### Introduction to Parallel Computing (CMSC416 / CMSC616)

# **Parallel CSE Applications**



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## Announcements

## Quiz 3 has been posted



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## Contact me

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## • CMSC416: If you are an undergrad interested in participating in International Student









motions



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## Calculate trajectories of atoms and molecules by solving Newton's equations of

- motions
- Force calculations
  - Bonded interactions: bonds, angles, dihedrals
  - Non-bonded interactions: van der Waal's and electrostatic forces



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- Simulation step: ~I femtosecond (10-15 s)
- Used for drug design, materials design







# Sequential Algorithm

- At every step, calculate forces on each atom
  - Calculate bonded and short-range forces every step
  - Calculate long-range non-bonded forces every few time steps (using PME or P3M etc.)
- Particle mesh Ewald (PME) summation:
  - Calculate long-range interactions in Fourier space
- Calculate velocities and new positions
- Repeat ...











## Atom decomposition:

• Partition the atoms across processes





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- Force decomposition:
  - Distribute the force matrix to processes
  - Matrix is sparse and non-uniform





- Atom decomposition:
  - Partition the atoms across processes
- Force decomposition:
  - Distribute the force matrix to processes
  - Matrix is sparse and non-uniform
- Spatial decomposition:
  - Assign a region of the 3D simulation space to each process







# Hybrid parallelization

- Hybrid of spatial and force decomposition
- Decouple assignment of data and work to processes
- Distribute both atoms and the force calculations to different processes









# Neutral territory (NT) methods

## Desmond's mid-point method





### SC23 Test-of-time award:

Shaw DE, RO Dror, JK Salmon, et al. 2009. "Millisecond-scale molecular dynamics simulations on Anton," In Proceedings of the Conference on High Performance Computing Networking, Storage and Analysis (SC09), Portland, OR, USA, pp. 1-11, doi: 10.1145/1654059.1654126



# Particle mesh Ewald

- Replace direct force calculations by:
  - Calculate short-range forces in real space
  - Calculate long-range forces in Fourier space
- Create a 3D mesh/grid representing charge densities of atoms
  - Compute a 3D Fast Fourier Transform (FFT)
- FFT computes the discrete Fourier transform (DFT) or inverse DFT
  - Reduces the complexity from  $O(N^2)$  to  $O(N \log N)$











• Bring all the data to one process





XA

- Bring all the data to one process
- ID or slab decomposition





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## 2D or pencil decomposition





# Measles killed 200,000 in 2020 alone!

### LARGE MEASLES OUTBREAKS Global trends in the estimated number of deaths The epidemic in the Democratic Republic of the Congo caused by TB and AIDS, 2000-2018 is the largest single-nation outbreak for decades. Shaded areas represent uncertainty intervals AIDS-related deaths Ukraine Tuberculosis deaths among (2017 - 20)deaths per year HIV-negative people 1.5 >115,000 cases >41 deaths Philippines ď Democratic (2018 - 20)Millions Republic 71,170 cases of the Congo 0.5 841 deaths (2019 - 20)Tuberculosis deaths among 348,158 cases\* people living with HIV 6,504 deaths 2000 2009 2018 Madagascar (2018-20) For AIDS, the latest estimates of the number of deaths in 2018 that have been published. 244.675 cases by UNAIDS are available at http://www.unaids.org/en/. For TB, the estimates for 2018 are >1,000 deaths these published in the Global Tuberculosis Report 2019. Deaths from TB among people living with HIV are officially classified as deaths caused by HIV/AIDS in the International Classification of Diseases. Source: Global tuberculosis report 2019. Geneva: World Health Organization, 2019. Data from March 2020. onature \*Suspected, not yet officially reported to WHO. Predictions say that 1.66 million

https://www.nature.com/articles/d41586-020-01011-6



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### people died of tuberculosis in 2020

### **TERRIBLE TOLL**

By the end of July 2020, there had been 646,949 COVID-19 deaths worldwide. In the 32 countries and 4 major cities with relevant data, there were more excess deaths than COVID-19 deaths, suggesting that some COVID-19 deaths are misclassified or that other causes of death have also risen.

Deaths attributed to COVID-19 Excess deaths



\*Cumulative deaths from outbreak onset to latest available data, as of 18 August 2020.

onature

https://www.nature.com/articles/d41586-020-02497-w



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# Societal challenge

- Controlling the spread of infectious diseases is important
- Computational and mathematical modeling of epidemics important to assist governments in responding to outbreaks
- Made challenging due to:
  - increased and denser urbanization
  - increased local and global travel
  - increasingly immuno-comprised population





# **Approach: individual-based simulation**

- Agent-based modeling to simulate epidemic diffusion
- Models agents (people) and interactions between them
- People interact when they visit the same location at the same time
- These "interactions" between pairs of people are represented as "visits" to locations
- Use a bi-partite graph of people and locations or a people-people interactivity graph





# Serial algorithm

- At each timestep (typically a day):
  - Determine which people visit which locations
  - "Send" people to those locations
  - At each location "interactions" happen and transmission happens
  - Update people's states at the end of the day and continue
- people's susceptibility, movements etc.



• Interventions (vaccinations, school closures) can be added on certain days to change

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Hybrid time-stepped and discrete-event simulation



https://sitn.hms.harvard.edu/flash/special-edition-on-infectiousdisease/2014/an-introduction-to-infectious-disease/ Abhinav Bhatele (CMSC416 / CMSC616)



Hybrid time-stepped and discrete-event simulation





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Hybrid time-stepped and discrete-event simulation







## Hybrid time-stepped and discrete-event simulation

while d < num days: for each person: Send visit messages to locations

for each location: Process all visit messages Run discrete event simulation Send interaction messages

for each person: Process interactions Update disease state



Normal (5.1) : 35%





# **Parallel simulation is challenging**

- Size and scale of the social contact network (6 billion agents for a global simulation)
  - Unstructured networks and complicated dependencies lead to high communication cost
- Individuals and their behaviors are not identical
- Co-evolving epidemics, public policies and agent behaviors make it impossible to apply standard model reduction techniques





# **Parallel implementation: Loimos**

- All the people and locations are distributed among all processes
- DES computation can be done locally in parallel
- Communication when sending visit and infection messages
- Uses Charm++, a message-driven model







# **Application software stack**

- Parallel programming model / runtime:
  - MPI, OpenMP, Charm++, CUDA, ...
- Libraries
  - Data and visualization libraries (mesh management, simulation output)
  - I/O libraries
  - Math/numerical libraries
  - Graph partitioning, load balancing ...





# Why use libraries?

- No need to reinvent the wheel
  - Libraries are highly optimized, have fewer bugs
- Avoids significant effort to write, optimize and maintain code
- Makes code more portable





# **Popular Libraries**

- Data/visualization and I/O libraries
  - I/O: HDF5, pNetCDF, ADIOS
- Numerical libraries:
  - Fast Fourier transforms: FFTW
  - Dense linear algebra: BLAS, LAPACK, Intel MKL
  - Solvers for sparse systems: Hypre, PETSc, Trilinos
- Graph partitioning/load balancing:
  - METIS, Scotch, Zoltan, Chaco



https://events.prace-ri.eu/event/176/contributions/38/attachments/154/305/HPC\_libraries.pdf



# **Domain-specific languages/frameworks**

- Structured grids: SAMRAI, Chombo, AMREx
- Unstructured grids: MFEM, Quinoa







# The *n*-body problem

- Simulate the motion of celestial objects interacting with one another due to gravitational forces
- Naive algorithm:  $O(n^2)$ 
  - Every body calculates forces pair-wise with every other body (particle)







- Naive approach: Assign n/p particles to each process
- Other approaches?





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https://en.wikipedia.org/wiki/Z-order\_curve





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http://datagenetics.com/blog/march22013/ https://en.wikipedia.org/wiki/Z-order\_curve





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http://charm.cs.uiuc.edu/workshops/charmWorkshop2011/slides/CharmWorkshop2011\_apps\_ChaNGa.pdf

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## • Let us consider a two-dimensional space with bodies/particles in it





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# **Different parallelization methods**

- Tree codes: Barnes-Hut simulations
- Fast multipole methods (FMM): Greengard and Rokhlin
- Particle mesh methods
- Particle-particle particle-mesh (P<sup>3</sup>M) methods





# **Barnes-Hut simulation**

- Represent the space containing the particles as an oct-tree
- Pairwise force calculations for nearby particles
- For tree nodes that are sufficiently far away, approximate the particles in the node by a single large particle at the center of mass
- O(N logN) algorithm





https://en.wikipedia.org/wiki/Barnes-Hut simulation





# Fast multipole methods

- Use multipole expansion for distant particles
- distant particles are similar
- Reduces the time complexity further to O(n)



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### • Takes advantage of the fact that for nearby particles, multipole-expanded forces from



# Particle-particle particle-mesh methods

- Explicit calculation of forces on nearby particles
- Fourier-based Ewald summation for calculating potentials on a grid
- Smoothed particle hydrodynamics









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