A Linear-Time Algorithm for Concave One-Dimensional Dynamic Programming

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A Linear-Time Algorithm for Concave One-Dimensional Dynamic Programming*

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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

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

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 *i*th row and the *j*th column. A[i:i', j:j'] denotes the submatrix of A that is the intersection of rows $i, i+1, \ldots, i'$ and columns $j, j+1, \ldots, j'$. We say that the cost function w is *concave* if it satisfies the quadrangle inequality [7]

$$w(a,c) + w(b,d) \le w(b,c) + w(a,d), \quad \text{for } a \le b \le c \le d.$$

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,

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

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

$$r(1) \le r(2) \le \dots \le r(m). \tag{1}$$

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 for $a \leq b \leq c \leq d$, 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 $1, 2, \ldots, i$ 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

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Figure 1. Matrix B at a typical iteration

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 (1)), 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 linear-time algorithm, which is quite different from ours.

The algorithm consists of a sequence of iterations. Figure 1 shows a typical iteration. We use N[j], $1 \le j \le n$, to store interim column minima 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) $0 \leq r$ 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 $j = c, c+1, \ldots$ 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 Figure 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) $B[r: j-2, j: n] (\alpha, \beta, \gamma, \alpha)$ and the part of G above β in Figure 1) and N[j: n] are dead. We start a new iteration with c = j + 1 and r = j 1.
- (2) If $F[j] \leq B[j-1,j]$, 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 Figure 1) is dead by relation (1). B[c: j-2, j:p] (β in Figure 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 $F[p] \leq B[j-1,p]$, B[j-1,j:p] (δ in Figure 1) is dead by relation (1) 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

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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
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Figure 2. The algorithm for concave 1D dynamic programming

E[j], the assumption holds at j + 1. We go on to column j + 1.

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 exited 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.

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