Administrivia

 Class Friday? (No — almost half the class is going to the ACM programming competition. Good luck!)

 Homework 4 to be on Web soon. Briefly — write a parallel version of Conway's Game of Life.

Slide 1

• Courses next term:

CSCI 3294 ("Unix Power Tools") will be similar to course last spring, syllabus/notes from under "Old course materials" on my Web page.

CSCI 3394 ("Topics in Parallel Computing") will be a follow-on to CSCI 3366.

Molecular Dynamics Example — Recap

- Previously discussed the problem (what we're computing and how) and sketched out how to decompose/analyze it.
- Also decided on overall algorithm structure of *Task Parallelism*. Pseudocode in next slide, again.

Pseudocode for Non-Bonded Force Computation

```
function non_bonded_forces (N, Atoms, neighbors, Forces)
  Int const N // number of atoms
  Array of Real :: atoms (3,N) //3D coordinates
  Array of Real :: forces (3,N) //force in each dimension
  Array of List :: neighbors (N) //atoms in cutoff volume
  Real :: forceX, forceY, forceZ

loop [i] over atoms
  loop [j] over neighbors (i)
        forceX = non_bond_force(atoms(1,i), atoms(1,j))
        forceY = non_bond_force(atoms(2,i), atoms(2,j))
        force(2,i) += forceX; force(1,i) -= forceX;
        force(2,i) += forceY; force(2,j) -= forceY;
        force(3,i) += forceZ; force(3,j) -= forceZ;
        end loop [j]
  end loop [i]
end function non bonded forces
```

Slide 3

Molecular Dynamics and Task Parallelism

- How to define tasks so we get "enough but not too many"?
 One task per atom pair is too many; one task per atom is probably right.
- How to manage data dependencies (if any)?
 Dependency involving forces array potentially any UE can write to any element, if we exploit symmetry resulting from Newton's third law. But computation is accumulation/reduction, so just give each UE a local copy and combine all copies at end.
- How to assign tasks to UEs? statically (at compile time) or dynamically (at runtime)?

Work per task can vary, since how many atoms are "close" varies. Decide at next level.

Molecular Dynamics and Task Parallelism, Continued

• How to structure program (i.e., which of *Supporting Structures* patterns to use)?

Obvious choices here *Loop Parallelism* and *SPMD*. Decide based on target platform.

Slide 5

Design of Program for Molecular Dynamics

- Finally, we turn the design into code, probably using patterns from *Supporting Structures* design space, and possibly some information/understanding from *Implementation Mechanisms*.
- Based on previous design steps, consider Loop Parallelism and/or SPMD.
 Decide based mostly on target platform. Tables in section 5.3 should be helpful . . .

Program Structures Versus Algorithm Structures

Geomet-Event-Recur-Task Divide Based ric Paraland Pipeline sive Decom-Coordilelism Conquer Data position nation SPMD **** **** *** *** Loop Parallelism Mas-*** ** ter/Worker **** *** *** **** Fork/Join

Slide 7

Program Structures Versus Programming Environments

| | MPI | OpenMP | Java |
|---------------------|-----|--------|------|
| SPMD | *** | *** | ** |
| Loop Parallelism | * | **** | *** |
| Mas- ter/Worker | *** | ** | *** |
| Fork/Join | | *** | *** |

Molecular Dynamics and *SPMD* — Key Design Decisions

 Only parallelize computation of non-bonded forces, since that's most of the computational load.

• Keep a copy of the full force and coordinate arrays on each node.

parallel and communicate the results).

- Have each UE redundantly update positions and velocities for the atoms (i.e., assume it's cheaper to redundantly compute these terms than to do them in
- Have each UE compute its contributions to the force array and then combine (or reduce) the UEs' contributions into a single global force array copied onto each UE.

Molecular Dynamics and SPMD — Code

- Slightly more detailed sequential pseudocode in figure 5.7 (p. 134).
- MPI main pseudocode in figure 5.8 (p. 135). Compare to figure 5.7.
- Pseudocode for computation of non-bonded forces in figure 5.9 (p. 136).
 Compare to sequential pseudocode in figure 4.4 (p. 72).
- Pseudocode for computation of neighbor list in figure 5.10 (p. 137). Notice that we exploit the symmetry resulting from Newton's third law.
- A remaining decision how to distribute atoms among UEs. Cyclic distribution is easy and will probably work okay. If not, could do something more complex — define "owner-computes filter" — boolean function of ID and loop iteration.
- Notice that we could do this in OpenMP too.

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Molecular Dynamics and *Loop Parallelism* — Key Design Decisions

- Parallelize computationally intensive loop only (the one for non-bonded forces).
- Figure out what to do about shared variables:
 - Make temporary variables used inside loop private.
 - Make forces array a reduction variable.
- Decide how to map iterations onto UEs. Dynamic schedule works well if available (as it is in OpenMP).
- OpenMP-based pseudocode as shown in figure 5.25 (p. 161) and following pragma omp directives). Compare to pseudocode in figure 4.4 (p. 72).

Minute Essay

• Do you have thoughts about a possible project? the two broad categories are (1) parallel applications, and (2) programs to determine something about a hardware/software environment (e.g., speed of message passing).

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