programming pearls

by Jon Bentley

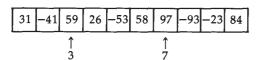
ALGORITHM DESIGN TECHNIQUES

The September 1983 column described the "everyday" impact that algorithm design can have on programmers: an algorithmic view of a problem gives insights that may make a program simpler to understand and to write. In this column we'll study a contribution of the field that is less frequent but more impressive: sophisticated algorithmic methods sometimes lead to dramatic performance improvements.

This column is built around one small problem, with an emphasis on the algorithms that solve it and the techniques used to design them. Some of the algorithms are a little complicated, but the complication is justified; while the first algorithm we'll study takes 39 days to solve a problem of size 10,000, the final algorithm solves the same problem in just a third of a second.

The Problem and a Simple Program

The problem arose in one-dimensional pattern recognition; I'll describe its history later. The input is a vector X of N real numbers; the output is the maximum sum found in any contiguous subvector of the input. For instance, if the input vector is



then the program returns the sum of X[3...7], or 187. The problem is easy when all the numbers are positive—the maximum subvector is the entire input vector. The rub comes when some of the numbers are negative. Should we include a negative number in hopes that the positive numbers to its sides will compensate for its negative contribution? To complete the definition of the problem, we'll say that when all inputs are negative the maximum sum subvector is the empty vector, which has sum zero.

The obvious program for this task is simple: for each pair of integers L and U (where $1 \le L \le U \le N$), compute the sum of X[L ... U] and check whether that sum is greater than the maximum sum so far. The pseudocode given in Algorithm 1 is short, straightforward, and easy to understand. Unfortunately, it has the severe disadvantage of being slow. On the computer I typically use, for instance, the code takes about an hour

if N is 1,000 and 39 days if N is 10,000 (we'll get to

timing details later).

Those times are anecdotal; we get a different kind of feeling for the algorithm's efficiency using "big-oh" notation. The statements in the outermost loop are executed exactly N times, and those in the middle loop are executed at most N times in each execution of the outer loop. Multiplying those two factors of N shows that the four lines contained in the middle loop are executed $O(N^2)$ times. The loop in those four lines is never executed more than N times, so its cost is O(N). Multiplying the cost per inner loop times its number of executions shows that the cost of the entire program is proportional to N cubed, so we'll refer to this as a cubic algorithm.

Those simple steps illustrate the technique of "bigoh" analysis of run time and many of its strengths and weaknesses. Its primary weakness is that we still don't really know the amount of time the program will take for any particular input; we just know that the number of steps it executes is $O(N^3)$. Two strong points of the method often compensate for that weakness. "Big-oh" analyses are usually easy to perform (as above), and the asymptotic run time is often sufficient for a "back-ofthe-envelope" calculation to decide whether or not a program is efficient enough for a given application.

The next several sections use asymptotic run time as the only measure of program efficiency. If that makes

```
MaxSoFar := 0.0
for L := 1 to N do
    for U := L to N do
        sum := 0.0
        for I := L to U do
            Sum := Sum + X[I]
          Sum now contains the
             sum of X[L..U] */
       MaxSoFar := max(MaxSoFar, Sum)
```

Algorithm 1. The cubic algorithm

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¹ The notation $O(N^2)$ can be thought of as "proportional to N^2 "; both $15N^2 + 100N$ and $N^2/2 - 10$ are $O(N^2)$. Informally, f(N) = O(g(N)) means that f(N) < cg(N) for some constant c and sufficiently large values of N. A formal definition of the notation can be found in most textbooks on algorithm design or discrete mathematics.

you uncomfortable, peek ahead to the section on "What Does It Matter?", which shows that for this problem such analyses are extremely informative. Before you read on, take a minute to try to find a faster algorithm for this problem.

Two Quadratic Algorithms

Most programmers have the same response to Algorithm 1: "There's an obvious way to make it a lot faster." There are two obvious ways, however, and if one is obvious to a given programmer then the other often isn't. Both algorithms are quadratic—they take $O(N^2)$ steps on an input of size N—and both achieve their run time by computing the sum of $X[L \ldots U]$ in a constant number of steps rather than in the U-L+1 steps of Algorithm 1. But the two quadratic algorithms use very different methods to compute the sum in constant time.

The first quadratic algorithm computes the sum quickly by noticing that the sum of X[L ... U] has an intimate relationship to the sum previously computed, that of X[L ... U - 1]. Exploiting that relationship leads to Algorithm 2. The statements inside the first loop are executed N times, and those inside the second loop are executed at most N times on each execution of the outer loop, so the total run time is $O(N^2)$.

An alternative quadratic algorithm computes the sum in the inner loop by accessing a data structure (called CumArray) built before the outer loop is ever executed. The I^{th} element of CumArray contains the cumulative sum of the values in X[1 ... I], so the sum of the values in X[L ... U] can be found by computing CumArray[U] - CumArray[U - 1]. This results in Algorithm 2b, which takes $O(N^2)$ time; the analysis is the same as for Algorithm 2.

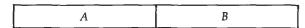
The algorithms we have seen so far inspect all possible pairs of starting and ending values of subvectors and consider the sum of the numbers in that subvector. Because there are $O(N^2)$ subvectors, any algorithm that inspects all such values must take at least quadratic time. Can you think of a way to sidestep this problem and achieve an algorithm that runs in less time?

A Divide-and-Conquer Algorithm

Our first subquadratic algorithm is complicated; if you get bogged down in its details, you won't lose much by skipping to the next section. It is based on the following divide-and-conquer schema:

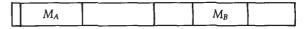
To solve a problem of size N, recursively solve two subproblems of size approximately N/2, and combine their solutions to yield a solution to the complete problem.

In this case the original problem deals with a vector of size N, so the most natural way to divide it into subproblems is to create two subvectors of approximately equal size, which we'll call A and B:

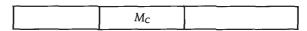


We then recursively find the maximum subvectors in A

and B, which we'll call M_A and M_B :



It is tempting to think that we have solved the problem because the maximum sum subvector of the entire vector must be either M_A or M_B , and that is almost right. In fact, the maximum is either entirely in A, entirely in B, or it crosses the border between A and B(which we'll call M_C for the maximum crossing the border):



Thus our divide-and-conquer algorithm will compute M_A and M_B recursively, compute M_C by some other means, and then return the maximum of the three.

That description is almost enough to write code. All we have left to describe is how we'll handle small vectors and how we'll compute M_C . The former is easy: the maximum of a one-element vector is the only value in the vector (or zero if that number is negative), and the maximum of a zero-element vector was previously defined to be zero. To compute M_C we observe that its component in A is the largest subvector starting at the boundary and reaching into A, and similarly for its component in B. Putting these facts together leads to Algorithm 3, which is originally invoked by the statement

```
Answer := MaxSum(1, N)
```

Although the code is complicated (and easy to get wrong), it does yield a substantial decrease in run time:

```
MaxSoFar := 0.0
for L := 1 to N do
    Sum := 0.0
    for U := L to N do
        Sum := Sum + X[I]
        /* Sum now contains the
        sum of X[L..U] */
        MaxSoFar := max(MaxSoFar, Sum)
```

Algorithm 2. The first quadratic algorithm

Algorithm 2b. An alternative quadratic algorithm

```
recursive function MaxSum(L, U)
                     /* Zero-element vector */
   if L > U then
        return 0.0
                    /* One-element vector */
   if L = U then
        return max(0.0, X[L])
                      /* A is X[L..M], B is X[M + 1..U] */
   M := (L + U)/2
    /* Find max crossing to left */
        Sum := 0.0; MaxToLeft := 0.0
        for I := M downto L do
            Sum := Sum + X[I]
           MaxToLeft := max(MaxToLeft, Sum)
    /* Find max crossing to right */
        Sum := 0.0; MaxToRight := 0.0
        for I := M + 1 to U do
            Sum := Sum + X[I]
            MaxToRight := max(MaxToRight, Sum)
    MaxCrossing := MaxToLeft + MaxToRight
   MaxInA := MaxSum(L, M)
   MaxInB := MaxSum(M + 1, U)
    return max(MaxCrossing, MaxInA, MaxInB)
```

Algorithm 3. A divide-and-conquer algorithm

it solves the problem in $O(N \log N)$ time. There are a number of ways of proving this fact. An informal argument observes that the algorithm does O(N) work on each of $O(\log N)$ levels of recursion. The argument can be made more precise by the use of recurrence relations; if T(N) denotes the time to solve a problem of size N, then we can show that T(1) = O(1) and that

$$T(N) = 2T(N/2) + O(N).$$

Most textbooks on algorithm design show that this recurrence has the solution $T(N) = O(N \log N)$.

A Scanning Algorithm

We'll now use the simplest kind of algorithm that operates on arrays: it starts at the left end (element X[1]) and scans through to the right end (element X[N]), keeping track of the maximum sum subvector seen so far. The maximum is initially zero. Suppose we've solved the problem for X[1 . I - 1]; how can we extend that to a solution for the first I elements? We use reasoning similar to that of the divide-and-conquer algorithm: the maximum sum in the first I elements is either the maximum sum in the first I - 1 elements (which we'll call MaxSoFar), or it is that of a subvector that ends in position I (which we'll call MaxEndingHere):

	MaxSoFar	MaxEndingHere
1 1		

Recomputing MaxEndingHere from scratch (using code like that in Algorithm 3) yields yet another quadratic algorithm. We can get around this by using the technique that led to Algorithm 2: instead of computing the maximum subvector ending in position I from scratch,

we'll use the maximum subvector that ends in position I-1. This results in Algorithm 4.

The key to understanding this program is the variable MaxEndingHere. Before the first assignment statement in the loop, MaxEndingHere contains the value of the maximum subvector ending in position I-1; the assignment statement modifies it to contain the value of the maximum subvector ending in position I. The statement increases it by the value X[I] so long as doing so keeps it positive; when it goes negative, it is reset to zero (that is, the maximum subvector ending at I is the empty vector). Although the code is subtle, it is short and fast: its running time is O(N) (so we'll refer to it as a linear algorithm). David Gries systematically derives and verifies this algorithm in his paper "A Note on the Standard Strategy for Developing Loop Invariants and Loops" (in Science of Computer Programming 2, pp. 207-214).

What Does It Matter?

So far I've played fast and loose with "big-ohs"; it's time for me to come clean and tell about the run times of the programs. I implemented the four primary algorithms (all except Algorithm 2b) in the C language on a Digital

Algorithm 4. The linear algorithm

TABLE I. Summary of the Algorithms

Algorithm		1	2	3	4
Lines of C Code	•	8	7	14	7
Run time in microseconds		3.4N³	13N²	46N log N	33 <i>N</i>
Time to solve	10 ²	3.4 secs	130 msecs	30 msecs	3.3 msecs
problem of size	10^{3}	.94 hrs	13 secs	.45 secs	33 msecs
	10⁴	39 days	22 mins	6.1 secs	.33 secs
	10 ⁵	108 yrs	1.5 days	1.3 min	3.3 secs
	10 ⁶	108 mill	5 mos	15 min	33 secs
Max problem	sec	67	280	2000	30,000
solved in one	min	260	2200	82,000	2,000,000
	hr	1000	17,000	3,500,000	120,000,000
	day	3000	81,000	73,000,000	2,800,000,000
If N multiplies by time multiplies b		1000	100	10+	10
If time multiplies 10, N multiplies	•	2.15	3.16	10-	10

Equipment Corporation VAX-11/750,² timed them, and extrapolated the run times to achieve Table I.

This table makes a number of points. The most important is that proper algorithm design can make a big difference in run time; that point is underscored by the middle rows. The table also shows something of the different character of cubic, quadratic, $N \log N$ and linear algorithms: the last two rows show how the problem size and run time vary as a function of each other.

Another important point is that when we're comparing cubic, quadratic, and linear algorithms with one another, the constant factors of the programs don't matter much. To underscore this point, I conducted an experiment in which I tried to make the constant factors of two algorithms differ by as much as possible. To achieve a huge constant factor I implemented Algorithm 4 on a BASIC interpreter on a Radio Shack TRS-80 Model III microcomputer. For the other end of the spectrum, Eric Grosse of AT&T Bell Laboratories and I implemented Algorithm 1 in fine-tuned FORTRAN on a CRAY-1 supercomputer. We got the disparity we wanted: the run time of the cubic algorithm was measured as $3.0N^3$ nanoseconds, while the run time of the linear algorithm was 19,500,000N nanoseconds. Table II

shows how those expressions translate to times for various problem sizes (the same data is displayed graphically in Figure 1.)

The difference in constants (a factor of six and a half million) allowed the cubic algorithm to start off faster, but the linear algorithm was bound to catch up. In this case, the break-even point for the two algorithms is around 2,500, where each takes about 50 seconds.

Principles

The history of the problem sheds light on the algorithm design techniques. The problem arose in a pattern-matching procedure designed by Ulf Grenander of Brown University in the two-dimensional form described in Problem 7. In that form, the maximum sum subarray was the maximum likelihood estimator of a certain kind of pattern in a digitized picture. Because the two-dimensional problem required too much time to solve, Grenander simplified it to one dimension to gain insight into its structure.

Grenander observed that the cubic time of Algorithm 1 was prohibitively slow, and derived Algorithm 2. In 1977 he described the problem to Michael Shamos of UNILOGIC, Ltd. (then of Carnegie-Mellon University)

TABLE II. The Tyranny of Asymptotics

N	Cray-1, FORTRAN, Cubic Algorithm	TRS-80, BASIC, Linear Algorithm
10	3.0 microsecs	200 millisecs
100	3.0 millisecs	2.0 secs
1000	3.0 secs	20 secs
10,000	49 mins	3.2 mins
100,000	35 days	32 mins
1,000,000	95 yrs	5.4 hrs

² VAX is a trademark of Digital Equipment Corporation.

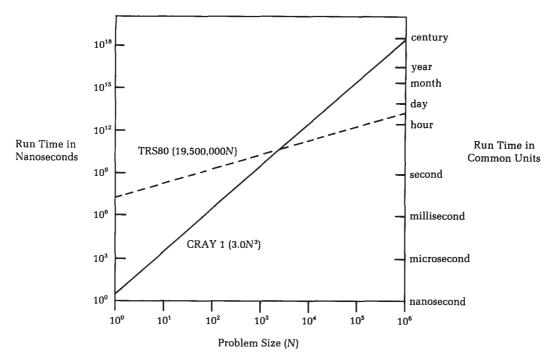


FIGURE 1. The Run Times of Two Programs

who overnight designed Algorithm 3. When Shamos showed me the problem shortly thereafter, we thought that it was probably the best possible; researchers had just shown that several similar problems require time proportional to $N \log N$. A few days later Shamos described the problem and its history at a seminar attended by Jay Kadane (a statistician at Carnegie-Mellon University), who designed the linear-time Algorithm 4 within a minute. Fortunately, we know that there can be no better algorithm: any algorithm must look at all N inputs.

Even though the one-dimensional problem is now completely solved, Grenander's original two-dimensional problem remains open. Because of the computational expense of all known algorithms, Grenander had to abandon that approach to the pattern-matching problem. Readers who feel that the linear-time algorithm for the one-dimensional problem is "obvious" are therefore urged to find an "obvious" algorithm for Problem 7!

Although the algorithms in this story were never incorporated into a system, they do illustrate several important algorithm design techniques that have had substantial impact on many systems (see the sidebar on page 870):

Save state to avoid recomputation. This simple form of dynamic programming arose in Algorithms 2 and 4. By using space to store results, we avoid using time to recompute them.

Preprocess information into data structures. This was the technique underlying Algorithm 2b: the CumArray structure allowed the sum of a subvector to be accessed in just a couple of operations.

Divide-and-conquer algorithms. Algorithm 3 uses a simple form of divide-and-conquer; textbooks on algorithm design describe more advanced forms.

Scanning algorithms. Problems on arrays can often be solved by asking "how can I extend a solution for X[1 ... I-1] to a solution for X[1 ... I]?" In Algorithm 4 we had to remember both the old answer and some auxiliary data to compute the new answer.

Cumulatives. Algorithm 2b uses a cumulative table in which the I^{th} element contains the sum of the first I values of X; such tables are common when dealing with ranges. In business data processing applications, for instance, one finds the sales from March to October by subtracting the February year-to-date sales from the October year-to-date sales.

Lower bounds. Algorithm designers sleep peacefully only when their algorithm is known to be the best possible because they have proved a matching lower bound. The linear lower bound for this problem was easy; more complex lower bounds can be difficult.

Problems

- Algorithms 3 and 4 use subtle code that is easy to get wrong. Use the program verification techniques described in the December 1983 column to argue the correctness of the code; specify the loop invariants carefully.
- 2. Our analysis of the four algorithms was done only at the "big-oh" level of detail. Analyze the number of additions and comparisons done by each algorithm as exactly as possible; does this exercise give any insight into the behavior of the algorithms?