A LINEAR-TIME ALGORITHM FOR CONCAVE ONE-DIMENSIONAL DYNAMIC PROGRAMMING *

Zvi GALIL

Department of Computer Science, 450 Computer Science Building, Columbia University, New York, NY 10027, USA Department of Computer Science, Tel-Aviv University, Tel-Aviv, Israel

Kunsoo PARK

Department of Computer Science, 450 Computer Science Building, Columbia University, New York, NY 10027, USA

Communicated by David Gries Received 2 May 1989 Revised 1 September 1989

Keywords: Dynamic programming, quadrangle inequality, total monotonicity

The one-dimensional dynamic programming problem is defined as follows: given a real-valued function w(i, j) for integers $0 \le i \le j \le n$ and E[0], compute

$$E[j] = \min_{0 \leq i < j} \{D[i] + w(i, j)\} \quad \text{for } 1 \leq j \leq n,$$

where D[i] is computed from E[i] in constant time. The least weight subsequence problem [4] is a special case of the problem where D[i] = E[i]. The modified edit distance problem [3], which arises in molecular biology, geology, and speech recognition, can be decomposed into 2n copies of the problem.

Let A be an $n \times m$ matrix. A[i, j] denotes the element in the ith row and the jth column. A[i:i',j:j'] denotes the submatrix of A that is the intersection of rows i, i+1,...,i' and columns j, j+1,...,j'. We say that the cost function w is concave if it satisfies the quadrangle inequality [7]

$$w(a, c) + w(b, d) \le w(b, c) + w(a, d),$$

for $a \le b \le c \le d.$

* Work supported in part by NSF Grants CCR-86-05353 and CCR-88-14977.

In the concave one-dimensional dynamic programming problem w is concave as defined above. A condition closely related to the quadrangle inequality was introduced by Aggarwal et al. [1]. An $n \times m$ matrix A is totally monotone if for all a < b and c < d,

$$A[a,c] > A[b,c] \Rightarrow A[a,d] > A[b,d].$$

Let r(j) be the smallest row index such that A[r(j), j] is the minimum value in column j. Then total monotonicity implies

$$r(1) \leqslant r(2) \leqslant \cdots \leqslant r(m).$$
 (*)

That is, the minimum row indices are nondecreasing. We say that an element A[i,j] is dead if $i \neq r(j)$. A submatrix of A is dead if all of its elements are dead. Note that the quadrangle inequality implies total monotonicity, but the converse is not true. Aggarwal et al. [1] show that the row maxima of a totally monotone $n \times m$ matrix A can be found in O(n+m) time if A[i,j] for any i,j can be computed in constant time. Their algorithm is easily adapted to find the column minima. We will refer to their algorithm as the SMAWK algorithm.

Let B[i,j] = D[i] + w(i,j) for $0 \le i < j \le n$. We say that B[i,j] is available if D[i] is known and

therefore B[i,j] can be computed in constant time. Then the problem is to find the column minima in the upper triangular matrix B with the restriction that B[i, j] is available only after the column minima for columns 1, 2, ..., i have been found. It is easy to see that when w satisfies the quadrangle inequality, B also satisfies the quadrangle inequality. For the concave problem Hirschberg and Larmore [4] and later Galil and Giancarlo [3] gave $O(n \log n)$ algorithms using queues. Wilber [6] proposed an O(n) time algorithm when D[i] = E[i]. However, his algorithm does not work if the availability of matrix B must be obeyed, which happens when many copies of the problem proceed simultaneously (i.e., the computation is interleaved among many copies) as in the modified edit distance problem [3] and the mixed convex and concave cost problem [2]. Eppstein [2] extended Wilber's algorithm for interleaved computation. Our algorithm is more general than Eppstein's; it works for any totally monotone matrix B (we use only relation (*)), whereas Eppstein's algorithm works only when B[i,j] = D[i] + w(i,j). Our algorithm is also simpler than both Wilber's and Eppstein's. Recently, Larmore and Schieber [5] reported another lineartime algorithm, which is quite different from ours.

The algorithm consists of a sequence of iterations. Fig. 1 shows a typical iteration. We use N[j], $1 \le j \le n$, to store interim column minima

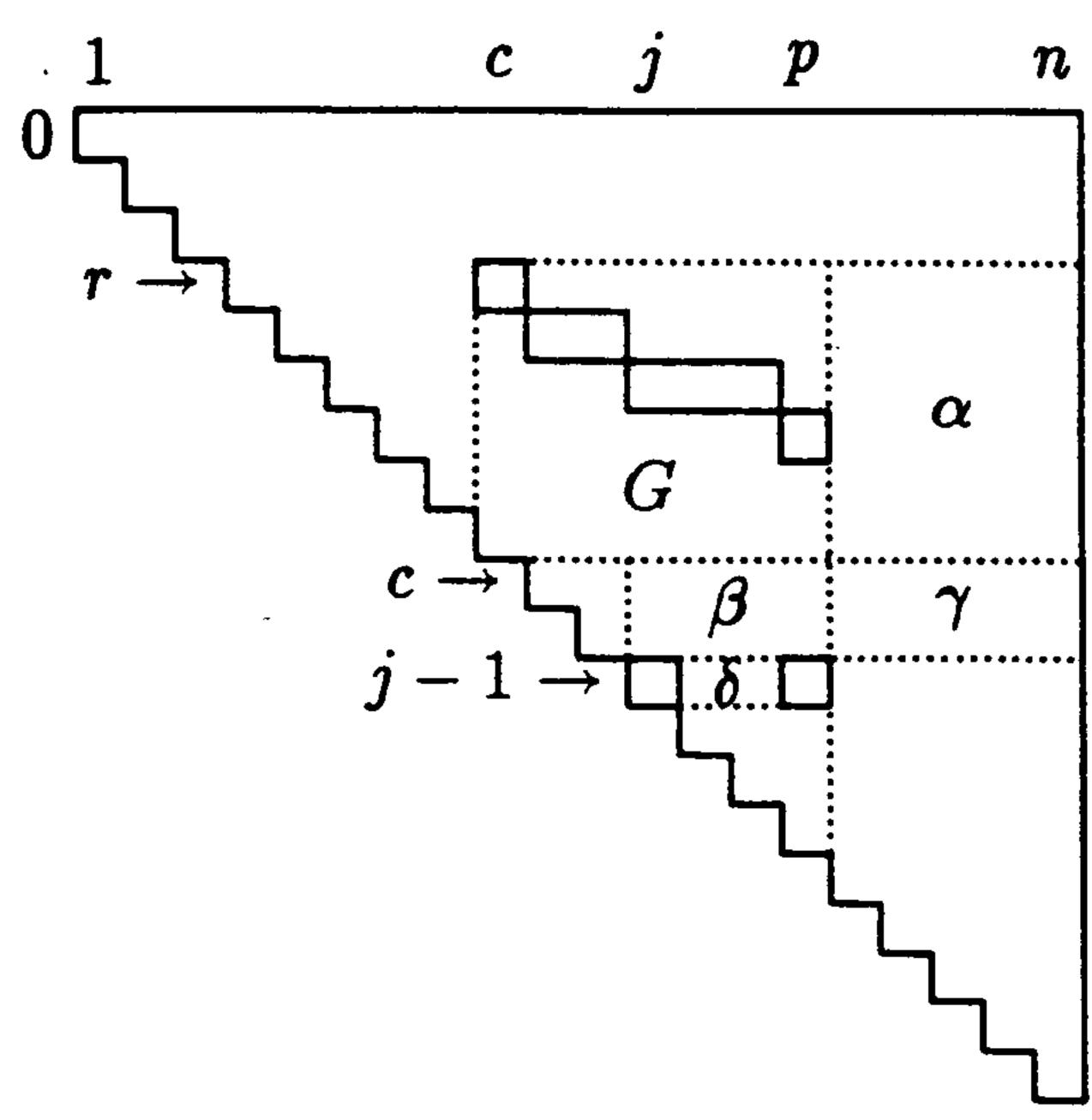


Fig. 1. Matrix B at a typical iteration.

before row r; N[j] = B[i, j] for some i < r (the usage will be clear shortly). At the beginning of each iteration the following invariants hold:

- (a) $0 \le r$ and r < c.
- (b) E[j] for all $1 \le j < c$ have been found.
- (c) E[j] for $j \ge c$ is $\min(N[j], \min_{i \ge r} B[i, j])$. Invariant (b) means that D[i] for all $0 \le i < c$ are known, and therefore B[i, j] for $0 \le i < c$ and $c \le j \le n$ is available. Initially, r = 0, c = 1, and all N[j] are $+\infty$.

Let $p = \min(2c - r, n)$, and let G be the union of N[c:p] and B[r:c-1, c:p], N[c:p] as its first row and B[r:c-1, c:p] as the other rows. G is a $(c-r+1)\times(c-r+1)$ matrix unless 2c-r>n. Let F[j], $c \le j \le p$, denote the column minima of G. Since matrix G is totally monotone, we use the SMAWK algorithm to find the column minima of G. Once F[c:p] are found, we compute E[j] for j=c, c+1,... as follows. Obviously, E[c]=F[c]. For $c+1 \le j \le p$, assume inductively that B[c:j-2,j:p] (β in Fig. 1) is dead and B[j-1,j:n] is available. It is trivially true when j=c+1. By the assumption $E[j]=\min(F[j],B[j-1,j])$.

- (1) If B[j-1,j] < F[j], then E[j] = B[j-1,j], and by relation (1) B[r:j-2,j:n] (α , β , γ , and the part of G above β in Fig. 1) and N[j:n] are dead. We start a new iteration with c=j+1 and r=j-1.
- (2) If $F[j] \le B[j-1,j]$, then E[j] = F[j]. We compare B[j-1,p] with F[p].
 - (2.1) If B[j-1,p] < F[p], B[r:j-2,p+1:n] (α and γ in Fig. 1) is dead by relation (*). B[c:j-2,j:p] (β in Fig. 1) is dead by the assumption. Thus only F[j+1:p] among B[0:j-2,j+1:n] may become column minima in the future computation. We store F[j+1:p] in N[j+1:p] and start a new iteration with c=j+1 and r=j-1.
 - (2.2) If $F[p] \le B[j-1,p]$, B[j-1,j:p](δ in Fig. 1) is dead by relation (*) in submatrix B[r:j-1,j:p] (β , δ , and the part of G above β). Since B[j,j+1:n] is available from E[j], the assumption holds at j+1. We go on to column j+1.

```
procedure concave 1D
    c \leftarrow 1;
    r \leftarrow 0;
    N[1:n] \leftarrow +\infty;
    while c \leq n do
         p \leftarrow \min(2c - r, n);
         use SMAWK to find column minima F[c:p] of G;
         E[c] \leftarrow F[c];
         for j \leftarrow c + 1 to p do
              if B[j-1,j] < F[j] then
                   E[j] \leftarrow B[j-1,j];
                   break
              else
                   E[j] \leftarrow F[j];
                  if B[j-1,p] < F[p] then
                       N[j+1:p] \leftarrow F[j+1:p];
                        break
                   end if
              end if
         end for
         if j \leq p then
             c \leftarrow j + 1;
              r \leftarrow j-1
         else
              c \leftarrow p + 1;
              r \leftarrow \max(r, \text{ row of } F[p])
         end if
    end while
end
```

Fig. 2. The algorithm for concave 1D dynamic programming.

If case (2.2) is repeated until j = p, we have found E[j] for $c \le j \le p$. We start a new iteration with c = p + 1. If the row of F[p] is greater than r, it becomes the new r (it may be smaller than r if it is the row of N[p]). Note that the three invariants hold at the beginning of new iterations. Figure 2 shows the algorithm, where the **break** statement causes the innermost enclosing loop to be excited immediately.

Each iteration takes time O(c-r). If either case (1) or case (2.1) happens, we charge the time to rows $r, \ldots, c-1$ because r is increased by $(j-1)-r \ge c-r$. If case (2.2) is repeated until j=p, there are two cases. If p < n, we charge the time to columns c, \ldots, p because c is increased by $(p+1)-c \ge c-r+1$. If p=n, we have finished the whole computation, and rows $r, \ldots, c-1$ (< n) have not been charged yet; we charge the time to the rows. Since c and r never decrease, only constant time is charged to each row or column. Thus the total time of the algorithm is linear in n.

References

- [1] A. Aggarwal, M.M. Klawe, S. Moran, P. Shor and R. Wilber, Geometric applications of a matrix-searching algorithm, *Algorithmica* 2 (1987) 195-208.
- [2] D. Eppstein, Sequence comparison with mixed convex and concave costs, *J. Algorithms*, to appear.
- [3] Z. Galil and R. Giancarlo, Speeding up dynamic programming with applications to molecular biology, *Theoret. Comput. Sci.* 64 (1989) 107–118.
- [4] D.S. Hirschberg and L.L. Larmore, The least weight subsequence problem, SIAM J. Comput. 16 (4) (1987) 628-638.
- [5] L.L. Larmore and B. Schieber, On-line dynamic programming with applications to the prediction of RNA secondary structure, to be presented at *1st Ann. ACM-SIAM Symp. on Discrete Algorithms*.
- [6] R. Wilber, The concave least-weight subsequence problem revisited, J. Algorithms 9 (1988) 418-425.
- [7] F.F. Yao, Speed-up in dynamic programming, SIAM J. Algebraic Discrete Methods 3 (1982) 532-540.