Shared-Memory Programming with OpenMP

Like Pthreads, OpenMP is an API for shared-memory parallel programming. The "MP" in OpenMP stands for "multiprocessing," a term that is synonymous with shared-memory parallel computing. Thus, OpenMP is designed for systems in which each thread or process can potentially have access to all available memory, and, when we're programming with OpenMP, we view our system as a collection of cores or CPUs, all of which have access to main memory, as in Figure 5.1.

Although OpenMP and Pthreads are both APIs for shared-memory programming, they have many fundamental differences. Pthreads requires that the programmer explicitly specify the behavior of each thread. OpenMP, on the other hand, sometimes allows the programmer to simply state that a block of code should be executed in parallel, and the precise determination of the tasks and which thread should execute them is left to the compiler and the run-time system. This suggests a further difference between OpenMP and Pthreads, that is, that Pthreads (like MPI) is a library of functions that can be linked to a C program, so any Pthreads program can be used with any C compiler, provided the system has a Pthreads library. OpenMP, on the other hand, requires compiler support for some operations, and hence it's entirely possible that you may run across a C compiler that can't compile OpenMP programs into parallel programs.

These differences also suggest why there are two standard APIs for shared-memory programming: Pthreads is lower level and provides us with the power to program virtually any conceivable thread behavior. This power, however, comes with some associated cost—it's up to us to specify every detail of the behavior of each thread. OpenMP, on the other hand, allows the compiler and run-time system to determine some of the details of thread behavior, so it can be simpler to code some parallel behaviors using OpenMP. The cost is that some low-level thread interactions can be more difficult to program.

OpenMP was developed by a group of programmers and computer scientists who believed that writing large-scale high-performance programs using APIs such as Pthreads was too difficult, and they defined the OpenMP specification so that shared-memory programs could be developed at a higher level. In fact, OpenMP was explicitly designed to allow programmers to *incrementally* parallelize

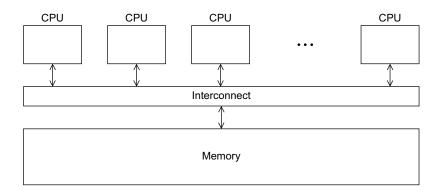


FIGURE 5.1

A shared-memory system

existing serial programs; this is virtually impossible with MPI and fairly difficult with Pthreads.

In this chapter, we'll learn the basics of OpenMP. We'll learn how to write a program that can use OpenMP, and we'll learn how to compile and run OpenMP programs. We'll then learn how to exploit one of the most powerful features of OpenMP: its ability to parallelize many serial for loops with only small changes to the source code. We'll then look at some other features of OpenMP: task-parallelism and explicit thread synchronization. We'll also look at some standard problems in shared-memory programming: the effect of cache memories on shared-memory programming and problems that can be encountered when serial code—especially a serial library—is used in a shared-memory program. Let's get started.

5.1 GETTING STARTED

OpenMP provides what's known as a "directives-based" shared-memory API. In C and C++, this means that there are special preprocessor instructions known as pragmas. Pragmas are typically added to a system to allow behaviors that aren't part of the basic C specification. Compilers that don't support the pragmas are free to ignore them. This allows a program that uses the pragmas to run on platforms that don't support them. So, in principle, if you have a carefully written OpenMP program, it can be compiled and run on any system with a C compiler, regardless of whether the compiler supports OpenMP.

Pragmas in C and C++ start with

#pragma

As usual, we put the pound sign, #, in column 1, and like other preprocessor directives, we shift the remainder of the directive so that it is aligned with the rest of the

code. Pragmas (like all preprocessor directives) are, by default, one line in length, so if a pragma won't fit on a single line, the newline needs to be "escaped"—that is, preceded by a backslash \. The details of what follows the #pragma depend entirely on which extensions are being used.

Let's take a look at a *very* simple example, a "hello, world" program that uses OpenMP. See Program 5.1.

```
#include <stdio.h>
   #include <stdlib.h>
   #include <omp.h>
5
   void Hello(void); /* Thread function */
6
7
   int main(int argc, char* argv[]) {
8
       /* Get number of threads from command line */
       int thread_count = strtol(argv[1], NULL, 10);
9
10
      pragma omp parallel num_threads(thread_count)
11
       Hello();
12
13
14
       return 0;
   } /* main */
15
16
   void Hello(void) {
17
       int my_rank = omp_get_thread_num();
18
19
       int thread_count = omp_get_num_threads();
20
21
       printf("Hello from thread %d of %d\n", my_rank, thread_count);
22
      /* Hello */
23
   }
```

Program 5.1: A "hello,world" program that uses OpenMP

5.1.1 Compiling and running OpenMP programs

To compile this with gcc we need to include the -fopenmp option:¹

```
$ gcc -g -Wall -fopenmp -o omp_hello omp_hello.c
```

To run the program, we specify the number of threads on the command line. For example, we might run the program with four threads and type

```
$ ./omp_hello 4
```

¹Some older versions of gcc may not include OpenMP support. Other compilers will, in general, use different command-line options to specify that the source is an OpenMP program. For details on our assumptions about compiler use, see Section 2.9.

If we do this, the output might be

```
Hello from thread 0 of 4
Hello from thread 1 of 4
Hello from thread 2 of 4
Hello from thread 3 of 4
```

However, it should be noted that the threads are competing for access to stdout, so there's no guarantee that the output will appear in thread-rank order. For example, the output might also be

```
Hello from thread 1 of 4
Hello from thread 2 of 4
Hello from thread 0 of 4
Hello from thread 3 of 4

Or

Hello from thread 3 of 4
Hello from thread 1 of 4
Hello from thread 2 of 4
Hello from thread 0 of 4
```

or any other permutation of the thread ranks.

If we want to run the program with just one thread, we can type

```
$ ./omp_hello 1
```

and we would get the output

```
Hello from thread 0 of 1
```

5.1.2 The program

Let's take a look at the source code. In addition to a collection of directives, OpenMP consists of a library of functions and macros, so we usually need to include a header file with prototypes and macro definitions. The OpenMP header file is omp.h, and we include it in Line 3.

In our Pthreads programs, we specified the number of threads on the command line. We'll also usually do this with our OpenMP programs. In Line 9 we therefore use the strtol function from stdlib.h to get the number of threads. Recall that the syntax of this function is

The first argument is a string—in our example, it's the command-line argument—and the last argument is the numeric base in which the string is represented—in our example, it's base 10. We won't make use of the second argument, so we'll just pass in a NULL pointer.

If you've done a little C programming, there's nothing really new up to this point. When we start the program from the command line, the operating system starts a single-threaded process and the process executes the code in the main function. However, things get interesting in Line 11. This is our first OpenMP directive, and we're using it to specify that the program should start some threads. Each thread that's forked should execute the Hello function, and when the threads return from the call to Hello, they should be terminated, and the process should then terminate when it executes the return statement.

That's a lot of bang for the buck (or code). If you studied the Pthreads chapter, you'll recall that we had to write a lot of code to fork and join multiple threads: we needed to allocate storage for a special struct for each thread, we used a for loop to start each thread, and we used another for loop to terminate the threads. Thus, it's immediately evident that OpenMP is higher-level than Pthreads.

We've already seen that pragmas in C and C++ start with

pragma

OpenMP pragmas always begin with

pragma omp

Our first directive is a parallel directive, and, as you might have guessed it specifies that the **structured block** of code that follows should be executed by multiple threads. A structured block is a C statement or a compound C statement with one point of entry and one point of exit, although calls to the function exit are allowed. This definition simply prohibits code that branches into or out of the middle of the structured block.

Recollect that **thread** is short for *thread of execution*. The name is meant to suggest a sequence of statements executed by a program. Threads are typically started or **forked** by a process, and they share most of the resources of the process that starts them—for example, access to stdin and stdout—but each thread has its own stack and program counter. When a thread completes execution it **joins** the process that started it. This terminology comes from diagrams that show threads as directed lines. See Figure 5.2. For more details see Chapters 2 and 4.

At its most basic the parallel directive is simply

pragma omp parallel

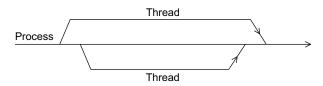


FIGURE 5.2

A process forking and joining two threads

and the number of threads that run the following structured block of code will be determined by the run-time system. The algorithm used is fairly complicated; see the OpenMP Standard [42] for details. However, if there are no other threads started, the system will typically run one thread on each available core.

As we noted earlier, we'll usually specify the number of threads on the command line, so we'll modify our parallel directives with the num_threads clause. A clause in OpenMP is just some text that modifies a directive. The num_threads clause can be added to a parallel directive. It allows the programmer to specify the number of threads that should execute the following block:

```
# pragma omp parallel num_threads(thread_count)
```

It should be noted that there may be system-defined limitations on the number of threads that a program can start. The OpenMP Standard doesn't guarantee that this will actually start thread_count threads. However, most current systems can start hundreds or even thousands of threads, so unless we're trying to start *a lot* of threads, we will almost always get the desired number of threads.

What actually happens when the program gets to the parallel directive? Prior to the parallel directive, the program is using a single thread, the process started when the program started execution. When the program reaches the parallel directive, the original thread continues executing and thread_count - 1 additional threads are started. In OpenMP parlance, the collection of threads executing the parallel block—the original thread and the new threads—is called a **team**, the original thread is called the **master**, and the additional threads are called **slaves**. Each thread in the team executes the block following the directive, so in our example, each thread calls the Hello function.

When the block of code is completed—in our example, when the threads return from the call to Hello—there's an **implicit barrier**. This means that a thread that has completed the block of code will wait for all the other threads in the team to complete the block—in our example, a thread that has completed the call to Hello will wait for all the other threads in the team to return. When all the threads have completed the block, the slave threads will terminate and the master thread will continue executing the code that follows the block. In our example, the master thread will execute the return statement in Line 14, and the program will terminate.

Since each thread has its own stack, a thread executing the Hello function will create its own private, local variables in the function. In our example, when the function is called, each thread will get its rank or id and the number of threads in the team by calling the OpenMP functions <code>omp_get_thread_num</code> and <code>omp_get_num_threads</code>, respectively. The rank or id of a thread is an int that is in the range $0,1,\ldots$, <code>thread_count -1</code>. The syntax for these functions is

```
int omp_get_thread_num(void);
int omp_get_num_threads(void);
```

Since stdout is shared among the threads, each thread can execute the printf statement, printing its rank and the number of threads. As we noted earlier, there is no scheduling of access to stdout, so the actual order in which the threads print their results is nondeterministic.

5.1.3 Error checking

In order to make the code more compact and more readable, our program doesn't do any error checking. Of course, this is dangerous, and, in practice, it's a *very* good idea—one might even say mandatory—to try to anticipate errors and check for them. In this example, we should definitely check for the presence of a command-line argument, and, if there is one, after the call to strtol we should check that the value is positive. We might also check that the number of threads actually created by the parallel directive is the same as thread_count, but in this simple example, this isn't crucial.

A second source of potential problems is the compiler. If the compiler doesn't support OpenMP, it will just ignore the parallel directive. However, the attempt to include omp.h and the calls to omp_get_thread_num and omp_get_num_threads will cause errors. To handle these problems, we can check whether the preprocessor macro _OPENMP is defined. If this is defined, we can include omp.h and make the calls to the OpenMP functions. We might make the following modifications to our program.

Instead of simply including omp.h in the line

```
#include <omp.h>
```

we can check for the definition of <code>_OPENMP</code> before trying to include it:

```
#ifdef _OPENMP
# include <omp.h>
#endif
```

Also, instead of just calling the OpenMP functions, we can first check whether _OPENMP is defined:

```
# ifdef _OPENMP
    int my_rank = omp_get_thread_num();
    int thread_count = omp_get_num_threads();
# else
    int my_rank = 0;
    int thread_count = 1;
# endif
```

Here, if OpenMP isn't available, we assume that the Hello function will be single-threaded. Thus, the single thread's rank will be 0 and the number of threads will be one.

The book's website contains the source for a version of this program that makes these checks. In order to make our code as clear as possible, we'll usually show little, if any, error checking in the code displayed in the text.

5.2 THE TRAPEZOIDAL RULE

Let's take a look at a somewhat more useful (and more complicated) example: the trapezoidal rule for estimating the area under a curve. Recall from Section 3.2 that if y = f(x) is a reasonably nice function, and a < b are real numbers, then we can estimate the area between the graph of f(x), the vertical lines x = a and x = b, and the x-axis by dividing the interval [a,b] into n subintervals and approximating the area over each subinterval by the area of a trapezoid. See Figure 5.3 for an example.

Also recall that if each subinterval has the same length and if we define h = (b - a)/n, $x_i = a + ih$, i = 0, 1, ..., n, then our approximation will be

$$h[f(x_0)/2+f(x_1)+f(x_2)+\cdots+f(x_{n-1})+f(x_n)/2].$$

Thus, we can implement a serial algorithm using the following code:

```
/* Input: a, b, n */
h = (b-a)/n;
approx = (f(a) + f(b))/2.0;
for (i = 1; i <= n-1; i++) {
    x_i = a + i*h;
    approx += f(x_i);
}
approx = h*approx;</pre>
```

See Section 3.2.1 for details.

5.2.1 A first OpenMP version

Recall that we applied Foster's parallel program design methodology to the trapezoidal rule as described in the following list (see Section 3.2.2).

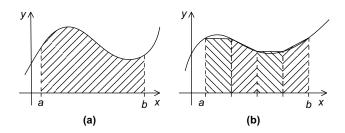


FIGURE 5.3

The trapezoidal rule: (a) area to be estimated and (b) approximate area using trapezoids

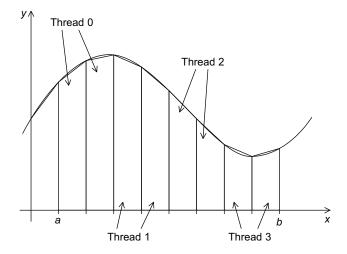


FIGURE 5.4

Assignment of trapezoids to threads

- **1.** We identified two types of tasks:
 - a. Computation of the areas of individual trapezoids, and
 - **b.** Adding the areas of trapezoids.
- **2.** There is no communication among the tasks in the first collection, but each task in the first collection communicates with task 1(b).
- **3.** We assumed that there would be many more trapezoids than cores, so we aggregated tasks by assigning a contiguous block of trapezoids to each thread (and a single thread to each core). Effectively, this partitioned the interval [a,b] into larger subintervals, and each thread simply applied the serial trapezoidal rule to its subinterval. See Figure 5.4 for an example.

We aren't quite done, however, since we still need to add up the threads' results. An obvious solution is to use a shared variable for the sum of all the threads' results, and each thread can add its (private) result into the shared variable. We would like to have each thread execute a statement that looks something like

```
global_result += my_result;
```

However, as we've already seen, this can result in an erroneous value for $global_result$ —if two (or more) threads attempt to simultaneously execute this statement, the result will be unpredictable. For example, suppose that $global_result$ has been initialized to 0, thread 0 has computed $my_result = 1$, and thread 1

²Since we were discussing MPI, we actually used *processes* instead of threads.

has computed my_result = 2. Furthermore, suppose that the threads execute the statement global_result += my_result according to the following timetable:

Time	Thread 0	Thread 1
0 1 2 3 4	<pre>global_result = 0 to register my_result = 1 to register add my_result to global_result store global_result = 1</pre>	<pre>finish my_result global_result = 0 to register my_result = 2 to register add my_result to global_result store global_result = 2</pre>

We see that the value computed by thread $0 \pmod{t} = 1$ is overwritten by thread 1.

Of course, the actual sequence of events might well be different, but unless one thread finishes the computation <code>global_result</code> += <code>my_result</code> before the other starts, the result will be incorrect. Recall that this is an example of a **race condition**: multiple threads are attempting to access a shared resource, at least one of the accesses is an update, and the accesses can result in an error. Also recall that the code that causes the race condition, <code>global_result</code> += <code>my_result</code>, is called a **critical section**. A critical section is code executed by multiple threads that updates a shared resource, and the shared resource can only be updated by one thread at a time.

We therefore need some mechanism to make sure that once one thread has started executing <code>global_result</code> += <code>my_result</code>, no other thread can start executing this code until the first thread has finished. In Pthreads we used mutexes or semaphores. In OpenMP we can use the <code>critical</code> directive

```
# pragma omp critical
  global_result += my_result;
```

This directive tells the compiler that the system needs to arrange for the threads to have **mutually exclusive** access to the following structured block of code. That is, only one thread can execute the following structured block at a time. The code for this version is shown in Program 5.2. We've omitted any error checking. We've also omitted code for the function f(x).

In the main function, prior to Line 16, the code is single-threaded, and it simply gets the number of threads and the input (a, b, and n). In Line 16 the parallel directive specifies that the Trap function should be executed by thread_count threads. After returning from the call to Trap, any new threads that were started by the parallel directive are terminated, and the program resumes execution with only one thread. The one thread prints the result and terminates.

In the Trap function, each thread gets its rank and the total number of threads in the team started by the parallel directive. Then each thread determines the following:

- **1.** The length of the bases of the trapezoids (Line 32)
- 2. The number of trapezoids assigned to each thread (Line 33)

```
#include <stdio.h>
   #include <stdlib.h>
   #include <omp.h>
   void Trap(double a, double b, int n, double* global_result_p);
5
   int main(int argc, char* argv[]) {
7
       double global_result = 0.0;
8
9
       double a, b;
10
       int
       int
               thread_count;
11
12
13
       thread\_count = strtol(argv[1], NULL, 10);
14
       printf("Enter a, b, and n\n");
15
       scanf("%lf %lf %d", &a, &b, &n);
   # pragma omp parallel num_threads(thread_count)
16
       Trap(a, b, n, &global_result);
17
18
       printf("With n = %d trapezoids, our estimate\n", n);
19
       printf("of the integral from %f to %f = \%.14e\n",
20
21
          a, b, global_result);
22
       return 0;
23 } /* main */
24
   void Trap(double a, double b, int n, double* global_result_p) {
25
26
       double h, x, my_result;
27
       double local_a, local_b;
       int i, local_n;
28
       int my_rank = omp_get_thread_num();
29
      int thread_count = omp_get_num_threads();
30
31
32
      h = (b-a)/n;
33
      local_n = n/thread_count;
34
       local_a = a + my_rank*local_n*h;
35
       local_b = local_a + local_n*h;
36
       my_result = (f(local_a) + f(local_b))/2.0;
       for (i = 1; i <= local_n-1; i++) {
37
        x = local_a + i*h;
38
39
        my_result += f(x);
40
41
      my_result = my_result*h;
42
43
   # pragma omp critical
44
       *global_result_p += my_result;
45
   } /* Trap */
```

Program 5.2: First OpenMP trapezoidal rule program

- **3.** The left and right endpoints of its interval (Lines 34 and 35, respectively)
- **4.** Its contribution to global_result (Lines 36–41)

The threads finish by adding in their individual results to global_result in Lines 43 and 44.

We use the prefix local for some variables to emphasize that their values may differ from the values of corresponding variables in the main function—for example, local a may differ from a, although it is the *thread's* left endpoint.

Notice that unless n is evenly divisible by thread_count, we'll use fewer than n trapezoids for global_result. For example, if n=14 and thread_count = 4, each thread will compute

```
local_n = n/thread_count = 14/4 = 3.
```

Thus each thread will only use 3 trapezoids, and global_result will be computed with $4 \times 3 = 12$ trapezoids instead of the requested 14. So in the error checking (which isn't shown) we check that n is evenly divisible by thread_count by doing something like this:

```
if (n % thread_count != 0) {
   fprintf(stderr, "n must be evenly divisible by thread_count\n");
   exit(0);
}
```

Since each thread is assigned a block of local_n trapezoids, the length of each thread's interval will be local_n*h, so the left endpoints will be

```
thread 0: a + 0*local_n*h
thread 1: a + 1*local_n*h
thread 2: a + 2*local_h*h
. . .
```

and in Line 34, we assign

```
local_a = a + my_rank*local_n*h;
```

Furthermore, since the length of each thread's interval will be local_n*h, its right endpoint will just be

```
local_b = local_a + local_n*h;
```

5.3 SCOPE OF VARIABLES

In serial programming, the *scope* of a variable consists of those parts of a program in which the variable can be used. For example, a variable declared at the beginning of a C function has "function-wide" scope, that is, it can only be accessed in the body of the function. On the other hand, a variable declared at the beginning of a .c file but outside any function has "file-wide" scope, that is, any function in the file

in which the variable is declared can access the variable. In OpenMP, the **scope** of a variable refers to the set of threads that can access the variable in a parallel block. A variable that can be accessed by all the threads in the team has **shared** scope, while a variable that can only be accessed by a single thread has **private** scope.

In the "hello, world" program, the variables used by each thread (my_rank and thread_count) were declared in the Hello function, which is called inside the parallel block. Consequently, the variables used by each thread are allocated from the thread's (private) stack, and hence all of the variables have private scope. This is *almost* the case in the trapezoidal rule program; since the parallel block is just a function call, all of the variables used by each thread in the Trap function are allocated from the thread's stack.

However, the variables that are declared in the main function (a, b, n, global_result, and thread_count) are all accessible to all the threads in the team started by the parallel directive. Hence, the *default* scope for variables declared before a parallel block is shared. In fact, we've made implicit use of this: each thread in the team gets the values of a, b, and n from the call to Trap. Since this call takes place in the parallel block, it's essential that each thread has access to a,b, and n when their values are copied into the corresponding formal arguments.

Furthermore, in the Trap function, although <code>global_result_p</code> is a private variable, it refers to the variable <code>global_result</code> which was declared in <code>main</code> before the <code>parallel</code> directive, and the value of <code>global_result</code> is used to store the result that's printed out after the <code>parallel</code> block. So in the code

```
*global_result_p += my_result;
```

it's essential that *global_result_p have shared scope. If it were private to each thread, there would be no need for the critical directive. Furthermore, if it were private, we would have a hard time determining the value of global_result in main after completion of the parallel block.

To summarize, then, variables that have been declared before a parallel directive have shared scope among the threads in the team, while variables declared in the block (e.g., local variables in functions) have private scope. Furthermore, the value of a shared variable at the beginning of the parallel block is the same as the value before the block, and, after completion of the parallel block, the value of the variable is the value at the end of the block.

We'll shortly see that the *default* scope of a variable can change with other directives, and that OpenMP provides clauses to modify the default scope.

5.4 THE REDUCTION CLAUSE

If we developed a serial implementation of the trapezoidal rule, we'd probably use a slightly different function prototype. Rather than

```
void Trap(double a, double b, int n, double* global_result_p);
```

```
we would probably define
```

```
double Trap(double a, double b, int n);
and our function call would be
global_result = Trap(a, b, n);
```

This is somewhat easier to understand and probably more attractive to all but the most fanatical believers in pointers.

We resorted to the pointer version because we needed to add each thread's local calculation to get global_result. However, we might prefer the following function prototype:

```
double Local_trap(double a, double b, int n);
```

With this prototype, the body of Local_trap would be the same as the Trap function in Program 5.2, except that there would be no critical section. Rather, each thread would return its part of the calculation, the final value of its my_result variable. If we made this change, we might try modifying our parallel block so that it looks like this:

```
global_result = 0.0;
# pragma omp parallel num_threads(thread_count)
{
# pragma omp critical
    global_result += Local_trap(double a, double b, int n);
}
```

Can you see a problem with this code? It should give the correct result. However, since we've specified that the critical section is

```
global_result += Local_trap(double a, double b, int n);
```

the call to Local_trap can only be executed by one thread at a time, and, effectively, we're forcing the threads to execute the trapezoidal rule sequentially. If we check the run-time of this version, it may actually be *slower* with multiple threads than one thread (see Exercise 5.3).

We can avoid this problem by declaring a private variable inside the parallel block and moving the critical section after the function call:

```
global_result = 0.0;
# pragma omp parallel num_threads(thread_count)
{
    double my_result = 0.0; /* private */
    my_result += Local_trap(double a, double b, int n);
# pragma omp critical
    global_result += my_result;
}
```

Now the call to Local_trap is outside the critical section, and the threads can execute their calls simultaneously. Furthermore, since my_result is declared in the

parallel block, it's private, and before the critical section each thread will store its part of the calculation in its my_result variable.

OpenMP provides a cleaner alternative that also avoids serializing execution of Local_trap: we can specify that global_result is a *reduction* variable. A **reduction operator** is a binary operation (such as addition or multiplication) and a **reduction** is a computation that repeatedly applies the same reduction operator to a sequence of operands in order to get a single result. Furthermore, all of the intermediate results of the operation should be stored in the same variable: the **reduction variable**. For example, if A is an array of n ints, the computation

```
int sum = 0;
for (i = 0; i < n; i++)
   sum += A[i];</pre>
```

is a reduction in which the reduction operator is addition.

In OpenMP it may be possible to specify that the result of a reduction is a reduction variable. To do this, a reduction clause can be added to a parallel directive. In our example, we can modify the code as follows:

```
global_result = 0.0;
# pragma omp parallel num_threads(thread_count) \
    reduction(+: global_result)
global_result += Local_trap(double a, double b, int n);
```

First note that the parallel directive is two lines long. Recall that C preprocessor directives are, by default, only one line long, so we need to "escape" the newline character by putting a backslash (\) immediately before it.

The code specifies that <code>global_result</code> is a reduction variable and the plus sign ("+") indicates that the reduction operator is addition. Effectively, OpenMP creates a private variable for each thread, and the run-time system stores each thread's result in this private variable. OpenMP also creates a critical section and the values stored in the private variables are added in this critical section. Thus, the calls to <code>Local_trap</code> can take place in parallel.

The syntax of the reduction clause is

```
reduction(<operator>: <variable list>)
```

In C, operator can be any one of the operators +, *, -, &, |, $^{\hat{}}$, &&, |, although the use of subtraction is a bit problematic, since subtraction isn't associative or commutative. For example, the serial code

```
result = 0;
for (i = 1; i <= 4; i++)
result -= i;
```

stores the value -10 in result. If, however, we split the iterations among two threads, with thread 0 subtracting 1 and 2 and thread 1 subtracting 3 and 4, then thread 0 will compute -3 and thread 1 will compute -7 and, of course, -3 - (-7) = 4.

In principle, the compiler should determine that the threads' individual results should actually be added (-3 + (-7) = -10), and, in practice, this seems to be the case. However, the OpenMP Standard [42] doesn't seem to guarantee this.

It should also be noted that if a reduction variable is a float or a double, the results may differ slightly when different numbers of threads are used. This is due to the fact that floating point arithmetic isn't associative. For example, if a, b, and c are **floats**, then (a+b)+c may not be exactly equal to a+(b+c). See Exercise 5.5.

When a variable is included in a reduction clause, the variable itself is shared. However, a private variable is created for each thread in the team. In the parallel block each time a thread executes a statement involving the variable, it uses the private variable. When the parallel block ends, the values in the private variables are combined into the shared variable. Thus, our latest version of the code

```
global_result = 0.0;
# pragma omp parallel num_threads(thread_count) \
    reduction(+: global_result)
global_result += Local_trap(double a, double b, int n);
```

effectively executes code that is identical to our previous version:

```
global_result = 0.0;
# pragma omp parallel num_threads(thread_count)
{
    double my_result = 0.0; /* private */
    my_result += Local_trap(double a, double b, int n);
# pragma omp critical
    global_result += my_result;
}
```

One final point to note is that the threads' private variables are initialized to 0. This is analogous to our initializing my_result to zero. In general, the private variables created for a reduction clause are initialized to the *identity value* for the operator. For example, if the operator is multiplication, the private variables would be initialized to 1.

5.5 THE parallel for DIRECTIVE

As an alternative to our explicit parallelization of the trapezoidal rule, OpenMP provides the parallel for directive. Using it, we can parallelize the serial trapezoidal rule

```
h = (b-a)/n;
approx = (f(a) + f(b))/2.0;
for (i = 1; i <= n-1; i++)
   approx += f(a + i*h);
approx = h*approx;
```

by simply placing a directive immediately before the for loop:

```
h = (b-a)/n;
approx = (f(a) + f(b))/2.0;
# pragma omp parallel for num_threads(thread_count) \
    reduction(+: approx)
for (i = 1; i <= n-1; i++)
    approx += f(a + i*h);
approx = h*approx;</pre>
```

Like the parallel directive, the parallel for directive forks a team of threads to execute the following structured block. However, the structured block following the parallel for directive must be a for loop. Furthermore, with the parallel for directive the system parallelizes the for loop by dividing the iterations of the loop among the threads. The parallel for directive is therefore very different from the parallel directive, because in a block that is preceded by a parallel directive, in general, the work must be divided among the threads by the threads themselves.

In a for loop that has been parallelized with a parallel for directive, the default partitioning, that is, of the iterations among the threads is up to the system. However, most systems use roughly a block partitioning, that is, if there are m iterations, then roughly the first $m/\text{thread_count}$ are assigned to thread 0, the next $m/\text{thread_count}$ are assigned to thread 1, and so on.

Note that it was essential that we made approx a reduction variable. If we hadn't, it would have been an ordinary shared variable, and the body of the loop

```
approx += f(a + i*h);
```

would be an unprotected critical section.

However, speaking of scope, the default scope for all variables in a parallel directive is shared, but in our parallel for if the loop variable i were shared, the variable update, i++, would also be an unprotected critical section. Hence, in a loop that is parallelized with a parallel for directive, the default scope of the loop variable is *private*; in our code, each thread in the team has its own copy of i.

5.5.1 Caveats

This is truly wonderful: It may be possible to parallelize a serial program that consists of one large for loop by just adding a single parallel for directive. It may be possible to incrementally parallelize a serial program that has many for loops by successively placing parallel for directives before each loop.

However, things may not be quite as rosy as they seem. There are several caveats associated with the use of the parallel for directive. First, OpenMP will only parallelize for loops. It won't parallelize while loops or do-while loops. This may not seem to be too much of a limitation, since any code that uses a while loop or a do-while loop can be converted to equivalent code that uses a for loop instead.

However, OpenMP will only parallelize for loops for which the number of iterations can be determined

- from the for statement itself (that is, the code for (. . . ; . . . ; . . .)), and
- prior to execution of the loop.

For example, the "infinite loop"

```
for (;;) {
    . . .
}
```

cannot be parallelized. Similarly, the loop

```
for (i = 0; i < n; i++) {
   if ( . . . ) break;
     . . .
}</pre>
```

cannot be parallelized, since the number of iterations can't be determined from the for statement alone. This for loop is also not a structured block, since the break adds another point of exit from the loop.

In fact, OpenMP will only parallelize for loops that are in **canonical form**. Loops in canonical form take one of the forms shown in Program 5.3. The variables and expressions in this template are subject to some fairly obvious restrictions:

- The variable index must have integer or pointer type (e.g., it can't be a **float**).
- The expressions start, end, and incr must have a compatible type. For example, if index is a pointer, then incr must have integer type.
- The expressions start, end, and incr must not change during execution of the loop.
- During execution of the loop, the variable index can only be modified by the "increment expression" in the for statement.

Program 5.3: Legal forms for parallelizable for statements

These restrictions allow the run-time system to determine the number of iterations prior to execution of the loop.

The sole exception to the rule that the run-time system must be able to determine the number of iterations prior to execution is that there *can* be a call to exit in the body of the loop.

5.5.2 Data dependences

If a for loop fails to satisfy one of the rules outlined in the preceding section, the compiler will simply reject it. For example, suppose we try to compile a program with the following linear search function:

```
int Linear_search(int key, int A[], int n) {
   int i;
   /* thread_count is global */

# pragma omp parallel for num_threads(thread_count)
for (i = 0; i < n; i++)
   if (A[i] == key) return i;
return -1; /* key not in list */
}</pre>
```

The gcc compiler reports:

```
Line 6: error: invalid exit from OpenMP structured block
```

A more insidious problem occurs in loops in which the computation in one iteration depends on the results of one or more previous iterations. As an example, consider the following code, which computes the first n fibonacci numbers:

```
fibo[0] = fibo[1] = 1;
for (i = 2; i < n; i++)
  fibo[i] = fibo[i-1] + fibo[i-2];</pre>
```

Although we may be suspicious that something isn't quite right, let's try parallellizing the for loop with a parallel for directive:

```
fibo[0] = fibo[1] = 1;

# pragma omp parallel for num_threads(thread_count)
for (i = 2; i < n; i++)
    fibo[i] = fibo[i-1] + fibo[i-2];</pre>
```

The compiler will create an executable without complaint. However, if we try running it with more than one thread, we may find that the results are, at best, unpredictable. For example, on one of our systems if we try using two threads to compute the first 10 Fibonacci numbers, we sometimes get

```
1 1 2 3 5 8 13 21 34 55,
```

which is correct. However, we also occasionally get

```
1 1 2 3 5 8 0 0 0 0.
```

What happened? It appears that the run-time system assigned the computation of fibo[2], fibo[3], fibo[4], and fibo[5] to one thread, while fibo[6], fibo[7], fibo[8], and fibo[9] were assigned to the other. (Remember the loop starts with i = 2.) In some runs of the program, everything is fine because the thread that was assigned fibo[2], fibo[3], fibo[4], and fibo[5] finishes its computations before the other thread starts. However, in other runs, the first thread has evidently not computed fibo[4] and fibo[5] when the second computes fibo[6]. It appears that the system has initialized the entries in fibo to 0, and the second thread is using the values fibo[4] = 0 and fibo[5] = 0 to compute fibo[6]. It then goes on to use fibo[5] = 0 and fibo[6] = 0 to compute fibo[7], and so on.

We see two important points here:

- 1. OpenMP compilers don't check for dependences among iterations in a loop that's being parallelized with a parallel for directive. It's up to us, the programmers, to identify these dependences.
- **2.** A loop in which the results of one or more iterations depend on other iterations *cannot*, in general, be correctly parallelized by OpenMP.

The dependence of the computation of fibo[6] on the computation of fibo[5] is called a **data dependence**. Since the value of fibo[5] is calculated in one iteration, and the result is used in a subsequent iteration, the dependence is sometimes called a **loop-carried dependence**.

5.5.3 Finding loop-carried dependences

Perhaps the first thing to observe is that when we're attempting to use a parallel for directive, we only need to worry about loop-carried dependences. We don't need to worry about more general data dependences. For example, in the loop

```
for (i = 0; i < n; i++) {
    x[i] = a + i*h;
    y[i] = exp(x[i]);
}</pre>
```

there is a data dependence between Lines 2 and 3. However, there is no problem with the parallelization

```
1  # pragma omp parallel for num_threads(thread_count)
2     for (i = 0; i < n; i++) {
3          x[i] = a + i*h;
4          y[i] = exp(x[i]);
5     }</pre>
```

since the computation of x[i] and its subsequent use will always be assigned to the same thread.

Also observe that at least one of the statements must write or update the variable in order for the statements to represent a dependence, so in order to detect a loop-carried dependence, we should only concern ourselves with variables that are updated

by the loop body. That is, we should look for variables that are read or written in one iteration, and written in another. Let's look at a couple of examples.

5.5.4 Estimating π

One way to get a numerical approximation to π is to use many terms in the formula³

$$\pi = 4\left[1 - \frac{1}{3} + \frac{1}{5} - \frac{1}{7} + \dots\right] = 4\sum_{k=0}^{\infty} \frac{(-1)^k}{2k+1}.$$

We can implement this formula in serial code with

(Why is it important that factor is a double instead of an int or a long?)

How can we parallelize this with OpenMP? We might at first be inclined to do something like this:

```
double factor = 1.0;
double sum = 0.0;

pragma omp parallel for num_threads(thread_count) \
reduction(+:sum)

for (k = 0; k < n; k++) {
    sum += factor/(2*k+1);
    factor = -factor;

pi_approx = 4.0*sum;</pre>
```

However, it's pretty clear that the update to factor in Line 7 in iteration k and the subsequent increment of sum in Line 6 in iteration k+1 is an instance of a loop-carried dependence. If iteration k is assigned to one thread and iteration k+1 is assigned to another thread, there's no guarantee that the value of factor in Line 6 will be correct. In this case we can fix the problem by examining the series

$$\sum_{k=0}^{\infty} \frac{(-1)^k}{2k+1}.$$

³This is by no means the best method for approximating π , since it requires a *lot* of terms to get a reasonably accurate result. However, we're more interested in the formula itself than the actual estimate.

We see that in iteration k the value of factor should be $(-1)^k$, which is +1 if k is even and -1 if k is odd, so if we replace the code

```
sum += factor/(2*k+1);
1
             factor = -factor;
by
             if (k \% 2 == 0)
1
2
                 factor = 1.0;
3
                 factor = -1.0;
4
             sum += factor/(2*k+1);
5
or, if you prefer the ?: operator,
              factor = (k \% 2 == 0) ? 1.0 : -1.0;
1
             sum += factor/(2*k+1);
```

we will eliminate the loop dependency.

However, things still aren't quite right. If we run the program on one of our systems with just two threads and n = 1000, the result is consistently wrong. For example,

```
With n = 1000 terms and 2 threads,
Our estimate of pi = 2.97063289263385
With n = 1000 terms and 2 threads,
Our estimate of pi = 3.22392164798593
```

On the other hand, if we run the program with only one thread, we always get

```
1 With n = 1000 terms and 1 threads,
2 Our estimate of pi = 3.14059265383979
```

What's wrong here?

Recall that in a block that has been parallelized by a parallel for directive, by default any variable declared before the loop—with the sole exception of the loop variable—is shared among the threads. So factor is shared and, for example, thread 0 might assign it the value 1, but before it can use this value in the update to sum, thread 1 could assign it the value -1. Therefore, in addition to eliminating the loop-carried dependence in the calculation of factor, we need to insure that each thread has its own copy of factor. That is, in order to make our code correct, we need to also insure that factor has private scope. We can do this by adding a private clause to the parallel for directive.

```
double sum = 0.0;
pragma omp parallel for num_threads(thread_count) \
reduction(+:sum) private(factor)
for (k = 0; k < n; k++) {
   if (k % 2 == 0)</pre>
```

The private clause specifies that for each variable listed inside the parentheses, a private copy is to be created for each thread. Thus, in our example, each of the thread_count threads will have its own copy of the variable factor, and hence the updates of one thread to factor won't affect the value of factor in another thread.

It's important to remember that the value of a variable with private scope is unspecified at the beginning of a parallel block or a parallel for block. Its value is also unspecified after completion of a parallel or parallel for block. So, for example, the output of the first printf statement in the following code is unspecified, since it prints the private variable x before it's explicitly initialized. Similarly, the output of the final printf is unspecified, since it prints x after the completion of the parallel block.

```
1
       int x = 5;
2
      pragma omp parallel num_threads(thread_count) \
3
         private(x)
4
5
         int my_rank = omp_get_thread_num();
         printf("Thread %d > before initialization, x = %d n",
6
7
                my_rank, x);
8
         x = 2*my_rank + 2;
9
         printf("Thread %d > after initialization, x = %d n",
10
                my_rank, x);
11
       printf("After parallel block, x = %d\n", x);
12
```

5.5.5 More on scope

Our problem with the variable factor is a common one. We usually need to think about the scope of each variable in a parallel block or a parallel for block. Therefore, rather than letting OpenMP decide on the scope of each variable, it's a very good practice for us as programmers to specify the scope of each variable in a block. In fact, OpenMP provides a clause that will explicitly require us to do this: the default clause. If we add the clause

```
default(none)
```

to our parallel or parallel for directive, then the compiler will require that we specify the scope of each variable we use in the block and that has been declared outside the block. (Variables that are declared within the block are always private, since they are allocated on the thread's stack.)

For example, using a default(none) clause, our calculation of π could be written as follows:

```
double sum = 0.0;
pragma omp parallel for num_threads(thread_count) \
    default(none) reduction(+:sum) private(k, factor) \
    shared(n)
for (k = 0; k < n; k++) {
    if (k % 2 == 0)
        factor = 1.0;
    else
        factor = -1.0;
    sum += factor/(2*k+1);
}</pre>
```

In this example, we use four variables in the for loop. With the default clause, we need to specify the scope of each. As we've already noted, sum is a reduction variable (which has properties of both private and shared scope). We've also already noted that factor and the loop variable k should have private scope. Variables that are never updated in the parallel or parallel for block, such as n in this example, can be safely shared. Recall that unlike private variables, shared variables have the same value in the parallel or parallel for block that they had before the block, and their value after the block is the same as their last value in the block. Thus, if n were initialized before the block to 1000, it would retain this value in the parallel for statement, and since the value isn't changed in the for loop, it would retain this value after the loop has completed.

5.6 MORE ABOUT LOOPS IN OPENMP: SORTING

5.6.1 Bubble sort

Recollect that the serial *bubble sort* algorithm for sorting a list of integers can be implemented as follows:

```
for (list_length = n; list_length >= 2; list_length—)
  for (i = 0; i < list_length-1; i++)
    if (a[i] > a[i+1]) {
        tmp = a[i];
        a[i] = a[i+1];
        a[i+1] = tmp;
    }
```

Here, a stores n ints and the algorithm sorts them in increasing order. The outer loop first finds the largest element in the list and stores it in a[n-1]; it then finds the next-to-the-largest element and stores it in a[n-2], and so on. So, effectively, the first pass is working with the full n-element list. The second is working with all of the elements, except the largest; it's working with an n-1-element list, and so on.

The inner loop compares consecutive pairs of elements in the current list. When a pair is out of order (a[i] > a[i+1]) it swaps them. This process of swapping will move the largest element to the last slot in the "current" list, that is, the list consisting of the elements

```
a[0], a[1], . . . , a[list_length-1]
```

It's pretty clear that there's a loop-carried dependence in the outer loop; in any iteration of the outer loop the contents of the current list depends on the previous iterations of the outer loop. For example, if at the start of the algorithm a=3, 4, 1, 2, then the second iteration of the outer loop should work with the list 3, 1, 2, since the 4 should be moved to the last position by the first iteration. But if the first two iterations are executing simultaneously, it's possible that the effective list for the second iteration will contain 4.

The loop-carried dependence in the inner loop is also fairly easy to see. In iteration i the elements that are compared depend on the outcome of iteration i-1. If in iteration i-1, a[i-1] and a[i] are not swapped, then iteration i should compare a[i] and a[i+1]. If, on the other hand, iteration i-1 swaps a[i-1] and a[i], then iteration i should be comparing the original a[i-1] (which is now a[i]) and a[i+1]. For example, suppose the current list is $\{3,1,2\}$. Then when i=1, we should compare 3 and 2, but if the i=0 and the i=1 iterations are happening simultaneously, it's entirely possible that the i=1 iteration will compare 1 and 2.

It's also not at all clear how we might remove either loop-carried dependence without completely rewriting the algorithm. It's important to keep in mind that even though we can always find loop-carried dependences, it may be difficult or impossible to remove them. The parallel for directive is not a universal solution to the problem of parallelizing for loops.

5.6.2 Odd-even transposition sort

Odd-even transposition sort is a sorting algorithm that's similar to bubble sort, but that has considerably more opportunities for parallelism. Recall from Section 3.7.1 that serial odd-even transposition sort can be implemented as follows:

```
for (phase = 0; phase < n; phase++)
  if (phase % 2 == 0)
     for (i = 1; i < n; i += 2)
        if (a[i-1] > a[i]) Swap(&a[i-1],&a[i]);
  else
     for (i = 1; i < n-1; i += 2)
        if (a[i] > a[i+1]) Swap(&a[i], &a[i+1]);
```

The list a stores n ints, and the algorithm sorts them into increasing order. During an "even phase" (phase % 2 == 0), each odd-subscripted element, a[i], is compared to the element to its "left," a[i-1], and if they're out of order, they're swapped. During an "odd" phase, each odd-subscripted element is compared to the element to its right, and if they're out of order, they're swapped. A theorem guarantees that after n phases, the list will be sorted.

Table 5.1 Serial Odd-Even Transposition Sort						
Subscript in Array						
0		1		2		3
9	\leftrightarrow	7		8	\leftrightarrow	6
7 7		9	\leftrightarrow	6		8 8
7		6		9		8
6	\leftrightarrow	-		-	\leftrightarrow	8
6		7	\leftrightarrow	8		9
6		7		8		9
	9 7 7 7 6 6	Sul 9 ↔ 7 7 7 ↔ 6 6 6	Subscr Subscr 0 1 9 ↔ 7 7 9 7 9 7 6 7 ↔ 6 6 7 6 7	sition Sort Subscript in 0 1 9 \leftrightarrow 7 7 9 7 9 \leftrightarrow 6 7 \leftrightarrow 6 6 7 \leftrightarrow 6 6 7 \leftrightarrow 6	Sition Sort Subscript in Ar 0 1 2 9 \rightarrow 7 8 7 9 6 7 9 \rightarrow 6 7 \rightarrow 6 9 7 \rightarrow 6 9 6 7 \rightarrow 8 6 7 \rightarrow 8	Subscript in Array 0 1 2 9 ↔ 7 8 ↔ 7 9 6 7 9 ↔ 6 7 6 9 7 ↔ 6 9 7 ↔ 6 9 6 7 8 6 7 ↔ 8

As a brief example, suppose a = {9, 7, 8, 6}. Then the phases are shown in Table 5.1. In this case, the final phase wasn't necessary, but the algorithm doesn't bother checking whether the list is already sorted before carrying out each phase.

It's not hard to see that the outer loop has a loop-carried dependence. As an example, suppose as before that $a = \{9, 7, 8, 6\}$. Then in phase 0 the inner loop will compare elements in the pairs (9,7) and (8,6), and both pairs are swapped. So for phase 1 the list should be $\{7, 9, 6, 8\}$, and during phase 1 the elements in the pair (9,6) should be compared and swapped. However, if phase 0 and phase 1 are executed simultaneously, the pair that's checked in phase 1 might be (7,8), which is in order. Furthermore, it's not clear how one might eliminate this loop-carried dependence, so it would appear that parallelizing the outer for loop isn't an option.

The *inner* for loops, however, don't appear to have any loop-carried dependences. For example, in an even phase loop, variable i will be odd, so for two distinct values of i, say i = j and i = k, the pairs $\{j - 1, j\}$ and $\{k - 1, k\}$ will be disjoint. The comparison and possible swaps of the pairs (a[j-1], a[j]) and (a[k-1], a[k]) can therefore proceed simultaneously.

Thus, we could try to parallelize odd-even transposition sort using the code shown in Program 5.4, but there are a couple of potential problems. First, although any iteration of, say, one even phase doesn't depend on any other iteration of that phase, we've already noted that this is not the case for iterations in phase p and phase p+1. We need to be sure that all the threads have finished phase p before any thread starts phase p+1. However, like the parallel directive, the parallel for directive has an implicit barrier at the end of the loop, so none of the threads will proceed to the next phase, phase p+1, until all of the threads have completed the current phase, phase p.

A second potential problem is the overhead associated with forking and joining the threads. The OpenMP implementation *may* fork and join thread_count threads on *each* pass through the body of the outer loop. The first row of Table 5.2 shows

```
1
       for (phase = 0; phase \langle n; phase++ \rangle {
          if (phase % 2 == 0)
2
   #
              pragma omp parallel for num_threads(thread_count) \
3
4
                 default(none) shared(a, n) private(i, tmp)
5
              for (i = 1; i < n; i += 2) {
 6
                 if (a[i-1] > a[i]) {
7
                    tmp = a[i-1];
8
                    a[i-1] = a[i];
9
                    a[i] = tmp;
10
11
12
          else
              pragma omp parallel for num_threads(thread_count) \
13
   #
14
                 default(none) shared(a, n) private(i, tmp)
15
              for (i = 1; i < n-1; i += 2) {
                 if (a[i] > a[i+1]) {
16
17
                    tmp = a[i+1];
18
                    a[i+1] = a[i];
19
                    a[i] = tmp;
20
21
              }
22
       }
```

Program 5.4: First OpenMP implementation of odd-even sort

Table 5.2 Odd-Even Sort with Two parallel for Directives and Two for Directives (times are in seconds)					
thread_count	1	2	3	4	
Two parallel for directives Two for directives	0.770 0.732			0.305 0.239	

run-times for 1, 2, 3, and 4 threads on one of our systems when the input list contained 20,000 elements.

These aren't terrible times, but let's see if we can do better. Each time we execute one of the inner loops, we use the same number of threads, so it would seem to be superior to fork the threads once and reuse the same team of threads for each execution of the inner loops. Not surprisingly, OpenMP provides directives that allow us to do just this. We can fork our team of thread_count threads *before* the outer loop with a parallel directive. Then, rather than forking a new team of threads with each execution of one of the inner loops, we use a for directive, which tells OpenMP to parallelize the for loop with the existing team of threads. This modification to the original OpenMP implementation is shown in Program 5.5

The for directive, unlike the parallel for directive, doesn't fork any threads. It uses whatever threads have already been forked in the enclosing parallel block.

```
pragma omp parallel num_threads(thread_count) \
          default(none) shared(a, n) private(i, tmp, phase)
2
3
       for (phase = 0; phase < n; phase++) {</pre>
          if (phase % 2 == 0)
4
5
    #
             pragma omp for
             for (i = 1; i < n; i += 2) {
6
                 if (a[i-1] > a[i]) {
7
8
                    tmp = a[i-1];
9
                    a[i-1] = a[i];
10
                    a[i] = tmp;
11
12
              }
13
          else
    #
14
             pragma omp for
              for (i = 1; i < n-1; i += 2) {
15
                 if (a[i] > a[i+1]) {
16
17
                    tmp = a[i+1];
                    a[i+1] = a[i];
18
19
                    a[i] = tmp;
20
21
              }
22
       }
```

Program 5.5: Second OpenMP implementation of odd-even sort

There *is* an implicit barrier at the end of the loop. The results of the code—the final list—will therefore be the same as the results obtained from the original parallelized code.

Run-times for this second version of odd-even sort are in the second row of Table 5.2. When we're using two or more threads, the version that uses two for directives is at least 17% faster than the version that uses two parallel for directives, so for this system the slight effort involved in making the change is well worth it.

5.7 SCHEDULING LOOPS

When we first encountered the parallel for directive, we saw that the exact assignment of loop iterations to threads is system dependent. However, most OpenMP implementations use roughly a block partitioning: if there are n iterations in the serial loop, then in the parallel loop the first $n/{\rm thread_count}$ are assigned to thread 0, the next $n/{\rm thread_count}$ are assigned to thread 1, and so on. It's not difficult to think of situations in which this assignment of iterations to threads would be less than optimal. For example, suppose we want to parallelize the loop

```
sum = 0.0;
for (i = 0; i <= n; i++)
   sum += f(i);
```

Also suppose that the time required by the call to f is proportional to the size of the argument i. Then a block partitioning of the iterations will assign much more work

to thread_count -1 than it will assign to thread 0. A better assignment of work to threads might be obtained with a **cyclic** partitioning of the iterations among the threads. In a cyclic partitioning, the iterations are assigned, one at a time, in a "round-robin" fashion to the threads. Suppose $t = \text{thread_count}$. Then a cyclic partitioning will assign the iterations as follows:

```
 \begin{array}{|c|c|c|c|c|c|} \hline \textbf{Thread} & \textbf{Iterations} \\ \hline 0 & 0, \, n/t, \, 2n/t, \, \dots \\ 1 & 1, \, n/t+1, \, 2n/t+1, \, \dots \\ \vdots & \vdots & \vdots \\ t-1 & t-1, \, n/t+t-1, \, 2n/t+t-1, \, \dots \\ \hline \end{array}
```

To get a feel for how drastically this can affect performance, we wrote a program in which we defined

```
double f(int i) {
  int j, start = i*(i+1)/2, finish = start + i;
  double return_val = 0.0;

  for (j = start; j <= finish; j++) {
     return_val += sin(j);
  }
  return return_val;
} /* f */</pre>
```

The call f(i) calls the sine function i times, and, for example, the time to execute f(2i) requires approximately twice as much time as the time to execute f(i).

When we ran the program with n = 10,000 and one thread, the run-time was 3.67 seconds. When we ran the program with two threads and the default assignment—iterations 0–5000 on thread 0 and iterations 5001–10,000 on thread 1—the run-time was 2.76 seconds. This is a speedup of only 1.33. However, when we ran the program with two threads and a cyclic assignment, the run-time was decreased to 1.84 seconds. This is a speedup of 1.99 over the one-thread run and a speedup of 1.5 over the two-thread block partition!

We can see that a good assignment of iterations to threads can have a very significant effect on performance. In OpenMP, assigning iterations to threads is called **scheduling**, and the schedule clause can be used to assign iterations in either a parallel for or a for directive.

5.7.1 The schedule clause

In our example, we already know how to obtain the default schedule: we just add a parallel for directive with a reduction clause:

```
sum = 0.0;
pragma omp parallel for num_threads(thread_count) \
    reduction(+:sum)
```

```
for (i = 0; i <= n; i++)
   sum += f(i);</pre>
```

To get a cyclic schedule, we can add a schedule clause to the parallel for directive:

```
sum = 0.0;
pragma omp parallel for num_threads(thread_count) \
    reduction(+:sum) schedule(static,1)
for (i = 0; i <= n; i++)
    sum += f(i);</pre>
```

In general, the schedule clause has the form

```
schedule(<type> [, <chunksize>])
```

The type can be any one of the following:

- static. The iterations can be assigned to the threads before the loop is executed.
- dynamic or guided. The iterations are assigned to the threads while the loop is
 executing, so after a thread completes its current set of iterations, it can request
 more from the run-time system.
- auto. The compiler and/or the run-time system determine the schedule.
- runtime. The schedule is determined at run-time.

The chunksize is a positive integer. In OpenMP parlance, a **chunk** of iterations is a block of iterations that would be executed consecutively in the serial loop. The number of iterations in the block is the chunksize. Only static, dynamic, and guided schedules can have a chunksize. This determines the details of the schedule, but its exact interpretation depends on the type.

5.7.2 The static schedule type

For a static schedule, the system assigns chunks of chunksize iterations to each thread in a round-robin fashion. As an example, suppose we have 12 iterations, $0,1,\ldots,11$, and three threads. Then if <code>schedule(static,1)</code> is used in the <code>parallel</code> for or for directive, we've already seen that the iterations will be assigned as

Thread 0: 0,3,6,9 Thread 1: 1,4,7,10 Thread 2: 2,5,8,11

If schedule(static,2) is used, then the iterations will be assigned as

Thread 0: 0,1,6,7 Thread 1: 2,3,8,9 Thread 2: 4,5,10,11 If schedule(static,4) is used, the iterations will be assigned as

Thread 0: 0,1,2,3 Thread 1: 4,5,6,7 Thread 2: 8,9,10,11

Thus the clause schedule(static, total_iterations/thread_count) is more or less equivalent to the default schedule used by most implementations of OpenMP.

The chunksize can be omitted. If it is omitted, the chunksize is approximately total_iterations/thread_count.

5.7.3 The dynamic and guided schedule types

In a dynamic schedule, the iterations are also broken up into chunks of chunksize consecutive iterations. Each thread executes a chunk, and when a thread finishes a chunk, it requests another one from the run-time system. This continues until all the iterations are completed. The chunksize can be omitted. When it is omitted, a chunksize of 1 is used.

In a guided schedule, each thread also executes a chunk, and when a thread finishes a chunk, it requests another one. However, in a guided schedule, as chunks are completed, the size of the new chunks decreases. For example, on one of our systems, if we run the trapezoidal rule program with the parallel for directive and a schedule(guided) clause, then when n=10,000 and thread_count = 2, the iterations are assigned as shown in Table 5.3. We see that the size of the chunk is approximately the number of iterations remaining divided by the number of threads. The first chunk has size $9999/2 \approx 5000$, since there are 9999 unassigned iterations. The second chunk has size $4999/2 \approx 2500$, and so on.

In a guided schedule, if no chunksize is specified, the size of the chunks decreases down to 1. If chunksize is specified, it decreases down to chunksize, with the exception that the very last chunk can be smaller than chunksize.

5.7.4 The runtime schedule type

To understand schedule(runtime) we need to digress for a moment and talk about **environment variables**. As the name suggests, environment variables are named values that can be accessed by a running program. That is, they're available in the program's *environment*. Some commonly used environment variables are PATH, HOME, and SHELL. The PATH variable specifies which directories the shell should search when it's looking for an executable. It's usually defined in both Unix and Windows. The HOME variable specifies the location of the user's home directory, and the SHELL variable specifies the location of the executable for the user's shell. These are usually defined in Unix systems. In both Unix-like systems (e.g., Linux and Mac OS X) and Windows, environment variables can be examined and specified on the command line. In Unix-like systems, you can use the shell's command line.

Table 5.3 Assignment of Trapezoidal Rule Iterations 1–9999 using a guided Schedule with Two Threads			
Thread	Chunk	Size of Chunk	Remaining Iterations
0	1-5000	5000	4999
1	5001-7500	2500	2499
1	7501-8750	1250	1249
1	8751–9375	625	624
0	9376–9687	312	312
1	9688–9843	156	156
0	9844-9921	78	78
1	9922-9960	39	39
1	9961–9980	20	19
1	9981–9990	10	9
1	9991–9995	5	4
0	9996–9997	2	2
1	9998–9998	1	1
0	9999-9999	1	0

In Windows systems, you can use the command line in an integrated development environment.

As an example, if we're using the bash shell, we can examine the value of an environment variable by typing

\$ echo \$PATH

and we can use the export command to set the value of an environment variable

\$ export TEST_VAR="hello"

For details about how to examine and set environment variables for your particular system, you should consult with your local expert.

When <code>schedule(runtime)</code> is specified, the system uses the environment variable <code>OMP_SCHEDULE</code> to determine at run-time how to schedule the loop. The <code>OMP_SCHEDULE</code> environment variable can take on any of the values that can be used for a static, dynamic, or guided schedule. For example, suppose we have a <code>parallel</code> for directive in a program and it has been modified by <code>schedule(runtime)</code>. Then if we use the bash shell, we can get a cyclic assignment of iterations to threads by executing the command

\$ export OMP_SCHEDULE="static,1"

Now, when we start executing our program, the system will schedule the iterations of the for loop as if we had the clause schedule(static,1) modifying the parallel for directive.

5.7.5 Which schedule?

If we have a for loop that we're able to parallelize, how do we decide which type of schedule we should use and what the chunksize should be? As you may have guessed, there is some overhead associated with the use of a schedule clause. Furthermore, the overhead is greater for dynamic schedules than static schedules, and the overhead associated with guided schedules is the greatest of the three. Thus, if we're getting satisfactory performance without a schedule clause, we should go no further. However, if we suspect that the performance of the default schedule can be substantially improved, we should probably experiment with some different schedules.

In the example at the beginning of this section, when we switched from the default schedule to schedule(static,1), the speedup of the two-threaded execution of the program increased from 1.33 to 1.99. Since it's *extremely* unlikely that we'll get speedups that are significantly better than 1.99, we can just stop here, at least if we're only going to use two threads with 10,000 iterations. If we're going to be using varying numbers of threads and varying numbers of iterations, we need to do more experimentation, and it's entirely possible that we'll find that the optimal schedule depends on both the number of threads and the number of iterations.

It can also happen that we'll decide that the performance of the default schedule isn't very good, and we'll proceed to search through a large array of schedules and iteration counts only to conclude that our loop doesn't parallelize very well and *no* schedule is going to give us much improved performance. For an example of this, see Programming Assignment 5.4.

There are some situations in which it's a good idea to explore some schedules before others:

- If each iteration of the loop requires roughly the same amount of computation, then it's likely that the default distribution will give the best performance.
- If the cost of the iterations decreases (or increases) linearly as the loop executes, then a static schedule with small chunksizes will probably give the best performance.
- If the cost of each iteration can't be determined in advance, then it may make sense to explore a variety of scheduling options. The schedule(runtime) clause can be used here, and the different options can be explored by running the program with different assignments to the environment variable OMP_SCHEDULE.

5.8 PRODUCERS AND CONSUMERS

Let's take a look at a parallel problem that isn't amenable to parallelization using a parallel for or for directive.

5.8.1 Queues

Recall that a **queue** is a list abstract datatype in which new elements are inserted at the "rear" of the queue and elements are removed from the "front" of the queue.

A queue can thus be viewed as an abstraction of a line of customers waiting to pay for their groceries in a supermarket. The elements of the list are the customers. New customers go to the end or "rear" of the line, and the next customer to check out is the customer standing at the "front" of the line.

When a new entry is added to the rear of a queue, we sometimes say that the entry has been "enqueued," and when an entry is removed from the front of a queue, we sometimes say that the entry has been "dequeued."

Queues occur frequently in computer science. For example, if we have a number of processes, each of which wants to store some data on a hard drive, then a natural way to insure that only one process writes to the disk at a time is to have the processes form a queue, that is, the first process that wants to write gets access to the drive first, the second process gets access to the drive next, and so on.

A queue is also a natural data structure to use in many multithreaded applications. For example, suppose we have several "producer" threads and several "consumer" threads. The producer threads might "produce" requests for data from a server—for example, current stock prices—while the consumer threads might "consume" the request by finding or generating the requested data—the current stock prices. The producer threads could enqueue the requested prices, and the consumer threads could dequeue them. In this example, the process wouldn't be completed until the consumer threads had given the requested data to the producer threads.

5.8.2 Message-passing

Another natural application would be implementing message-passing on a shared-memory system. Each thread could have a shared message queue, and when one thread wanted to "send a message" to another thread, it could enqueue the message in the destination thread's queue. A thread could receive a message by dequeuing the message at the head of its message queue.

Let's implement a relatively simple message-passing program in which each thread generates random integer "messages" and random destinations for the messages. After creating the message, the thread enqueues the message in the appropriate message queue. After sending a message, a thread checks its queue to see if it has received a message. If it has, it dequeues the first message in its queue and prints it out. Each thread alternates between sending and trying to receive messages. We'll let the user specify the number of messages each thread should send. When a thread is done sending messages, it receives messages until all the threads are done, at which point all the threads quit. Pseudocode for each thread might look something like this:

```
for (sent_msgs = 0; sent_msgs < send_max; sent_msgs++) {
    Send_msg();
    Try_receive();
}
while (!Done())
    Try_receive();</pre>
```

5.8.3 Sending messages

Note that accessing a message queue to enqueue a message is probably a critical section. Although we haven't looked into the details of the implementation of the message queue, it seems likely that we'll want to have a variable that keeps track of the rear of the queue. For example, if we use a singly linked list with the tail of the list corresponding to the rear of the queue, then, in order to efficiently enqueue, we would want to store a pointer to the rear. When we enqueue a new message, we'll need to check and update the rear pointer. If two threads try to do this simultaneously, we may lose a message that has been enqueued by one of the threads. (It might help to draw a picture!) The results of the two operations will conflict, and hence enqueueing a message will form a critical section.

Pseudocode for the Send_msg() function might look something like this:

```
mesg = random();
  dest = random() % thread_count;
# pragma omp critical
Enqueue(queue, dest, my_rank, mesg);
```

Note that this allows a thread to send a message to itself.

5.8.4 Receiving messages

The synchronization issues for receiving a message are a little different. Only the owner of the queue (that is, the destination thread) will dequeue from a given message queue. As long as we dequeue one message at a time, if there are at least two messages in the queue, a call to Dequeue can't possibly conflict with any calls to Enqueue, so if we keep track of the size of the queue, we can avoid any synchronization (for example, critical directives), as long as there are at least two messages.

Now you may be thinking, "What about the variable storing the size of the queue?" This would be a problem if we simply store the size of the queue. However, if we store two variables, enqueued and dequeued, then the number of messages in the queue is

```
queue_size = enqueued - dequeued
```

and the only thread that will update dequeued is the owner of the queue. Observe that one thread can update enqueued at the same time that another thread is using it to compute queue_size. To see this, let's suppose thread q is computing queue_size. It will either get the old value of enqueued or the new value. It may therefore compute a queue_size of 0 or 1 when queue_size should actually be 1 or 2, respectively, but in our program this will only cause a modest delay. Thread q will try again later if queue_size is 0 when it should be 1, and it will execute the critical section directive unnecessarily if queue_size is 1 when it should be 2.

Thus, we can implement Try_receive as follows:

```
queue_size = enqueued - dequeued;
if (queue_size == 0) return;
```

```
else if (queue_size == 1)
#    pragma omp critical
    Dequeue(queue, &src, &mesg);
else
    Dequeue(queue, &src, &mesg);
Print_message(src, mesg);
```

5.8.5 Termination detection

We also need to think about implementation of the Done function. First note that the following "obvious" implementation will have problems:

```
queue_size = enqueued - dequeued;
if (queue_size == 0)
   return TRUE;
else
   return FALSE:
```

If thread u executes this code, it's entirely possible that some thread—call it thread v—will send a message to thread u after u has computed queue_size = 0. Of course, after thread u computes queue_size = 0, it will terminate and the message sent by thread v will never be received.

However, in our program, after each thread has completed the for loop, it won't send any new messages. Thus, if we add a counter done_sending, and each thread increments this after completing its for loop, then we *can* implement Done as follows:

```
queue_size = enqueued - dequeued;
if (queue_size == 0 && done_sending == thread_count)
   return TRUE;
else
   return FALSE;
```

5.8.6 Startup

When the program begins execution, a single thread, the master thread, will get command-line arguments and allocate an array of message queues, one for each thread. This array needs to be shared among the threads, since any thread can send to any other thread, and hence any thread can enqueue a message in any of the queues. Given that a message queue will (at a minimum) store

- a list of messages,
- a pointer or index to the rear of the queue,
- a pointer or index to the front of the queue,
- · a count of messages enqueued, and
- a count of messages dequeued,

it makes sense to store the queue in a struct, and in order to reduce the amount of copying when passing arguments, it also makes sense to make the message queue

an array of pointers to structs. Thus, once the array of queues is allocated by the master thread, we can start the threads using a parallel directive, and each thread can allocate storage for its individual queue.

An important point here is that one or more threads may finish allocating their queues before some other threads. If this happens, the threads that finish first could start trying to enqueue messages in a queue that hasn't been allocated and cause the program to crash. We therefore need to make sure that none of the threads starts sending messages until all the queues are allocated. Recall that we've seen that several OpenMP directives provide implicit barriers when they're completed, that is, no thread will proceed past the end of the block until all the threads in the team have completed the block. In this case, though, we'll be in the middle of a parallel block, so we can't rely on an implicit barrier from some other OpenMP construct—we need an *explicit* barrier. Fortunately, OpenMP provides one:

```
# pragma omp barrier
```

When a thread encounters the barrier, it blocks until all the threads in the team have reached the barrier. After all the threads have reached the barrier, all the threads in the team can proceed.

5.8.7 The atomic directive

After completing its sends, each thread increments done_sending before proceeding to its final loop of receives. Clearly, incrementing done_sending is a critical section, and we could protect it with a critical directive. However, OpenMP provides a potentially higher performance directive: the atomic directive:

```
# pragma omp atomic
```

Unlike the critical directive, it can only protect critical sections that consist of a single C assignment statement. Further, the statement must have one of the following forms:

```
x <op>= <expression>;
x++;
++x;
x—;
--x;
```

Here <op> can be one of the binary operators

```
+, *, -, /, \&, ^, |, <<, or >>.
```

It's also important to remember that <expression> must not reference x.

It should be noted that only the load and store of x are guaranteed to be protected. For example, in the code

```
# pragma omp atomic
x += y++;
```

a thread's update to \times will be completed before any other thread can begin updating \times . However, the update to y may be unprotected and the results may be unpredictable.

The idea behind the atomic directive is that many processors provide a special load-modify-store instruction, and a critical section that only does a load-modify-store can be protected much more efficiently by using this special instruction rather than the constructs that are used to protect more general critical sections.

5.8.8 Critical sections and locks

To finish our discussion of the message-passing program, we need to take a more careful look at OpenMP's specification of the critical directive. In our earlier examples, our programs had at most one critical section, and the critical directive forced mutually exclusive access to the section by all the threads. In this program, however, the use of critical sections is more complex. If we simply look at the source code, we'll see three blocks of code preceded by a critical or an atomic directive:

done_sending++;Enqueue(q_p, my_rank, mesg);Dequeue(q_p, &src, &mesg);

However, we don't need to enforce exclusive access across all three of these blocks of code. We don't even need to enforce completely exclusive access within the second and third blocks. For example, it would be fine for, say, thread 0 to enqueue a message in thread 1's queue at the same time that thread 1 is enqueuing a message in thread 2's queue. But for the second and third blocks—the blocks protected by critical directives—this is exactly what OpenMP does. From OpenMP's point of view our program has two distinct critical sections: the critical section protected by the atomic directive, done_sending++, and the "composite" critical section in which we enqueue and dequeue messages.

Since enforcing mutual exclusion among threads serializes execution, this default behavior of OpenMP—treating all critical blocks as part of one composite critical section—can be highly detrimental to our program's performance. OpenMP *does* provide the option of adding a name to a critical directive:

```
# pragma omp critical(name)
```

When we do this, two blocks protected with critical directives with different names *can* be executed simultaneously. However, the names are set during compilation, and we want a different critical section for each thread's queue. Therefore, we need to set the names at run-time, and in our setting, when we want to allow simultaneous access to the same block of code by threads accessing different queues, the named critical directive isn't sufficient.

The alternative is to use **locks**.⁴ A lock consists of a data structure and functions that allow the programmer to explicitly enforce mutual exclusion in a critical section. The use of a lock can be roughly described by the following pseudocode:

```
/* Executed by one thread */
Initialize the lock data structure;
...
/* Executed by multiple threads */
Attempt to lock or set the lock data structure;
Critical section;
Unlock or unset the lock data structure;
...
/* Executed by one thread */
Destroy the lock data structure;
```

The lock data structure is shared among the threads that will execute the critical section. One of the threads (e.g., the master thread) will initialize the lock, and when all the threads are done using the lock, one of the threads should destroy it.

Before a thread enters the critical section, it attempts to *set* or lock the lock data structure by calling the lock function. If no other thread is executing code in the critical section, it *obtains* the lock and proceeds into the critical section past the call to the lock function. When the thread finishes the code in the critical section, it calls an unlock function, which *relinquishes* or *unsets* the lock and allows another thread to obtain the lock.

While a thread owns the lock, no other thread can enter the critical section. If another thread attempts to enter the critical section, it will *block* when it calls the lock function. If multiple threads are blocked in a call to the lock function, then when the thread in the critical section relinquishes the lock, one of the blocked threads returns from the call to the lock, and the others remain blocked.

OpenMP has two types of locks: **simple** locks and **nested** locks. A simple lock can only be set once before it is unset, while a nested lock can be set multiple times by the same thread before it is unset. The type of an OpenMP simple lock is omplock_t, and the simple lock functions that we'll be using are

```
void omp_init_lock(omp_lock_t* lock_p /* out */);
void omp_set_lock(omp_lock_t* lock_p /* in/out */);
void omp_unset_lock(omp_lock_t* lock_p /* in/out */);
void omp_destroy_lock(omp_lock_t* lock_p /* in/out */);
```

The type and the functions are specified in omp.h. The first function initializes the lock so that it's unlocked, that is, no thread owns the lock. The second function attempts to set the lock. If it succeeds, the calling thread proceeds; if it fails, the calling thread blocks until the lock becomes available. The third function unsets the

⁴If you've studied the Pthreads chapter, you've already learned about locks, and you can skip ahead to the syntax for OpenMP locks.

lock so another thread can obtain it. The fourth function makes the lock uninitialized. We'll only use simple locks. For information about nested locks, see [8, 10], or [42].

5.8.9 Using locks in the message-passing program

In our earlier discussion of the limitations of the critical directive, we saw that in the message-passing program, we wanted to insure mutual exclusion in each individual message queue, not in a particular block of source code. Locks allow us to do this. If we include a data member with type <code>omp_lock_t</code> in our queue struct, we can simply call <code>omp_set_lock</code> each time we want to insure exclusive access to a message queue. So the code

```
# pragma omp critical
    /* q-p = msg_queues[dest] */
    Enqueue(q_p, my_rank, mesg);

can be replaced with

/* q_p = msg_queues[dest] */
    omp_set_lock(&q_p->lock);
    Enqueue(q_p, my_rank, mesg);
    omp_unset_lock(&q_p->lock);

Similarly, the code

# pragma omp critical
    /* q_p = msg_queues[my_rank] */
    Dequeue(q_p, &src, &mesg);

can be replaced with

/* q_p = msg_queues[my_rank] */
    omp_set_lock(&q_p->lock);
```

Dequeue(q_p, &src, &mesg);
omp_unset_lock(&q_p->lock);

Now when a thread tries to send or receive a message, it can only be blocked by a thread attempting to access the same message queue, since different message queues have different locks. In our original implementation, only one thread could send at a time, regardless of the destination.

Note that it would also be possible to put the calls to the lock functions in the queue functions Enqueue and Dequeue. However, in order to preserve the performance of Dequeue, we would also need to move the code that determines the size of the queue (enqueued – dequeued) to Dequeue. Without it, the Dequeue function will lock the queue every time it is called by Try_receive. In the interest of preserving the structure of the code we've already written, we'll leave the calls to omp_set_lock and omp_unset_lock in the Send and Try_receive functions.

Since we're now including the lock associated with a queue in the queue struct, we can add initialization of the lock to the function that initializes an empty queue.

Destruction of the lock can be done by the thread that owns the queue before it frees the queue.

5.8.10 critical directives, atomic directives, or locks?

Now that we have three mechanisms for enforcing mutual exclusion in a critical section, it's natural to wonder when one method is preferable to another. In general, the atomic directive has the potential to be the fastest method of obtaining mutual exclusion. Thus, if your critical section consists of an assignment statement having the required form, it will probably perform at least as well with the atomic directive as the other methods. However, the OpenMP specification [42] allows the atomic directive to enforce mutual exclusion across *all* atomic directives in the program—this is the way the unnamed critical directive behaves. If this might be a problem—for example, you have multiple different critical sections protected by atomic directives—you should use named critical directives or locks. For example, suppose we have a program in which it's possible that one thread will execute the code on the left while another executes the code on the right.

Even if x and y are unrelated memory locations, it's possible that if one thread is executing x++, then no thread can simultaneously execute y++. It's important to note that the standard doesn't require this behavior. If two statements are protected by atomic directives and the two statements modify different variables, then there are implementations that treat the two statements as different critical sections. See Exercise 5.10. On the other hand, different statements that modify the same variable *will* be treated as if they belong to the same critical section, regardless of the implementation.

We've already seen some limitations to the use of critical directives. However, both named and unnamed critical directives are very easy to use. Furthermore, in the implementations of OpenMP that we've used there doesn't seem to be a very large difference between the performance of critical sections protected by a critical directive, and critical sections protected by locks, so if you can't use an atomic directive, but you can use a critical directive, you probably should. Thus, the use of locks should probably be reserved for situations in which mutual exclusion is needed for a data structure rather than a block of code.

5.8.11 Some caveats

You should exercise caution when you're using the mutual exclusion techniques we've discussed. They can definitely cause serious programming problems. Here are a few things to be aware of:

1. You shouldn't mix the different types of mutual exclusion for a single critical section. For example, suppose a program contains the following two segments.

```
# pragma omp atomic # pragma omp critical x += f(y); x = g(x);
```

The update to x on the right doesn't have the form required by the atomic directive, so the programmer used a critical directive. However, the critical directive won't exclude the action executed by the atomic block, and it's possible that the results will be incorrect. The programmer needs to either rewrite the function g so that its use can have the form required by the atomic directive, or she needs to protect both blocks with a critical directive.

2. There is no guarantee of **fairness** in mutual exclusion constructs. This means that it's possible that a thread can be blocked forever in waiting for access to a critical section. For example, in the code

it's possible that, for example, thread 1 can block forever waiting to execute $x = g(my_rank)$, while the other threads repeatedly execute the assignment. Of course, this wouldn't be an issue if the loop terminated. Also note that many implementations give threads access to the critical section in the order in which they reach it, and for these implementations, this won't be an issue.

3. It can be dangerous to "nest" mutual exclusion constructs. As an example, suppose a program contains the following two segments.

```
# pragma omp critical
    y = f(x);
    ...
    double f(double x) {
        pragma omp critical
        z = g(x); /* z is shared */
        ...
}
```

This is guaranteed to **deadlock**. When a thread attempts to enter the second critical section, it will block forever. If thread u is executing code in the first critical block, no thread can execute code in the second block. In particular, thread u can't execute this code. However, if thread u is blocked waiting to enter the second critical block, then it will never leave the first, and it will stay blocked forever.

In this example, we can solve the problem by using named critical sections. That is, we could rewrite the code as

```
# pragma omp critical(one)
    y = f(x);
    ...
    double f(double x) {
        pragma omp critical(two)
        z = g(x); /* z is global */
        ...
}
```

However, it's not difficult to come up with examples when naming won't help. For example, if a program has two named critical sections—say one and two—and threads can attempt to enter the critical sections in different orders, then deadlock can occur. For example, suppose thread u enters one at the same time that thread v enters two and u then attempts to enter two while v attempts to enter one:

Time	Thread u	Thread v		
0	Enter crit. sect. one	Enter crit. sect. two		
1	Attempt to enter two	Attempt to enter one		
2	Block	Block		

Then both u and v will block forever waiting to enter the critical sections. So it's not enough to just use different names for the critical sections—the programmer must insure that different critical sections are always entered in the same order.

5.9 CACHES, CACHE COHERENCE, AND FALSE SHARING⁵

Recall that for a number of years now, processors have been able to execute operations much faster than they can access data in main memory, so if a processor must read data from main memory for each operation, it will spend most of its time simply waiting for the data from memory to arrive. Also recall that in order to address this problem, chip designers have added blocks of relatively fast memory to processors. This faster memory is called **cache memory**.

The design of cache memory takes into consideration the principles of *temporal* and spatial locality: if a processor accesses main memory location x at time t, then it is likely that at times close to t, it will access main memory locations close to x. Thus, if a processor needs to access main memory location x, rather than transferring only the contents of x to/from main memory, a block of memory containing x is transferred from/to the processor's cache. Such a block of memory is called a **cache line** or **cache block**.

We've already seen in Section 2.3.4 that the use of cache memory can have a huge impact on shared memory. Let's recall why. First, consider the following situation. Suppose \times is a shared variable with the value 5, and both thread 0 and thread 1 read \times from memory into their (separate) caches, because both want to execute the statement

 $my_y = x;$

⁵This material is also covered in Chapter 4. So if you've already read that chapter, you may want to just skim this section.

Here, my_y is a private variable defined by both threads. Now suppose thread 0 executes the statement

x++;

Finally, suppose that thread 1 now executes

$$my_z = x$$
;

where my_z is another private variable.

What's the value in my_z ? Is it 5? Or is it 6? The problem is that there are (at least) three copies of x: the one in main memory, the one in thread 0's cache, and the one in thread 1's cache. When thread 0 executed x++, what happened to the values in main memory and thread 1's cache? This is the **cache coherence** problem, which we discussed in Chapter 2. We saw there that most systems insist that the caches be made aware that changes have been made to data they are caching. The line in the cache of thread 1 would have been marked *invalid* when thread 0 executed x++, and before assigning $my_z = x$, the core running thread 1 would see that it's value of x was out of date. Thus, the core running thread 0 would have to update the copy of x in main memory (either now or earlier), and the core running thread 1 would get the line with the updated value of x from main memory. For further details, see Chapter 2.

The use of cache coherence can have a dramatic effect on the performance of shared-memory systems. To illustrate this, let's take a look at matrix-vector multiplication. Recall that if $A = (a_{ij})$ is an $m \times n$ matrix and \mathbf{x} is a vector with n components, then their product $\mathbf{y} = A\mathbf{x}$ is a vector with m components, and its ith component y_i is found by forming the dot product of the ith row of A with \mathbf{x} :

$$y_i = a_{i0}x_0 + a_{i1}x_1 + \dots + a_{i,n-1}x_{n-1}.$$

See Figure 5.5.

So if we store A as a two-dimensional array and \mathbf{x} and \mathbf{y} as one-dimensional arrays, we can implement serial matrix-vector multiplication with the following code:

a_{00}	a_{01}	• • • •	$a_{0,n-1}$		
a_{10}	a_{11}		$a_{1,n-1}$		
:	:		:		
a_{i0}	a_{i1}		$a_{i,n-1}$		
<i>a</i> _{i0} :	<i>a_{i1}</i> :	•••	$a_{i,n-1}$		

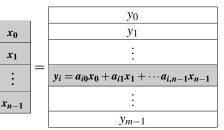


FIGURE 5.5

Matrix-vector multiplication

```
for (j = 0; j < n; j++)
    y[i] += A[i][j]*x[j];
}</pre>
```

There are no loop-carried dependences in the outer loop, since A and X are never updated and iteration i only updates y[i]. Thus, we can parallelize this by dividing the iterations in the outer loop among the threads:

If T_{serial} is the run-time of the serial program and T_{parallel} is the run-time of the parallel program, recall that the *efficiency E* of the parallel program is the speedup S divided by the number of threads, t:

$$E = \frac{S}{t} = \frac{\left(\frac{T_{\text{serial}}}{T_{\text{parallel}}}\right)}{t} = \frac{T_{\text{serial}}}{t \times T_{\text{parallel}}}.$$

Since $S \le t$, $E \le 1$. Table 5.4 shows the run-times and efficiencies of our matrix-vector multiplication with different sets of data and differing numbers of threads.

In each case, the total number of floating point additions and multiplications is 64,000,000. An analysis that only considers arithmetic operations would predict that a single thread running the code would take the same amount of time for all three inputs. However, it's clear that this is *not* the case. The $8,000,000 \times 8$ system requires about 22% more time than the 8000×8000 system, and the $8 \times 8,000,000$ system requires about 26% more time than the 8000×8000 system. Both of these differences are at least partially attributable to cache performance.

Recall that a **write-miss** occurs when a core tries to update a variable that's not in cache, and it has to access main memory. A cache profiler (such as Valgrind [49]) shows that when the program is run with the $8,000,000 \times 8$ input, it has far more

Table 5.4 Run-Times and Efficiencies of Matrix-Vector
Multiplication (times in seconds)

	Matrix Dimension						
	8,000,000 × 8		8000 × 8000		8 × 8,000,000		
Threads	Time	Eff.	Time	Eff.	Time	Eff.	
1	0.322	1.000	0.264	1.000	0.333	1.000	
2	0.219	0.735	0.189	0.698	0.300	0.555	
4	0.141	0.571	0.119	0.555	0.303	0.275	

cache write-misses than either of the other inputs. The bulk of these occur in Line 4. Since the number of elements in the vector y is far greater in this case (8,000,000 vs. 8000 or 8), and each element must be initialized, it's not surprising that this line slows down the execution of the program with the $8,000,000 \times 8$ input.

Also recall that a **read-miss** occurs when a core tries to read a variable that's not in cache, and it has to access main memory. A cache profiler shows that when the program is run with the $8 \times 8,000,000$ input, it has far more cache read-misses than either of the other inputs. These occur in Line 6, and a careful study of this program (see Exercise 5.12) shows that the main source of the differences is due to the reads of x. Once again, this isn't surprising, since for this input, x has 8,000,000 elements, versus only 8000 or 8 for the other inputs.

It should be noted that there may be other factors that affect the relative performance of the single-threaded program with differing inputs. For example, we haven't taken into consideration whether virtual memory (see Section 2.2.4) has affected the performance of the program with the different inputs. How frequently does the CPU need to access the page table in main memory?

Of more interest to us, though, are the differences in efficiency as the number of threads is increased. The two-thread efficiency of the program with the $8\times8,000,000$ input is more than 20% less than the efficiency of the program with the $8,000,000\times8$ and the 8000×8000 inputs. The four-thread efficiency of the program with the $8\times8,000,000$ input is more than 50% less than the program's efficiency with the $8,000,000\times8$ and the 8000×8000 inputs. Why, then, is the multithreaded performance of the program so much worse with the $8\times8,000,000$ input?

In this case, once again, the answer has to do with cache. Let's take a look at the program when we run it with four threads. With the $8,000,000 \times 8$ input, y has 8,000,000 components, so each thread is assigned 2,000,000 components. With the 8000×8000 input, each thread is assigned 2000 components of y, and with the $8 \times 8,000,000$ input, each thread is assigned two components. On the system we used, a cache line is 64 bytes. Since the type of y is double, and a double is 8 bytes, a single cache line will store eight doubles.

Cache coherence is enforced at the "cache-line level." That is, each time any value in a cache line is written, if the line is also stored in another core's cache, the entire *line* will be invalidated—not just the value that was written. The system we're using has two dual-core processors and each processor has its own cache. Suppose for the moment that threads 0 and 1 are assigned to one of the processors and threads 2 and 3 are assigned to the other. Also suppose that for the $8 \times 8,000,000$ problem all of y is stored in a single cache line. Then every write to some element of y will invalidate the line in the other processor's cache. For example, each time thread 0 updates y[0] in the statement

```
y[i] += A[i][j]*x[j];
```

if thread 2 or 3 is executing this code, it will have to reload y. Each thread will update each of its components 8,000,000 times. We see that with this assignment of threads to processors and components of y to cache lines, all the threads will

have to reload y many times. This is going to happen in spite of the fact that only one thread accesses any one component of y—for example, only thread 0 accesses y[0].

Each thread will update its assigned components of y a total of 16,000,000 times. It appears that many, if not most, of these updates are forcing the threads to access main memory. This is called **false sharing**. Suppose two threads with separate caches access different variables that belong to the same cache line. Further suppose at least one of the threads updates its variable. Even though neither thread has written to a shared variable, the cache controller invalidates the entire cache line and forces the other threads to get the values of the variables from main memory. The threads aren't sharing anything (except a cache line), but the behavior of the threads with respect to memory access is the same as if they were sharing a variable. Hence the name *false sharing*.

Why is false sharing not a problem with the other inputs? Let's look at what happens with the 8000×8000 input. Suppose thread 2 is assigned to one of the processors and thread 3 is assigned to another. (We don't actually know which threads are assigned to which processors, but it turns out—see Exercise 5.13—that it doesn't matter.) Thread 2 is responsible for computing

```
y[4000], y[4001], . . . , y[5999],
```

and thread 3 is responsible for computing

```
y[6000], y[6001], . . . , y[7999]
```

If a cache line contains eight consecutive doubles, the only possibility for false sharing is on the interface between their assigned elements. If, for example, a single cache line contains

```
y[5996], y[5997], y[5998], y[5999], y[6000], y[6001], y[6002], y[6003],
```

then it's conceivable that there might be false sharing of this cache line. However, thread 2 will access

```
y[5996], y[5997], y[5998], y[5999]
```

at the end of its iterations of the for i loop, while thread 3 will access

```
y[6000], y[6001], y[6002], y[6003]
```

at the *beginning* of its iterations. So it's very likely that when thread 2 accesses, say, y[5996], thread 3 will be long done with all four of

```
y[6000], y[6001], y[6002], y[6003].
```

Similarly, when thread 3 accesses, say, y[6003], it's very likely that thread 2 won't be anywhere near starting to access

```
y[5996], y[5997], y[5998], y[5999].
```

It's therefore unlikely that false sharing of the elements of y will be a significant problem with the 8000×8000 input. Similar reasoning suggests that false sharing of y is unlikely to be a problem with the $8,000,000 \times 8$ input. Also note that we don't need to worry about false sharing of A or X, since their values are never updated by the matrix-vector multiplication code.

This brings up the question of how we might avoid false sharing in our matrix-vector multiplication program. One possible solution is to "pad" the y vector with dummy elements in order to insure that any update by one thread won't affect another thread's cache line. Another alternative is to have each thread use its own private storage during the multiplication loop, and then update the shared storage when they're done (see Exercise 5.15).

5.10 THREAD-SAFETY⁶

Let's look at another potential problem that occurs in shared-memory programming: *thread-safety*. A block of code is **thread-safe** if it can be simultaneously executed by multiple threads without causing problems.

As an example, suppose we want to use multiple threads to "tokenize" a file. Let's suppose that the file consists of ordinary English text, and that the tokens are just contiguous sequences of characters separated from the rest of the text by white space—spaces, tabs, or newlines. A simple approach to this problem is to divide the input file into lines of text and assign the lines to the threads in a round-robin fashion: the first line goes to thread 0, the second goes to thread $1, \ldots$, the tth goes to thread t, the t+ 1st goes to thread 0, and so on.

We'll read the text into an array of strings, with one line of text per string. Then we can use a parallel for directive with a schedule(static,1) clause to divide the lines among the threads.

One way to tokenize a line is to use the strtok function in string.h. It has the following prototype:

Its usage is a little unusual: the first time it's called, the string argument should be the text to be tokenized, so in our example it should be the line of input. For subsequent calls, the first argument should be NULL. The idea is that in the first call, strtok caches a pointer to string, and for subsequent calls it returns successive tokens taken from the cached copy. The characters that delimit tokens should be passed in separators, so we should pass in the string " \t\n" as the separators argument.

⁶This material is also covered in Chapter 4, so if you've already read that chapter, you may want to just skim this section.

```
void Tokenize(
                               /* in/out */,
         char* lines[]
2
3
          int
                 line_count /* in
                                      */.
4
         int
                 thread_count /* in
                                         */) {
5
       int my_rank, i, j;
6
      char *my_token;
7
8
      pragma omp parallel num_threads(thread_count) \
9
          default(none) private(my_rank, i, j, my_token) \
          shared(lines, line_count)
10
11
12
         my_rank = omp_get_thread_num();
         pragma omp for schedule(static, 1)
13
14
          for (i = 0; i < line_count; i++) {
15
            printf("Thread %d > line %d = %s", my_rank, i,
                   lines[i]);
16
             j = 0;
17
            my_token = strtok(lines[i], " \t\n");
18
            while ( my_token != NULL ) {
               printf("Thread %d > token %d = %s\n", my_rank, j,
19
                      my_token);
                my_token = strtok(NULL, " \t\n");
20
21
                j++;
22
23
          } /* for i */
         /* omp parallel */
24
25
26
   } /* Tokenize */
```

Program 5.6: A first attempt at a multi threaded tokenizer

Given these assumptions, we can write the Tokenize function shown in Program 5.6. The main function has initialized the array lines so that it contains the input text, and line_count is the number of strings stored in lines. Although for our purposes, we only need the lines argument to be an input argument, the strtok function modifies its input. Thus, when Tokenize returns, lines will be modified. When we run the program with a single thread, it correctly tokenizes the input stream. The first time we run it with two threads and the input

Pease porridge hot.

Pease porridge cold.

Pease porridge in the pot

Nine days old.

the output is also correct. However, the second time we run it with this input, we get the following output.

```
Thread 0 > line 0 = Pease porridge hot.
Thread 1 > line 1 = Pease porridge cold.
Thread 0 > token 0 = Pease
```

```
Thread 1 > token 0 = Pease
Thread 0 > token 1 = porridge
Thread 1 > token 1 = cold.
Thread 0 > line 2 = Pease porridge in the pot
Thread 1 > line 3 = Nine days old.
Thread 0 > token 0 = Pease
Thread 1 > token 0 = Nine
Thread 0 > token 1 = days
Thread 1 > token 1 = old.
```

What happened? Recall that strtok caches the input line. It does this by declaring a variable to have static storage class. This causes the value stored in this variable to persist from one call to the next. Unfortunately for us, this cached string is shared, not private. Thus, it appears that thread 1's call to strtok with the second line has apparently overwritten the contents of thread 0's call with the first line. Even worse, thread 0 has found a token ("days") that should be in thread 1's output.

The strtok function is therefore *not* thread-safe: if multiple threads call it simultaneously, the output it produces may not be correct. Regrettably, it's not uncommon for C library functions to fail to be thread-safe. For example, neither the random number generator random in stdlib.h nor the time conversion function localtime in time.h is thread-safe. In some cases, the C standard specifies an alternate, thread-safe, version of a function. In fact, there is a thread-safe version of strtok:

The "_r" is supposed to suggest that the function is *re-entrant*, which is sometimes used as a synonym for thread-safe. The first two arguments have the same purpose as the arguments to strtok. The saveptr_p argument is used by strtok_r for keeping track of where the function is in the input string; it serves the purpose of the cached pointer in strtok. We can correct our original Tokenize function by replacing the calls to strtok with calls to strtok_r. We simply need to declare a char* variable to pass in for the third argument, and replace the calls in Line 17 and Line 20 with the calls

```
my_token = strtok_r(lines[i], " \t\n", &saveptr);
...
my_token = strtok_r(NULL, " \t\n", &saveptr);
respectively.
```

5.10.1 Incorrect programs can produce correct output

Notice that our original version of the tokenizer program shows an especially insidious form of program error: The first time we ran it with two threads, the program produced correct output. It wasn't until a later run that we saw an error. This,

unfortunately, is not a rare occurrence in parallel programs. It's especially common in shared-memory programs. Since, for the most part, the threads are running independently of each other, as we noted back at the beginning of the chapter, the exact sequence of statements executed is nondeterministic. For example, we can't say when thread 1 will first call strtok. If its first call takes place after thread 0 has tokenized its first line, then the tokens identified for the first line should be correct. However, if thread 1 calls strtok before thread 0 has finished tokenizing its first line, it's entirely possible that thread 0 may not identify all the tokens in the first line, so it's especially important in developing shared-memory programs to resist the temptation to assume that since a program produces correct output, it must be correct. We always need to be wary of race conditions.

5.11 SUMMARY

OpenMP is a standard for programming shared-memory systems. It uses both special functions and preprocessor directives called **pragmas**, so unlike Pthreads and MPI, OpenMP requires compiler support. One of the most important features of OpenMP is that it was designed so that developers could *incrementally* parallelize existing serial programs, rather than having to write parallel programs from scratch.

OpenMP programs start multiple **threads** rather than multiple processes. Threads can be much lighter weight than processes; they can share almost all the resources of a process, except each thread must have its own stack and program counter.

To get OpenMP's function prototypes and macros, we include the omp.h header in OpenMP programs. There are several OpenMP directives that start multiple threads; the most general is the parallel directive:

```
# pragma omp parallel
    structured block
```

This directive tells the run-time system to execute the following structured block of code in parallel. It may **fork** or start several threads to execute the structured block. A **structured block** is a block of code with a single entry point and a single exit point, although calls to the C library function exit are allowed within a structured block. The number of threads started is system dependent, but most systems will start one thread for each available core. The collection of threads executing block of code is called a **team**. One of the threads in the team is the thread that was executing the code before the parallel directive. This thread is called the **master**. The additional threads started by the parallel directive are called **slaves**. When all of the threads are finished, the slave threads are terminated or **joined** and the master thread continues executing the code beyond the structured block.

Many OpenMP directives can be modified by **clauses**. We made frequent use of the num_threads clause. When we use an OpenMP directive that starts a team of threads, we can modify it with the num_threads clause so that the directive will start the number of threads we desire.

When OpenMP starts a team of threads, each of the threads is assigned a rank or an id in the range $0,1,\ldots,$ thread_count -1. The OpenMP library function omp_get_thread_num then returns the calling thread's rank. The function omp_get_num_threads returns the number of threads in the current team.

A major problem in the development of shared-memory programs is the possibility of **race conditions**. A race condition occurs when multiple threads attempt to access a shared resource, at least one of the accesses is an update, and the accesses can result in an error. Code that is executed by multiple threads that update a shared resource that can only be updated by one thread at a time is called a **critical section**. Thus, if multiple threads try to update a shared variable, the program has a race condition and the code that updates the variable is a critical section. OpenMP provides several mechanisms for insuring **mutual exclusion** in critical sections. We examined four of them:

- 1. Critical directives insure that only one thread at a time can execute the structured block. If multiple threads try to execute the code in the critical section, all but one of them will block before the critical section. As threads finish the critical section, other threads will be unblocked and enter the code.
- 2. Named critical directives can be used in programs having different critical sections that can be executed concurrently. Multiple threads trying to execute code in critical section(s) with the same name will be handled in the same way as multiple threads trying to execute an unnamed critical section. However, threads entering critical sections with different names can execute concurrently.
- **3.** An atomic directive can only be used when the critical section has the form x = (expression), x++, ++x, x--, or --x. It's designed to exploit special hardware instructions, so it can be much faster than an ordinary critical section.
- **4.** Simple locks are the most general form of mutual exclusion. They use function calls to restrict access to a critical section:

```
omp_set_lock(&lock);
critical section
omp_unset_lock(&lock);
```

When multiple threads call <code>omp_set_lock</code>, only one of them will proceed to the critical section. The others will block until the first thread calls <code>omp_unset_lock</code>. Then one of the blocked threads can proceed.

All of the mutual exclusion mechanisms can cause serious program problems such as deadlock, so they need to be used with great care.

A for directive can be used to partition the iterations in a for loop among the threads. This directive doesn't start a team of threads, it divides the iterations in a for loop among the threads in an existing team. If we want to also start a team of threads, we can use the parallel for directive. There are a number of restrictions on the form of a for loop that can be parallelized; basically, the run-time system must

be able to determine the total number of iterations through the loop body before the loop begins execution. For details, see Program 5.3.

It's not enough, however, to insure that our for loop has one of the canonical forms. It must also not have any **loop-carried dependences**. A loop-carried dependence occurs when a memory location is read or written in one iteration and written in another iteration. OpenMP won't detect loop-carried dependences; it's up to us, the programmers, to detect them and eliminate them. It may, however, be impossible to eliminate them, in which case, the loop isn't a candidate for parallelization.

By default, most systems use a **block partitioning** of the iterations in a parallelized for loop. If there are n iterations, this means that roughly the first $n/\text{thread_count}$ are assigned to thread 0, the next $n/\text{thread_count}$ are assigned to thread 1, and so on. However, there are a variety of **scheduling** options provided by OpenMP. The schedule clause has the form

```
schedule(<type> [,<chunksize>])
```

The type can be static, dynamic, guided, auto, or runtime. In a static schedule, the iterations can be assigned to the threads before the loop starts execution. In dynamic and guided schedules the iterations are assigned on the fly. When a thread finishes a chunk of iterations—a contiguous block of iterations—it requests another chunk. If auto is specified, the schedule is determined by the compiler or run-time system, and if runtime is specified, the schedule is determined at run-time by examining the environment variable OMP_SCHEDULE.

Only static, dynamic, and guided schedules can have a chunksize. In a static schedule, the chunks of chunksize iterations are assigned in round robin fashion to the threads. In a dynamic schedule, each thread is assigned chunksize iterations, and when a thread completes its chunk, it requests another chunk. In a quided schedule, the size of the chunks decreases as the iteration proceeds.

In OpenMP the **scope** of a variable is the collection of threads to which the variable is accessible. Typically, any variable that was defined before the OpenMP directive has **shared** scope within the construct. That is, all the threads have access to it. The principal exception to this is that the loop variable in a for or parallel for construct is **private**, that is, each thread has its own copy of the variable. Variables that are defined within an OpenMP construct have private scope, since they will be allocated from the executing thread's stack.

As a rule of thumb, it's a good idea to explicitly assign the scope of variables. This can be done by modifying a parallel or parallel for directive with the *scoping* clause:

```
default(none)
```

This tells the system that the scope of every variable that's used in the OpenMP construct must be explicitly specified. Most of the time this can be done with private or shared clauses.

The only exceptions we encountered were **reduction variables**. A **reduction operator** is a binary operation (such as addition or multiplication) and a **reduction** is a computation that repeatedly applies the same reduction operator to a sequence of operands in order to get a single result. Furthermore, all of the intermediate results of the operation should be stored in the same variable: the **reduction variable**. For example, if *A* is an array with *n* elements, then the code

```
int sum = 0;
for (i = 0; i < n; i++)
   sum += A[i];
```

is a reduction. The reduction operator is addition and the reduction variable is sum. If we try to parallelize this loop, the reduction variable should have properties of both private and shared variables. Initially we would like each thread to add its array elements into its own private sum, but when the threads are done, we want the private sum's combined into a single, shared sum. OpenMP therefore provides the reduction clause for identifying reduction variables and operators.

A barrier directive will cause the threads in a team to block until all the threads have reached the directive. We've seen that the parallel, parallel for, and for directives have implicit barriers at the end of the structured block.

We recalled that modern microprocessor architectures use caches to reduce memory access times, so typical architectures have special hardware to insure that the caches on the different chips are **coherent**. Since the unit of cache coherence, a **cache line** or **cache block**, is usually larger than a single word of memory, this can have the unfortunate side effect that two threads may be accessing different memory locations, but when the two locations belong to the same cache line, the cache-coherence hardware acts as if the threads were accessing the same memory location—if one of the threads updates its memory location, and then the other thread tries to read its memory location, it will have to retrieve the value from main memory. That is, the hardware is forcing the thread to act as if it were actually sharing the memory location. Hence, this is called **false sharing**, and it can seriously degrade the performance of a shared-memory program.

Some C functions cache data between calls by declaring variables to be static. This can cause errors when multiple threads call the function; since static storage is shared among the threads, one thread can overwrite another thread's data. Such a function is not **thread-safe**, and, unfortunately, there are several such functions in the C library. Sometimes, however, the library has a thread-safe variant of a function that isn't thread-safe.

In one of our programs we saw a particularly insidious problem: when we ran the program with multiple threads and a fixed set of input, it sometimes produced correct output, even though it had an error. Producing correct output during testing doesn't guarantee that the program is in fact correct. It's up to us to identify possible race condtions.

5.12 EXERCISES

- **5.1.** If it's defined, the _OPENMP macro is a decimal int. Write a program that prints its value. What is the significance of the value?
- **5.2.** Download omp_trap_1.c from the book's website, and delete the critical directive. Now compile and run the program with more and more threads and larger and larger values of *n*. How many threads and how many trapezoids does it take before the result is incorrect?
- **5.3.** Modify omp_trap_1.c so that
 - a. it uses the first block of code on page 222, and
 - **b.** the time used by the parallel block is timed using the OpenMP function omp_get_wtime(). The syntax is

```
double omp_get_wtime(void)
```

It returns the number of seconds that have passed since some time in the past. For details on taking timings, see Section 2.6.4. Also recall that OpenMP has a barrier directive:

```
# pragma omp barrier
```

Now find a system with at least two cores and time the program with

- **c.** one thread and a large value of n, and
- **d.** two threads and the same value of n.

What happens? Download omp_trap_2.c from the book's website. How does its performance compare? Explain your answers.

- **5.4.** Recall that OpenMP creates private variables for reduction variables, and these private variables are initialized to the identity value for the reduction operator. For example, if the operator is addition, the private variables are initialized to 0, while if the operator is multiplication, the private variables are initialized to 1. What are the identity values for the operators &&, ||, &, |, ^?
- **5.5.** Suppose that on the amazing Bleeblon computer, variables with type float can store three decimal digits. Also suppose that the Bleeblon's floating point registers can store four decimal digits, and that after any floating point operation, the result is rounded to three decimal digits before being stored. Now suppose a C program declares an array a as follows:

```
float a[] = \{4.0, 3.0, 3.0, 1000.0\};
```

a. What is the output of the following block of code if it's run on the Bleeblon?

```
int i;
float sum = 0.0;
```

```
for (i = 0; i < 4; i++)
    sum += a[i];
printf("sum = %4.1f\n", sum);</pre>
```

b. Now consider the following code:

```
int i;
  float sum = 0.0;
# pragma omp parallel for num_threads(2) \
    reduction(+:sum)
  for (i = 0; i < 4; i++)
    sum += a[i];
  printf("sum = %4.1f\n", sum);</pre>
```

Suppose that the run-time system assigns iterations i = 0, 1 to thread 0 and i = 2, 3 to thread 1. What is the output of this code on the Bleeblon?

5.6. Write an OpenMP program that determines the default scheduling of parallel for loops. Its input should be the number of iterations, and its output should be which iterations of a parallelized for loop are executed by which thread. For example, if there are two threads and four iterations, the output might be:

```
Thread 0: Iterations 0 - 1 Thread 1: Iterations 2 - 3
```

- **5.7.** In our first attempt to parallelize the program for estimating π , our program was incorrect. In fact, we used the result of the program when it was run with one thread as evidence that the program run with two threads was incorrect. Explain why we could "trust" the result of the program when it was run with one thread.
- **5.8.** Consider the loop

```
a[0] = 0;

for (i = 1; i < n; i++)

a[i] = a[i-1] + i;
```

There's clearly a loop-carried dependence, as the value of a[i] can't be computed without the value of a[i-1]. Can you see a way to eliminate this dependence and parallelize the loop?

5.9. Modify the trapezoidal rule program that uses a parallel for directive (omp_trap_3.c) so that the parallel for is modified by a schedule(runtime) clause. Run the program with various assignments to the environment variable OMP_SCHEDULE and determine which iterations are assigned to which thread. This can be done by allocating an array iterations of *n* ints and in the Trap function assigning omp_get_thread_num() to iterations[i] in the *i*th iteration of the for loop. What is the default assignment of iterations on your system? How are guided schedules determined?

5.10. Recall that all structured blocks modified by an unnamed critical directive form a single critical section. What happens if we have a number of atomic directives in which different variables are being modified? Are they all treated as a single critical section?

We can write a small program that tries to determine this. The idea is to have all the threads simultaneously execute something like the following code

```
int i;
  double my_sum = 0.0;
  for (i = 0; i < n; i++)
#     pragma omp atomic
     my_sum += sin(i);</pre>
```

We can do this by modifying the code by a parallel directive:

```
# pragma omp parallel num_threads(thread_count)
{
    int i;
    double my_sum = 0.0;
    for (i = 0; i < n; i++)
        pragma omp atomic
        my_sum += sin(i);
}</pre>
```

Note that since my_sum and i are declared in the parallel block, each thread has its own private copy. Now if we time this code for large n when thread_count = 1 and we also time it when thread_count > 1, then as long as thread_count is less than the number of available cores, the run-time for the single-threaded run should be roughly the same as the time for the multithreaded run if the different threads' executions of my_sum += sin(i) are treated as different critical sections. On the other hand, if the different executions of my_sum += sin(i) are all treated as a single critical section, the multithreaded run should be much slower than the single-threaded run. Write an OpenMP program that implements this test. Does your implementation of OpenMP allow simultaneous execution of updates to different variables when the updates are protected by atomic directives?

- **5.11.** Recall that in C, a function that takes a two-dimensional array argument must specify the number of columns in the argument list, so it is quite common for C programmers to only use one-dimensional arrays, and to write explicit code for converting pairs of subscripts into a single dimension. Modify the OpenMP matrix-vector multiplication so that it uses a one-dimensional array for the matrix.
- **5.12.** Download the source file omp_mat_vect_rand_split.c from the book's website. Find a program that does cache profiling (e.g., Valgrind [49]) and compile the program according to the instructions in the cache profiler documentation. (For example, with Valgrind you will want a symbol table and full optimization. (With gcc use, gcc -g -02 . . .). Now run the program according to the instructions in the cache profiler documentation, using input $k \times (k \cdot 10^6)$,

- $(k \cdot 10^3) \times (k \cdot 10^3)$, and $(k \cdot 10^6) \times k$. Choose k so large that the number of level 2 cache misses is of the order 10^6 for at least one of the input sets of data
- **a.** How many level 1 cache write-misses occur with each of the three inputs?
- **b.** How many level 2 cache write-misses occur with each of the three inputs?
- **c.** Where do most of the write-misses occur? For which input data does the program have the most write-misses? Can you explain why?
- **d.** How many level 1 cache read-misses occur with each of the three inputs?
- **e.** How many level 2 cache read-misses occur with each of the three inputs?
- **f.** Where do most of the read-misses occur? For which input data does the program have the most read-misses? Can you explain why?
- **g.** Run the program with each of the three inputs, but without using the cache profiler. With which input is the program the fastest? With which input is the program the slowest? Can your observations about cache misses help explain the differences? How?
- **5.13.** Recall the matrix-vector multiplication example with the 8000 × 8000 input. Suppose that thread 0 and thread 2 are assigned to different processors. If a cache line contains 64 bytes or 8 doubles, is it possible for false sharing between threads 0 and 2 to occur for any part of the vector y? Why? What about if thread 0 and thread 3 are assigned to different processors; is it possible for false sharing to occur between them for any part of y?
- **5.14.** Recall the matrix-vector multiplication example with an $8 \times 8,000,000$ matrix. Suppose that doubles use 8 bytes of memory and that a cache line is 64 bytes. Also suppose that our system consists of two dual-core processors.
 - **a.** What is the minimum number of cache lines that are needed to store the vector y?
 - **b.** What is the maximum number of cache lines that are needed to store the vector *y*?
 - **c.** If the boundaries of cache lines always coincide with the boundaries of 8-byte doubles, in how many different ways can the components of y be assigned to cache lines?
 - **d.** If we only consider which pairs of threads share a processor, in how many different ways can four threads be assigned to the processors in our computer? Here, we're assuming that cores on the same processor share cache.
 - **e.** Is there an assignment of components to cache lines and threads to processors that will result in no false-sharing in our example? In other words, is it possible that the threads assigned to one processor will have their components of *y* in one cache line, and the threads assigned to the other processor will have their components in a different cache line?
 - **f.** How many assignments of components to cache lines and threads to processors are there?
 - **g.** Of these assignments, how many will result in no false sharing?

- **5.15. a.** Modify the matrix-vector multiplication program so that it pads the vector y when there's a possibility of false sharing. The padding should be done so that if the threads execute in lock-step, there's no possibility that a single cache line containing an element of y will be shared by two or more threads. Suppose, for example, that a cache line stores eight doubles and we run the program with four threads. If we allocate storage for at least 48 doubles in y, then, on each pass through the for i loop, there's no possibility that two threads will simultaneously access the same cache line.
 - **b.** Modify the matrix-vector multiplication program so that each thread uses private storage for its part of y during the for i loop. When a thread is done computing its part of y, it should copy its private storage into the shared variable.
 - **c.** How does the performance of these two alternatives compare to the original program. How do they compare to each other?
- **5.16.** Although strtok_r is thread-safe, it has the rather unfortunate property that it gratuitously modifies the input string. Write a tokenizer that is thread-safe and doesn't modify the input string.

5.13 PROGRAMMING ASSIGNMENTS

- **5.1.** Use OpenMP to implement the parallel histogram program discussed in Chapter 2.
- **5.2.** Suppose we toss darts randomly at a square dartboard, whose bullseye is at the origin, and whose sides are 2 feet in length. Suppose also that there's a circle inscribed in the square dartboard. The radius of the circle is 1 foot, and it's area is π square feet. If the points that are hit by the darts are uniformly distributed (and we always hit the square), then the number of darts that hit inside the circle should approximately satisfy the equation

```
\frac{\text{number in circle}}{\text{total number of tosses}} = \frac{\pi}{4},
```

since the ratio of the area of the circle to the area of the square is $\pi/4$.

We can use this formula to estimate the value of π with a random number generator:

```
number_in_circle = 0;
for (toss = 0; toss < number_of_tosses; toss++) {
    x = random double between -1 and 1;
    y = random double between -1 and 1;
    distance_squared = x*x + y*y;
    if (distance_squared <= 1) number_in_circle++;
}
pi_estimate = 4*number_in_circle/((double) number_of_tosses);</pre>
```

This is called a "Monte Carlo" method, since it uses randomness (the dart tosses).

Write an OpenMP program that uses a Monte Carlo method to estimate π . Read in the total number of tosses before forking any threads. Use a reduction clause to find the total number of darts hitting inside the circle. Print the result after joining all the threads. You may want to use long long ints for the number of hits in the circle and the number of tosses, since both may have to be very large to get a reasonable estimate of π .

5.3. Count sort is a simple serial sorting algorithm that can be implemented as follows:

```
void Count_sort(int a[], int n) {
   int i, j, count;
   int* temp = malloc(n*sizeof(int));

for (i = 0; i < n; i++) {
     count = 0;
     for (j = 0; j < n; j++)
        if (a[j] < a[i])
          count++;
     else if (a[j] == a[i] && j < i)
          count++;
     temp[count] = a[i];
}

memcpy(a, temp, n*sizeof(int));
   free(temp);
} /* Count_sort */</pre>
```

The basic idea is that for each element a[i] in the list a, we count the number of elements in the list that are less than a[i]. Then we insert a[i] into a temporary list using the subscript determined by the count. There's a slight problem with this approach when the list contains equal elements, since they could get assigned to the same slot in the temporary list. The code deals with this by incrementing the count for equal elements on the basis of the subscripts. If both a[i] = a[j] and $j \le i$, then we count a[j] as being "less than" a[i].

After the algorithm has completed, we overwrite the original array with the temporary array using the string library function memcpy.

- **a.** If we try to parallelize the for i loop (the outer loop), which variables should be private and which should be shared?
- **b.** If we parallelize the for i loop using the scoping you specified in the previous part, are there any loop-carried dependences? Explain your answer.
- **c.** Can we parallelize the call to memcpy? Can we modify the code so that this part of the function will be parallelizable?
- **d.** Write a C program that includes a parallel implementation of Count_sort.
- **e.** How does the performance of your parallelization of Count_sort compare to serial Count_sort? How does it compare to the serial qsort library function?

- **5.4.** Recall that when we solve a large linear system, we often use Gaussian elimination followed by *backward substitution*. Gaussian elimination converts an $n \times n$ linear system into an *upper triangular* linear system by using the "row operations."
 - Add a multiple of one row to another row
 - · Swap two rows
 - Multiply one row by a nonzero constant

An upper triangular system has zeroes below the "diagonal" extending from the upper left-hand corner to the lower right-hand corner.

For example, the linear system

$$2x_0 - 3x_1 = 3$$

$$4x_0 - 5x_1 + x_2 = 7$$

$$2x_0 - x_1 - 3x_2 = 5$$

can be reduced to the upper triangular form

$$2x_0 - 3x_1 = 3
x_1 + x_2 = 1 ,
-5x_2 = 0$$

and this system can be easily solved by first finding x_2 using the last equation, then finding x_1 using the second equation, and finally finding x_0 using the first equation.

We can devise a couple of serial algorithms for back substitution. The "row-oriented" version is

```
for (row = n-1; row >= 0; row---) {
    x[row] = b[row];
    for (col = row+1; col < n; col++)
        x[row] -= A[row][col]*x[col];
    x[row] /= A[row][row];
}</pre>
```

Here the "right-hand side" of the system is stored in array b, the two-dimensional array of coefficients is stored in array A, and the solutions are stored in array A. An alternative is the following "column-oriented" algorithm:

a. Determine whether the outer loop of the row-oriented algorithm can be parallelized.

- **b.** Determine whether the inner loop of the row-oriented algorithm can be parallelized.
- **c.** Determine whether the (second) outer loop of the column-oriented algorithm can be parallelized.
- **d.** Determine whether the inner loop of the column-oriented algorithm can be parallelized.
- **e.** Write one OpenMP program for each of the loops that you determined could be parallelized. You may find the single directive useful—when a block of code is being executed in parallel and a sub-block should be executed by only one thread, the sub-block can be modified by a #pragma omp single directive. The threads in the executing team will block at the end of the directive until all of the threads have completed it.
- **f.** Modify your parallel loop with a schedule(runtime) clause and test the program with various schedules. If your upper triangular system has 10,000 variables, which schedule gives the best performance?
- **5.5.** Use OpenMP to implement a program that does Gaussian elimination. (See the preceding problem.) You can assume that the input system doesn't need any row-swapping.
- **5.6.** Use OpenMP to implement a producer-consumer program in which some of the threads are producers and others are consumers. The producers read text from a collection of files, one per producer. They insert lines of text into a single shared queue. The consumers take the lines of text and tokenize them. Tokens are "words" separated by white space. When a consumer finds a token, it writes it to stdout.

Parallel Program Development

In the last three chapters we haven't just learned about parallel APIs, we've also developed a number of small parallel programs, and each of these programs has involved the implementation of a parallel algorithm. In this chapter, we'll look at a couple of larger examples: solving *n*-body problems and solving the traveling salesperson problem. For each problem, we'll start by looking at a serial solution and examining modifications to the serial solution. As we apply Foster's methodology, we'll see that there are some striking similarities between developing shared- and distributed-memory programs. We'll also see that in parallel programming there are problems that we need to solve for which there is no serial analog. We'll see that there are instances in which, as parallel programmers, we'll have to start "from scratch."

6.1 TWO n-BODY SOLVERS

In an *n*-body problem, we need to find the positions and velocities of a collection of interacting particles over a period of time. For example, an astrophysicist might want to know the positions and velocities of a collection of stars, while a chemist might want to know the positions and velocities of a collection of molecules or atoms. An *n*-body solver is a program that finds the solution to an *n*-body problem by simulating the behavior of the particles. The input to the problem is the mass, position, and velocity of each particle at the start of the simulation, and the output is typically the position and velocity of each particle at a sequence of user-specified times, or simply the position and velocity of each particle at the end of a user-specified time period.

Let's first develop a serial *n*-body solver. Then we'll try to parallelize it for both shared- and distributed-memory systems.

6.1.1 The problem

For the sake of explicitness, let's write an n-body solver that simulates the motions of planets or stars. We'll use Newton's second law of motion and his law of universal gravitation to determine the positions and velocities. Thus, if particle q has position $\mathbf{s}_q(t)$ at time t, and particle k has position $\mathbf{s}_k(t)$, then the force on particle q exerted by

particle k is given by

$$\mathbf{f}_{qk}(t) = -\frac{Gm_q m_k}{\left|\mathbf{s}_q(t) - \mathbf{s}_k(t)\right|^3} \left[\mathbf{s}_q(t) - \mathbf{s}_k(t)\right]. \tag{6.1}$$

Here, G is the gravitational constant $(6.673 \times 10^{-11} \text{m}^3/(\text{kg} \cdot \text{s}^2))$, and m_q and m_k are the masses of particles q and k, respectively. Also, the notation $|\mathbf{s}_q(t) - \mathbf{s}_k(t)|$ represents the distance from particle k to particle q. Note that in general the positions, the velocities, the accelerations, and the forces are vectors, so we're using boldface to represent these variables. We'll use an italic font to represent the other, scalar, variables, such as the time t and the gravitational constant G.

We can use Formula 6.1 to find the total force on any particle by adding the forces due to all the particles. If our n particles are numbered 0, 1, 2, ..., n-1, then the total force on particle q is given by

$$\mathbf{F}_{q}(t) = \sum_{\substack{k=0\\k\neq q}}^{n-1} \mathbf{f}_{qk} = -Gm_{q} \sum_{\substack{k=0\\k\neq q}}^{n-1} \frac{m_{k}}{|\mathbf{s}_{q}(t) - \mathbf{s}_{k}(t)|^{3}} \left[\mathbf{s}_{q}(t) - \mathbf{s}_{k}(t) \right]. \tag{6.2}$$

Recall that the acceleration of an object is given by the second derivative of its position and that Newton's second law of motion states that the force on an object is given by its mass multiplied by its acceleration, so if the acceleration of particle q is $\mathbf{a}_q(t)$, then $\mathbf{F}_q(t) = m_q \mathbf{a}_q(t) = m_q \mathbf{s}_q''(t)$, where $\mathbf{s}_q''(t)$ is the second derivative of the position $\mathbf{s}_q(t)$. Thus, we can use Formula 6.2 to find the acceleration of particle q:

$$\mathbf{s}_{q}''(t) = -G \sum_{\substack{j=0\\ i \neq q}}^{n-1} \frac{m_{j}}{\left|\mathbf{s}_{q}(t) - \mathbf{s}_{j}(t)\right|^{3}} \left[\mathbf{s}_{q}(t) - \mathbf{s}_{j}(t)\right]. \tag{6.3}$$

Thus Newton's laws give us a system of *differential* equations—equations involving derivatives—and our job is to find at each time t of interest the position $\mathbf{s}_q(t)$ and velocity $\mathbf{v}_q(t) = \mathbf{s}_q'(t)$.

We'll suppose that we either want to find the positions and velocities at the times

$$t = 0, \Delta t, 2\Delta t, \dots, T\Delta t,$$

or, more often, simply the positions and velocities at the final time $T\Delta t$. Here, Δt and T are specified by the user, so the input to the program will be n, the number of particles, Δt , T, and, for each particle, its mass, its initial position, and its initial velocity. In a fully general solver, the positions and velocities would be three-dimensional vectors, but in order to keep things simple, we'll assume that the particles will move in a plane, and we'll use two-dimensional vectors instead.

The output of the program will be the positions and velocities of the n particles at the timesteps $0, \Delta t, 2\Delta t, \ldots$, or just the positions and velocities at $T\Delta t$. To get the output at only the final time, we can add an input option in which the user specifies whether she only wants the final positions and velocities.

6.1.2 Two serial programs

In outline, a serial *n*-body solver can be based on the following pseudocode:

We can use our formula for the total force on a particle (Formula 6.2) to refine our pseudocode for the computation of the forces in Lines 4–5:

```
for each particle q {
   for each particle k != q {
     x_diff = pos[q][X] - pos[k][X];
     y_diff = pos[q][Y] - pos[k][Y];
     dist = sqrt(x_diff*x_diff + y_diff*y_diff);
     dist_cubed = dist*dist*dist;
     forces[q][X] -= G*masses[q]*masses[k]/dist_cubed * x_diff;
     forces[q][Y] -= G*masses[q]*masses[k]/dist_cubed * y_diff;
}
}
```

Here, we're assuming that the forces and the positions of the particles are stored as two-dimensional arrays, forces and pos, respectively. We're also assuming we've defined constants X = 0 and Y = 1. So the x-component of the force on particle q is forces[q][X] and the y-component is forces[q][Y]. Similarly, the components of the position are pos[q][X] and pos[q][Y]. (We'll take a closer look at data structures shortly.)

We can use Newton's third law of motion, that is, for every action there is an equal and opposite reaction, to halve the total number of calculations required for the forces. If the force on particle q due to particle k is \mathbf{f}_{qk} , then the force on k due to q is $-\mathbf{f}_{qk}$. Using this simplification we can modify our code to compute forces, as shown in Program 6.1. To better understand this pseudocode, imagine the individual forces as a two-dimensional array:

$$\begin{bmatrix} 0 & \mathbf{f}_{01} & \mathbf{f}_{02} & \cdots & \mathbf{f}_{0,n-1} \\ -\mathbf{f}_{01} & 0 & \mathbf{f}_{12} & \cdots & \mathbf{f}_{1,n-1} \\ -\mathbf{f}_{02} & -\mathbf{f}_{12} & 0 & \cdots & \mathbf{f}_{2,n-1} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ -\mathbf{f}_{0,n-1} & -\mathbf{f}_{1,n-1} & -\mathbf{f}_{2,n-1} & \cdots & 0 \end{bmatrix}.$$

(Why are the diagonal entries 0?) Our original solver simply adds all of the entries in row q to get forces[q]. In our modified solver, when q = 0, the body of the loop

```
for each particle q
   forces[q] = 0;
for each particle q {
        for each particle k > q {
            x_diff = pos[q][X] - pos[k][X];
            y_diff = pos[q][Y] - pos[k][Y];
            dist = sqrt(x_diff*x_diff + y_diff*y_diff);
            dist_cubed = dist*dist*dist;
            force_qk[X] = G*masses[q]*masses[k]/dist_cubed * x_diff;
            force_qk[Y] = G*masses[q]*masses[k]/dist_cubed * y_diff

            forces[q][X] += force_qk[X];
            forces[q][Y] += force_qk[Y];
            forces[k][X] -= force_qk[Y];
            forces[k][Y] -= force_qk[Y];
        }
}
```

Program 6.1: A reduced algorithm for computing *n*-body forces

for each particle q will add the entries in row 0 into forces[0]. It will also add the kth entry in column 0 into forces[k] for k = 1, 2, ..., n-1. In general, the qth iteration will add the entries to the right of the diagonal (that is, to the right of the 0) in row q into forces[q], and the entries below the diagonal in column q will be added into their respective forces, that is, the kth entry will be added in to forces[k].

Note that in using this modified solver, it's necessary to initialize the forces array in a separate loop, since the qth iteration of the loop that calculates the forces will, in general, add the values it computes into forces[k] for k = q + 1, q + 2, ..., n - 1, not just forces[q].

In order to distinguish between the two algorithms, we'll call the n-body solver with the original force calculation, the basic algorithm, and the solver with the number of calculations reduced, the reduced algorithm.

The position and the velocity remain to be found. We know that the acceleration of particle q is given by

$$\mathbf{a}_q(t) = \mathbf{s}_q''(t) = \mathbf{F}_q(t)/m_q,$$

where $\mathbf{s}_q''(t)$ is the second derivative of the position $\mathbf{s}_q(t)$ and $\mathbf{F}_q(t)$ is the force on particle q. We also know that the velocity $\mathbf{v}_q(t)$ is the first derivative of the position $\mathbf{s}_q'(t)$, so we need to integrate the acceleration to get the velocity, and we need to integrate the velocity to get the position.

We might at first think that we can simply find an antiderivative of the function in Formula 6.3. However, a second look shows us that this approach has problems: the right-hand side contains unknown functions \mathbf{s}_q and \mathbf{s}_k —not just the variable t—so we'll instead use a **numerical** method for *estimating* the position and the velocity. This means that rather than trying to find simple closed formulas, we'll approximate

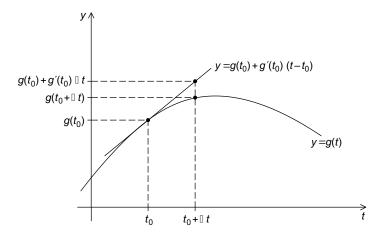


FIGURE 6.1

Using the tangent line to approximate a function

the values of the position and velocity at the times of interest. There are *many* possible choices for numerical methods, but we'll use the simplest one: Euler's method, which is named after the famous Swiss mathematician Leonhard Euler (1707–1783). In Euler's method, we use the tangent line to approximate a function. The basic idea is that if we know the value of a function $g(t_0)$ at time t_0 and we also know its derivative $g'(t_0)$ at time t_0 , then we can approximate its value at time $t_0 + \Delta t$ by using the tangent line to the graph of $g(t_0)$. See Figure 6.1 for an example. Now if we know a point $(t_0, g(t_0))$ on a line, and we know the slope of the line $g'(t_0)$, then an equation for the line is given by

$$y = g(t_0) + g'(t_0)(t - t_0).$$

Since we're interested in the time $t = t_0 + \Delta t$, we get

$$g(t + \Delta t) \approx g(t_0) + g'(t_0)(t + \Delta t - t) = g(t_0) + \Delta t g'(t_0).$$

Note that this formula will work even when g(t) and y are vectors: when this is the case, g'(t) is also a vector and the formula just adds a vector to a vector multiplied by a scalar, Δt .

Now we know the value of $\mathbf{s}_q(t)$ and $\mathbf{s}_q'(t)$ at time 0, so we can use the tangent line and our formula for the acceleration to compute $\mathbf{s}_q(\Delta t)$ and $\mathbf{v}_q(\Delta t)$:

$$\mathbf{s}_q(\Delta t) \approx \mathbf{s}_q(0) + \Delta t \mathbf{s}'_q(0) = \mathbf{s}_q(0) + \Delta t \mathbf{v}_q(0),$$

$$\mathbf{v}_q(\Delta t) \approx \mathbf{v}_q(0) + \Delta t \, \mathbf{v}_q'(0) = \mathbf{v}_q(0) + \Delta t \, \mathbf{a}_q(0) = \mathbf{v}_q(0) + \Delta t \frac{1}{m_q} \mathbf{F}_q(0).$$

When we try to extend this approach to the computation of $\mathbf{s}_q(2\Delta t)$ and $\mathbf{s}_q'(2\Delta t)$, we see that things are a little bit different, since we don't know the exact value of $\mathbf{s}_q(\Delta t)$

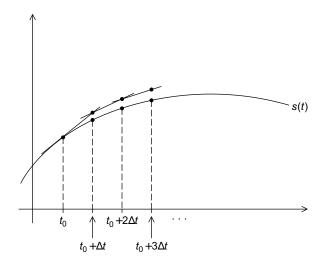


FIGURE 6.2

Euler's method

and $\mathbf{s}_q'(\Delta t)$. However, if our approximations to $\mathbf{s}_q(\Delta t)$ and $\mathbf{s}_q'(\Delta t)$ are good, then we should be able to get a reasonably good approximation to $\mathbf{s}_q(2\Delta t)$ and $\mathbf{s}_q'(2\Delta t)$ using the same idea. This is what Euler's method does (see Figure 6.2).

Now we can complete our pseudocode for the two *n*-body solvers by adding in the code for computing position and velocity:

```
pos[q][X] += delta_t*vel[q][X];
pos[q][Y] += delta_t*vel[q][Y];
vel[q][X] += delta_t/masses[q]*forces[q][X];
vel[q][Y] += delta_t/masses[q]*forces[q][Y];
```

Here, we're using pos[q], vel[q], and forces[q] to store the position, the velocity, and the force, respectively, of particle q.

Before moving on to parallelizing our serial program, let's take a moment to look at data structures. We've been using an array type to store our vectors:

```
#define DIM 2
typedef double vect_t[DIM];
```

A struct is also an option. However, if we're using arrays and we decide to change our program so that it solves three-dimensional problems, in principle, we only need to change the macro DIM. If we try to do this with structs, we'll need to rewrite the code that accesses individual components of the vector.

For each particle, we need to know the values of

- its mass,
- its position,

- · its velocity,
- · its acceleration, and
- the total force acting on it.

Since we're using Newtonian physics, the mass of each particle is constant, but the other values will, in general, change as the program proceeds. If we examine our code, we'll see that once we've computed a new value for one of these variables for a given timestep, we never need the old value again. For example, we don't need to do anything like this

```
new_pos_q = f(old_pos_q);
new_vel_q = g(old_pos_q, new_pos_q);
```

Also, the acceleration is only used to compute the velocity, and its value can be computed in one arithmetic operation from the total force, so we only need to use a local, temporary variable for the acceleration.

For each particle it suffices to store its mass and the current value of its position, velocity, and force. We could store these four variables as a struct and use an array of structs to store the data for all the particles. Of course, there's no reason that all of the variables associated with a particle need to be grouped together in a struct. We can split the data into separate arrays in a variety of different ways. We've chosen to group the mass, position, and velocity into a single struct and store the forces in a separate array. With the forces stored in contiguous memory, we can use a fast function such as memset to quickly assign zeroes to all of the elements at the beginning of each iteration:

```
#include <string.h> /* For memset */
...
vect_t* forces = malloc(n*sizeof(vect_t));
...
for (step = 1; step <= n_steps; step++) {
...
    /* Assign 0 to each element of the forces array */
    forces = memset(forces, 0, n*sizeof(vect_t);
    for (part = 0; part < n-1; part++)
        Compute_force(part, forces, ...)
...
}</pre>
```

If the force on each particle were a member of a struct, the force members wouldn't occupy contiguous memory in an array of structs, and we'd have to use a relatively slow for loop to assign zero to each element.

6.1.3 Parallelizing the n-body solvers

Let's try to apply Foster's methodology to the *n*-body solver. Since we initially want *lots* of tasks, we can start by making our tasks the computations of the positions, the velocities, and the total forces at each timestep. In the basic algorithm, the algorithm in which the total force on each particle is calculated directly from Formula 6.2, the



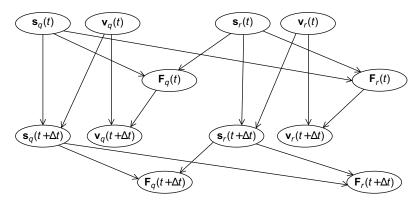


FIGURE 6.3

Communications among tasks in the basic *n*-body solver

computation of $\mathbf{F}_q(t)$, the total force on particle q at time t, requires the positions of each of the particles $\mathbf{s}_r(t)$, for each r. The computation of $\mathbf{v}_q(t+\Delta t)$ requires the velocity at the previous timestep, $\mathbf{v}_q(t)$, and the force, $\mathbf{F}_q(t)$, at the previous timestep. Finally, the computation of $\mathbf{s}_q(t+\Delta t)$ requires $\mathbf{s}_q(t)$ and $\mathbf{v}_q(t)$. The communications among the tasks can be illustrated as shown in Figure 6.3. The figure makes it clear that most of the communication among the tasks occurs among the tasks associated with an individual particle, so if we agglomerate the computations of $\mathbf{s}_q(t)$, $\mathbf{v}_q(t)$, and $\mathbf{F}_q(t)$, our intertask communication is greatly simplified (see Figure 6.4). Now the tasks correspond to the particles and, in the figure, we've labeled the communications with the data that's being communicated. For example, the arrow from particle q at timestep t to particle r at timestep t is labeled with \mathbf{s}_q , the position of particle q.

For the reduced algorithm, the "intra-particle" communications are the same. That is, to compute $\mathbf{s}_q(t+\Delta t)$ we'll need $\mathbf{s}_q(t)$ and $\mathbf{v}_q(t)$, and to compute $\mathbf{v}_q(t+\Delta t)$, we'll need $\mathbf{v}_q(t)$ and $\mathbf{F}_q(t)$. Therefore, once again it makes sense to agglomerate the computations associated with a single particle into a composite task.

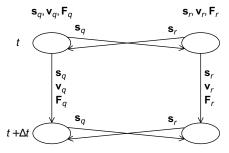


FIGURE 6.4

Communications among agglomerated tasks in the basic *n*-body solver

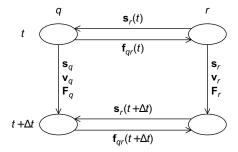


FIGURE 6.5

Communications among agglomerated tasks in the reduced *n*-body solver (q < r)

Recollect that in the reduced algorithm, we make use of the fact that the force $\mathbf{f}_{rq} = -\mathbf{f}_{qr}$. So if q < r, then the communication *from* task r to task q is the same as in the basic algorithm—in order to compute $\mathbf{F}_q(t)$, task/particle q will need $\mathbf{s}_r(t)$ from task/particle r. However, the communication from task q to task r is no longer $s_q(t)$, it's the force on particle q due to particle r, that is, $\mathbf{f}_{qr}(t)$. See Figure 6.5.

The final stage in Foster's methodology is mapping. If we have n particles and T timesteps, then there will be nT tasks in both the basic and the reduced algorithm. Astrophysical *n*-body problems typically involve thousands or even millions of particles, so n is likely to be several orders of magnitude greater than the number of available cores. However, T may also be much larger than the number of available cores. So, in principle, we have two "dimensions" to work with when we map tasks to cores. However, if we consider the nature of Euler's method, we'll see that attempting to assign tasks associated with a single particle at different timesteps to different cores won't work very well. Before estimating $\mathbf{s}_q(t+\Delta t)$ and $\mathbf{v}_q(t+\Delta t)$, Euler's method must "know" $\mathbf{s}_q(t)$, $\mathbf{v}_q(t)$, and $\mathbf{a}_q(t)$. Thus, if we assign particle q at time t to core c_0 , and we assign particle q at time $t + \Delta t$ to core $c_1 \neq c_0$, then we'll have to communicate $\mathbf{s}_q(t)$, $\mathbf{v}_q(t)$, and $\mathbf{F}_q(t)$ from c_0 to c_1 . Of course, if particle q at time t and particle q at time $t + \Delta t$ are mapped to the same core, this communication won't be necessary, so once we've mapped the task consisting of the calculations for particle q at the first timestep to core c_0 , we may as well map the subsequent computations for particle q to the same cores, since we can't simultaneously execute the computations for particle q at two different timesteps. Thus, mapping tasks to cores will, in effect, be an assignment of particles to cores.

At first glance, it might seem that any assignment of particles to cores that assigns roughly $n/\text{thread_count}$ particles to each core will do a good job of balancing the workload among the cores, and for the basic algorithm this is the case. In the basic algorithm the work required to compute the position, velocity, and force is the same for every particle. However, in the reduced algorithm the work required in the forces computation loop is much greater for lower-numbered iterations than the work required for higher-numbered iterations. To see this, recall the pseudocode that computes the total force on particle q in the reduced algorithm:

```
for each particle k > q {
    x_diff = pos[q][X] - pos[k][X];
    y_diff = pos[q][Y] - pos[k][Y];
    dist = sqrt(x_diff*x_diff + y_diff*y_diff);
    dist_cubed = dist*dist*dist;
    force_qk[X] = G*masses[q]*masses[k]/dist_cubed * x_diff;
    force_qk[Y] = G*masses[q]*masses[k]/dist_cubed * y_diff;

    forces[q][X] += force_qk[X];
    forces[q][Y] += force_qk[Y];
    forces[k][X] -= force_qk[Y];
}
```

Then, for example, when q=0, we'll make n-1 passes through the for each particle k > q loop, while when q=n-1, we won't make any passes through the loop. Thus, for the reduced algorithm we would expect that a cyclic partition of the particles would do a better job than a block partition of evenly distributing the *computation*.

However, in a shared-memory setting, a cyclic partition of the particles among the cores is almost certain to result in a much higher number of cache misses than a block partition, and in a distributed-memory setting, the overhead involved in communicating data that has a cyclic distribution will probably be greater than the overhead involved in communicating data that has a block distribution (see Exercises 6.8 and 6.9).

Therefore with a composite task consisting of all of the computations associated with a single particle throughout the simulation, we conclude the following:

- 1. A block distribution will give the best performance for the basic *n*-body solver.
- **2.** For the reduced *n*-body solver, a cyclic distribution will best distribute the workload in the computation of the forces. However, this improved performance *may* be offset by the cost of reduced cache performance in a shared-memory setting and additional communication overhead in a distributed-memory setting.

In order to make a final determination of the optimal mapping of tasks to cores, we'll need to do some experimentation.

6.1.4 A word about I/O

You may have noticed that our discussion of parallelizing the *n*-body solver hasn't touched on the issue of I/O, even though I/O can figure prominently in both of our serial algorithms. We've discussed the problem of I/O several times in earlier chapters. Recall that different parallel systems vary widely in their I/O capabilities, and with the very basic I/O that is commonly available it is very difficult to obtain high performance. This basic I/O was designed for use by single-process, single-threaded programs, and when multiple processes or multiple threads attempt to access the I/O buffers, the system makes no attempt to schedule their access. For example, if multiple threads attempt to execute

```
printf("Hello from thread %d of %d\n", my_rank, thread_count);
```

more or less simultaneously, the order in which the output appears will be unpredictable. Even worse, one thread's output may not even appear as a single line. It can happen that the output from one thread appears as multiple segments, and the individual segments are separated by output from other threads.

Thus, as we've noted earlier, except for debug output, we generally assume that one process/thread does all the I/O, and when we're timing program execution, we'll use the option to only print output for the final timestep. Furthermore, we won't include this output in the reported run-times.

Of course, even if we're ignoring the cost of I/O, we can't ignore its existence. We'll briefly discuss its implementation when we discuss the details of our parallel implementations.

6.1.5 Parallelizing the basic solver using OpenMP

How can we use OpenMP to map tasks/particles to cores in the basic version of our *n*-body solver? Let's take a look at the pseudocode for the serial program:

```
for each timestep {
   if (timestep output) Print positions and velocities of particles;
   for each particle q
      Compute total force on q;
   for each particle q
      Compute position and velocity of q;
}
```

The two inner loops are both iterating over particles. So, in principle, parallelizing the two inner for loops will map tasks/particles to cores, and we might try something like this:

```
for each timestep {
    if (timestep output) Print positions and velocities of
        particles;

# pragma omp parallel for
    for each particle q
        Compute total force on q;

# pragma omp parallel for
    for each particle q
        Compute position and velocity of q;
}
```

We may not like the fact that this code could do a lot of forking and joining of threads, but before dealing with that, let's take a look at the loops themselves: we need to see if there are any race conditions caused by loop-carried dependences.

In the basic version the first loop has the following form:

```
# pragma omp parallel for
for each particle q {
  forces[q][X] = forces[q][Y] = 0;
  for each particle k != q {
    x_diff = pos[q][X] - pos[k][X];
    y_diff = pos[q][Y] - pos[k][Y];
```

```
dist = sqrt(x_diff*x_diff + y_diff*y_diff);
    dist_cubed = dist*dist*dist;
    forces[q][X] -= G*masses[q]*masses[k]/dist_cubed * x_diff;
    forces[q][Y] -= G*masses[q]*masses[k]/dist_cubed * y_diff;
}
}
```

Since the iterations of the for each particle q loop are partitioned among the threads, only one thread will access forces[q] for any q. Different threads do access the same elements of the pos array and the masses array. However, these arrays are only read in the loop. The remaining variables are used for temporary storage in a single iteration of the inner loop, and they can be private. Thus, the parallelization of the first loop in the basic algorithm won't introduce any race conditions.

The second loop has the form:

```
# pragma omp parallel for
for each particle q {
    pos[q][X] += delta_t*vel[q][X];
    pos[q][Y] += delta_t*vel[q][Y];
    vel[q][X] += delta_t/masses[q]*forces[q][X];
    vel[q][Y] += delta_t/masses[q]*forces[q][Y];
}
```

Here, a single thread accesses pos[q], vel[q], masses[q], and forces[q] for any particle q, and the scalar variables are only read, so parallelizing this loop also won't introduce any race conditions.

Let's return to the issue of repeated forking and joining of threads. In our pseudocode, we have

```
for each timestep {
    if (timestep output) Print positions and velocities of
        particles;

# pragma omp parallel for
    for each particle q
        Compute total force on q;

pragma omp parallel for
    for each particle q
        Compute position and velocity of q;
}
```

We encountered a similar issue when we parallelized odd-even transposition sort (see Section 5.6.2). In that case, we put a parallel directive before the outermost loop and used OpenMP for directives for the inner loops. Will a similar strategy work here? That is, can we do something like this?

```
# pragma omp parallel
   for each timestep {
      if (timestep output) Print positions and velocities of
          particles;
# pragma omp for
   for each particle q
```

```
Compute total force on q;

# pragma omp for
for each particle q
Compute position and velocity of q;
}
```

This will have the desired effect on the two for each particle loops: the same team of threads will be used in both loops and for every iteration of the outer loop. However, we have a clear problem with the output statement. As it stands now, every thread will print all the positions and velocities, and we only want one thread to do the I/O. However, OpenMP provides the single directive for exactly this situation: we have a team of threads executing a block of code, but a part of the code should only be executed by one of the threads. Adding the single directive gives us the following pseudocode:

```
# pragma omp parallel
    for each timestep {
        if (timestep output) {
            pragma omp single
                Print positions and velocities of particles;
        }
# pragma omp for
        for each particle q
            Compute total force on q;
# pragma omp for
        for each particle q
            Compute position and velocity of q;
}
```

There are still a few issues that we need to address. The most important has to do with possible race conditions introduced in the transition from one statement to another. For example, suppose thread 0 completes the first for each particle loop before thread 1, and it then starts updating the positions and velocities of its assigned particles in the second for each particle loop. Clearly, this could cause thread 1 to use an updated position in the first for each particle loop. However, recall that there is an implicit barrier at the end of each structured block that has been parallelized with a for directive. So, if thread 0 finishes the first inner loop before thread 1, it will block until thread 1 (and any other threads) finish the first inner loop, and it won't start the second inner loop until all the threads have finished the first. This will also prevent the possibility that a thread might rush ahead and print positions and velocities before they've all been updated by the second loop.

There's also an implicit barrier after the single directive, although in this program the barrier isn't necessary. Since the output statement won't update any memory locations, it's OK for some threads to go ahead and start executing the next iteration before output has been completed. Furthermore, the first inner for loop in the next iteration only updates the forces array, so it can't cause a thread executing the output statement to print incorrect values, and because of the barrier at the end of the first inner loop, no thread can race ahead and start updating positions and velocities in

the second inner loop before the output has been completed. Thus, we could modify the single directive with a nowait clause. If the OpenMP implementation supports it, this simply eliminates the implied barrier associated with the single directive. It can also be used with for, parallel for, and parallel directives. Note that in this case, addition of the nowait clause is unlikely to have much effect on performance, since the two for each particle loops have implied barriers that will prevent any one thread from getting more than a few statements ahead of any other.

Finally, we may want to add a schedule clause to each of the for directives in order to insure that the iterations have a block partition:

```
# pragma omp for schedule(static, n/thread_count)
```

6.1.6 Parallelizing the reduced solver using OpenMP

The reduced solver has an additional inner loop: the initialization of the forces array to 0. If we try to use the same parallelization for the reduced solver, we should also parallelize this loop with a for directive. What happens if we try this? That is, what happens if we try to parallelize the reduced solver with the following pseudocode?

```
# pragma omp parallel
   for each timestep {
      if (timestep output) {
#
         pragma omp single
         Print positions and velocities of particles;
      pragma omp for
#
      for each particle q
         forces[q] = 0.0;
      pragma omp for
      for each particle q
         Compute total force on q;
      pragma omp for
      for each particle q
         Compute position and velocity of q;
```

Parallelization of the initialization of the forces should be fine, as there's no dependence among the iterations. The updating of the positions and velocities is the same in both the basic and reduced solvers, so if the computation of the forces is OK, then this should also be OK.

How does parallelization affect the correctness of the loop for computing the forces? Recall that in the reduced version, this loop has the following form:

```
# pragma omp for /* Can be faster than memset */
for each particle q {
   force_qk[X] = force_qk[Y] = 0;
   for each particle k > q {
```

```
x_diff = pos[q][X] - pos[k][X];
y_diff = pos[q][Y] - pos[k][Y];
dist = sqrt(x_diff*x_diff + y_diff*y_diff);
dist_cubed = dist*dist*dist;
force_qk[X] = G*masses[q]*masses[k]/dist_cubed * x_diff;
force_qk[Y] = G*masses[q]*masses[k]/dist_cubed * y_diff;

forces[q][X] += force_qk[X];
forces[q][Y] += force_qk[Y];
forces[k][X] -= force_qk[Y];
forces[k][Y] -= force_qk[Y];
}
```

As before, the variables of interest are pos, masses, and forces, since the values in the remaining variables are only used in a single iteration, and hence, can be private. Also, as before, elements of the pos and masses arrays are only read, not updated. We therefore need to look at the elements of the forces array. In this version, unlike the basic version, a thread *may* update elements of the forces array other than those corresponding to its assigned particles. For example, suppose we have two threads and four particles and we're using a block partition of the particles. Then the total force on particle 3 is given by

$$\mathbf{F}_3 = -\mathbf{f}_{03} - \mathbf{f}_{13} - \mathbf{f}_{23}$$
.

Furthermore, thread 0 will compute \mathbf{f}_{03} and \mathbf{f}_{13} , while thread 1 will compute \mathbf{f}_{23} . Thus, the updates to forces[3] do create a race condition. In general, then, the updates to the elements of the forces array introduce race conditions into the code.

A seemingly obvious solution to this problem is to use a critical directive to limit access to the elements of the forces array. There are at least a couple of ways to do this. The simplest is to put a critical directive before all the updates to forces

```
# pragma omp critical
{
    forces[q][X] += force_qk[X];
    forces[q][Y] += force_qk[Y];
    forces[k][X] -= force_qk[X];
    forces[k][Y] -= force_qk[Y];
}
```

However, with this approach access to the elements of the forces array will be effectively serialized. Only one element of forces can be updated at a time, and contention for access to the critical section is actually likely to seriously degrade the performance of the program. See Exercise 6.3.

An alternative would be to have one critical section for each particle. However, as we've seen, OpenMP doesn't readily support varying numbers of critical sections, so we would need to use one lock for each particle instead and our updates would

look something like this:

```
omp_set_lock(&locks[q]);
forces[q][X] += force_qk[X];
forces[q][Y] += force_qk[Y];
omp_unset_lock(&locks[q]);

omp_set_lock(&locks[k]);
forces[k][X] -= force_qk[X];
forces[k][Y] -= force_qk[Y];
omp_unset_lock(&locks[k]);
```

This assumes that the master thread will create a shared array of locks, one for each particle, and when we update an element of the forces array, we first set the lock corresponding to that particle. Although this approach performs much better than the single critical section, it still isn't competitive with the serial code. See Exercise 6.4.

Another possible solution is to carry out the computation of the forces in two phases. In the first phase, each thread carries out exactly the same calculations it carried out in the erroneous parallelization. However, now the calculations are stored in its *own* array of forces. Then, in the second phase, the thread that has been assigned particle q will add the contributions that have been computed by the different threads. In our example above, thread 0 would compute $-\mathbf{f}_{03} - \mathbf{f}_{13}$, while thread 1 would compute $-\mathbf{f}_{23}$. After each thread was done computing its contributions to the forces, thread 1, which has been assigned particle 3, would find the total force on particle 3 by adding these two values.

Let's look at a slightly larger example. Suppose we have three threads and six particles. If we're using a block partition of the particles, then the computations in the first phase are shown in Table 6.1. The last three columns of the table show each thread's contribution to the computation of the total forces. In phase 2 of the computation, the thread specified in the first column of the table will add the contents of each of its assigned rows—that is, each of its assigned particles.

Note that there's nothing special about using a block partition of the particles. Table 6.2 shows the same computations if we use a cyclic partition of the particles.

Table 6.1 First-Phase Computations for a Reduced Algorithm with Block Partition							
		Threa	ad				
Thread	Particle	0	1	2			
0	0	$\mathbf{f}_{01} + \mathbf{f}_{02} + \mathbf{f}_{03} + \mathbf{f}_{04} + \mathbf{f}_{05}$	0	0			
	1	$-\mathbf{f}_{01} + \mathbf{f}_{12} + \mathbf{f}_{13} + \mathbf{f}_{14} + \mathbf{f}_{15}$	0	0			
1	2	$-\mathbf{f}_{02} - \mathbf{f}_{12}$	$\mathbf{f}_{23} + \mathbf{f}_{24} + \mathbf{f}_{25}$	0			
	3	$-\mathbf{f}_{03} - \mathbf{f}_{13}$	$-\mathbf{f}_{23} + \mathbf{f}_{34} + \mathbf{f}_{35}$	0			
2	4	$-\mathbf{f}_{04} - \mathbf{f}_{14}$	$-\mathbf{f}_{24} - \mathbf{f}_{34}$	f ₄₅			
	5	$-\mathbf{f}_{05} - \mathbf{f}_{15}$	$-\mathbf{f}_{25} - \mathbf{f}_{35}$	- f ₄₅			

Table 6	Table 6.2 First-Phase Computations for a Reduced Algorithm with Cyclic Partition							
			Thread					
Thread	Particle	0	1	2				
0	0	$\mathbf{f}_{01} + \mathbf{f}_{02} + \mathbf{f}_{03} + \mathbf{f}_{04} + \mathbf{f}_{05}$	0	0				
1	1	- f ₀₁	$\mathbf{f}_{12} + \mathbf{f}_{13} + \mathbf{f}_{14} + \mathbf{f}_{15}$	0				
2	2	- f ₀₂	$-\mathbf{f}_{12}$	$\mathbf{f}_{23} + \mathbf{f}_{24} + \mathbf{f}_{25}$				
0	3	$-\mathbf{f}_{03} + \mathbf{f}_{34} + \mathbf{f}_{35}$	- f ₁₃	$-\mathbf{f}_{23}$				
1	4	$-\mathbf{f}_{04} - \mathbf{f}_{34} \\ -\mathbf{f}_{05} - \mathbf{f}_{35}$	$-\mathbf{f}_{14} + \mathbf{f}_{45}$	$-\mathbf{f}_{24}$				
2	5	$-\mathbf{f}_{05}-\mathbf{f}_{35}$	$-\mathbf{f}_{15} - \mathbf{f}_{45}$	−f ₂₅				

Note that if we compare this table with the table that shows the block partition, it's clear that the cyclic partition does a better job of balancing the load.

To implement this, during the first phase our revised algorithm proceeds as before, except that each thread adds the forces it computes into its own subarray of loc_forces:

```
pragma omp for
for each particle q {
   force_qk[X] = force_qk[Y] = 0;
   for each particle k > q {
      x_diff = pos[q][X] - pos[k][X];
      y_diff = pos[q][Y] - pos[k][Y];
      dist = sqrt(x_diff*x_diff + y_diff*y_diff);
      dist_cubed = dist*dist*dist;
       force_qk[X] = G*masses[q]*masses[k]/dist_cubed * x_diff;
       force_qk[Y] = G*masses[q]*masses[k]/dist_cubed * y_diff;
       loc_forces[my_rank][q][X] += force_qk[X];
       loc_forces[my_rank][q][Y] += force_qk[Y];
      loc_forces[my_rank][k][X] -= force_qk[X];
      loc_forces[my_rank][k][Y] -= force_qk[Y];
   }
}
```

During the second phase, each thread adds the forces computed by all the threads for its assigned particles:

```
# pragma omp for
    for (q = 0; q < n; q++) {
        forces[q][X] = forces[q][Y] = 0;
        for (thread = 0; thread < thread_count; thread++) {
            forces[q][X] += loc_forces[thread][q][X];
            forces[q][Y] += loc_forces[thread][q][Y];
        }
}</pre>
```

Before moving on, we should make sure that we haven't inadvertently introduced any new race conditions. During the first phase, since each thread writes to its own subarray, there isn't a race condition in the updates to loc_forces . Also, during the second phase, only the "owner" of thread q writes to $loc_forces[q]$, so there are no race conditions in the second phase. Finally, since there is an implied barrier after each of the parallelized for loops, we don't need to worry that some thread is going to race ahead and make use of a variable that hasn't been properly initialized, or that some slow thread is going to make use of a variable that has had its value changed by another thread.

6.1.7 Evaluating the OpenMP codes

Before we can compare the basic and the reduced codes, we need to decide how to schedule the parallelized for loops. For the basic code, we've seen that any schedule that divides the iterations equally among the threads should do a good job of balancing the computational load. (As usual, we're assuming no more than one thread/core.) We also observed that a block partitioning of the iterations would result in fewer cache misses than a cyclic partition. Thus, we would expect that a block schedule would be the best option for the basic version.

In the reduced code, the amount of work done in the first phase of the computation of the forces decreases as the for loop proceeds. We've seen that a cyclic schedule should do a better job of assigning more or less equal amounts of work to each thread. In the remaining parallel for loops—the initialization of the loc_forces array, the second phase of the computation of the forces, and the updating of the positions and velocities—the work required is roughly the same for all the iterations. Therefore, taken out of context each of these loops will probably perform best with a block schedule. However, the schedule of one loop can affect the performance of another (see Exercise 6.10), so it may be that choosing a cyclic schedule for one loop and block schedules for the others will degrade performance.

With these choices, Table 6.3 shows the performance of the *n*-body solvers when they're run on one of our systems with no I/O. The solver used 400 particles for 1000 timesteps. The column labeled "Default Sched" gives times for the OpenMP reduced solver when all of the inner loops use the default schedule, which, on our system, is a block schedule. The column labeled "Forces Cyclic" gives times when the first phase of the forces computation uses a cyclic schedule and the other inner loops use the default schedule. The last column, labeled "All Cyclic," gives times when all of

Table 6.3 Run-Times of the <i>n</i> -Body Solvers Parallelized with OpenMP (times are in seconds)							
Reduced Reduced Reduced Threads Basic Default Sched Forces Cyclic All Cy							
1	7.71	3.90	3.90	3.90			
2 4	3.87 1.95	2.94 1.73	1.98 1.01	2.01 1.08			
8	0.99	0.95	0.54	0.61			

the inner loops use a cyclic schedule. The run-times of the serial solvers differ from those of the single-threaded solvers by less than 1%, so we've omitted them from the table.

Notice that with more than one thread the reduced solver, using all default schedules, takes anywhere from 50 to 75% longer than the reduced solver with the cyclic forces computation. Using the cyclic schedule is clearly superior to the default schedule in this case, and any loss in time resulting from cache issues is more than made up for by the improved load balance for the computations.

For only two threads there is very little difference between the performance of the reduced solver with only the first forces loop cyclic and the reduced solver with all loops cyclic. However, as we increase the number of threads, the performance of the reduced solver that uses a cyclic schedule for all of the loops does start to degrade. In this particular case, when there are more threads, it appears that the overhead involved in changing distributions is less than the overhead incurred from false sharing.

Finally, notice that the basic solver takes about twice as long as the reduced solver with the cyclic scheduling of the forces computation. So if the extra memory is available, the reduced solver is clearly superior. However, the reduced solver increases the memory requirement for the storage of the forces by a factor of thread_count, so for very large numbers of particles, it may be impossible to use the reduced solver.

6.1.8 Parallelizing the solvers using pthreads

Parallelizing the two *n*-body solvers using Pthreads is very similar to parallelizing them using OpenMP. The differences are only in implementation details, so rather than repeating the discussion, we will point out some of the principal differences between the Pthreads and the OpenMP implementations. We will also note some of the more important similarities.

- By default local variables in Pthreads are private, so all shared variables are global in the Pthreads version.
- The principal data structures in the Pthreads version are identical to those in the OpenMP version: vectors are two-dimensional arrays of doubles, and the mass, position, and velocity of a single particle are stored in a struct. The forces are stored in an array of vectors.
- Startup for Pthreads is basically the same as the startup for OpenMP: the main thread gets the command-line arguments, and allocates and initializes the principal data structures.
- The main difference between the Pthreads and the OpenMP implementations is in the details of parallelizing the inner loops. Since Pthreads has nothing analogous to a parallel for directive, we must explicitly determine which values of the loop variables correspond to each thread's calculations. To facilitate this, we've written a function Loop_schedule, which determines
 - the initial value of the loop variable,
 - the final value of the loop variable, and
 - the increment for the loop variable.

The input to the function is

- the calling thread's rank,
- the number of threads,
- the total number of iterations, and
- an argument indicating whether the partitioning should be block or cyclic.
- Another difference between the Pthreads and the OpenMP versions has to do with barriers. Recall that the end of a parallel for directive in OpenMP has an implied barrier. As we've seen, this is important. For example, we don't want a thread to start updating its positions until all the forces have been calculated, because it could use an out-of-date force and another thread could use an out-of-date position. If we simply partition the loop iterations among the threads in the Pthreads version, there won't be a barrier at the end of an inner for loop and we'll have a race condition. Thus, we need to add explicit barriers after the inner loops when a race condition can arise. The Pthreads standard includes a barrier. However, some systems don't implement it, so we've defined a function that uses a Pthreads condition variable to implement a barrier. See Subsection 4.8.3 for details.

6.1.9 Parallelizing the basic solver using MPI

With our composite tasks corresponding to the individual particles, it's fairly straightforward to parallelize the basic algorithm using MPI. The only communication among the tasks occurs when we're computing the forces, and, in order to compute the forces, each task/particle needs the position and mass of every other particle. MPI_Allgather is expressly designed for this situation, since it collects on each process the same information from every other process. We've already noted that a block distribution will probably have the best performance, so we should use a block mapping of the particles to the processes.

In the shared-memory implementations, we collected most of the data associated with a single particle (mass, position, and velocity) into a single struct. However, if we use this data structure in the MPI implementation, we'll need to use a derived datatype in the call to MPI_Allgather, and communications with derived datatypes tend to be slower than communications with basic MPI types. Thus, it will make more sense to use individual arrays for the masses, positions, and velocities. We'll also need an array for storing the positions of all the particles. If each process has sufficient memory, then each of these can be a separate array. In fact, if memory isn't a problem, each process can store the entire array of masses, since these will never be updated and their values only need to be communicated during the initial setup.

On the other hand, if memory is short, there is an "in-place" option that can be used with some MPI collective communications. For our situation, suppose that the array pos can store the positions of all n particles. Further suppose that $vect_mpi_t$ is an MPI datatype that stores two contiguous doubles. Also suppose that n is evenly divisible by comm_sz and $loc_n = n/comm_sz$. Then, if we store the local positions in a separate array, loc_pos , we can use the following call to collect all of the positions

on each process:

If we can't afford the extra storage for loc_pos , then we can have each process q store its local positions in the qth block of pos. That is, the local positions of each process should be stored in the appropriate block of each process' pos array:

With the pos array initialized this way on each process, we can use the following call to MPI_Allgather:

```
MPI_Allgather(MPI_IN_PLACE, loc_n, vect_mpi_t,
pos, loc_n, vect_mpi_t, comm);
```

In this call, the first loc_n and vect_mpi_t arguments are ignored. However, it's not a bad idea to use arguments whose values correspond to the values that will be used, just to increase the readability of the program.

In the program we've written, we made the following choices with respect to the data structures:

- Each process stores the entire global array of particle masses.
- Each process only uses a single *n*-element array for the positions.
- Each process uses a pointer loc_pos that refers to the start of its block of pos. Thus, on process, 0 local_pos = pos, on process 1 local_pos = pos + loc_n, and, so on.

With these choices, we can implement the basic algorithm with the pseudocode shown in Program 6.2. Process 0 will read and broadcast the command line arguments. It will also read the input and print the results. In Line 1, it will need to distribute the input data. Therefore, Get input data might be implemented as follows:

```
if (my_rank == 0) {
   for each particle
     Read masses[particle], pos[particle], vel[particle];
}
MPI_Bcast(masses, n, MPI_DOUBLE, 0, comm);
MPI_Bcast(pos, n, vect_mpi_t, 0, comm);
MPI_Scatter(vel, loc_n, vect_mpi_t, loc_vel, loc_n, vect_mpi_t, 0, comm);
```

So process 0 reads all the initial conditions into three *n*-element arrays. Since we're storing all the masses on each process, we broadcast masses. Also, since each process

```
Get input data;
   for each timestep {
3
      if (timestep output)
         Print positions and velocities of particles;
4
      for each local particle loc_q
5
         Compute total force on loc_q;
6
7
      for each local particle loc_q
         Compute position and velocity of loc_q;
8
9
      Allgather local positions into global pos array;
10
   Print positions and velocities of particles;
11
```

Program 6.2: Pseudocode for the MPI version of the basic *n*-body solver

will need the global array of positions for the first computation of forces in the main for loop, we just broadcast pos. However, velocities are only used locally for the updates to positions and velocities, so we scatter vel.

Notice that we gather the updated positions in Line 9 at the end of the body of the outer for loop of Program 6.2. This insures that the positions will be available for output in both Line 4 and Line 11. If we're printing the results for each timestep, this placement allows us to eliminate an expensive collective communication call: if we simply gathered the positions onto process 0 before output, we'd have to call MPI_Allgather before the computation of the forces. With this organization of the body of the outer for loop, we can implement the output with the following pseudocode:

```
Gather velocities onto process 0;
if (my_rank == 0) {
   Print timestep;
   for each particle
       Print pos[particle] and vel[particle]
}
```

6.1.10 Parallelizing the reduced solver using MPI

The "obvious" implementation of the reduced algorithm is likely to be extremely complicated. Before computing the forces, each process will need to gather a subset of the positions, and after the computation of the forces, each process will need to scatter some of the individual forces it has computed and add the forces it receives. Figure 6.6 shows the communications that would take place if we had three processes, six particles, and used a block partitioning of the particles among the processes. Not suprisingly, the communications are even more complex when we use a cyclic distribution (see Exercise 6.13). Certainly it would be possible to implement these communications. However, unless the implementation were *very* carefully done, it would probably be *very* slow.

Fortunately, there's a much simpler alternative that uses a communication structure that is sometimes called a **ring pass**. In a ring pass, we imagine the processes

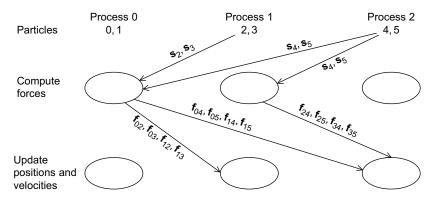


FIGURE 6.6

Communication in a possible MPI implementation of the reduced *n*-body solver

as being interconnected in a ring (see Figure 6.7). Process 0 communicates directly with processes 1 and communicates 1 communicates with processes 0 and 2, and so on. The communication in a ring pass takes place in phases, and during each phase each process sends data to its "lower-ranked" neighbor, and receives data from its "higher-ranked" neighbor. Thus, 0 will send to communicates q = 1 and receive from 1.1 will send to 0 and receive from 2, and so on. In general, process q = 1 + communicates 2 % communicates directly with process q = 1 + communicates 3 % communicates directly with process q = 1 + communicates 3 % communicates directly with process q = 1 + communicates 3 % communicates q = 1 + commun

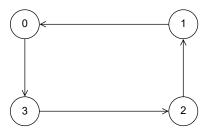


FIGURE 6.7

A ring of processes

By repeatedly sending and receiving data using this ring structure, we can arrange that each process has access to the positions of all the particles. During the first phase, each process will send the positions of its assigned particles to its "lower-ranked" neighbor and receive the positions of the particles assigned to its higher-ranked neighbor. During the next phase, each process will forward the positions it received in the first phase. This process continues through $comm_sz - 1$ phases until each process has received the positions of all of the particles. Figure 6.8 shows the three phases if there are four processes and eight particles that have been cyclically distributed.

Of course, the virtue of the reduced algorithm is that we don't need to compute all of the inter-particle forces since $\mathbf{f}_{kq} = -\mathbf{f}_{qk}$, for every pair of particles q and k. To see



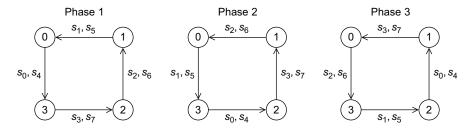


FIGURE 6.8

Ring pass of positions

how to exploit this, first observe that using the reduced algorithm, the interparticle forces can be divided into those that are *added* into and those that are subtracted from the total forces on the particle. For example, if we have six particles, then the reduced algorithm will compute the force on particle 3 as

$$\mathbf{F}_3 = -\mathbf{f}_{03} - \mathbf{f}_{13} - \mathbf{f}_{23} + \mathbf{f}_{34} + \mathbf{f}_{35}.$$

The key to understanding the ring pass computation of the forces is to observe that the interparticle forces that are *subtracted* are computed by another task/particle, while the forces that are *added* are computed by the owning task/particle. Thus, the computations of the interparticle forces on particle 3 are assigned as follows:

Force	f 03	f ₁₃	f ₂₃	f ₃₄	f ₃₅
Task/Particle	0	1	2	3	3

So, suppose that for our ring pass, instead of simply passing $loc_n = n/comm_sz$ positions, we also pass loc_n forces. Then in each phase, a process can

- 1. compute interparticle forces resulting from interaction between its assigned particles and the particles whose positions it has received, and
- **2.** once an interparticle force has been computed, the process can add the force into a local array of forces corresponding to its particles, *and* it can subtract the interparticle force from the received array of forces.

See, for example, [15, 34] for further details and alternatives.

Let's take a look at how the computation would proceed when we have four particles, two processes, and we're using a cyclic distribution of the particles among the processes (see Table 6.4). We're calling the arrays that store the local positions and local forces loc_pos and loc_forces, respectively. These are not communicated among the processes. The arrays that are communicated among the processes are tmp_pos and tmp_forces.

Before the ring pass can begin, both arrays storing positions are initialized with the positions of the local particles, and the arrays storing the forces are set to 0. Before the ring pass begins, each process computes those forces that are due to interaction

Table 6.	Table 6.4 Computation of Forces in Ring Pass						
Time	Variable	Process 0	Process 1				
Start		s ₀ , s ₂ 0, 0 s ₀ , s ₂ 0, 0	s ₁ , s ₃ 0, 0 s ₁ , s ₃ 0, 0				
	loc_forces tmp_pos	s ₀ , s ₂ f ₀₂ , 0 s ₀ , s ₂ 0, - f ₀₂	s ₁ , s ₃ f ₁₃ , 0 s ₁ , s ₃ 0, - f ₁₃				
After First Comm	loc_forces tmp_pos	s ₀ , s ₂ f ₀₂ , 0 s ₁ , s ₃ 0, - f ₁₃	s ₁ , s ₃ f ₁₃ , 0 s ₀ , s ₂ 0, - f ₀₂				
After Comp of Forces	loc_forces tmp_pos	$\mathbf{s}_{0}, \mathbf{s}_{2}$ $\mathbf{f}_{01} + \mathbf{f}_{02} + \mathbf{f}_{03}, \mathbf{f}_{23}$ $\mathbf{s}_{1}, \mathbf{s}_{3}$ $-\mathbf{f}_{01}, -\mathbf{f}_{03} - \mathbf{f}_{13} - \mathbf{f}_{23}$	$\mathbf{s}_{1}, \mathbf{s}_{3}$ $\mathbf{f}_{12} + \mathbf{f}_{13}, 0$ $\mathbf{s}_{0}, \mathbf{s}_{2}$ $0, -\mathbf{f}_{02} - \mathbf{f}_{12}$				
After Second Comm	loc_forces tmp_pos	$\mathbf{s}_{0}, \mathbf{s}_{2}$ $\mathbf{f}_{01} + \mathbf{f}_{02} + \mathbf{f}_{03}, \mathbf{f}_{23}$ $\mathbf{s}_{0}, \mathbf{s}_{2}$ $0, -\mathbf{f}_{02} - \mathbf{f}_{12}$	$\mathbf{s}_{1}, \mathbf{s}_{3}$ $\mathbf{f}_{12} + \mathbf{f}_{13}, 0$ $\mathbf{s}_{1}, \mathbf{s}_{3}$ $-\mathbf{f}_{01}, -\mathbf{f}_{03} - \mathbf{f}_{13} - \mathbf{f}_{23}$				
After Comp of Forces	loc_forces	s ₀ , s ₂	$\begin{array}{l} \textbf{s}_{1}, \textbf{s}_{3} \\ -\textbf{f}_{01} + \textbf{f}_{12} + \textbf{f}_{13}, -\textbf{f}_{03} - \textbf{f}_{13} - \textbf{f}_{23} \\ \textbf{s}_{1}, \textbf{s}_{3} \\ -\textbf{f}_{01}, -\textbf{f}_{03} - \textbf{f}_{13} - \textbf{f}_{23} \end{array}$				

among its assigned particles. Process 0 computes \mathbf{f}_{02} and process 1 computes \mathbf{f}_{13} . These values are added into the appropriate locations in loc_forces and subtracted from the appropriate locations in tmp_forces.

Now, the two processes exchange tmp_pos and tmp_forces and compute the forces due to interaction among their local particles and the received particles. In the reduced algorithm, the lower ranked task/particle carries out the computation. Process 0 computes \mathbf{f}_{01} , \mathbf{f}_{03} , and \mathbf{f}_{23} , while process 1 computes \mathbf{f}_{12} . As before, the newly computed forces are added into the appropriate locations in loc_forces and subtracted from the appropriate locations in tmp_forces.

To complete the algorithm, we need to exchange the tmp arrays one final time. Once each process has received the updated tmp_forces, it can carry out a simple vector sum

```
loc_forces += tmp_forces
```

to complete the algorithm.

¹Actually, we only need to exchange tmp_forces for the final communication.

```
source = (my\_rank + 1) \% comm\_sz;
   dest = (my\_rank - 1 + comm\_sz) \% comm\_sz;
3
   Copy loc_pos into tmp_pos;
   loc_forces = tmp_forces = 0;
   Compute forces due to interactions among local particles;
   for (phase = 1; phase < comm_sz; phase++) {</pre>
8
      Send current tmp_pos and tmp_forces to dest;
9
      Receive new tmp_pos and tmp_forces from source;
10
       /* Owner of the positions and forces we're receiving */
       owner = (my_rank + phase) % comm_sz;
11
12
      Compute forces due to interactions among my particles
13
         and owner's particles;
14
   Send current tmp_pos and tmp_forces to dest;
   Receive new tmp_pos and tmp_forces from source;
```

Program 6.3: Pseudocode for the MPI implementation of the reduced *n*-body solver

Thus, we can implement the computation of the forces in the reduced algorithm using a ring pass with the pseudocode shown in Program 6.3. Recall that using MPI_Send and MPI_Recv for the send-receive pairs in Lines 8 and 9 and 15 and 16 is *unsafe* in MPI parlance, since they can hang if the system doesn't provide sufficient buffering. In this setting, recall that MPI provides MPI_Sendrecv and MPI_Sendrecv_replace. Since we're using the same memory for both the outgoing and the incoming data, we can use MPI_Sendrecv_replace.

Also recall that the time it takes to start up a message is substantial. We can probably reduce the cost of the communication by using a single array to store both tmp_pos and tmp_forces. For example, we could allocate storage for an array tmp_data that can store $2 \times loc_n$ objects with type vect_t and use the first loc_n for tmp_pos and the last loc_n for tmp_forces. We can continue to use tmp_pos and tmp_forces by making these pointers to tmp_data[0] and tmp_data[loc_n], respectively.

The principal difficulty in implementing the actual computation of the forces in Lines 12 and 13 lies in determining whether the current process should compute the force resulting from the interaction of a particle q assigned to it and a particle r whose position it has received. If we recall the reduced algorithm (Program 6.1), we see that task/particle q is responsible for computing \mathbf{f}_{qr} if and only if q < r. However, the arrays loc_pos and tmp_pos (or a larger array containing tmp_pos and tmp_forces) use local subscripts, not global subscripts. That is, when we access an element of (say) loc_pos, the subscript we use will lie in the range $0, 1, \ldots, loc_n - 1$, not $0, 1, \ldots, n - 1$; so, if we try to implement the force interaction with the following pseudocode, we'll run into (at least) a couple of problems:

```
for (loc_part1 = 0; loc_part1 < loc_n-1; loc_part1++)
   for (loc_part2 = loc_part1+1; loc_part2 < loc_n; loc_part2++)</pre>
```

The first, and most obvious, is that masses is a global array and we're using local subscripts to access its elements. The second is that the relative sizes of loc_part1 and loc_part2 don't tell us whether we should compute the force due to their interaction. We need to use global subscripts to determine this. For example, if we have four particles and two processes, and the preceding code is being run by process 0, then when loc_part1 = 0, the inner loop will skip loc_part2 = 0 and start with loc_part2 = 1; however, if we're using a cyclic distribution, loc_part1 = 0 corresponds to global particle 0 and loc_part2 = 0 corresponds to global particle 1, and we *should* compute the force resulting from interaction between these two particles.

Clearly, the problem is that we shouldn't be using local particle indexes, but rather we should be using *global* particle indexes. Thus, using a cyclic distribution of the particles, we could modify our code so that the loops also iterate through global particle indexes:

The function First_index should determine a global index glb_part2 with the following properties:

- 1. The particle glb_part2 is assigned to the process with rank owner.
- **2.** glb_part1 < glb_part2 < glb_part1 + comm_sz.

The function Global_to_local should convert a global particle index into a local particle index, and the function Compute_force should compute the force resulting from the interaction of two particles. We already know how to implement Compute_force. See Exercises 6.15 and 6.16 for the other two functions.

6.1.11 Performance of the MPI solvers

Table 6.5 shows the run-times of the two *n*-body solvers when they're run with 800 particles for 1000 timesteps on an Infiniband-connected cluster. All the timings were taken with one process per cluster node. The run-times of the serial solvers differed from the single-process MPI solvers by less than 1%, so we haven't included them.

Clearly, the performance of the reduced solver is much superior to the performance of the basic solver, although the basic solver achieves higher efficiencies.

Table 6.5 Performance of the MPI <i>n</i> -Body Solvers (times in seconds)						
Processes Basic Reduced						
1	17.30	8.68				
2	8.65	4.45				
4	4.35	2.30				
8	2.20	1.26				
16	1.13	0.78				

Table 6.6 Run-Times for OpenMP and MPI *n*-Body Solvers (times in seconds)

Processes/	Ор	enMP	MPI		
Threads	Basic	Reduced	Basic	Reduced	
1	15.13	8.77	17.30	8.68	
2	7.62	4.42	8.65	4.45	
4	3.85	2.26	4.35	2.30	

For example, the efficiency of the basic solver on 16 nodes is about 0.95, while the efficiency of the reduced solver on 16 nodes is only about 0.70.

A point to stress here is that the reduced MPI solver makes much more efficient use of memory than the basic MPI solver; the basic solver must provide storage for all n positions on each process, while the reduced solver only needs extra storage for n/comm_sz positions and n/comm_sz forces. Thus, the extra storage needed on each process for the basic solver is nearly comm_sz/2 times greater than the storage needed for the reduced solver. When n and comm_sz are very large, this factor can easily make the difference between being able to run a simulation only using the process' main memory and having to use secondary storage.

The nodes of the cluster on which we took the timings have four cores, so we can compare the performance of the OpenMP implementations with the performance of the MPI implementations (see Table 6.6). We see that the basic OpenMP solver is a good deal faster than the basic MPI solver. This isn't surprising since MPI_Allgather is such an expensive operation. Perhaps surprisingly, though, the reduced MPI solver is quite competitive with the reduced OpenMP solver.

Let's take a brief look at the amount of memory required by the MPI and OpenMP reduced solvers. Say that there are n particles and p threads or processes. Then each solver will allocate the same amount of storage for the local velocities and the local positions. The MPI solver allocates n doubles per process for the masses. It also allocates 4n/p doubles for the tmp_pos and tmp_forces arrays, so in addition to the

local velocities and positions, the MPI solver stores

$$n + 4n/p$$

doubles per process. The OpenMP solver allocates a total of 2pn + 2n doubles for the forces and n doubles for the masses, so in addition to the local velocities and positions, the OpenMP solver stores

$$3n/p + 2n$$

doubles per thread. Thus, the difference in the local storage required for the OpenMP version and the MPI version is

$$n - n/p$$

doubles. In other words, if *n* is large, the local storage required for the MPI version is substantially less than the local storage required for the OpenMP version. So, for a fixed number of processes or threads, we should be able to run much larger simulations with the MPI version than the OpenMP version. Of course, because of hardware considerations, we're likely to be able to use many more MPI processes than OpenMP threads, so the size of the largest possible MPI simulations should be *much* greater than the size of the largest possible OpenMP simulations. The MPI version of the reduced solver is much more scalable than any of the other versions, and the "ring pass" algorithm provides a genuine breakthrough in the design of *n*-body solvers.

6.2 TREE SEARCH

Many problems can be solved using a tree search. As a simple example, consider the traveling salesperson problem, or TSP. In TSP, a salesperson is given a list of cities she needs to visit and a cost for traveling between each pair of cities. Her problem is to visit each city once, returning to her hometown, and she must do this with the least possible cost. A route that starts in her hometown, visits each city once and returns to her hometown is called a *tour*; thus, the TSP is to find a minimum-cost tour.

Unfortunately, TSP is what's known as an **NP-complete** problem. From a practical standpoint, this means that there is no algorithm known for solving it that, in all cases, is significantly better than exhaustive search. Exhaustive search means examining all possible solutions to the problem and choosing the best. The number of possible solutions to TSP grows exponentially as the number of cities is increased. For example, if we add one additional city to an n-city problem, we'll increase the number of possible solutions by a factor of n-1. Thus, although there are only six possible solutions to a four-city problem, there are $4 \times 6 = 24$ to a five-city problem, $5 \times 24 = 120$ to a six-city problem, $6 \times 120 = 720$ to a seven-city problem, and so on. In fact, a 100-city problem has far more possible solutions than the number of atoms in the universe!

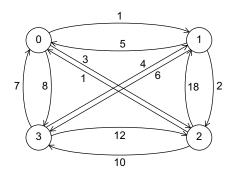


FIGURE 6.9

A four-city TSP

Furthermore, if we could find a solution to TSP that's significantly better in all cases than exhaustive search, then there are literally hundreds of other very hard problems for which we could find fast solutions. Not only is there no known solution to TSP that is better in all cases than exhaustive search, it's very unlikely that we'll find one.

So how can we solve TSP? There are a number of clever solutions. However, let's take a look at an especially simple one. It's a very simple form of tree search. The idea is that in searching for solutions, we build a *tree*. The leaves of the tree correspond to tours, and the other tree nodes correspond to "partial" tours—routes that have visited some, but not all, of the cities.

Each node of the tree has an associated cost, that is, the cost of the partial tour. We can use this to eliminate some nodes of the tree. Thus, we want to keep track of the cost of the best tour so far, and, if we find a partial tour or node of the tree that couldn't possibly lead to a less expensive complete tour, we shouldn't bother searching the children of that node (see Figures 6.9 and 6.10).

In Figure 6.9 we've represented a four-city TSP as a labeled, directed graph. A **graph** (not to be confused with a graph in calculus) is a collection of vertices and edges or line segments joining pairs of vertices. In a **directed graph** or **digraph**, the edges are oriented—one end of each edge is the tail, and the other is the head. A graph or digraph is **labeled** if the vertices and/or edges have labels. In our example, the vertices of the digraph correspond to the cities in an instance of the TSP, the edges correspond to routes between the cities, and the labels on the edges correspond to the costs of the routes. For example, there's a cost of 1 to go from city 0 to city 1 and a cost of 5 to go from city 1 to city 0.

If we choose vertex 0 as the salesperson's home city, then the initial partial tour consists only of vertex 0, and since we've gone nowhere, it's cost is 0. Thus, the root of the tree in Figure 6.10 has the partial tour consisting only of the vertex 0 with cost 0. From 0 we can first visit 1, 2, or 3, giving us three two-city partial tours with costs 1, 3, and 8, respectively. In Figure 6.10 this gives us three children of the root. Continuing, we get six three-city partial tours, and since there are

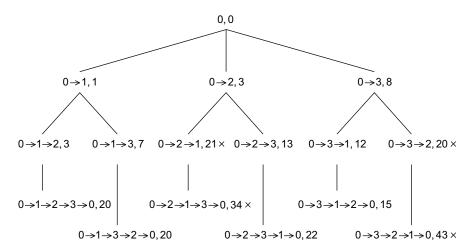


FIGURE 6.10

Search tree for four-city TSP

only four cities, once we've chosen three of the cities, we know what the complete tour is.

Now, to find a least-cost tour, we should search the tree. There are many ways to do this, but one of the most commonly used is called **depth-first search**. In depth-first search, we probe as deeply as we can into the tree. After we've either reached a leaf or found a tree node that can't possibly lead to a least-cost tour, we back up to the deepest "ancestor" tree node with unvisited children, and probe one of its children as deeply as possible.

In our example, we'll start at the root, and branch left until we reach the leaf labeled

$$0 \rightarrow 1 \rightarrow 2 \rightarrow 3 \rightarrow 0$$
, Cost 20.

Then we back up to the tree node labeled $0 \rightarrow 1$, since it is the deepest ancestor node with unvisited children, and we'll branch down to get to the leaf labeled

$$0 \rightarrow 1 \rightarrow 3 \rightarrow 2 \rightarrow 0$$
, Cost 20.

Continuing, we'll back up to the root and branch down to the node labeled $0 \rightarrow 2$. When we visit its child, labeled

$$0 \rightarrow 2 \rightarrow 1$$
, Cost 21,

we'll go no further in this subtree, since we've already found a complete tour with cost less than 21. We'll back up to $0 \rightarrow 2$ and branch down to its remaining unvisited child. Continuing in this fashion, we eventually find the least-cost tour

$$0 \rightarrow 3 \rightarrow 1 \rightarrow 2 \rightarrow 0$$
, Cost 15.

6.2.1 Recursive depth-first search

Using depth-first search we can systematically visit each node of the tree that could possibly lead to a least-cost solution. The simplest formulation of depth-first search uses recursion (see Program 6.4). Later on it will be useful to have a definite order in which the cities are visited in the for loop in Lines 8 to 13, so we'll assume that the cities are visited in order of increasing index, from city 1 to city n-1.

The algorithm makes use of several global variables:

- n: the total number of cities in the problem
- digraph: a data structure representing the input digraph
- hometown: a data structure representing vertex or city 0, the salesperson's hometown
- best_tour: a data structure representing the best tour so far

The function City_count examines the partial tour to see if there are n cities on the partial tour. If there are, we know that we simply need to return to the hometown to complete the tour, and we can check to see if the complete tour has a lower cost than the current "best tour" by calling Best_tour. If it does, we can replace the current best tour with this tour by calling the function Update_best_tour. Note that before the first call to Depth_first_search, the best_tour variable should be initialized so that its cost is greater than the cost of any possible least-cost tour.

If the partial tour tour hasn't visited n cities, we can continue branching down in the tree by "expanding the current node," in other words, by trying to visit other cities from the city last visited in the partial tour. To do this we simply loop through the cities. The function Feasible checks to see if the city or vertex has already been visited, and, if not, whether it can possibly lead to a least-cost tour. If the city is feasible, we add it to the tour, and recursively call Depth_first_search. When

```
void Depth_first_search(tour_t tour) {
      city_t city;
2
3
4
       if (City_count(tour) == n) {
5
          if (Best_tour(tour))
             Update_best_tour(tour);
6
7
       } else {
8
          for each neighboring city
9
             if (Feasible(tour, city)) {
10
                Add_city(tour, city);
11
                Depth_first_search(tour);
12
                Remove_last_city(tour, city);
13
14
       /* Depth_first_search */
15
   }
```

Program 6.4: Pseudocode for a recursive solution to TSP using depth-first search

we return from Depth_first_search, we remove the city from the tour, since it shouldn't be included in the tour used in subsequent recursive calls.

6.2.2 Nonrecursive depth-first search

Since function calls are expensive, recursion can be slow. It also has the disadvantage that at any given instant of time only the current tree node is accessible. This could be a problem when we try to parallelize tree search by dividing tree nodes among the threads or processes.

It is possible to write a nonrecursive depth-first search. The basic idea is modeled on recursive implementation. Recall that recursive function calls can be implemented by pushing the current state of the recursive function onto the run-time stack. Thus, we can try to eliminate recursion by pushing necessary data on our own stack before branching deeper into the tree, and when we need to go back up the tree—either because we've reached a leaf or because we've found a node that can't lead to a better solution—we can pop the stack.

This outline leads to the implementation of iterative depth-first search shown in Program 6.5. In this version, a stack record consists of a single city, the city that will be added to the tour when its record is popped. In the recursive version we continue to make recursive calls until we've visited every node of the tree that corresponds to a feasible partial tour. At this point, the stack won't have any more activation records for calls to Depth_first_search, and we'll return to the function that made the

```
for (city = n-1; city >= 1; city—)
2
      Push(stack, city);
3
   while (!Empty(stack)) {
4
      city = Pop(stack);
5
       if (city == NO_CITY) // End of child list, back up
          Remove_last_city(curr_tour);
6
7
8
          Add_city(curr_tour, city);
          if (City_count(curr_tour) == n) {
9
10
             if (Best_tour(curr_tour))
11
                Update_best_tour(curr_tour);
12
             Remove_last_city(curr_tour);
13
          } else {
14
             Push(stack, NO_CITY);
15
             for (nbr = n-1; nbr >= 1; nbr---)
16
                if (Feasible(curr_tour, nbr))
17
                   Push(stack, nbr);
18
19
          /* if Feasible */
       /* while !Empty */
```

Program 6.5: Pseudocode for an implementation of a depth-first solution to TSP that doesn't use recursion

original call to Depth_first_search. The main control structure in our iterative version is the while loop extending from Line 3 to Line 20, and the loop termination condition is that our stack is empty. As long as the search needs to continue, we need to make sure the stack is nonempty, and, in the first two lines, we add each of the non-hometown cities. Note that this loop visits the cities in decreasing order, from n-1 down to 1. This is because of the order created by the stack, whereby the stack pops the top cities first. By reversing the order, we can insure that the cities are visited in the same order as the recursive function.

Also notice that in Line 5 we check whether the city we've popped is the constant NO_CITY. This constant is used so that we can tell when we've visited all of the children of a tree node; if we didn't use it, we wouldn't be able to tell when to back up in the tree. Thus, before pushing all of the children of a node (Lines 15–17), we push the NO_CITY marker.

An alternative to this iterative version uses partial tours as stack records (see Program 6.6). This gives code that is closer to the recursive function. However, it also results in a slower version, since it's necessary for the function that pushes onto the stack to create a copy of the tour before actually pushing it on to the stack. To emphasize this point, we've called the function Push_copy. (What happens if we simply push a pointer to the current tour onto the stack?) The extra memory required will probably not be a problem. However, allocating storage for a new tour and copying the existing tour is time-consuming. To some degree we can mitigate these costs by saving freed tours in our own data structure, and when a freed tour is available we can use it in the Push_copy function instead of calling malloc.

On the other hand, this version has the virtue that the stack is more or less independent of the other data structures. Since entire tours are stored, multiple threads or processes can "help themselves" to tours, and, if this is done reasonably carefully,

```
Push_copy(stack, tour); // Tour that visits only the hometown
2
   while (!Emptv(stack)) {
       curr_tour = Pop(stack);
3
       if (City\_count(curr\_tour) == n) {
4
5
          if (Best_tour(curr_tour))
             Update_best_tour(curr_tour);
6
7
       } else {
          for (nbr = n-1; nbr >= 1; nbr--)
Q
             if (Feasible(curr_tour, nbr)) {
10
                Add_city(curr_tour, nbr);
11
                Push_copy(stack, curr_tour);
                Remove_last_city(curr_tour);
12
13
14
       Free_tour(curr_tour);
15
16
   }
```

Program 6.6: Pseudocode for a second solution to TSP that doesn't use recursion

it won't destroy the correctness of the program. With the original iterative version, a stack record is just a city and it doesn't provide enough information by itself to show where we are in the tree.

6.2.3 Data structures for the serial implementations

Our principal data structures are the tour, the digraph, and, in the iterative implementations, the stack. The tour and the stack are essentially list structures. In problems that we're likely to be able to tackle, the number of cities is going to be small—certainly less than 100—so there's no great advantage to using a linked list to represent the tours and we've used an array that can store n+1 cities. We repeatedly need both the number of cities in the partial tour and the cost of the partial tour. Therefore, rather than just using an array for the tour data structure and recomputing these values, we use a struct with three members: the array storing the cities, the number of cities, and the cost of the partial tour.

To improve the readability and the performance of the code, we can use preprocessor macros to access the members of the struct. However, since macros can be a nightmare to debug, it's a good idea to write "accessor" functions for use during initial development. When the program with accessor functions is working, they can be replaced with macros. As an example, we might start with the function

```
/* Find the ith city on the partial tour */
int Tour_city(tour_t tour, int i) {
   return tour—>cities[i];
} /* Tour_city */
```

When the program is working, we could replace this with the macro

```
/* Find the ith city on the partial tour */
#define Tour_city(tour, i) (tour->cities[i])
```

The stack in the original iterative version is just a list of cities or ints. Furthermore, since there can't be more than $n^2/2$ records on the stack (see Exercise 6.17) at any one time, and n is likely to be small, we can just use an array, and like the tour data structure, we can store the number of elements on the stack. Thus, for example, Push can be implemented with

```
void Push(my_stack_t stack, int city) {
  int loc = stack->list_sz;
  stack->list[loc] = city;
  stack->list_sz++;
} /* Push */
```

In the second iterative version, the version that stores entire tours in the stack, we can probably still use an array to store the tours on the stack. Now the push function will look something like this:

```
void Push_copy(my_stack_t stack, tour_t tour) {
  int loc = stack->list_sz;
```

```
tour_t tmp = Alloc_tour();
Copy_tour(tour, tmp);
stack->list[loc] = tmp;
stack->list_sz++;
/* Push */
```

Once again, element access for the stack can be implemented with macros.

There are many possible representations for digraphs. When the digraph has relatively few edges, list representations are preferred. However, in our setting, if vertex i is different from vertex j, there are directed, weighted edges from i to j and from j to i, so we need to store a weight for each ordered pair of distinct vertices i and j. Thus, in our setting, an **adjacency matrix** is almost certainly preferable to a list structure. This is an $n \times n$ matrix, in which the weight of the edge from vertex i to vertex j can be the entry in the ith row and jth column of the matrix. We can access this weight directly, without having to traverse a list. The diagonal elements (row i and column i) aren't used, and we'll set them to 0.

6.2.4 Performance of the serial implementations

The run-times of the three serial implementations are shown in Table 6.7. The input digraph contained 15 vertices (including the hometown), and all three algorithms visited approximately 95,000,000 tree nodes. The first iterative version is less than 5% faster than the recursive version, and the second iterative version is about 8% slower than the recursive version. As expected, the first iterative solution eliminates some of the overhead due to repeated function calls, while the second iterative solution is slower because of the repeated copying of tour data structures. However, as we'll see, the second iterative solution is relatively easy to parallelize, so we'll be using it as the basis for the parallel versions of tree search.

6.2.5 Parallelizing tree search

Let's take a look at parallelizing tree search. The tree structure suggests that we identify tasks with tree nodes. If we do this, the tasks will communicate down the tree edges: a parent will communicate a new partial tour to a child, but a child, except for terminating, doesn't communicate directly with a parent.

We also need to take into consideration the updating and use of the best tour. Each task examines the best tour to determine whether the current partial tour is feasible or the current complete tour has lower cost. If a leaf task determines its tour is a better tour, then it will also update the best tour. Although all of the actual computation can

Table 6.7 Run-Times of the Three SerialImplementations of Tree Search (times in seconds)						
Recursive First Iterative Second Iterative						
30.5 29.2 32.9						

be considered to be carried out by the tree node tasks, we need to keep in mind that the best tour data structure requires additional communication that is not explicit in the tree edges. Thus, it's convenient to add an additional task that corresponds to the best tour. It "sends" data to every tree node task, and receives data from some of the leaves. This latter view is convenient for shared-memory, but not so convenient for distributed-memory.

A natural way to agglomerate and map the tasks is to assign a subtree to each thread or process, and have each thread/process carry out all the tasks in its subtree. For example, if we have three threads or processes, as shown earlier in Figure 6.10, we might map the subtree rooted at $0 \rightarrow 1$ to thread/process 0, the subtree rooted at $0 \rightarrow 2$ to thread/process 1, and the subtree rooted at $0 \rightarrow 3$ to thread/process 2.

Mapping details

There are many possible algorithms for identifying which subtrees we assign to the processes or threads. For example, one thread or process could run the last version of serial depth-first search until the stack stores one partial tour for each thread or process. Then it could assign one tour to each thread or process. The problem with depth-first search is that we expect a subtree whose root is deeper in the tree to require less work than a subtree whose root is higher up in the tree, so we would probably get better load balance if we used something like **breadth-first search** to identify the subtrees.

As the name suggests, breadth-first search searches as widely as possible in the tree before going deeper. So if, for example, we carry out a breadth-first search until we reach a level of the tree that has at least thread_count or comm_sz nodes, we can then divide the nodes at this level among the threads or processes. See Exercise 6.18 for implementation details.

The best tour data structure

On a shared-memory system, the best tour data structure can be shared. In this setting, the Feasible function can simply examine the data structure. However, updates to the best tour will cause a race condition, and we'll need some sort of locking to prevent errors. We'll discuss this in more detail when we implement the parallel version.

In the case of a distributed-memory system, there are a couple of choices that we need to make about the best tour. The simplest option would be to have the processes operate independently of each other until they have completed searching their subtrees. In this setting, each process would store its own *local* best tour. This local best tour would be used by the process in Feasible and updated by the process each time it calls <code>Update_best_tour</code>. When all the processes have finished searching, they can perform a global reduction to find the tour with the *global* least cost.

This approach has the virtue of simplicity, but it also suffers from the problem that it's entirely possible for a process to spend most or all of its time searching through partial tours that couldn't possibly lead to a global best tour. Thus, we should

probably try using an approach that makes the current global best tour available to all the processes. We'll take a look at details when we discuss the MPI implementation.

Dynamic mapping of tasks

A second issue we should consider is the problem of load imbalance. Although the use of breadth-first search ensures that all of our subtrees have approximately the same number of nodes, there is no guarantee that they all have the same amount of work. It's entirely possible that one process or thread will have a subtree consisting of very expensive tours, and, as a consequence, it won't need to search very deeply into its assigned subtree. However, with our current, *static* mapping of tasks to threads/processes, this one thread or process will simply have to wait until the other threads/processes are done.

An alternative is to implement a **dynamic** mapping scheme. In a dynamic scheme, if one thread/process runs out of useful work, it can obtain additional work from another thread/process. In our final implementation of serial depth-first search, each stack record contains a partial tour. With this data structure a thread or process can give additional work to another thread/process by dividing the contents of its stack. This might at first seem to have the potential for causing problems with the program's correctness, since if we give part of one thread's or one process' stack to another, there's a good chance that the order in which the tree nodes will be visited will be changed.

However, we're already going to do this; when we assign different subtrees to different threads/processes, the order in which the tree nodes are visited is no longer the serial depth-first ordering. In fact, in principle, there's no reason to visit any node before any other node as long as we make sure we visit "ancestors" before "descendants." But this isn't a problem since a partial tour isn't added to the stack until after all its ancestors have been visited. For example, in Figure 6.10 the node consisting of the tour $0 \to 2 \to 1$ will be pushed onto the stack when the node consisting of the tour $0 \to 2$ is the currently active node, and consequently the two nodes won't be on the stack simultaneously. Similarly, the parent of $0 \to 2$, the root of the tree, 0, is no longer on the stack when $0 \to 2$ is visited.

A second alternative for dynamic load balancing—at least in the case of shared memory—would be to have a shared stack. However, we couldn't simply dispense with the local stacks. If a thread needed to access the shared stack every time it pushed or popped, there would be a tremendous amount of contention for the shared stack and the performance of the program would probably be worse than a serial program. This is exactly what happened when we parallelized the reduced *n*-body solver with mutexes/locks protecting the calculations of the total forces on the various particles. If every call to Push or Pop formed a critical section, our program would grind to nearly a complete halt. Thus, we would want to retain local stacks for each thread, with only occasional accesses to the shared stack. We won't pursue this alternative. See Programming Assignment 6.7 for further details.

6.2.6 A static parallelization of tree search using pthreads

In our static parallelization, a single thread uses breadth-first search to generate enough partial tours so that each thread gets at least one partial tour. Then each thread takes its partial tours and runs iterative tree search on them. We can use the pseudocode shown in Program 6.7 on each thread. Note that most of the function calls—for example, Best_tour, Feasible, Add_city—need to access the adjacency matrix representing the digraph, so all the threads will need to access the digraph. However, since these are only *read* accesses, this won't result in a race condition or contention among the threads.

There are only four potential differences between this pseudocode and the pseudocode we used for the second iterative serial implementation:

- The use of my_stack instead of stack; since each thread has its own, private stack, we use my_stack as the identifier for the stack object instead of stack.
- Initialization of the stack.
- Implementation of the Best_tour function.
- Implementation of the Update_best_tour function.

In the serial implementation, the stack is initialized by pushing the partial tour consisting only of the hometown onto the stack. In the parallel version we need to generate at least thread_count partial tours to distribute among the threads. As we discussed earlier, we can use breadth-first search to generate a list of at least thread_count tours by having a single thread search the tree until it reaches a level with at least thread_count tours. (Note that this implies that the number of threads should be less than (n-1)!, which shouldn't be a problem). Then the threads can

```
Partition_tree(my_rank, my_stack);
while (!Empty(my_stack)) {
    curr_tour = Pop(my_stack);
    if (City_count(curr_tour) == n) {
        if (Best_tour(curr_tour)) Update_best_tour(curr_tour);
    } else {
        for (city = n-1; city >= 1; city—)
            if (Feasible(curr_tour, city)) {
                Add_city(curr_tour, city);
                Push_copy(my_stack, curr_tour);
                 Remove_last_city(curr_tour)
            }
    }
    Free_tour(curr_tour);
}
```

Program 6.7: Pseudocode for a Pthreads implementation of a statically parallelized solution to TSP

use a block partition to divide these tours among themselves and push them onto their private stacks. Exercise 6.18 looks into the details.

To implement the Best_tour function, a thread should compare the cost of its current tour with the cost of the global best tour. Since multiple threads may be simultaneously accessing the global best cost, it might at first seem that there will be a race condition. However, the Best_tour function only reads the global best cost, so there won't be any conflict with threads that are also checking the best cost. If a thread is updating the global best cost, then a thread that is just checking it will either read the old value or the new, updated value. While we would prefer that it get the new value, we can't insure this without using some very costly locking strategy. For example, threads wanting to execute Best_tour or Update_best_tour could wait on a single mutex. This would insure that no thread is updating while another thread is only checking, but would have the unfortunate side effect that only one thread could check the best cost at a time. We could improve on this by using a read-write lock, but this would have the side effect that the readers—the threads calling Best_tour—would all block while a thread updated the best tour. In principle, this doesn't sound too bad, but recall that in practice read-write locks can be quite slow. So it seems pretty clear that the "no contention" solution of possibly getting a best tour cost that's out-of-date is probably better, as the next time the thread calls Best_tour, it will get the updated value of the best tour cost.

On the other hand, we call <code>Update_best_tour</code> with the intention of writing to the best tour structure, and this clearly can cause a race condition if two threads call it simultaneously. To avoid this problem, we can protect the body of the <code>Update_best_tour</code> function with a mutex. This isn't enough, however; between the time a thread completes the test in <code>Best_tour</code> and the time it obtains the lock in <code>Update_best_tour</code>, another thread may have obtained the lock and updated the best tour cost, which now may be less than the best tour cost that the first thread found in <code>Best_tour</code>. Thus, correct pseudocode for <code>Update_best_tour</code> should look something like this:

```
pthread_mutex_lock(best_tour_mutex);
/* We've already checked Best_tour, but we need to check it
    again */
if (Best_tour(tour))
    Replace old best tour with tour;
pthread_mutex_unlock(best_tour_mutex).
```

This may seem wasteful, but if updates to the best tour are infrequent, then most of the time Best_tour will return false and it will only be rarely necessary to make the "double" call.

6.2.7 A dynamic parallelization of tree search using pthreads

If the initial distribution of subtrees doesn't do a good job of distributing the work among the threads, the static parallelization provides no means of redistributing work. The threads with "small" subtrees will finish early, while the threads with large subtrees will continue to work. It's not difficult to imagine that one thread gets the lion's

share of the work because the edges in its initial tours are very cheap, while the edges in the other threads' initial tours are very expensive. To address this issue, we can try to dynamically redistribute the work as the computation proceeds.

To do this, we can replace the test !Empty(my_stack) controlling execution of the while loop with more complex code. The basic idea is that when a thread runs out of work—that is, !Empty(my_stack) becomes false—instead of immediately exiting the while loop, the thread waits to see if another thread can provide more work. On the other hand, if a thread that still has work in its stack finds that there is at least one thread without work, and its stack has at least two tours, it can "split" its stack and provide work for one of the threads.

Pthreads condition variables provide a natural way to implement this. When a thread runs out of work it can call pthread_cond_wait and go to sleep. When a thread with work finds that there is at least one thread waiting for work, after splitting its stack, it can call pthread_cond_signal. When a thread is awakened it can take one of the halves of the split stack and return to work.

This idea can be extended to handle termination. If we maintain a count of the number of threads that are in pthread_cond_wait, then when a thread whose stack is empty finds that thread_count -1 threads are already waiting, it can call pthread_cond_broadcast and as the threads awaken, they'll see that all the threads have run out of work and quit.

Termination

Thus, we can use the pseudocode shown in Program 6.8 for a Terminated function that would be used instead of Empty for the while loop implementing tree search.

There are several details that we should look at more closely. Notice that the code executed by a thread before it splits its stack is fairly complicated. In Lines 1–2 the thread

- checks that it has at least two tours in its stack,
- · checks that there are threads waiting, and
- checks whether the new_stack variable is NULL.

The reason for the check that the thread has enough work should be clear: if there are fewer than two records on the thread's stack, "splitting" the stack will either do nothing or result in the active thread's trading places with one of the waiting threads.

It should also be clear that there's no point in splitting the stack if there aren't any threads waiting for work. Finally, if some thread has already split its stack, but a waiting thread hasn't retrieved the new stack, that is, new_stack != NULL, then it would be disastrous to split a stack and overwrite the existing new stack. Note that this makes it essential that after a thread retrieves new_stack by, say, copying new_stack into its private my_stack variable, the thread must set new_stack to NULL.

If all three of these conditions hold, then we can try splitting our stack. We can acquire the mutex that protects access to the objects controlling termination (threads_in_cond_wait, new_stack, and the condition variable). However, the condition

```
threads_in_cond_wait > 0 && new_stack == NULL
```

```
if (my_stack_size >= 2 && threads_in_cond_wait > 0 &&
          new_stack == NULL) {
3
       lock term_mutex;
       if (threads_in_cond_wait > 0 && new_stack == NULL) {
4
5
          Split my_stack creating new_stack;
          pthread_cond_signal(&term_cond_var);
6
7
8
       unlock term_mutex;
9
       return 0; /* Terminated = false; don't quit */
10
   } else if (!Empty(my_stack)) /* Keep working */
       return 0; /* Terminated = false; don't quit */
11
    } else { /* My stack is empty */
12
       lock term_mutex;
13
14
       if (threads_in_cond_wait == thread_count-1)
15
                                            /* Last thread running */
16
          threads_in_cond_wait++;
17
          pthread_cond_broadcast(&term_cond_var);
18
          unlock term_mutex;
19
          return 1; /* Terminated = true; quit */
20
       } else { /* Other threads still working, wait for work */
2.1
          threads_in_cond_wait++;
22
          while (pthread_cond_wait(&term_cond_var, &term_mutex) != 0);
23
          /* We've been awakened */
          if (threads_in_cond_wait < thread_count) { /* We got work */</pre>
24
25
             my_stack = new_stack;
26
             new_stack = NULL;
27
             threads_in_cond_wait—;
28
             unlock term_mutex;
29
             return 0; /* Terminated = false */
          } else { /* All threads done */
30
31
             unlock term_mutex;
             return 1; /* Terminated = true; quit */
32
33
34
          /* else wait for work */
35
      /* else my_stack is empty */
```

Program 6.8: Pseudocode for Pthreads Terminated function

can change between the time we start waiting for the mutex and the time we actually acquire it, so as with <code>Update_best_tour</code>, we need to confirm that this condition is still true after acquiring the mutex (Line 4). Once we've verified that these conditions still hold, we can split the stack, awaken one of the waiting threads, unlock the mutex, and return to work.

If the test in Lines 1 and 2 is false, we can check to see if we have any work at all—that is, our stack is nonempty. If it is, we return to work. If it isn't, we'll start the termination sequence by waiting for and acquiring the termination mutex in Line 13. Once we've acquired the mutex, there are two possibilities:

- We're the last thread to enter the termination sequence, that is, threads_in_cond_wait == thread_count-1.
- · Other threads are still working.

In the first case, we know that since all the other threads have run out of work, and we have also run out of work, the tree search should terminate. We therefore signal all the other threads by calling pthread_cond_broadcast and returning true. Before executing the broadcast, we increment threads_in_cond_wait, even though the broadcast is telling all the threads to return from the condition wait. The reason is that threads_in_cond_wait is serving a dual purpose: When it's less than thread_count, it tells us how many threads are waiting. However, when it's equal to thread_count, it tells us that all the threads are out of work, and it's time to quit.

In the second case—other threads are still working—we call pthread_cond_wait (Line 22) and wait to be awakened. Recall that it's possible that a thread could be awakened by some event other than a call to pthread_cond_signal or pthread_cond_broadcast. So, as usual, we put the call to pthread_cond_wait in a while loop, which will immediately call pthread_cond_wait again if some other event (return value not 0) awakens the thread.

Once we've been awakened, there are also two cases to consider:

- threads_in_cond_wait < thread_count
- threads_in_cond_wait == thread_count

In the first case, we know that some other thread has split its stack and created more work. We therefore copy the newly created stack into our private stack, set the new_stack variable to NULL, and decrement threads_in_cond_wait (i.e., Lines 25–27). Recall that when a thread returns from a condition wait, it obtains the mutex associated with the condition variable, so before returning, we also unlock the mutex (i.e., Line 28). In the second case, there's no work left, so we unlock the mutex and return true.

In the actual code, we found it convenient to group the termination variables together into a single struct. Thus, we defined something like

```
typedef struct {
   my_stack_t new_stack;
   int threads_in_cond_wait;
   pthread_cond_t term_cond_var;
   pthread_mutex_t term_mutex;
} term_struct;
typedef term_struct* term_t;
```

and we defined a couple of functions, one for initializing the term variable and one for destroying/freeing the variable and its members.

Before discussing the function that splits the stack, note that it's possible that a thread with work can spend a lot of time waiting for term_mutex before being able

to split its stack. Other threads may be either trying to split their stacks, or preparing for the condition wait. If we suspect that this is a problem, Pthreads provides a nonblocking alternative to pthread_mutex_lock called pthread_mutex_trylock:

This function attempts to acquire mutex_p. However, if it's locked, instead of waiting, it returns immediately. The return value will be zero if the calling thread has successfully acquired the mutex, and nonzero if it hasn't. As an alternative to waiting on the mutex before splitting its stack, a thread can call pthread_mutex_trylock. If it acquires term_mutex, it can proceed as before. If not, it can just return. Presumably on a subsequent call it can successfully acquire the mutex.

Splitting the stack

Since our goal is to balance the load among the threads, we would like to insure that the amount of work in the new stack is roughly the same as the amount remaining in the original stack. We have no way of knowing in advance of searching the subtree rooted at a partial tour how much work is actually associated with the partial tour, so we'll never be able to guarantee an equal division of work. However, we can use the same strategy that we used in our original assignment of subtrees to threads: that the subtrees rooted at two partial tours with the same number of cities have identical structures. Since on average two partial tours with the same number of cities are equally likely to lead to a "good" tour (and hence more work), we can try splitting the stack by assigning the tours on the stack on the basis of their numbers of edges. The tour with the least number of edges remains on the original stack, the tour with the next to the least number of edges goes to the new stack, the tour with the next number of edges remains on the original, and so on.

This is fairly simple to implement, since the tours on the stack have an increasing number of edges. That is, as we proceed from the bottom of the stack to the top of the stack, the number of edges in the tours increases. This is because when we push a new partial tour with k edges onto the stack, the tour that's immediately "beneath" it on the stack either has k edges or k-1 edges. We can implement the split by starting at the bottom of the stack, and alternately leaving partial tours on the old stack and pushing partial tours onto the new stack, so tour 0 will stay on the old stack, tour 1 will go to the new stack, tour 2 will stay on the old stack, and so on. If the stack is implemented as an array of tours, this scheme will require that the old stack be "compressed" so that the gaps left by removing alternate tours are eliminated. If the stack is implemented as a linked list of tours, compression won't be necessary.

This scheme can be further refined by observing that partial tours with lots of cities won't provide much work, since the subtrees that are rooted at these trees are very small. We could add a "cutoff size" and not reassign a tour unless its number of cities was less than the cutoff. In a shared-memory setting with an array-based stack, reassigning a tour when a stack is split won't increase the cost of the split, since the tour (which is a pointer) will either have to be copied to the new stack or a new

Table 6.8 Run-Times of Pthreads Tree-Search Programs (times in seconds)								
	First Problem Second Problem							
Threads	Serial	Static	Dynamic		Serial	Static	Dynamic	
1 2 4 8	32.9	32.7 27.9 25.7 23.8	34.7 28.9 25.9 22.4	(0) (7) (47) (180)	26.0	25.8 25.8 25.8 24.0	27.5 19.2 9.3 5.7	(0) (6) (49) (256)

location in the old stack. We'll defer exploration of this alternative to Programming Assignment 6.6.

6.2.8 Evaluating the Pthreads tree-search programs

Table 6.8 shows the performance of the two Pthreads programs on two fifteen-city problems. The "Serial" column gives the run-time of the second iterative solution—the solution that pushes a copy of each new tour onto the stack. For reference, the first problem in Table 6.8 is the same as the problem the three serial solutions were tested with in Table 6.7, and both the Pthreads and serial implementations were tested on the same system. Run-times are in seconds, and the numbers in parentheses next to the run-times of the program that uses dynamic partitioning give the total number of times the stacks were split.

From these numbers, it's apparent that different problems can result in radically different behaviors. For example, the program that uses static partitioning generally performs better on the first problem than the program that uses dynamic partitioning. However, on the second problem, the performance of the static program is essentially independent of the number of threads, while the dynamic program obtains excellent performance. In general, it appears that the dynamic program is more scalable than the static program.

As we increase the number of threads, we would expect that the size of the local stacks will decrease, and hence threads will run out of work more often. When threads are waiting, other threads will split their stacks, so as the number of threads is increased, the total number of stack splits should increase. Both problems confirm this prediction.

It should be noted that if the input problem has more than one possible solution—that is, different tours with the same minimum cost—then the results of both of the programs are nondeterministic. In the static program, the sequence of best tours depends on the speed of the threads, and this sequence determines which tree nodes are examined. In the dynamic program, we also have nondeterminism because different runs may result in different places where a thread splits its stack and variation in which thread receives the new work. This can also result in run-times, especially dynamic run-times, that are *highly* variable.

6.2.9 Parallelizing the tree-search programs using OpenMP

The issues involved in implementing the static and dynamic parallel tree-search programs using OpenMP are the same as the issues involved in implementing the programs using Pthreads.

There are almost no substantive differences between a static implementation that uses OpenMP and one that uses Pthreads. However, a couple of points should be mentioned:

1. When a single thread executes some code in the Pthreads version, the test

```
if (my_rank == whatever)
```

can be replaced by the OpenMP directive

```
# pragma omp single
```

This will insure that the following structured block of code will be executed by one thread in the team, and the other threads in the team will wait in an implicit barrier at the end of the block until the executing thread is finished.

When whatever is 0 (as it is in each test in the Pthreads program), the test can also be replaced by the OpenMP directive

```
# pragma omp master
```

This will insure that thread 0 executes the following structured block of code. However, the master directive doesn't put an implicit barrier at the end of the block, so it may be necessary to also add a barrier directive after a structured block that has been modified by a master directive.

2. The Pthreads mutex that protects the best tour can be replaced by a single critical directive placed either inside the <code>Update_best_tour</code> function or immediately before the call to <code>Update_best_tour</code>. This is the only potential source of a race condition after the distribution of the initial tours, so the simple <code>critical</code> directive won't cause a thread to block unnecessarily.

The dynamically load-balanced Pthreads implementation depends heavily on Pthreads condition variables, and OpenMP doesn't provide a comparable object. The rest of the Pthreads code can be easily converted to OpenMP. In fact, OpenMP even provides a nonblocking version of <code>omp_set_lock</code>. Recall that OpenMP provides a lock object <code>omp_lock_t</code> and the following functions for acquiring and relinquishing the lock, respectively:

```
void omp_set_lock(omp_lock_t* lock_p /* in/out */);
void omp_unset_lock(omp_lock_t* lock_p /* in/out */);
It also provides the function
int omp_test_lock(omp_lock_t* lock_p /* in/out */);
```

which is analogous to pthread_mutex_trylock; it attempts to acquire the lock *lock_p, and if it succeeds it returns true (or nonzero). If the lock is being used by some other thread, it returns immediately with return value false (or zero).

If we examine the pseudocode for the Pthreads Terminated function in Program 6.8, we see that in order to adapt the Pthreads version to OpenMP, we need to emulate the functionality of the Pthreads function calls

```
pthread_cond_signal(&term_cond_var);
pthread_cond_broadcast(&term_cond_var);
pthread_cond_wait(&term_cond_var, &term_mutex);
```

in Lines 6, 17, and 22, respectively.

Recall that a thread that has entered the condition wait by calling

```
pthread_cond_wait(&term_cond_var, &term_mutex);
```

is waiting for either of two events:

- Another thread has split its stack and created work for the waiting thread.
- All of the threads have run out of work.

Perhaps the simplest solution to emulating a condition wait in OpenMP is to use busy-waiting. Since there are two conditions a waiting thread should test for, we can use two different variables in the busy-wait loop:

```
/* Global variables */
int awakened_thread = -1;
int work_remains = 1; /* true */
. . .
while (awakened_thread != my_rank && work_remains);
```

Initialization of the two variables is crucial: If awakened_thread has the value of some thread's rank, that thread will exit immediately from the while, but there may be no work available. Similarly, if work_remains is initialized to 0, all the threads will exit the while loop immediately and quit.

Now recall that when a thread enters a Pthreads condition wait, it relinquishes the mutex associated with the condition variable so that another thread can also enter the condition wait or signal the waiting thread. Thus, we should relinquish the lock used in the Terminated function before starting the while loop.

Also recall that when a thread returns from a Pthreads condition wait, it reacquires the mutex associated with the condition variable. This is especially important in this setting since if the awakened thread has received work, it will need to access the shared data structures storing the new stack. Thus, our complete emulated condition wait should look something like this:

```
/* Global vars */
int awakened_thread = -1;
work_remains = 1; /* true */
```

```
omp_unset_lock(&term_lock);
while (awakened_thread != my_rank && work_remains);
omp_set_lock(&term_lock);
```

If you recall the discussion of busy-waiting in Section 4.5 and Exercise 4.3 of Chapter 4, you may be concerned about the possibility that the compiler might reorder the code around the busy-wait loop. The compiler should not reorder across calls to omp_set_lock or omp_unset_lock. However, the updates to the variables *could* be reordered, so if we're going to be using compiler optimization, we should declare both with the volatile keyword.

Emulating the condition broadcast is straightforward: When a thread determines that there's no work left (Line 14 in Program 6.8), then the condition broadcast (Line 17) can be replaced with the assignment

```
work_remains = 0; /* Assign false to work_remains */
```

The "awakened" threads can check if they were awakened by some thread's setting work_remains to false, and, if they were, return from Terminated with the value true.

Emulating the condition signal requires a little more work. The thread that has split its stack needs to choose one of the sleeping threads and set the variable awakened_thread to the chosen thread's rank. Thus, at a minimum, we need to keep a list of the ranks of the sleeping threads. A simple way to do this is to use a shared queue of thread ranks. When a thread runs out of work, it enqueues its rank before entering the busy-wait loop. When a thread splits its stack, it can choose the thread to awaken by dequeuing the queue of waiting threads:

```
got_lock = omp_test_lock(&term_lock);
if (got_lock != 0) {
   if (waiting_threads > 0 && new_stack == NULL) {
      Split my_stack creating new_stack;
      awakened_thread = Dequeue(term_queue);
   }
   omp_unset_lock(&term_lock);
}
```

The awakened thread needs to reset awakened_thread to -1 before it returns from its call to the Terminated function.

Note that there is no danger that some other thread will be awakened before the awakened thread reacquires the lock. As long as new_stack is not NULL, no thread will attempt to split its stack, and hence no thread will try to awaken another thread. So if several threads call Terminated before the awakened thread reacquires the lock, they'll either return if their stacks are nonempty, or they'll enter the wait if their stacks are empty.

6.2.10 Performance of the OpenMP implementations

Table 6.9 shows run-times of the two OpenMP implementations on the same two fifteen-city problems that we used to test the Pthreads implementations. The programs

Search (times in seconds)														
	First Problem						Second Problem							
	Static Dynamic				Sta	tic	Dynamic							
Th	OMP	Pth	Ol	ЛP	P	th	OMP	Pth	OMP		Pth			
1	32.5	32.7	33.7	(O)	34.7	(O)	25.6	25.8	26.6	(O)	27.5	(O)		
2	27.7	27.9	28.0	(6)	28.9	(7)	25.6	25.8	18.8	(9)	19.2	(6)		
4	25.4	25.7	33.1	(75)	25.9	(47)	25.6	25.8	9.8	(52)	9.3	(49)		
8	28.0	23.8	19.2	(134)	22.4	(180)	23.8	24.0	6.3	(163)	5.7	(256)		

Table 6.9 Performance of OpenMP and Pthreads Implementations of Tree

were also run on the same system we used for the Pthreads and serial tests. For ease of comparison, we also show the Pthreads run-times. Run-times are in seconds and the numbers in parentheses show the total number of times stacks were split in the dynamic implementations.

For the most part, the OpenMP implementations are comparable to the Pthreads implementations. This isn't surprising since the system on which the programs were run has eight cores, and we wouldn't expect busy-waiting to degrade overall performance unless we were using more threads than cores.

There are two notable exceptions for the first problem. The performance of the static OpenMP implementation with eight threads is much worse than the Pthreads implementation, and the dynamic implementation with four threads is much worse than the Pthreads implementation. This could be a result of the nondeterminism of the programs, but more detailed profiling will be necessary to determine the cause with any certainty.

6.2.11 Implementation of tree search using MPI and static partitioning

The vast majority of the code used in the static parallelizations of tree search using Pthreads and OpenMP is taken straight from the second implementation of serial, iterative tree search. In fact, the only differences are in starting the threads, the initial partitioning of the tree, and the Update_best_tour function. We might therefore expect that an MPI implementation would also require relatively few changes to the serial code, and this is, in fact, the case.

There is the usual problem of distributing the input data and collecting the results. In order to construct a complete tour, a process will need to choose an edge into each vertex and out of each vertex. Thus, each tour will require an entry from each row and each column for each city that's added to the tour, so it would clearly be advantageous for each process to have access to the entire adjacency matrix. Note that the adjacency matrix is going to be relatively small. For example, even if we have 100 cities, it's unlikely that the matrix will require more than 80,000 bytes of storage, so it makes sense to simply read in the matrix on process 0 and broadcast it to all the processes.

Once the processes have copies of the adjacency matrix, the bulk of the tree search can proceed as it did in the Pthreads and OpenMP implementations. The principal differences lie in

- partitioning the tree,
- · checking and updating the best tour, and
- after the search has terminated, making sure that process 0 has a copy of the best tour for output.

We'll discuss each of these in turn.

Partitioning the tree

In the Pthreads and OpenMP implementations, thread 0 uses breadth-first search to search the tree until there are at least thread_count partial tours. Each thread then determines which of these initial partial tours it should get and pushes its tours onto its local stack. Certainly MPI process 0 can also generate a list of comm_sz partial tours. However, since memory isn't shared, it will need to send the initial partial tours to the appropriate process. We could do this using a loop of sends, but distributing the initial partial tours looks an awful lot like a call to MPI_Scatter. In fact, the only reason we can't use MPI_Scatter is that the number of initial partial tours may not be evenly divisible by comm_sz. When this happens, process 0 won't be sending the same number of tours to each process, and MPI_Scatter requires that the source of the scatter send the same number of objects to each process in the communicator.

Fortunately, there is a variant of MPI_Scatter, MPI_Scatterv, which *can* be used to send different numbers of objects to different processes. First recall the syntax of MPI_Scatter:

```
int MPI_Scatter(
                              /* in */.
      void
                   sendbuf
      int
                   sendcount
                              /* in */,
      MPI_Datatype sendtype
                              /* in */,
      void*
                              /* out */,
                   recybuf
      int
                   recvcount
                              /* in */.
      MPI_Datatype recvtype
                              /* in */,
      int
                   root
                               /* in */,
      MPI_Comm
                   comm
                               /* in */);
```

Process root sends sendcount objects of type sendtype from sendbuf to each process in comm. Each process in comm receives recovcount objects of type recovtype into recovbuf. Most of the time, sendtype and recovtype are the same and sendcount and recovcount are also the same. In any case, it's clear that the root process must send the same number of objects to each process.

MPI_Scatterv, on the other hand, has syntax

```
int MPI_Scatterv(
    void* sendbuf /* in */,
```

```
int*
           sendcounts
                          /* in */,
           displacements /* in */,
int*
MPI_Datatype sendtype
                          /* in */.
void* recvbuf
                          /* out */,
                         /* in */,
          recvcount
MPI_Datatype recvtype
                          /* in */,
                          /* in */,
int.
          root
MPI_Comm
           comm
                          /* in */);
```

The single sendcount argument in a call to MPI_Scatter is replaced by two array arguments: sendcounts and displacements. Both of these arrays contain comm_sz elements: sendcounts[q] is the number of objects of type sendtype being sent to process q. Furthermore, displacements[q] specifies the start of the block that is being sent to process q. The displacement is calculated in units of type sendtype. So, for example, if sendtype is MPI_INT, and sendbuf has type int*, then the data that is sent to process q will begin in location

```
sendbuf + displacements[q]
```

In general, displacements[q] specifies the offset into sendbuf of the data that will go to process q. The "units" are measured in blocks with extent equal to the extent of sendtype.

Similarly, MPI_Gatherv generalizes MPI_Gather:

```
int MPI_Gatherv(
     void*
                 sendbuf
                                /* in */.
     int
                sendcount
                                /* in
     MPI_Datatype sendtype
                                /* in */.
     void* recvbuf
                                /* out */,
                recvcounts
                               /* in */,
     int*
                displacements
                                /* in */,
     int*
     MPI_Datatype recvtype
                                /* in */,
                                /* in */.
     int
                root
     MPI_Comm
                 comm
                                /* in */);
```

Maintaining the best tour

As we observed in our earlier discussion of parallelizing tree search, having each process use its own best tour is likely to result in a lot of wasted computation since the best tour on one process may be much more costly than most of the tours on another process (see Exercise 6.21). Therefore, when a process finds a new best tour, it should send it to the other processes.

First note that when a process finds a new best tour, it really only needs to send its *cost* to the other processes. Each process only makes use of the cost of the current best tour when it calls Best_tour. Also, when a process updates the best tour, it doesn't care what the actual cities on the former best tour were; it only cares that the cost of the former best tour is greater than the cost of the new best tour.

During the tree search, when one process wants to communicate a new best cost to the other processes, it's important to recognize that we can't use MPI_Bcast,

for recall that MPI_Bcast is blocking and every process in the communicator must call MPI_Bcast. However, in parallel tree search the only process that will know that a broadcast should be executed is the process that has found a new best cost. If it tries to use MPI_Bcast, it will probably block in the call and never return, since it will be the only process that calls it. We therefore need to arrange that the new tour is sent in such a way that the sending process won't block indefinitely.

MPI provides several options. The simplest is to have the process that finds a new best cost use MPI_Send to send it to all the other processes:

Here, we're using a special tag defined in our program, NEW_COST_TAG. This will tell the receiving process that the message is a new cost—as opposed to some other type of message—for example, a tour.

The destination processes can periodically check for the arrival of new best tour costs. We can't use MPI_Recv to check for messages since it's blocking; if a process calls

the process will block until a matching message arrives. If no message arrives—for example, if no process finds a new best cost—the process will hang. Fortunately, MPI provides a function that only *checks* to see if a message is available; it doesn't actually try to receive a message. It's called MPI_Iprobe, and its syntax is

It checks to see if a message from process rank source in communicator comm and with tag tag is available. If such a message is available, *msg_avail_p will be assigned the value true and the members of *status_p will be assigned the appropriate values. For example, status_p->MPI_SOURCE will be assigned the rank of the source of the message that's been received. If no message is available, *msg_avail_p will be assigned the value false. The source and tag arguments can be the wildcards MPI_ANY_SOURCE and MPI_ANY_TAG, respectively. So, to check for a message with a new cost from any process, we can call

```
MPI_Iprobe(MPI_ANY_SOURCE, NEW_COST_TAG, comm, &msg_avail, &status);
```

Program 6.9: MPI code to check for new best tour costs

If msg_avail is true, then we can receive the new cost with a call to MPI_Recv:

A natural place to do this is in the Best_tour function. Before checking whether our new tour is the best tour, we can check for new tour costs from other processes with the code in Program 6.9.

This code will continue to receive messages with new costs as long as they're available. Each time a new cost is received that's better than the current best cost, the variable best_tour_cost will be updated.

Did you spot the potential problem with this scheme? If there is no buffering available for the sender, then the loop of calls to MPI_Send can cause the sending process to block until a matching receive is posted. If all the other processes have completed their searches, the sending process will hang. The loop of calls to MPI_Send is therefore unsafe.

There are a couple of alternatives provided by MPI: **buffered sends** and **non-blocking sends**. We'll discuss buffered sends here. See Exercise 6.22 for a discussion of nonblocking operations in MPI.

Modes and Buffered Sends

MPI provides four **modes** for sends: **standard**, **synchronous**, **ready**, and **buffered**. The various modes specify different semantics for the sending functions. The send that we first learned about, MPI_Send, is the standard mode send. With it, the MPI implementation can decide whether to copy the contents of the message into its own storage or to block until a matching receive is posted. Recall that in synchronous mode, the send will block until a matching receive is posted. In ready mode, the send is erroneous unless a matching receive is posted *before* the send is started. In buffered mode, the MPI implementation must copy the message into local temporary storage if a matching receive hasn't been posted. The local temporary storage must be provided by the user program, not the MPI implementation.

Each mode has a different function: MPI_Send, MPI_Ssend, MPI_Rsend, and MPI_Bsend, respectively, but the argument lists are identical to the argument lists for MPI_Send:

The buffer that's used by MPI_Bsend must be turned over to the MPI implementation with a call to MPI_Buffer_attach:

```
int MPI_Buffer_attach(
    void* buffer /* in */,
    int buffer_size /* in */);
```

The buffer argument is a pointer to a block of memory allocated by the user program and buffer_size is its size in bytes. A previously "attached" buffer can be reclaimed by the program with a call to

```
int MPI_Buffer_detach(
    void* buf_p /* out */,
    int* buf_size_p /* out */);
```

The *buf_p argument returns the address of the block of memory that was previously attached, and *buf_size_p gives its size in bytes. A call to MPI_Buffer_detach will block until all messages that have been stored in the buffer are transmitted. Note that since buf_p is an output argument, it should probably be passed in with the ampersand operator. For example:

```
char buffer[1000];
char* buf;
int buf_size;
...
MPI_Buffer_attach(buffer, 1000);
...
/* Calls to MPI_Bsend */
...
MPI_Buffer_detach(&buf, &buf_size);
```

At any point in the program only one user-provided buffer can be attached, so if there may be multiple buffered sends that haven't been completed, we need to estimate the amount of data that will be buffered. Of course, we can't know this with any certainty, but we do know that in any "broadcast" of a best tour, the process doing the broadcast will make comm_sz - 1 calls to MPI_Bsend, and each of these calls will send a single int. We can thus determine the size of the buffer needed for a single broadcast. The amount of storage that's needed for the *data* that's transmitted can be determined with a call to MPI_Pack_size:

The output argument gives an upper bound on the number of bytes needed to store the data in a message. This won't be enough, however. Recall that in addition to the data, a message stores information such as the destination, the tag, and the communicator, so for each message there is some additional overhead. An upper bound on this additional overhead is given by the MPI constant MPI_BSEND_OVERHEAD. For a single broadcast, the following code determines the amount of storage needed:

```
int data_size;
int message_size;
int bcast_buf_size;

MPI_Pack_size(1, MPI_INT, comm, &data_size);
message_size = data_size + MPI_BSEND_OVERHEAD;
bcast_buf_size = (comm_sz - 1)*message_size;
```

We should guess a generous upper bound on the number of broadcasts and multiply that by bcast_buf_size to get the size of the buffer to attach.

Printing the best tour

When the program finishes, we'll want to print out the actual tour as well as its cost, so we do need to get the tour to process 0. It might at first seem that we could arrange this by having each process store its local best tour—the best tour that it finds—and when the tree search has completed, each process can check its local best tour cost and compare it to the global best tour cost. If they're the same, the process could send its local best tour to process 0. There are, however, several problems with this. First, it's entirely possible that there are multiple "best" tours in the TSP digraph, tours that all have the same cost, and different processes may find these different tours. If this happens, multiple processes will try to send their best tours to process 0, and all but one of the threads could hang in a call to MPI_Send. A second problem is that it's possible that one or more processes never received the best tour cost, and they may try to send a tour that isn't optimal.

We can avoid these problems by having each process store its local best tour, but after all the processes have completed their searches, they can all call MPI_All reduce and the process with the global best tour can then send it to process 0 for output. The following pseudocode provides some details:

```
struct {
   int cost;
   int rank;
} loc_data, global_data;

loc_data.cost = Tour_cost(loc_best_tour);
loc_data.rank = my_rank;
```

The key here is the operation we use in the call to MPI_Allreduce. If we just used MPI_MIN, we would know what the cost of the global best tour was, but we wouldn't know who owned it. However, MPI provides a predefined operator, MPI_MINLOC, which operates on pairs of values. The first value is the value to be minimized—in our setting, the cost of the tour—and the second value is the *location* of the minimum—in our setting, the rank of the process that actually owns the best tour. If more than one process owns a tour with minimum cost, the location will be the lowest of the ranks of the processes that own a minimum cost tour. The input and the output buffers in the call to MPI_Allreduce are two-member structs. Since both the cost and the rank are ints, both members are ints. Note that MPI also provides a predefined type MPI_2INT for this type. When the call to MPI_Allreduce returns, we have two alternatives:

- If process 0 already has the best tour, we simply return.
- Otherwise, the process owning the best tour sends it to process 0.

Unreceived messages

As we noted in the preceding discussion, it is possible that some messages won't be received during the execution of the parallel tree search. A process may finish searching its subtree before some other process has found a best tour. This won't cause the program to print an incorrect result; the call to MPI_Allreduce that finds the process with the best tour won't return until every process has called it, and some process will have the best tour. Thus, it will return with the correct least-cost tour, and process 0 will receive this tour.

However, unreceived messages can cause problems with the call to MPI_Buffer_detach or the call to MPI_Finalize. A process can hang in one of these calls if it is storing buffered messages that were never received, so before we attempt to shut down MPI, we can try to receive any outstanding messages by using MPI_Iprobe. The code is very similar to the code we used to check for new best tour costs. See Program 6.9. In fact, the only messages that are not sent in collectives are the "best tour" message sent to process 0, and the best tour cost broadcasts. The MPI collectives will hang if some process doesn't participate, so we only need to look for unreceived best tours.

In the dynamically load-balanced code (which we'll discuss shortly) there are other messages, including some that are potentially quite large. To handle this situation, we can use the status argument returned by MPI_Iprobe to determine the size of the message and allocate additional storage as necessary (see Exercise 6.23).

6.2.12 Implementation of tree search using MPI and dynamic partitioning

In an MPI program that dynamically partitions the search tree, we can try to emulate the dynamic partitioning that we used in the Pthreads and OpenMP programs. Recall that in those programs, before each pass through the main while loop in the search function, a thread called a boolean-valued function called Terminated. When a thread ran out of work—that is, its stack was empty—it went into a condition wait (Pthreads) or a busy-wait (OpenMP) until it either received additional work or it was notified that there was no more work. In the first case, it returned to searching for a best tour. In the second case, it quit. A thread that had at least two records on its stack would give half of its stack to one of the waiting threads.

Much of this can be emulated in a distributed-memory setting. When a process runs out of work, there's no condition wait, but it can enter a busy-wait, in which it waits to either receive more work or notification that the program is terminating. Similarly, a process with work can split its stack and send work to an idle process.

The key difference is that there is no central repository of information on which processes are waiting for work, so a process that splits its stack can't just dequeue a queue of waiting processes or call a function such as pthread_cond_signal. It needs to "know" a process that's waiting for work so it can send the waiting process more work. Thus, rather than simply going into a busy-wait for additional work or termination, a process that has run out of work should send a request for work to another process. If it does this, then, when a process enters the Terminated function, it can check to see if there's a request for work from some other process. If there is, and the process that has just entered Terminated has work, it can send part of its stack to the requesting process. If there is a request, and the process has no work available, it can send a rejection. Thus, when we have distributed-memory, pseudocode for our Terminated function can look something like the pseudocode shown in Program 6.10.

Terminated begins by checking on the number of tours that the process has in its stack (Line 1); if it has at least two that are "worth sending," it calls Fulfill_request (Line 2). Fulfill_request checks to see if the process has received a request for work. If it has, it splits its stack and sends work to the requesting process. If it hasn't received a request, it just returns. In either case, when it returns from Fulfill_request it returns from Terminated and continues searching.

If the calling process doesn't have at least two tours worth sending, Terminated calls Send_rejects (Line 5), which checks for any work requests from other processes and sends a "no work" reply to each requesting process. After this, Terminated checks to see if the calling process has any work at all. If it does—that is, if its stack isn't empty—it returns and continues searching.

Things get interesting when the calling process has no work left (Line 9). If there's only one process in the communicator (comm_sz = 1), then the process returns from Terminated and quits. If there's more than one process, then the process "announces" that it's out of work in Line 11. This is part of the implementation

```
if (My_avail_tour_count(my_stack) >= 2) {
2
       Fulfill_request(my_stack);
3
       return false; /* Still more work */
   } else { /* At most 1 available tour */
4
5
       Send_rejects(); /* Tell everyone who's requested */
                         /* work that I have none
6
7
       if (!Empty_stack(my_stack)) {
          return false; /* Still more work */
8
       } else { /* Empty stack */
9
10
          if (comm_sz == 1) return true;
11
          Out_of_work();
12
          work_request_sent = false;
13
          while (1) {
             Clear_msgs(); /* Msgs unrelated to work, termination */
14
             if (No_work_left()) {
15
16
                return true; /* No work left. Quit */
17
             } else if (!work_request_sent) {
18
                Send_work_request(); /* Request work from someone */
19
                work_request_sent = true;
20
             } else {
21
                Check_for_work(&work_request_sent, &work_avail);
22
                if (work_avail) {
23
                   Receive_work(my_stack);
24
                   return false;
25
                }
26
             }
27
             /* while */
          }
28
       } /* Empty stack */
29
   } /* At most 1 available tour */
```

Program 6.10: Terminated function for a dynamically partitioned TSP solver that uses MPI

of a "distributed termination detection algorithm," which we'll discuss shortly. For now, let's just note that the termination detection algorithm that we used with shared-memory may not work, since it's impossible to guarantee that a variable storing the number of processes that have run out of work is up to date.

Before entering the apparently infinite while loop (Line 13), we set the variable work_request_sent to false (Line 12). As its name suggests, this variable tells us whether we've sent a request for work to another process; if we have, we know that we should wait for work or a message saying "no work available" from that process before sending out a request to another process.

The while(1) loop is the distributed-memory version of the OpenMP busy-wait loop. We are essentially waiting until we either receive work from another process or we receive word that the search has been completed.

When we enter the while(1) loop, we deal with any outstanding messages in Line 14. We may have received updates to the best tour cost and we may have received requests for work. It's essential that we tell processes that have requested

work that we have none, so that they don't wait forever when there's no work available. It's also a good idea to deal with updates to the best tour cost, since this will free up space in the sending process' message buffer.

After clearing out outstanding messages, we iterate through the possibilities:

- The search has been completed, in which case we quit (Lines 15–16).
- We don't have an outstanding request for work, so we choose a process and send it a request (Lines 17–19). We'll take a closer look at the problem of which process should be sent a request shortly.
- We do have an outstanding request for work (Lines 21–25). So we check whether the request has been fulfilled or rejected. If it has been fulfilled, we receive the new work and return to searching. If we received a rejection, we set work_request_sent to false and continue in the loop. If the request was neither fulfilled nor rejected, we also continue in the while(1) loop.

Let's take a closer look at some of these functions.

My_avail_tour_count

The function My_avail_tour_count can simply return the size of the process' stack. It can also make use of a "cutoff length." When a partial tour has already visited most of the cities, there will be very little work associated with the subtree rooted at the partial tour. Since sending a partial tour is likely to be a relatively expensive operation, it may make sense to only send partial tours with fewer than some cutoff number of edges. In Exercise 6.24 we take a look at how such a cutoff affects the overall run-time of the program.

Fulfill_request

If a process has enough work so that it can usefully split its stack, it calls Fulfill_request (Line 2). Fulfill_request uses MPI_Iprobe to check for a request for work from another process. If there is a request, it receives it, splits its stack, and sends work to the requesting process. If there isn't a request for work, the process just returns.

Splitting the stack

A Split_stack function is called by Fulfill_request. It uses the same basic algorithm as the Pthreads and OpenMP functions, that is, alternate partial tours with fewer than split_cutoff cities are collected for sending to the process that has requested work. However, in the shared-memory programs, we simply copy the tours (which are pointers) from the original stack to a new stack. Unfortunately, because of the pointers involved in the new stack, such a data structure cannot be simply sent to another process (see Exercise 6.25). Thus, the MPI version of Split_stack packs the contents of the new stack into contiguous memory and sends the block of contiguous memory, which is unpacked by the receiver into a new stack.

MPI provides a function, MPI_Pack, for packing data into a buffer of contiguous memory. It also provides a function, MPI_Unpack, for unpacking data from a buffer

of contiguous memory. We took a brief look at them in Exercise 6.20 of Chapter 3. Recall that their syntax is

```
int MPI_Pack(
      void*
                    data_to_be_packed
                                          /* in
      int.
                   to_be_packed_count
                                         /* in
                                                    */,
      MPI_Datatype datatype
                                          /* in
                                                    */.
      void*
                   contig_buf
                                          /* out
                                                    */,
      int
                    contig_buf_size
                                         /* in
                                                    */,
      int*
                    position_p
                                          /* in/out */,
      MPI_Comm
                                          /* in
                    comm
                                                    */);
int MPI_Unpack(
      void*
                    contig_buf
                                          /* in
      int
                    contig_buf_size
                                          /* in
                                          /* in/out */,
      int.*
                    position_p
      void*
                    unpacked_data
                                          /* out
                                                    */.
                    unpack_count
                                          /* in
      int
                                                    */,
      MPI_Datatype datatype
                                          /* in
                                                    */,
                                          /* in
      MPI_Comm
                                                    */);
                    comm
```

MPI_Pack takes the data in data_to_be_packed and packs it into contig_buf. The *position_p argument keeps track of where we are in contig_buf. When the function is called, it should refer to the first available location in contig_buf before data_to_be_packed is added. When the function returns, it should refer to the first available location in contig_buf after data_to_be_packed has been added.

MPI_Unpack reverses the process. It takes the data in <code>contig_buf</code> and unpacks it into <code>unpacked_data</code>. When the function is called, <code>*position_p</code> should refer to the first location in <code>contig_buf</code> that hasn't been unpacked. When it returns, <code>*position_p</code> should refer to the next location in <code>contig_buf</code> after the data that was just unpacked.

As an example, suppose that a program contains the following definitions:

Then we can send a variable with type tour_t using the following code:

```
&position, comm);
MPI_Send(contig_buf, position, MPI_PACKED, dest, 0, comm);
} /* Send_tour */
```

Similarly, we can receive a variable of type tour_t using the following code:

Note that the MPI datatype that we use for sending and receiving packed buffers is MPI_PACKED.

Send_rejects

The Send_rejects function (Line 5) is similar to the function that looks for new best tours. It uses MPI_Iprobe to search for messages that have requested work. Such messages can be identified by a special tag value, for example, WORK_REQ_TAG. When such a message is found, it's received, and a reply is sent indicating that there is no work available. Note that both the request for work and the reply indicating there is no work can be messages with zero elements, since the tag alone informs the receiver of the message's purpose. Even though such messages have no content outside of the envelope, the envelope does take space and they need to be received.

Distributed termination detection

The functions <code>Out_of_work</code> and <code>No_work_left</code> (Lines 11 and 15) implement the termination detection algorithm. As we noted earlier, an algorithm that's modeled on the termination detection algorithm we used in the shared-memory programs will have problems. To see this, suppose each process stores a variable <code>oow</code>, which stores the number of processes that are out of work. The variable is set to 0 when the program starts. Each time a process runs out of work, it sends a message to all the other processes saying it's out of work so that all the processes will increment their copies of <code>oow</code>. Similarly, when a process receives work from another process, it sends a message to every process informing them of this, and each process will decrement its copy of <code>oow</code>. Now suppose we have three process, and process 2 has work but processes 0 and 1 have run out of work. Consider the sequence of events shown in Table 6.10.

The error here is that the work sent from process 1 to process 0 is lost. The reason is that process 0 receives the notification that process 2 is out of work before it receives the notification that process 1 has received work. This may seem improbable,

Table 6.10 Termination Events that Result in an Error									
Time	Process 0	Process 1	Process 2						
0	Out of Work Notify 1, 2 00W = 1	Out of Work Notify 0, 2 00w = 1	Working oow = 0						
1	Send request to 1 oow = 1	Send Request to 2 oow = 1	Recv notify fr 1 oow = 1						
2	oow = 1	Recv notify fr 0 oow = 2	Recv request fr 1 oow = 1						
3	oow = 1	oow = 2	Send work to 1 $00W = 0$						
4	oow = 1	Recv work fr 2 00w = 1	Recv notify fr 0 oow = 1						
5	oow = 1	Notify 0 $00W = 1$	Working oow = 1						
6	oow = 1	Recv request fr 0 oow = 1	Out of work Notify 0, 1 oow = 2						
7	Recv notify fr 2 oow = 2	Send work to 0 00w = 0	Send request to 1 oow = 2						
8	Recv 1st notify fr 1 00w = 3	Recv notify fr 2	oow = 2						
9	Quit	Recv request fr 2 00w = 1	00W = 2						

but it's not improbable that process 1 was, for example, interrupted by the operating system and its message wasn't transmitted until after the message from process 2 was transmitted.

Although MPI guarantees that two messages sent from process A to process B will, in general, be received in the order in which they were sent, it makes no guarantee about the order in which messages will be received if they were sent by different processes. This is perfectly reasonable in light of the fact that different processes will, for various reasons, work at different speeds.

Distributed termination detection is a challenging problem, and much work has gone into developing algorithms that are guaranteed to correctly detect it. Conceptually, the simplest of these algorithms relies on keeping track of a quantity that is conserved and can be measured precisely. Let's call it *energy*, since, of course, energy is conserved. At the start of the program, each process has 1 unit of energy. When a process runs out of work, it sends its energy to process 0. When a process fulfills a request for work, it divides its energy in half, keeping half for itself, and sending half to the process that's receiving the work. Since energy is conserved and since the program started with comm_sz units, the program should terminate when process 0 finds that it has received a total of comm_sz units.

The <code>Out_of_work</code> function when executed by a process other than 0 sends its energy to process 0. Process 0 can just add its energy to a <code>received_energy</code> variable. The <code>No_work_left</code> function also depends on whether process 0 or some other process is calling. If process 0 is calling, it can receive any outstanding messages sent by <code>Out_of_work</code> and add the energy into <code>received_energy</code>. If <code>received_energy</code> equals <code>comm_sz</code>, process 0 can send a termination message (with a special tag) to every process. On the other hand, a process other than 0 can just check to see if there's a message with the termination tag.

The tricky part here is making sure that no energy is inadvertently lost; if we try to use floats or doubles, we'll almost certainly run into trouble since at some point dividing by two will result in underflow. Since the amount of energy in exact arithmetic can be represented by a common fraction, we can represent the amount of energy on each process exactly by a pair of fixed-point numbers. The denominator will always be a power of two, so we can represent it by its base-two logarithm. For a large problem it is possible that the numerator could overflow. However, if this becomes a problem, there are libraries that provide arbitrary precision rational numbers (e.g, GMP [21]). An alternate solution is explored in Exercise 6.26.

Sending requests for work

Once we've decided on which process we plan to send a request to, we can just send a zero-length message with a "request for work" tag. However, there are many possibilities for choosing a destination:

- 1. Loop through the processes in round-robin fashion. Start with (my_rank + 1) % comm_sz and increment this destination (modulo comm_sz) each time a new request is made. A potential problem here is that two processes can get "in synch" and request work from the same destination repeatedly.
- 2. Keep a global destination for requests on process 0. When a process runs out of work, it first requests the current value of the global destination from 0. Process 0 can increment this value (modulo comm_sz) each time there's a request. This avoids the issue of multiple processes requesting work from the same destination, but clearly process 0 can become a bottleneck.
- **3.** Each process uses a random number generator to generate destinations. While it can still happen that several processes may simultaneously request work from the same process, the random choice of successive process ranks should reduce the chance that several processes will make repeated requests to the same process.

These are three possible options. We'll explore these options in Exercise 6.29. Also see [22] for an analysis of the options.

Checking for and receiving work

Once a request is sent for work, it's critical that the sending process repeatedly check for a response from the destination. In fact, a subtle point here is that it's critical that the sending process check for a message from the destination process with a "work available tag" or a "no work available tag." If the sending process simply checks

for a message from the destination, it may be "distracted" by other messages from the destination and never receive work that's been sent. For example, there might be a message from the destination requesting work that would mask the presence of a message containing work.

The Check_for_work function should therefore first probe for a message from the destination indicating work is available, and, if there isn't such a message, it should probe for a message from the destination saying there's no work available. If there is work available, the Receive_work function can receive the message with work and unpack the contents of the message buffer into the process' stack. Note also that it needs to unpack the energy sent by the destination process.

Performance of the MPI programs

Table 6.11 shows the performance of the two MPI programs on the same two fifteen-city problems on which we tested the Pthreads and the OpenMP implementations. Run-times are in seconds and the numbers in parentheses show the total number of times stacks were split in the dynamic implementations. These results were obtained on a different system from the system on which we obtained the Pthreads results. We've also included the Pthreads results for this system, so that the two sets of results can be compared. The nodes of this system only have four cores, so the Pthreads results don't include times for 8 or 16 threads. The cutoff number of cities for the MPI runs was 12.

The nodes of this system are small shared-memory systems, so communication through shared variables should be much faster than distributed-memory communication, and it's not surprising that in every instance the Pthreads implementation outperforms the MPI implementation.

The cost of stack splitting in the MPI implementation is quite high; in addition to the cost of the communication, the packing and unpacking is very time-consuming. It's also therefore not surprising that for relatively small problems with few processes, the static MPI parallelization outperforms the dynamic parallelization. However, the

Search (times in seconds)													
	First Problem						Second Problem						
Static				Dynamic			Sta	atic	Dynamic				
Th/Pr	Pth	MPI	Pt	h	М	'PI	Pth	MPI	MPI Pth		М	'PI	
1 2 4 8 16	35.8 29.9 27.2	40.9 34.9 31.7 35.7 20.1	41.9 34.3 30.2	(0) (9) (55)	56.5 55.6 52.6 45.5 10.5	(0) (5) (85) (165) (441)	27.4 27.4 27.4	31.5 31.5 31.5 35.7 17.8	32.3 22.0 10.7	(0) (8) (44)	43.8 37.4 21.8 16.5 0.1	(0) (9) (76) (161) (173)	

8- and 16-process results suggest that if a problem is large enough to warrant the use of many processes, the dynamic MPI program is much more scalable, and it can provide far superior performance. This is borne out by examination of a 17-city problem run with 16 processes: the dynamic MPI implementation has a run-time of 296 seconds, while the static implementation has a run-time of 601 seconds.

Note that times such as 0.1 second for the second problem running with 16 processes don't really show superlinear speedup. Rather, the initial distribution of work has allowed one of the processes to find the best tour much faster than the initial distributions with fewer processes, and the dynamic partitioning has allowed the processes to do a much better job of load balancing.

6.3 A WORD OF CAUTION

In developing our solutions to the *n*-body problem and TSP, we chose our serial algorithms because they were easy to understand and their parallelization was relatively straightforward. In no case did we choose a serial algorithm because it was the fastest or because it could solve the largest problem. Thus, it should not be assumed that either the serial or the parallel solutions are the best available. For information on "state-of-the-art" algorithms, see the bibliography, especially [12] for the *n*-body problem and [22] for parallel tree search.

6.4 WHICH API?

How can we decide which API, MPI, Pthreads, or OpenMP is best for our application? In general, there are many factors to consider, and the answer may not be at all clear cut. However, here are a few points to consider.

As a first step, decide whether to use distributed-memory, or shared-memory. In order to do this, first consider the amount of memory the application will need. In general, distributed-memory systems can provide considerably more main memory than shared-memory systems, so if the memory requirements are very large, you may need to write the application using MPI.

If the problem will fit into the main memory of your shared-memory system, you may still want to consider using MPI. Since the total available cache on a distributed-memory system will probably be much greater than that available on a shared-memory system, it's conceivable that a problem that requires lots of main memory accesses on a shared-memory system will mostly access cache on a distributed-memory system, and, consequently, have much better overall performance.

However, even if you'll get a big performance improvement from the large aggregate cache on a distributed-memory system, if you already have a large and complex serial program, it often makes sense to write a shared-memory program. It's often

possible to reuse considerably more serial code in a shared-memory program than a distributed-memory program. It's more likely that the serial data structures can be easily adapted to a shared-memory system. If this is the case, the development effort for the shared-memory program will probably be much less. This is especially true for OpenMP programs, since some serial programs can be parallelized by simply inserting some OpenMP directives.

Another consideration is the communication requirements of the parallel algorithm. If the processes/threads do little communication, an MPI program should be fairly easy to develop, and very scalable. At the other extreme, if the processes/threads need to be very closely coordinated, a distributed-memory program will probably have problems scaling to large numbers of processes, and the performance of a shared-memory program should be better.

If you decided that shared-memory is preferable, you will need to think about the details of parallelizing the program. As we noted earlier, if you already have a large, complex serial program, you should see if it lends itself to OpenMP. For example, if large parts of the program can be parallelized with parallel for directives, OpenMP will be much easier to use than Pthreads. On the other hand, if the program involves complex synchronization among the threads—for example, read-write locks or threads waiting on signals from other threads—then Pthreads will be much easier to use.

6.5 SUMMARY

In this chapter, we've looked at serial and parallel solutions to two very different problems: the *n*-body problem and solving the traveling salesperson problem using tree search. In each case we began by studying the problem and looking at serial algorithms for solving the problem. We continued by using Foster's methodology for devising a parallel solution, and then, using the designs developed with Foster's methodology, we implemented parallel solutions using Pthreads, OpenMP, and MPI. In developing the reduced MPI solution to the *n*-body problem, we determined that the "obvious" solution would be extremely difficult to implement correctly and would require a huge amount of communication. We therefore turned to an alternative "ring pass" algorithm, which proved to be much easier to implement and is probably more scalable.

In the dynamically partitioned solutions for parallel tree search, we used different methods for the three APIs. With Pthreads, we used a condition variable both for communicating new work among the threads and for termination. OpenMP doesn't provide an analog to Pthreads condition variables, so we used busy-waiting instead. In MPI, since all data is local, we needed to use a more complicated scheme to redistribute work, in which a process that runs out of work chooses a destination process and requests work from that process. To implement this correctly, a process that runs out of work enters a busy-wait loop in which it requests work, looks for a response to the work request, and looks for a termination message.

We saw that in a distributed-memory environment in which processes send each other work, determining when to terminate is a nontrivial problem. We also looked at a relatively straightforward solution to the problem of distributed termination detection, in which there is a fixed amount of "energy" throughout the execution of the program. When processes run out of work they send their energy to process 0, and when processes send work to other processes, they also send half of their current energy. Thus, when process 0 finds that it has all the energy, there is no more work, and it can send a termination message.

In closing, we looked briefly at the problem of deciding which API to use. The first consideration is whether to use shared-memory or distributed-memory. To decide this, we should look at the memory requirements of the application and the amount of communication among the processes/threads. If the memory requirements are great or the distributed-memory version can work mainly with cache, then a distributed-memory program is likely to be much faster. On the other hand, if there is considerable communication, a shared-memory program will probably be faster.

In choosing between OpenMP and Pthreads, if there's an existing serial program and it can be parallelized by the insertion of OpenMP directives, then OpenMP is probably the clear choice. However, if complex thread synchronization is needed—for example, read-write locks or thread signaling—then Pthreads will be easier to use. In the course of developing these programs, we also learned some more about Pthreads, OpenMP, and MPI.

6.5.1 Pthreads and OpenMP

In tree search, we need to check the cost of the current best tour before updating the best tour. In the Pthreads and OpenMP implementations of parallel tree search, updating the best tour introduces a race condition. A thread that wants to update the best tour must therefore first acquire a lock. The combination of "test lock condition" and "update lock condition" can cause a problem: the lock condition (e.g. the cost of the best tour) can change between the time of the first test and the time that the lock is acquired. Thus, the threads also need to check the lock condition *after* they acquire the lock, so pseudocode for updating the best tour should look something like this:

```
if (new_tour_cost < best_tour_cost) {
   Acquire lock protecting best tour;
   if (new_tour_cost < best_tour_cost)
        Update best tour;
   Relinquish lock;
}</pre>
```

Remember that we have also learned that Pthreads has a *nonblocking* version of pthreads_mutex_lock called pthread_mutex_trylock. This function checks to see if the mutex is available. If it is, it acquires the mutex and returns the value 0. If the mutex isn't available, instead of waiting for it to become available, it will return a nonzero value.

The analog of pthread_mutex_trylock in OpenMP is omp_test_lock. However, its return values are the opposite of those for pthread_mutex_trylock: it returns a nonzero value if the lock is acquired and a zero value if the lock is not acquired.

When a single thread should execute a structured block, OpenMP provides a couple of alternatives to the test, and:

```
if (my_rank == special_rank) {
   Execute action;
}
```

With the single directive

```
# pragma omp single
Execute action;
Next action;
```

the run-time system will choose a single thread to execute the action. The other threads will wait in an implicit barrier before proceeding to Next action. With the master directive

```
# pragma omp master
Execute action;
Next action;
```

the master thread (thread 0) will execute the action. However, unlike the single directive, there is no implicit barrier after the block Execute action, and the other threads in the team will proceed immediately to execute Next action. Of course, if we need a barrier before proceeding, we can add an explicit barrier after completing the structured block Execute action. In Exercise 6.6 we see that OpenMP provides a nowait clause which can modify a single directive:

```
# pragma omp single nowait
Execute action;
Next action:
```

When this clause is added, the thread selected by the run-time system to execute the action will execute it as before. However, the other threads in the team won't wait, they'll proceed immediately to execute Next action. The nowait clause can also be used to modify parallel for and for directives.

6.5.2 MPI

We learned quite a bit more about MPI. We saw that in some of the collective communication functions that use an input and an output buffer, we can use the argument MPI_IN_PLACE so that the input and output buffers are the same. This can save on memory and the implementation may be able to avoid copying from the input buffer to the output buffer.

The functions MPI_Scatter and MPI_Gather can be use to split an array of data among processes and collect distributed data into a single array, respectively. However, they can only be used when the amount of data going to or coming from each process is the same for each process. If we need to assign different amounts of data to each process, or to collect different amounts of data from each process, we can use MPI_Scatterv and MPI_Gatherv, respectively:

```
int MPI_Scatterv(
         void*
                      sendbuf
                                        /* in */.
         int*
                      sendcounts
                                        /* in */,
         int*
                      displacements
                                        /* in */,
         MPI_Datatype sendtype
                                        /* in */.
         void*
                      recvbuf
                                        /* out */,
         int
                      recvcount
                                        /* in */,
                                        /* in */,
         MPI_Datatype recvtype
                                        /* in */,
                      root
         int
         MPI_Comm
                      comm
                                        /* in */);
int MPI_Gatherv(
         void*
                      sendbuf
                                        /* in */,
         int
                      sendcount
                                        /* in */.
         MPI_Datatype sendtype
                                        /* in */,
         void*
                      recvbuf
                                        /* out */,
                                        /* in */,
         int*
                      recvcounts
         int.*
                      displacements
                                        /* in */.
         MPI_Datatype recytype
                                        /* in */.
                      root
                                        /* in */,
         MPI\_Comm
                                        /* in */);
                      comm
```

The arguments sendcounts for MPI_Scatterv and recvcounts for MPI_Gatherv are arrays with comm_sz elements. They specify the amount of data (in units of sendtype/recvtype) going to or coming from each process. The displacements arguments are also arrays with comm_sz elements. They specify the offsets (in units of sendtype/recvtype) of the data going to or coming from each process.

We saw that there is a special operator, MPI_MIN_LOC, that can be used in calls to MPI_Reduce and MPI_All reduce. It operates on pairs of values and returns a pair of values. If the pairs are

```
(a_0,b_0),(a_1,b_1),\ldots,(a_{\text{COMM\_S}\,Z-1},b_{\text{COMM\_S}\,Z-1}),
```

suppose that a is the minimum of the a_i 's and q is the smallest process rank at which a occurs. Then the MPI_MIN_LOC operator will return the pair (a_q,b_q) . We used this to find not only the cost of the minimum-cost tour, but by making the b_i 's the process ranks, we determined which process owned the minimum-cost tour.

In our development of two MPI implementations of parallel tree search, we made repeated use of MPI_Iprobe:

It checks to see if there is a message from source with tag tag available to be received. If such a message is available, msg_avail_p will be given the value true. Note that MPI_Iprobe doesn't actually receive the message, but if such a message is available, a call to MPI_Recv will receive it. Both the source and tag arguments can be the wildcards MPI_ANY_SOURCE and MPI_ANY_TAG, respectively. For example, we often wanted to check whether any process had sent a message with a new best cost. We checked for the arrival of such a message with the call

If such a message is available, its source will be returned in the *status_p argument. Thus, status.MPI_SOURCE can be used to receive the message:

There were several occasions when we wanted a send function to return immediately, regardless of whether the message had actually been sent. One way to arrange this in MPI is to use **buffered send mode**. In a buffered send, the user program provides storage for messages with a call to MPI_Buffer_attach. Then when the program sends the message with MPI_Bsend, the message is either transmitted immediately or copied to the user-program-provided buffer. In either case the call returns without blocking. When the program no longer needs to use buffered send mode, the buffer can be recovered with a call to MPI_Buffer_detach.

We also saw that MPI provides three other modes for sending: **synchronous**, **standard**, and **ready**. Synchronous sends won't buffer the data; a call to the synchronous send function MPI_Ssend won't return until the receiver has begun receiving the data. Ready sends (MPI_Rsend) are erroneous unless the matching receive has already been started when MPI_Rsend is called. The ordinary send MPI_Send is called the standard mode send.

In Exercise 6.22 we explore an alternative to buffered mode: nonblocking sends. As the name suggests, a nonblocking send returns regardless of whether the message has been transmitted. However, the send must be completed by calling one of several functions that *wait* for completion of the nonblocking operation. There is also a nonblocking receive function.

Since addresses on one system will, in general, have no relation to addresses on another system, pointers should not be sent in MPI messages. If you're using data structures that have embedded pointers, MPI provides the function MPI_Pack for storing a data structure in a single, contiguous buffer before sending. Similarly, the function MPI_Unpack can be used to take data that's been received into a single contiguous buffer and unpack it into a local data structure. Their syntax is

```
int MPI_Pack(
      void*
                    data_to_be_packed
                                          /* in
      int
                    to_be_packed_count
                                          /* in
                                                     */.
      MPI_Datatype datatype
                                          /* in
                                                     */.
                    contig_buf
                                          /* out
      void*
                                                     */,
      int
                    contig_buf_size
                                          /* in
                                                     */,
                    position_p
                                          /* in/out */,
      int.*
      MPI_Comm
                                          /* in
                    comm
                                                     */);
int MPI_Unpack(
      void*
                    contig_buf
                                          /* in
                                                     */,
      int
                    contig_buf_size
                                          /* in
                                                     */,
                                          /* in/out */,
      int*
                    position_p
      void*
                    unpacked_data
                                          /* out
                                                     */,
      int
                    unpack_count
                                          /* in
                                                     */,
                                          /* in
      MPI_Datatype
                    datatype
                                                     */,
      {\tt MPI\_Comm}
                    comm
                                          /* in
                                                     */);
```

The key to their use is the position_p argument. When MPI_Pack is called, it should reference the first available location in contig_buf. So, for example, when we start packing the data *position_p should be set to 0. When MPI_Pack returns, *position_p will refer to the first available location following the data that was just packed. Thus, successive elements of a data structure can be packed into a single buffer by repeated calls to MPI_Pack. When a packed buffer is received, the data can be unpacked in a completely analogous fashion. Note that when a buffer packed with MPI_Pack is sent, the datatype for both the send and the receive should be MPI_PACKED.

6.6 EXERCISES

6.1. In each iteration of the serial *n*-body solver, we first compute the total force on each particle, and then we compute the position and velocity of each particle. Would it be possible to reorganize the calculations so that in each iteration we did all of the calculations for each particle before proceeding to the next particle? That is, could we use the following pseudocode?

```
for each timestep
  for each particle {
    Compute total force on particle;
    Find position and velocity of particle;
    Print position and velocity of particle;
}
```

If so, what other modifications would we need to make to the solver? If not, why not?

6.2. Run the basic serial n-body solver for 1000 timesteps with a stepsize of 0.05, no output, and internally generated initial conditions. Let the number

of particles range from 500 to 2000. How does the run-time change as the number of particles increases? Can you extrapolate and predict how many particles the solver could handle if it ran for 24 hours?

- **6.3.** Parallelize the reduced version of the *n*-body solver with OpenMP or Pthreads and a single critical directive (OpenMP) or a single mutex (Pthreads) to protect access to the forces array. Parallelize the rest of the solver by parallelizing the inner for loops. How does the performance of this code compare with the performance of the serial solver? Explain your answer.
- **6.4.** Parallelize the reduced version of the *n*-body solver with OpenMP or Pthreads and a lock/mutex for each particle. The locks/mutexes should be used to protect access to updates to the forces array. Parallelize the rest of the solver by parallelizing the inner for loops. How does the performance compare with the performance of the serial solver? Explain your answer.
- **6.5.** In the shared-memory reduced *n*-body solver, if we use a block partition in both phases of the calculation of the forces, the loop in the second phase can be changed so that the for thread loop only goes up to my_rank instead of thread_count. That is, the code

```
# pragma omp for
for (part = 0; part < n; part++) {
    forces[part][X] = forces[part][Y] = 0.0;
    for (thread = 0; thread < thread_count; thread++) {
        forces[part][X] += loc_forces[thread][part][X];
        forces[part][Y] += loc_forces[thread][part][Y];
    }
}</pre>
```

can be changed to

```
# pragma omp for
for (part = 0; part < n; part++) {
    forces[part][X] = forces[part][Y] = 0.0;
    for (thread = 0; thread < my_rank; thread++) {
        forces[part][X] += loc_forces[thread][part][X];
        forces[part][Y] += loc_forces[thread][part][Y];
    }
}</pre>
```

Explain why this change is OK. Run the program with this modification and compare its performance with the original code with block partitioning and the code with a cyclic partitioning of the first phase of the forces calculation. What conclusions can you draw?

6.6. In our discussion of the OpenMP implementation of the basic *n*-body solver, we observed that the implied barrier after the output statement wasn't necessary. We could therefore modify the single directive with a nowait clause. It's possible to also eliminate the implied barriers at the ends of the two for

each particle q loops by modifying for directives with nowait clauses. Would doing this cause any problems? Explain your answer.

- **6.7.** For the shared-memory implementation of the reduced *n*-body solver, we saw that a cyclic schedule for the computation of the forces outperformed a block schedule, in spite of the reduced cache performance. By experimenting with the OpenMP or the Pthreads implementation, determine the performance of various block-cyclic schedules. Is there an optimal block size for your system?
- **6.8.** If **x** and **y** are double-precision *n*-dimensional vectors and α is a double-precision scalar, the assignment

$$\mathbf{y} \longleftarrow \alpha \mathbf{x} + \mathbf{y}$$

is called a DAXPY. DAXPY is an abbreviation of "Double precision Alpha times **X** Plus **Y**." Write a Pthreads or OpenMP program in which the master thread generates two large, random *n*-dimensional arrays and a random scalar, all of which are doubles. The threads should then carry out a DAXPY on the randomly generated values. For large values of *n* and various numbers of threads compare the performance of the program with a block partition and a cyclic partition of the arrays. Which partitioning scheme performs better? Why?

- **6.9.** Write an MPI program in which each process generates a large, initialized, *m*-dimensional array of doubles. Your program should then repeatedly call MPI_Allgather on the *m*-dimensional arrays. Compare the performance of the calls to MPI_Allgather when the global array (the array that's created by the call to MPI_Allgather) has
 - a. a block distribution, and
 - **b.** a cyclic distribution.

To use a cyclic distribution, download the code cyclic_derived.c from the book's web site, and use the MPI datatype created by this code for the *destination* in the calls to MPI_Allgather. For example, we might call

if the new MPI datatype were called cyclic_mpi_t.

Which distribution performs better? Why? Don't include the overhead involved in building the derived datatype.

6.10. Consider the following code:

```
int n, thread_count, i, chunksize;
double x[n], y[n], a;
    . . .
# pragma omp parallel num_threads(thread_count) \
    default(none) private(i) \
    shared(x, y, a, n, thread_count, chunksize)
```

```
# pragma omp for schedule(static, n/thread_count)
    for (i = 0; i < n; i++) {
        x[i] = f(i); /* f is a function */
        y[i] = g(i); /* g is a function */
    }
# pragma omp for schedule(static, chunksize)
    for (i = 0; i < n; i++)
        y[i] += a*x[i];
} /* omp parallel */</pre>
```

Suppose n = 64, thread_count = 2, the cache-line size is 8 doubles, and each core has an L2 cache that can store 131,072 doubles. If chunksize = n/thread_count, how many L2 cache misses do you expect in the second loop? If chunksize = 8, how many L2 misses do you expect in the second loop? You can assume that both x and y are aligned on a cache-line boundary. That is, both x[0] and y[0] are the first elements in their respective cache lines.

- **6.11.** Write an MPI program that compares the performance of MPI_Allgather using MPI_IN_PLACE with the performance of MPI_Allgather when each process uses separate send and receive buffers. Which call to MPI_Allgather is faster when run with a single process? What if you use multiple processes?
- **6.12. a.** Modify the basic MPI implementation of the *n*-body solver so that it uses a separate array for the local positions. How does its performance compare with the performance of the original *n*-body solver? (Look at performance with I/O turned off.)
 - **b.** Modify the basic MPI implementation of the *n*-body solver so that it distributes the masses. What changes need to be made to the communications in the program? How does the performance compare with the original solver?
- **6.13.** Using Figure 6.6 as a guide, sketch the communications that would be needed in an "obvious" MPI implementation of the reduced *n*-body solver if there were three processes, six particles, and the solver used a cyclic distribution of the particles.
- **6.14.** Modify the MPI version of the reduced *n*-body solver so that it uses two calls to MPI_Sendrecv_replace for each phase of the ring pass. How does the performance of this implementation compare to the implementation that uses a single call to MPI_Sendrecv_replace?
- **6.15.** A common problem in MPI programs is converting global array indexes to local array indexes and vice-versa.
 - **a.** Find a formula for determining a global index from a local index if the array has a block distribution.
 - **b.** Find a formula for determining a local index from a global index if the array has a block distribution.

- **c.** Find a formula for determining a global index from a local index if the array has a cyclic distribution.
- **d.** Find a formula for determining a local index from a global index if the array has cyclic distribution.

You can assume that the number of processes evenly divides the number of elements in the global array. Your solutions should only use basic arithmetic operators (+, -, *, /). They shouldn't use any loops or branches.

- **6.16.** In our implementation of the reduced *n*-body solver, we make use of a function First_index which, given a global index of a particle assigned to one process, determines the "next higher" global index of a particle assigned to another process. The input arguments to the function are the following:
 - a. The global index of the particle assigned to the first process
 - **b.** The rank of the first process
 - **c.** The rank of the second process
 - **d.** The number of processes

The return value is the global index of the second particle. The function assumes that the particles have a cyclic distribution among the processes. Write C-code for First_index. (*Hint*: Consider two cases: the rank of the first process is less than the rank of the second, and the rank of the first is greater than or equal to the rank of the second).

- **6.17. a.** Use Figure 6.10 to determine the maximum number of records that would be on the stack at any one time in solving a four-city TSP. (*Hint*: Look at the stack after branching as far as possible to the left).
 - **b.** Draw the tree structure that would be generated in solving a five-city TSP.
 - **c.** Determine the maximum number of records that would be on the stack at any one time during a search of this tree.
 - **d.** Use your answers to the preceding parts to determine a formula for the maximum number of records that would be on the stack at any one time in solving an *n*-city TSP.
- **6.18.** Breadth-first search can be implemented as an iterative algorithm using a **queue**. Recall that a queue is a "first-in first-out" list data structure, in which objects are removed, or *dequeued*, in the same order in which they're added, or *enqueued*. We can use a queue to solve TSP and implement breadth-first search as follows:

```
for each neighboring city
    if (Feasible(tour, city)) {
        Add_city(tour, city);
        Enqueue(tour);
        Remove_last_city(tour);
    }
}
Free_tour(tour);
/* while !Empty */
```

This algorithm, although correct, will have serious difficulty if it's used on a problem with more than 10 or so cities. Why?

In the shared-memory implementations of our solutions to TSP, we use breadth-first search to create an initial list of tours that can be divided among the threads.

- a. Modify this code so that it can be used by thread 0 to generate a queue of at least thread_count tours.
- **b.** Once the queue has been generated by thread 0, write pseudocode that shows how the threads can initialize their stacks with a block of tours stored in the queue.
- **6.19.** Modify the Pthreads implementation of static tree search so that it uses a readwrite lock to protect the examination of the best tour. Read-lock the best tour when calling Best_tour, and write-lock it when calling Update_best_tour. Run the modified program with several input sets. How does the change affect the overall run-time?
- **6.20.** Suppose the stack on process/thread A contains k tours.
 - **a.** Perhaps the simplest strategy for implementing stack splitting in TSP is to pop k/2 tours from A's existing stack and push them onto the new stack. Explain why this is unlikely to be a good strategy.
 - **b.** Another simple strategy is to split the stack on the basis of the cost of the partial tours on the stack. The least-cost partial tour goes to A. The second cheapest tour goes to new_stack. The third cheapest goes to A, and so on. Is this likely to be a good strategy? Explain your answer.
 - c. A variation on the strategy outlined in the preceding problem is to use average cost per edge. In average cost per edge, the partial tours on A's stack are ordered according to their cost divided by the number of edges in the partial tour. Then the tours are assigned in round-robin fashion to the stacks, that is, the cheapest cost per edge to A, the next cheapest cost per edge to new_stack, and so on. Is this likely to be a good strategy? Explain your answer.

Implement the three strategies outlined here in one of the dynamic loadbalancing codes. How do these strategies compare to each other and the strategy outlined in the text? How did you collect your data?

6.21. a. Modify the static MPI TSP program so that each process uses a local best tour data structure until it has finished searching. When all the processes

have finished executing, the processes should execute a global reduction to find the least-cost tour. How does the performance of this implementation compare to the static implementation? Can you find input problems for which its performance is competitive with the original static implementation?

- **b.** Create a TSP digraph in which the initial tours assigned to processes 1,2,...,comm_sz -1 all have an edge that has a cost that is much greater than the total cost of any complete tour that will be examined by process 0. How do the various implementations perform on this problem when comm_sz processes are used?
- **6.22.** MPI_Recv and each of the sends we've studied is blocking. MPI_Recv won't return until the message is received, and the various sends won't return until the message is sent or buffered. Thus, when one of these operations returns, you know the status of the message buffer argument. For MPI_Recv, the message buffer contains the received message—at least if there's been no error—and for the send operations, the message buffer can be reused. MPI also provides a **nonblocking** version of each of these functions, that is, they return as soon as the MPI run-time system has registered the operation. Furthermore, when they return, the message buffer argument cannot be accessed by the user program: the MPI run-time system can use the actual user message buffer to store the message. This has the virtue that the message doesn't have to be copied into or from an MPI-supplied storage location.

When the user program wants to reuse the message buffer, she can force the operation to complete by calling one of several possible MPI functions. Thus, the nonblocking operations split a communication into two phases:

- Begin the communication by calling one of the nonblocking functions
- Complete the communication by calling one of the completion functions

 Each of the nonblocking send initiation functions has the same syntax
 as the blocking function, except that there is a final *request* argument. For example,

```
int MPI_Isend(
     void*
                   msa
                              /* in
                   count
                              /* in
     int.
     MPI_Datatype datatype
                              /* in */.
                              /* in */,
     int
                   dest
                              /* in */,
     int.
                   tag
     MPI_Comm
                   comm
                              /* in */,
     MPI_Request* request_p /* out */);
```

The nonblocking receive replaces the status argument with a request argument. The request arguments identify the operation to the run-time system, so that when a program wishes to complete the operation, the completion function takes a request argument.

The simplest completion function is MPI_Wait:

```
int MPI_Wait(
          MPI_Request* request_p /* in/out */
          MPI_Status* status_p /* out */);
```

When this returns, the operation that created *request_p will have completed. In our setting, *request_p will be set to MPI_REQUEST_NULL, and *status_p will store information on the completed operation.

Note that nonblocking receives can be matched to blocking sends and nonblocking sends can be matched to blocking receives.

We can use nonblocking sends to implement our broadcast of the best tour. The basic idea is that we create a couple of arrays containing comm_sz elements. The first stores the cost of the new best tour, the second stores the requests, so the basic broadcast looks something like this:

When this loop is completed, the sends will have been started, and they can be matched by ordinary calls to MPI_Recv.

There are a variety of ways to deal with subsequent broadcasts. Perhaps the simplest is to wait on all the previous nonblocking sends with the function MPI_Waitall:

When this returns, all of the operations will have completed (assuming there are no errors). Note that it's OK to call MPI_Wait and MPI_Waitall if a request has the value MPI_REQUEST_NULL.

Use nonblocking sends to implement a broadcast of best tour costs in the static MPI implementation of the TSP program. How does its performance compare to the performance of the implementation that uses buffered sends?

6.23. Recall that an MPI_Status object is a struct with members for the source, the tag, and any error code for the associated message. It also stores information

on the size of the message. However, this isn't directly accessible as a member, it is only accessible through the MPI function MPI_Get_count:

```
int MPI_Get_count(
    MPI_Status* status_p /* in */,
    MPI_Datatype datatype /* in */,
    int* count_p /* out */);
```

When MPI_Get_count is passed the status of a message and a datatype, it returns the number of objects of the given datatype in the message. Thus, MPI_Iprobe and MPI_Get_count can be used to determine the size of an incoming message before the message is received. Use these to write a Cleanup_messages function that can be called before an MPI program quits. The purpose of the function is to receive any unreceived messages so that functions such as MPI_Buffer_detach won't hang.

- **6.24.** The program mpi_tsp_dyn.c takes a command-line argument split_cutoff. If a partial tour has visited split_cutoff or more cities, it's not considered a candidate for sending to another process. The Fulfill_request function will therefore only send partial tours with fewer than split_cutoff cities. How does split_cutoff affect the overall run-time of the program? Can you find a reasonably good rule of thumb for deciding on the split_cutoff? How does changing the number of processes (that is, changing comm_sz) affect the best value for split_cutoff?
- **6.25.** Pointers cannot be sent by MPI programs since an address that is valid on the sending process may cause a segmentation violation on the receiving process, or, perhaps worse, refer to memory that's already being used by the receiving process. There are a couple of alternatives that can be used to address this problem:
 - **a.** The object that uses pointers can be packed into contiguous memory by the sender and unpacked by the receiver.
 - **b.** The sender and the receiver can build MPI derived datatypes that map the memory used by the sender and valid memory on the receiver.

Write two Send_linked_list functions and two matching Recv_linked_list functions. The first pair of send-receive functions should use MPI_Pack and MPI_Unpack. The second pair should use derived datatypes. Note that the second pair may need to send two messages: the first will tell the receiver how many nodes are in the linked list, and the second will send the actual list. How does the performance of the two pairs of functions compare? How does their performance compare to the cost of sending a block of contiguous memory of the same size as the packed list?

6.26. The dynamically partitioned MPI implementation of the TSP solver uses a termination detection algorithm that may require the use of very high-precision

rational arithmetic (that is, common fractions with very large numerators and/or denominators).

- **a.** If the total amount of energy is comm_sz, explain why the amount of energy stored by any process other than zero will have the form $1/2^k$ for some nonnegative integer k. Thus, the amount of energy stored by any process other than zero can be represented by k, an unsigned integer.
- **b.** Explain why the representation in the first part is extremely unlikely to overflow or underflow.
- **c.** Process 0, on the other hand, will need to store fractions with a numerator other than one. Explain how to implement such a fraction using an unsigned integer for the denominator and a bit array for the numerator. How can this implementation deal with overflow of the numerator?
- **6.27.** If there are many processes and many redistributions of work in the dynamic MPI implementation of the TSP solver, process 0 could become a bottleneck for energy returns. Explain how one could use a spanning tree of processes in which a child sends energy to its parent rather than process 0.
- **6.28.** Modify the implementation of the TSP solver that uses MPI and dynamic partitioning of the search tree so that each process reports the number of times it sends an "out of work" message to process 0. Speculate about how receiving and handling the out of work messages affects the overall run-time for process 0.
- **6.29.** The C source file mpi_tsp_dyn.c contains the implementation of the MPI TSP solver that uses dynamic partitioning of the search tree. The online version uses the first of the three methods outlined in Section 6.2.12 for determining to which process a request for work should be sent. Implement the other two methods and compare the performance of the three. Does one method consistently outperform the other two?
- **6.30.** Determine which of the three APIs is preferable for the *n*-body solvers and solving TSP.
 - **a.** How much memory is required for each of the serial programs? When the parallel programs solve large problems, will they fit into the memory of your shared-memory system? What about your distributed-memory system?
 - **b.** How much communication is required by each of the parallel algorithms?
 - **c.** Can the serial programs be easily parallelized by the use of OpenMP directives? Do they need synchronization constructs such as condition variables or read-write locks?

Compare your decisions with the actual performance of the programs. Did you make the right decisions?

6.7 PROGRAMMING ASSIGNMENTS

6.1. Look up the classical fourth-order Runge Kutta method for solving an ordinary differential equation. Use this method instead of Euler's method to estimate

the values of $\mathbf{s}_q(t)$ and $\mathbf{s}_q'(t)$. Modify the reduced versions of the serial *n*-body solver, either the Pthreads or the OpenMP *n*-body solver, and the MPI *n*-body solver. How does the output compare to the output using Euler's method? How does the performance of the two methods compare?

- **6.2.** Modify the basic MPI *n*-body solver so that it uses a ring pass the instead of a call to MPI_Allgather. When a process receives the positions of particles assigned to another process, it computes *all* the forces resulting from interactions between its assigned particles and the received particles. After receiving comm_sz 1 sets of positions, each process should be able to compute the total force on each of its particles. How does the performance of this solver compare with the original basic MPI solver? How does its performance compare with the reduced MPI solver?
- **6.3.** We can simulate a ring pass using shared-memory:

To implement this, the main thread can allocate n storage locations for the total forces and n locations for the "temp" forces. Each thread will operate on the appropriate subset of locations in the two arrays. It's easiest to implement "notify" and "wait" using semaphores. The main thread can allocate a semaphore for each source-dest pair and initialize each semaphore to 0 (or "locked"). After a thread has computed the forces, it can call <code>sem_post</code> to notify the dest thread, and a thread can block in a call to <code>sem_wait</code> to wait for the availability of the next set of forces. Implement this scheme in Pthreads. How does its performance compare with the performance of the original reduced OpenMP/Pthreads solver? How does its performance compare with the reduced OpenMP/Pthreads solver? The reduced MPI solver?

6.4. The storage used in the reduced MPI *n*-body solver can be further reduced by having each process store only its *n*/comm_sz masses and communicating masses as well as positions and forces. This can be implemented by adding storage for an additional *n*/comm_sz doubles to the tmp_data array. How does this change affect the performance of the solver? How does the memory required by this program compare with the memory required for the original MPI *n*-body solver? How does its memory usage compare with the reduced OpenMP solver?

- **6.5.** The Terminated function in the OpenMP dynamic implementation of tree search uses busy-waiting, which can be very wasteful of system resources. Ask a system guru if your Pthreads and OpenMP implementations can be used together in a single program. If so, modify the solution in the OpenMP dynamic implementation so that it uses Pthreads and condition variables for work redistribution and termination. How does the performance of this implementation compare with the performance of the original implementation?
- **6.6.** The implementations of iterative tree search that we discussed used an array-based stack. Modify the implementation of either the Pthreads or OpenMP dynamic tree search program so that it uses a linked-list-based stack. How does the use of the linked list affect the performance?

Add a command-line argument for the "cutoff size" that was discussed briefly in the text. How does the use of a cutoff size affect the performance?

- **6.7.** Use Pthreads or OpenMP to implement tree search in which there's a shared stack. As we discussed in the text, it would be very inefficient to have all calls to Push and Pop access a shared stack, so the program should also use a local stack for each thread. However, the Push function can occasionally push partial tours onto the shared stack, and the Pop function can pop several tours from the shared stack and push them onto the local stack, if the calling thread has run out of work. Thus, the program will need some additional input arguments:
 - **a.** The frequency with which tours are pushed onto the shared stack. This can be an int. For example, if every 10th tour generated by a thread should be pushed onto the shared stack, then the command-line argument would be 10.
 - **b.** A blocksize for pushing. There may be less contention if, rather than pushing a single tour onto the shared stack, a block of several tours is pushed.
 - **c.** A blocksize for popping. If we pop a single tour from the shared stack when we run out of work, we may get too much contention for the shared stack.

How can a thread determine whether the program has terminated?

Implement this design and your termination detection with Pthreads or OpenMP. How do the various input arguments affect its performance? How does the optimal performance of this program compare with the optimal performance of the dynamically load-balanced code we discussed in the text?