E7441: Scientific computing in biology and biomedicine

Introduction to parallel computing

Vlad Popovici, Ph.D.

Fac. of Science - RECETOX

◆□▶ ◆□▶ ◆□▶ ◆□▶ ●□ ● ●

Outline





Principles of parallel computing

- Introduction
- Programming models
- Implementations

- "I think there is a world market for maybe five computers." (Thomas Watson, chairman of IBM, 1943)
- "There is no reason for any individual to have a computer in their home." (Ken Olson, founder Digital Equipment Corporation, 1977)
- "640K of memory ought to be enough for anybody." Bill Gates, chairman of Microsoft, 1981

- \sim 2500 BC: Babylon the first abacus
- ~ 100 BC: Antikythera device believed to be the first mechanical computer
- first half of the 19th century: Charles Babbage's differential machine (to tabulate polynomials) and analytical machine (only design)
- 1941: Z3 computer by Konrad Zuse: first programmable, fully automatic computing machine





source: Wikipedia

 \sim 1840 Charles Babbage produces the differential machine, a mechanical computer.



source: Wikipedia

< A > <

∃ →

1941: Z3 computer: electro-mechanical computer, \sim 2000 relays, 22-bit words, operating at 5-10 Hz.



source: Wikipedia

イロト イボト イヨト イヨト

1946: ENIAC - Electronic Numerical Integrator And Computer used initially by US Army to compute tables for artilery. Uses vacuum tubes as switches.



source: Wikipedia

1976: Cray-1 - the first successful supercomputer



Vlad Popovici, Ph.D. (Fac. of Science - RECET(E7441: Scientific computing in biology and biom

...fast forward: Tianhe-2 (top supercomputer as Nov. 2013): 33.86 PFlop/s, 3,120,000 cores; 1,024,000 GB, CPU: Intel Xeon



Moore's law

Gordon E. Moore (co-founder Intel): "Cramming More Components onto Integrated Circuits", Electronics Magazine, 1965

16-Core SPARC TS Ske-Core Core i7 2,600,000,000 Six-Core Xeon 7400 tanium 2 🖷 1,000,000,000 - Core Opteron 2400 Farium 2-100.000.000 AMD KI MD KT curve shows transistor AND K6 Transistor count 10.000.000 Pentium II count doubling every AND KS 1,000,000 A0096 100.000 66000 e 10.000 2,300 1971 1980 1990 2000 2011 Date of introduction

Microprocessor Transistor Counts 1971-2011 & Moore's Law

source: Wikipedia

A 10

Software and hardware

Software crises:

- '60s-'70s: assembly language difficult to use for large complex problems → Fortran, C: provide abstraction and portability for uniprocessors
- '80s-'90s: problems in maintaining complex systems → object-oriented programming (C++, Java)
- ~ 2000s: sequential performance lags behind Moore's law →
 programmers are oblivious to hardware better compilers, higher level
 languages, virtual machines

- parallel computing: using multiple execution units concurrently to solve a problem
- examples:
 - multi-core processors: several processors (cores) in a chip
 - shared memory processors (SMP): several processors interconnected through a shared memory
 - cluster computer: several computers interconnected through high-speed network

Issues with the traditional model: power density

(Ross: Why CPU Frequency Stalled, IEEE Spectrum Magazine, 2008)





イロト イポト イヨト イヨト

Issues cont'd: gains from implicit parallelism tapped out

Example: instruction-level parallelism. Machine instruction: decomposed into 4-stages: fetch, decode, execute and write-back



Other issues:

- increase in production costs (decrease in "chip yield")
- increase in amount of data to be processed
- Solution: explicit parallelism
 - multi-core
 - multi-processor
 - multi-machine

Principles

- identifying parallelism
- granularity: more smaller or fewer larger tasks?
- locality: data and instruction location
- load balance: aim: no lost CPU cycles
- synchronization
- overhead

Identifying parallelism

Amdahl's law:

$$S_n = \frac{T_1}{T_n} \le \frac{1}{\alpha + (1 - \alpha)/n} \le \frac{1}{\alpha}$$

where α is the fraction of the program that is strictly sequential, T_i is the execution time on *i* processors and S_i is the speed-up obtained by using *i* processors instead of 1.



Identifying parallelism

- implicit parallelism
 - hardware level: superscalar processors, multi-core, cluster computing
 - compiler level: parallelizing compilers
- explicit parallelism
 - programming language level
 - library level

Processing architectures

Flynn's taxonomy ("old way"): Single/Multiple Instruction \times Single/Multiple Data



Source: Wikipedia

Examples: SISD: mainframes; SIMD: GPUs; MISD: fault tolerant systems; MIMD: most computers nowadays

Locality: a box in a box in a box...



э

・ロト ・ 御 ト ・ ヨ ト ・ ヨ ト ・

Computing topologies







Figure 2.15 Linear arrays: (a) with no wraparound links; (b) with wraparound link.



Figure 2.16 Two and three dimensional meshes: (a) 2-D mesh with no wraparound; (b) 2-D mesh with wraparound link (2-D torus); and (c) a 3-D mesh with no wraparound.

Source: Grama - Introduction to Parallel Computing

イロト イボト イヨト イヨト

Shared memory: multicore or multi-CPU machines



Distributed memory: clusters with single CPUs nodes



Hybrid systems



- a limited number of CPUs have access to a pooled memory
- using more CPUs implies communication over network through message-passing

Hybrid systems with multicore CPUs



- extension of the hybrid model
- communication becomes increasingly complex
- many levels in the memory hierachy: cache(s), local main memory, other node's memory, etc
- you can add accelerators: e.g, GPUs
- requires a new programming model, and different communication protocols

Load balancing

- aim: distribute evenly the load (work) on all available resources...
- ...and thus minimize the time a resource is idle
- causes of imbalanced load:
 - insufficient paralelism
 - unequal task size (poor design?)

Types of parallelism

- data parallelism: each processor performs the same task on different data (h/w: SIMD, MIMD)
- task parallelism: each processor performs a different task on the same data (h/w: MISD, MIMD)
- usually, both types of parallelism are present

Example: re-annotation of a microarray chip

(embarrassingly parallel problem)

Problem: map (BLAST) each probe from a microarray against the latest version of the human genome (RefSeq).

Naive implementation on 2 CPUs:

program :	program :	program :
if CPU == 'CPU1' then idx = 1Np/2	idx = 1,,Np/2	idx = Np/2 + 1,, N
elseif CPU == 'CPU2' then idx = Np/2 + 1,,N endif	BLAST(Probes[idx])	BLAST(Probes[idx])
BLAST(Probes[idx])		

Better ways of distributing the data exists for this problem! Ex: distribute also the RefSeq...

Problem decomposition

- split the computations into concurrent tasks
- build the task-dependency graph
- there is no one-size-fits-all technique
- some methods: recursive decomposition, data-decomposition, exploratory decomposition and speculative decomposition

Recursive decomposition: example

Problem: find the minimum of a vector

```
proc serial min(A, n)
                                                       proc rec min(A, i, j)
  min = A[1]
                                                         if i == i
  for i = 2 to n do
                                                        then min = A[i]
    if A[i] < min
                                                         else
    then min = A[i]
                                                          lmin = rec min(A, i, j/2)
  end for
                                                          rmin = rec_min(A, j/2+1, j)
  return min
                                                           if Imin < rmin
end serial min
                                                           then min = Imin
                                                           else min = rmin
                                                          end if
                                                        end if
                                                         return min
                                                      end rec min
```

< ロ > < 同 > < 回 > < 回 > < 回 > <

Data decomposition: example

Matrix multiplication: $\mathbf{A} \cdot \mathbf{B} = \mathbf{C}$. Write it as

$$\begin{bmatrix} \mathbf{A}_{11} & \mathbf{A}_{12} \\ \mathbf{A}_{21} & \mathbf{A}_{22} \end{bmatrix} \cdot \begin{bmatrix} \mathbf{B}_{11} & \mathbf{B}_{12} \\ \mathbf{B}_{21} & \mathbf{B}_{22} \end{bmatrix} = \begin{bmatrix} \mathbf{C}_{11} & \mathbf{C}_{12} \\ \mathbf{C}_{21} & \mathbf{C}_{22} \end{bmatrix}$$

and distribute the four tasks:

Task 1:
$$C_{11} = A_{11}B_{11} + A_{12}B_{21}$$

Task 2: $C_{12} = A_{11}B_{12} + A_{12}B_{22}$
Task 3: $C_{21} = A_{21}B_{11} + A_{22}B_{21}$
Task 4: $C_{22} = A_{21}B_{12} + A_{22}B_{22}$

Other decompositions

- exploratory decomposition: decompose the search space for the solution and search for a solution in each subspace; then choose among the solutions
- *speculative decomposition*: launch alternative computation branches in parallel while waiting for input for deciding which branch to use
- hybrid decompositions

Mapping techniques

- problem decomposition \rightarrow tasks
- the tasks need to be allocated (mapped) to processors/processes
- objective: minimize the execution time
- *overheads*: time spent for everything else but actually solving the problem:
 - inter-process interaction synchronization and control
 - time spent being idle poor load balancing
- reduce the process inter-dependencies and communication: e.g. maximize data locality
- improve load balancing
- reduce blocking operations

Implementations on multihtread/multicore machines

- POSIX threads (pthreads): OS-level paralelism.
 - threads: lightweight processes
 - the same program runs on single- or multi-core machines
 - OS has the responsibility of mapping the threads
 - needs low-level programming, dedicated library
- OpenMP: built on top of pthreads for SIMD-kind of parallelism
 - implemented through compiler directives
 - easier to use than pthreads
 - performance depends on compiler's 'intelligence'

OpenMP: how does it look like? $(\sum_i a_i b_i)$

```
double a[N];
  double sum = 0.0;
  int i. n. tid:
#pragma omp parallel shared(a) private(i)
    tid = omp_get_thread_num();
    /* Only one of the threads do this */
#pragma omp single
      n = omp get num threads(); printf("Number_of_threads_=_%d\n", n);
    /* Initialize a */
#pragma omp for
    for (i=0; i < N; i++) {
      a[i] = 1.0:
    }
    /* Parallel for loop computing the sum of a[i] */
#pragma omp for reduction (+:sum)
    for (i=0; i < N; i++) {
      sum = sum + (a[i]);
  } /* End of parallel region */
```

・ロト ・同ト ・ヨト ・ヨト

Implementations on distributed-memory systems

• MPI: Message Passing Interface

- de facto standard for distributed memory programming (clusters)
- data must be manually decomposed
- use special libraries
- based on sending and receiving messages: data and synchronization
- PVM: Parallel Virtual Machine
 - previous library for cluster programming
 - based on message-passing principle
 - supplanted by MPI

MPI: how does it look like?

```
#include <mpi.h>
int main(int argc, char *argv[])
{
    int numprocs, myid;
    MPl_Init(&argc,&argv);
    MPL_comm_size(MPL_COMM_WORLD,&numprocs);
    MPL_comm_rank(MPL_COMM_WORLD,&myid);
    /* print out my rank and this run's PE size */
    printf("Hello_from_%d_of_%d\n", myid, numprocs);
    MPI_Finalize();
}
```

・ロト ・同ト ・ヨト ・ヨト

Implementations in R

- parallelism came as an after thought
- target: massive data applications
- tries to bring to R some of the libraries existing to other languages
- snow: for traditional clusters, supports PVM, MPI,...; is portable (UNIX, Windows)
- multicore: targets multi-core/-CPU machines; simple; does not run on Windows; does not handle parallel RNGs
- parallel: snow+multicore in new R (>=2.14); strange interactions with OS
- *R+Hadoop*: based on Hadoop cluster
- *RHIPE*: based on Hadoop, targets map-reduce operations
- Segue: APPLY-like calculations on Hadoop clusters, using Amazon's Elastic MapReduce

Questions?

Vlad Popovici, Ph.D. (Fac. of Science - RECET(E7441: Scientific computing in biology and biom

글 🖌 🔺 글 🕨

A 10