2(p_i)$ and obtain $T2_left$ and $T2_right$. Note that $MAX_2(DOM_2(p_i))$ are available in sorted order at the root of $T2_right$. Again, this takes time $O(\log^2 n)$.
- (iv) Obtain SR_i and $BEST(i)$ from $MAX_1(DOM_1(p_i))$ and $MAX_2(DOM_2(p_i))$, in time $O(|SR_i|)$ (Observation 3.1).

(v) Reconstruct $T1$ by concatenating $T1_left$ and $T1_right$. Similarly reconstruct $T2$ from $T2_left$ and $T2_right$. This takes time $O(\log^2 n)$.

End of Algorithm MER1

The cost of *MER1* is dominated by that of Step 2, whose total time is

$$O\left(\sum_{i=1}^n (|SR_i| + \log^2 n)\right) = O(s + n \log^2 n).$$

This implies that algorithm *MER* itself takes $O(s + n \log^2 n)$ time.

4. Conclusion

We have given an $O(s + n \log^2 n)$ time algorithm for solving the maximum empty rectangle problem, where s is the number of *RR*'s. Our algorithm can be combined with the algorithms of Naamad et. al. and Chazelle et. al. to give an $O(\min(s + n \log^2 n, s \log n, n \log^3 n))$ time algorithm for this problem.

References

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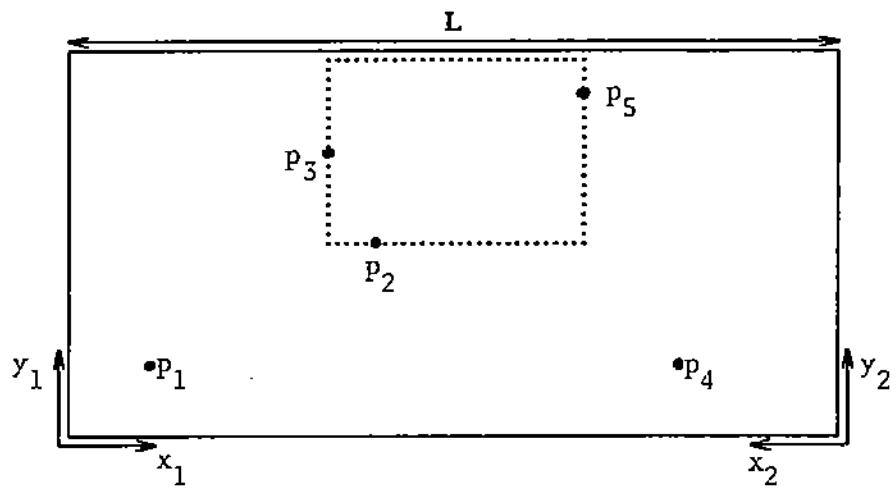


Figure 1

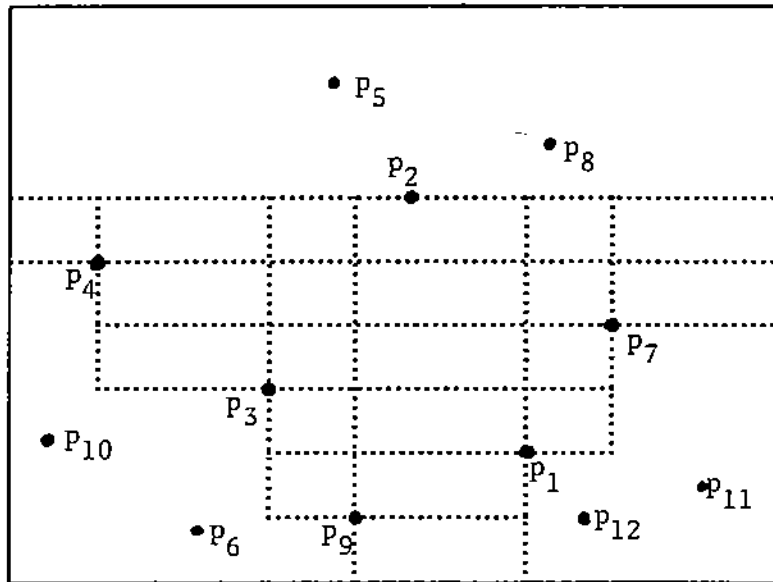


Figure 2