

Sackler Course BMSC-GA 4448

# High Performance Computing in Biomedical Informatics

Class 2: Friday February 14<sup>th</sup>, 2014  
2:30PM – 5:30PM

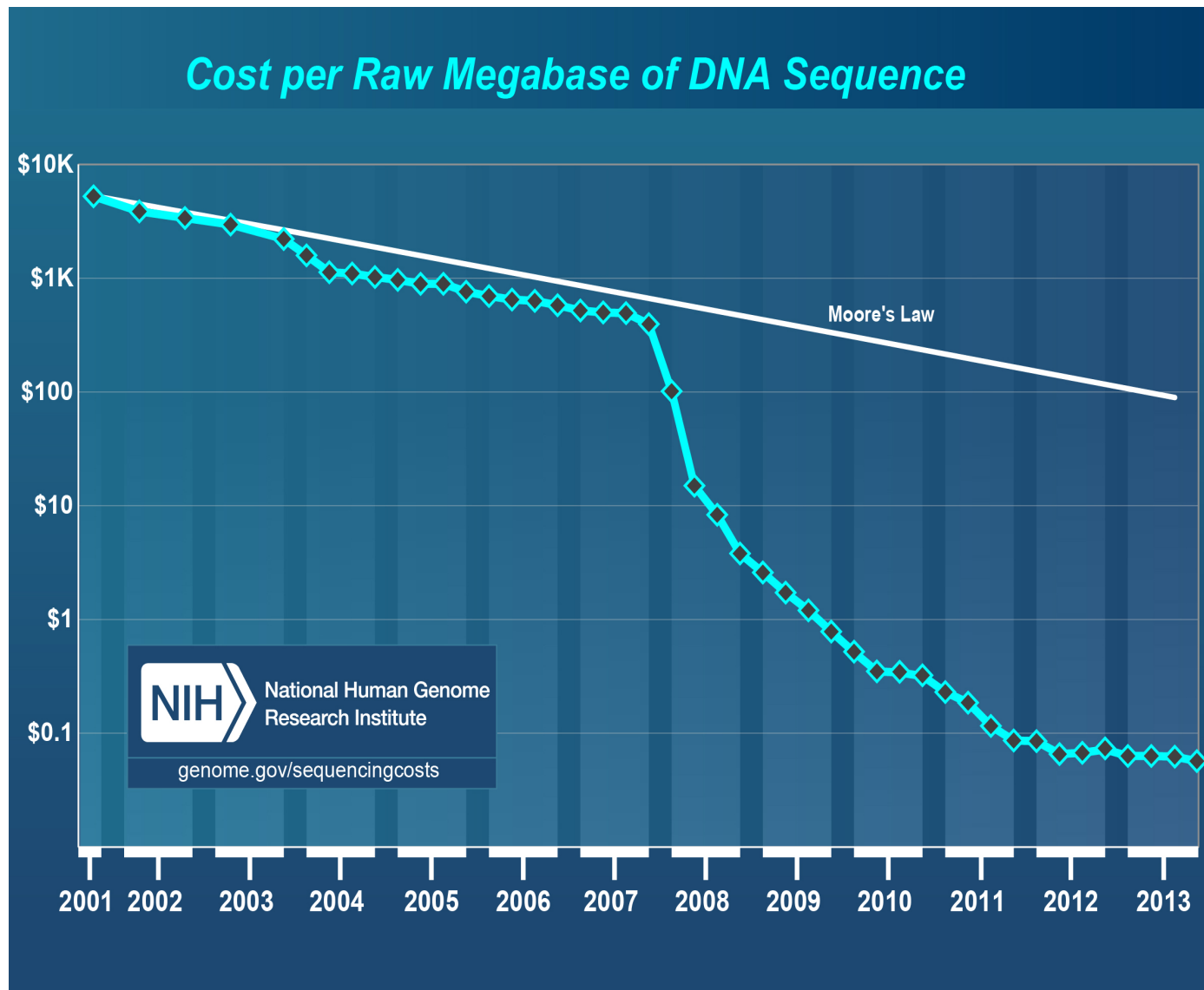
## AGENDA

- Recap 1<sup>st</sup> class & Homework discussion.
- Fundamentals of Parallel Computing: Multi-Processing
- Unix Hands-On:
  - Data sharing and File Permissions
- Student Projects

# Single-Processor Performance Growth

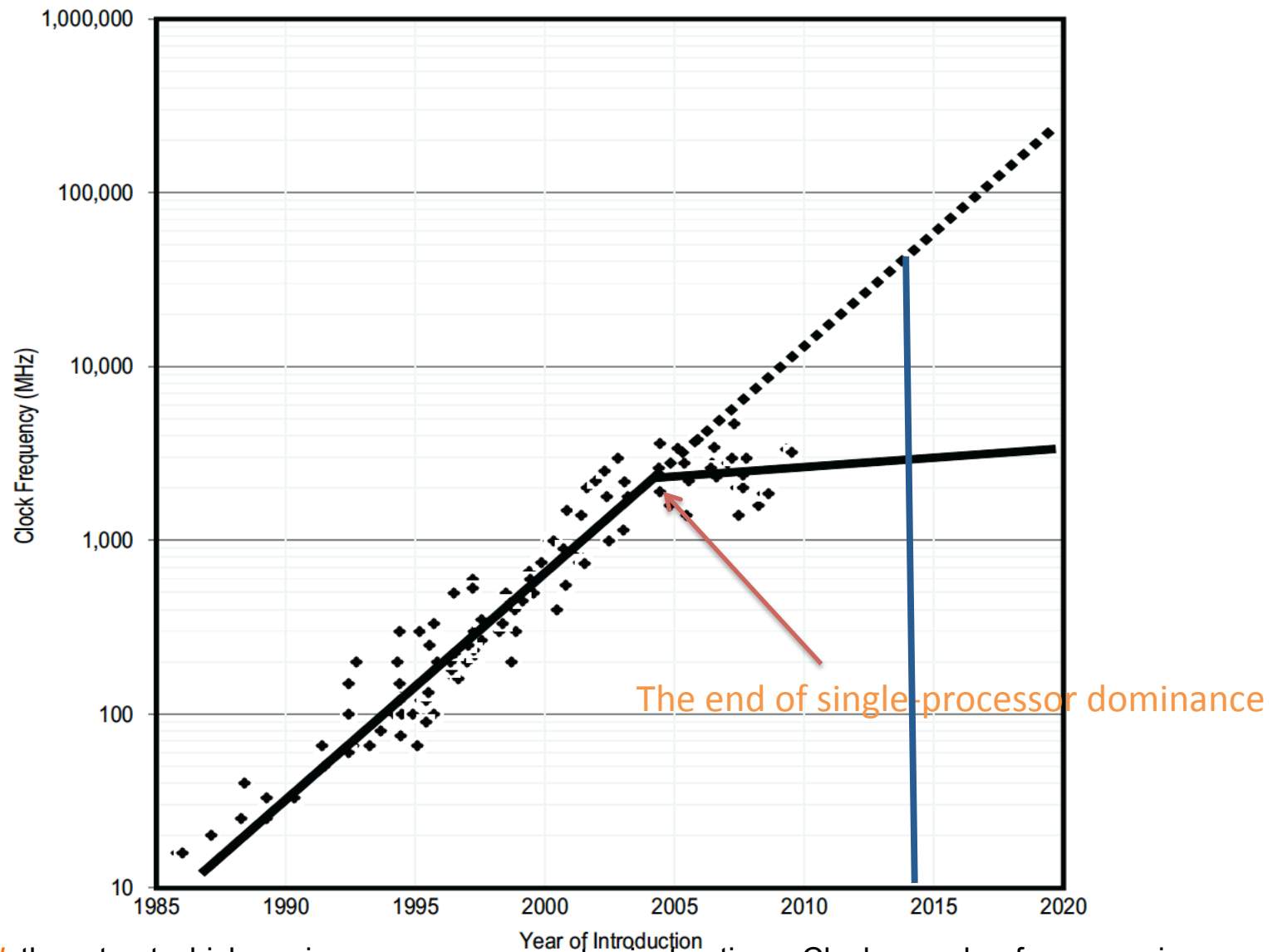


- **Moore's Law:** Processor performance doubles every 18 months  
(Gordon E. Moore, Intel co-founder, 1965)
  - (1) Increased transistor density
  - (2) Higher clock rates
- Exponential growth of Single-Processor performance for over 40 years
  - 1971: 0.7 MHz (Intel 4004)
  - 2001: 1,600.0 MHz
- Maintain compatibility of specific instruction sets.
- Exponential growth in computing power transformed the way we do business, interact and socialize.



Advances in DNA sequencing technologies vs. Moore's Law

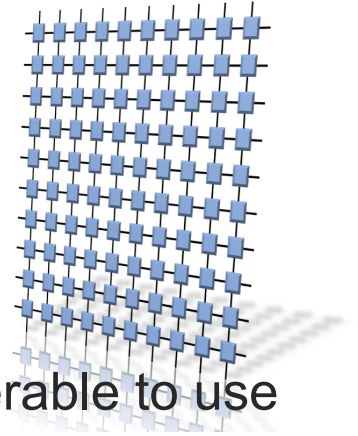
# Single-Processor Clock Frequency Growth



**Clock speed:** the rate at which a microprocessor executes instructions. Clock speed or frequency is expressed in MHz or GHz.

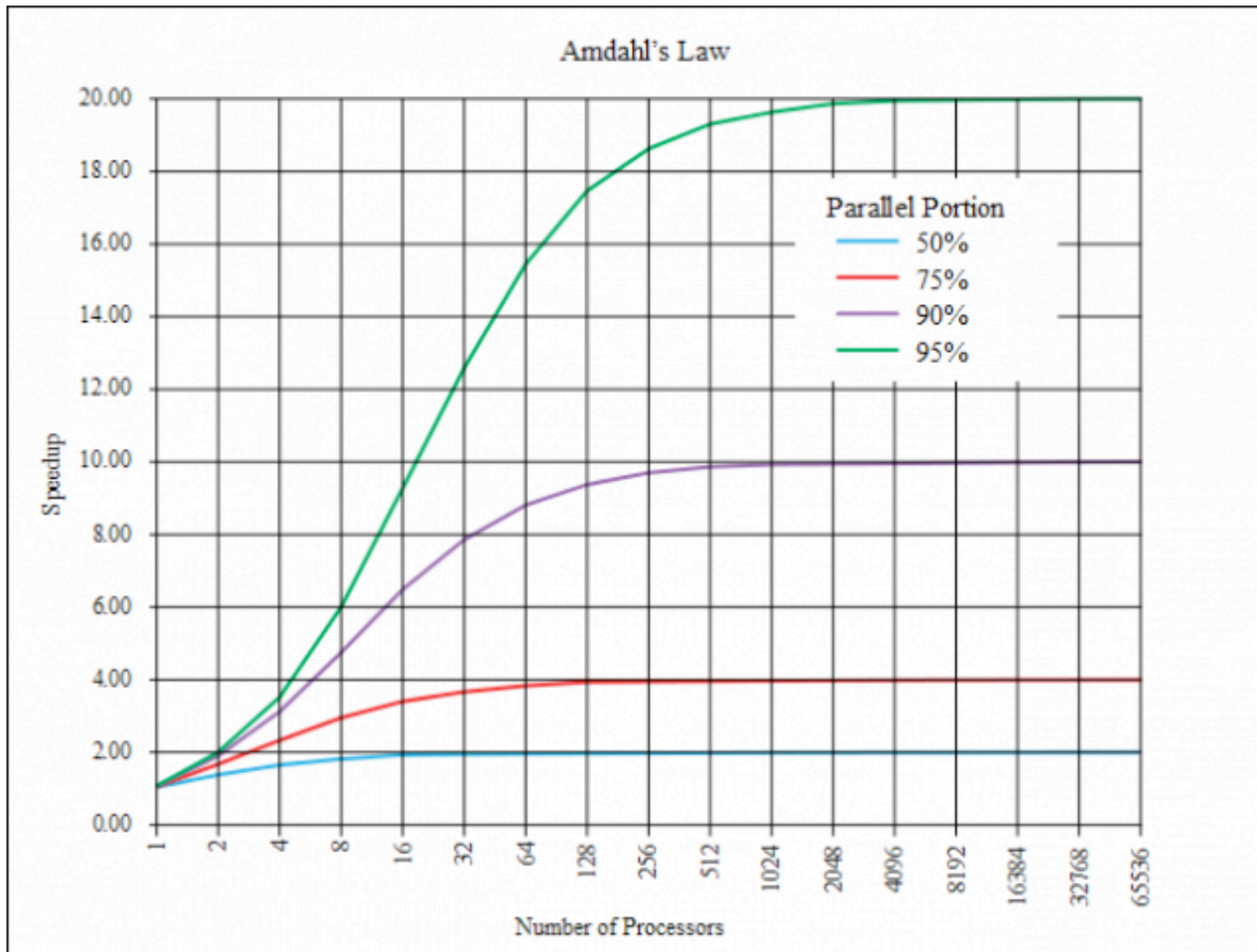
# Multi-core Processors

$$\text{Power} \sim \text{Frequency}^3$$



- Horizontal growth: **More processing cores per chip**. It is preferable to use many slower devices than a single ultrafast device.
- **Energy efficient cores** to turn more transistors into more performance.
- **New challenges for software developers:**  
Sequential (serial) programs do not automatically benefit from the use of multiple cores. Examples.
- **Parallel computing** enters **mainstream computing**.
- Can innovation in **software and algorithms** enable ongoing performance growth?

3 GHz + 3 GHz < 6 GHz



In a parallelized program, performance is limited by the amount of serial code

$$SpeedUp = \frac{1}{(1 - P) + \frac{P}{N}}$$

P: Parallel Portion

N: Processing Elements

**Cache:** memory set aside as a specialized buffer storage memory that is continually updated ; Used to optimize data transfers between system elements with different characteristics.

**Cache** memory bridges the speed gap between the processor and the memory.

Figure 2a. Multi-core processor separate L2

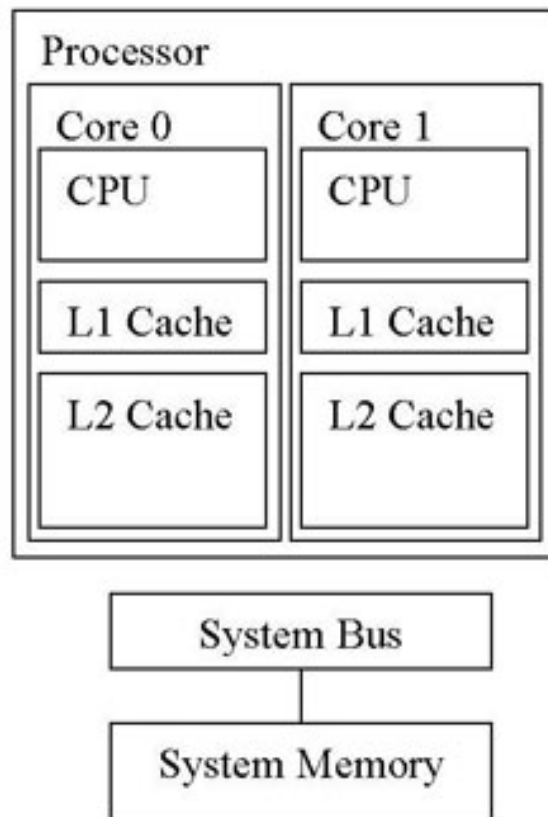
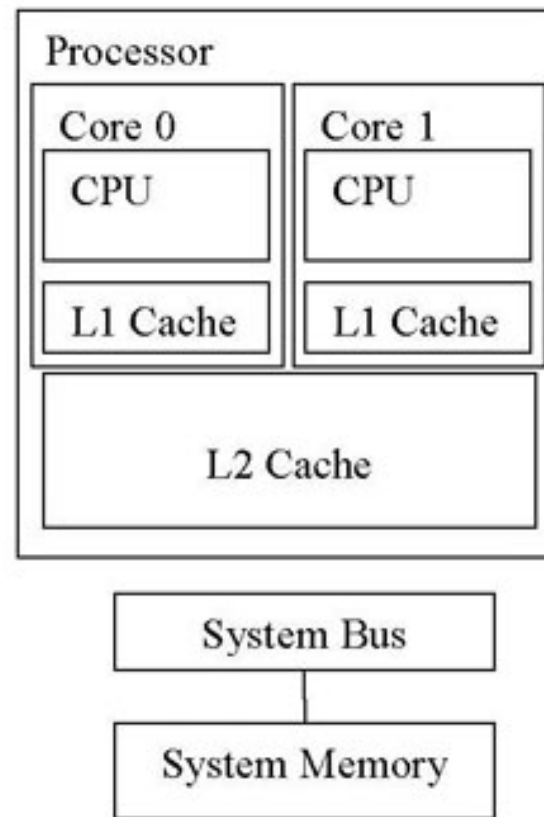
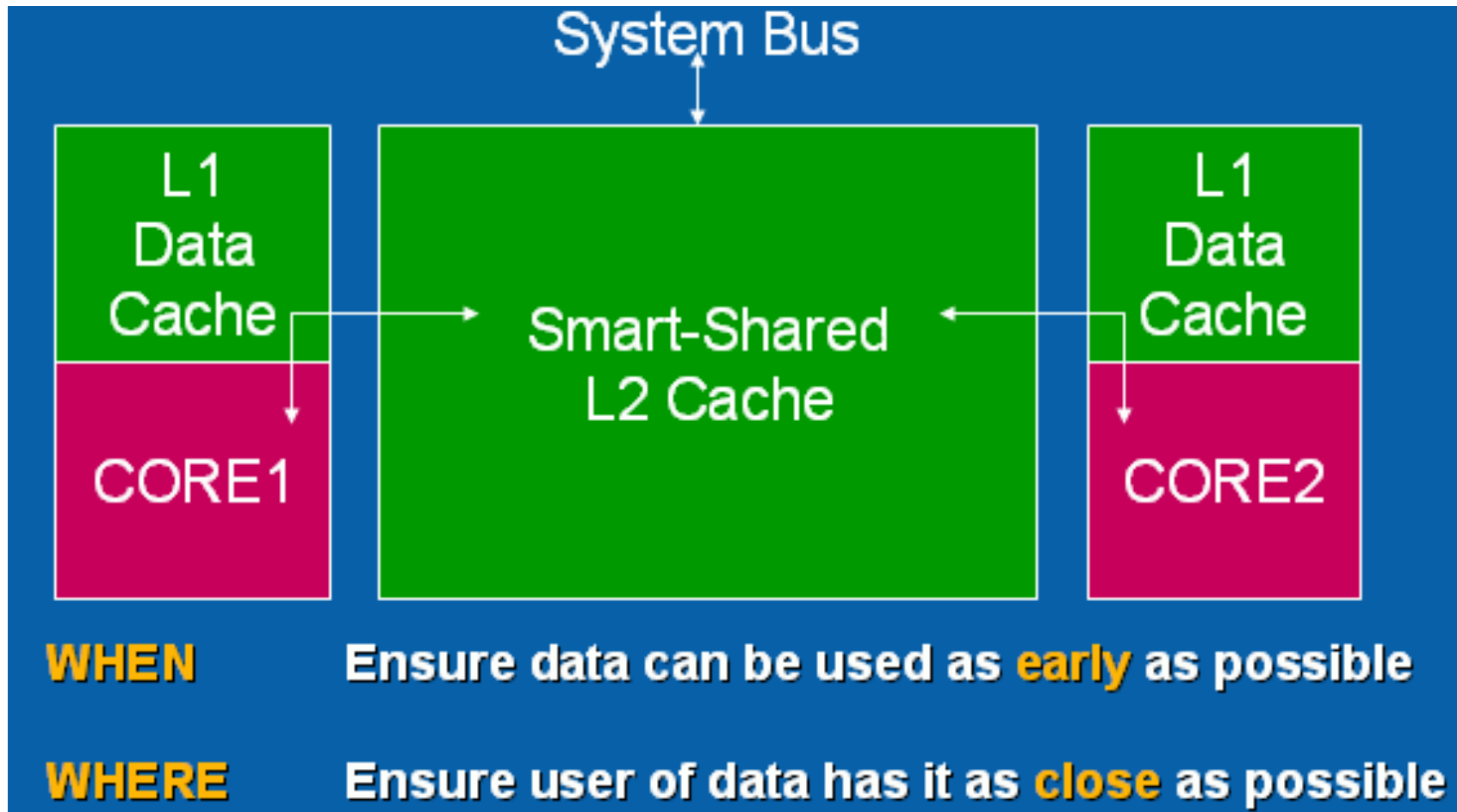


Figure 2b. Multi-core processor shared L2





Advantages of smart/shared L2 Cache:

- Size allocated per core can be dynamically adjusted with the potential of the total level 2 cache being available to applications that require it.
- Sharing of items between cores. Threads on separate cores can synchronize through the faster level 2 cache as opposed to main memory or the next level of cache.



## Questions/Discussion

### BREAK

- How many CPUs, cores are in your laptop? A cluster compute node?
- What is the clock speed of the processor on your laptop? A cluster compute node?
- How much system RAM is in your laptop? A cluster compute node?
- How much L2 cache is on the processors of your laptop? A cluster compute node

Hint: On the cluster, use:

`cat /proc/cpuinfo`

`cat /proc/meminfo`

# Computer Processes

Problem: We need to find out how many words are in a book.

*Serial Job* : A single task (*a process*)

Serial pseudo-code:

```
sum = 0;

Open (BOOK);

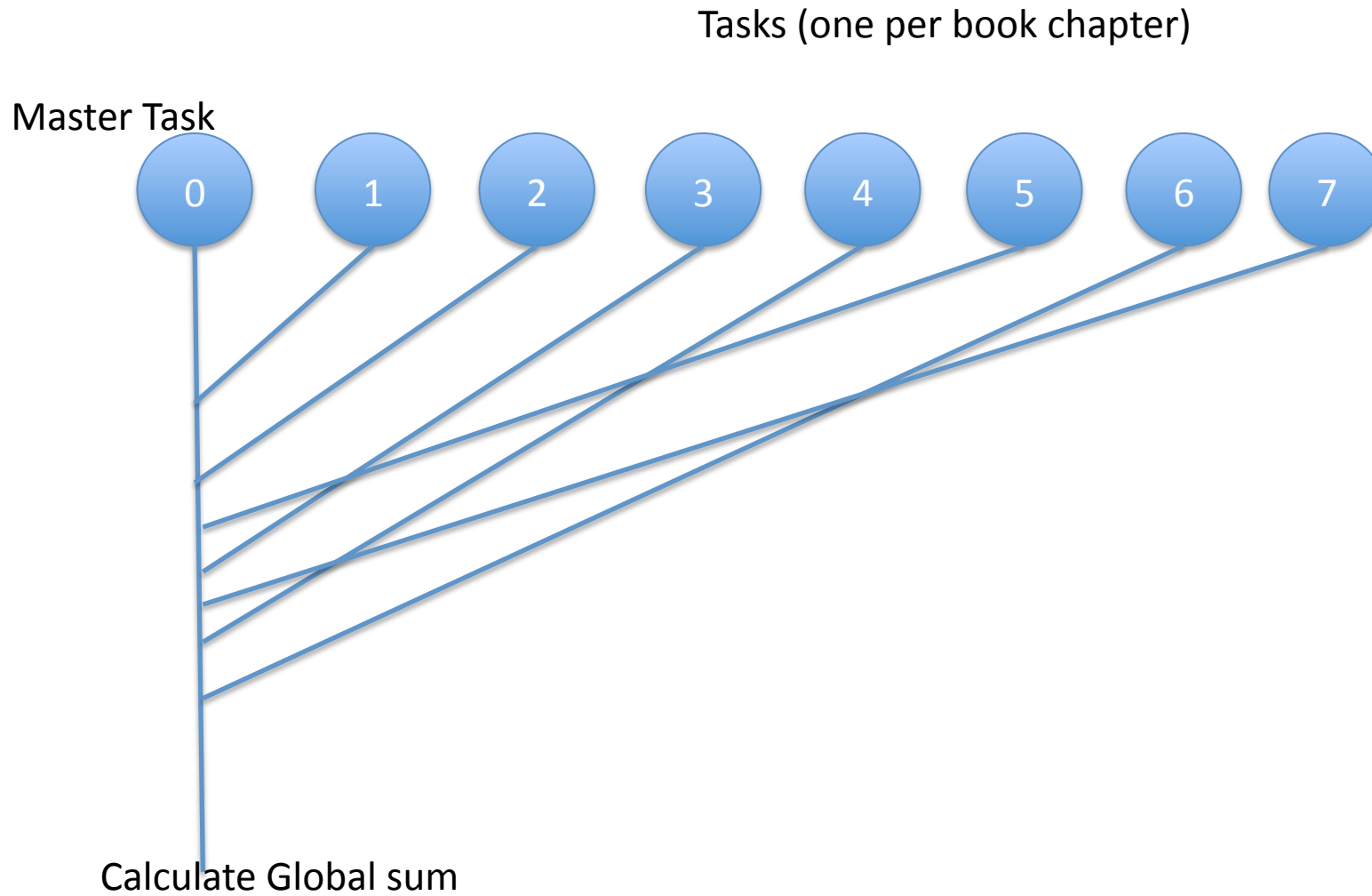
while (in BOOK) {
    Read( something, BOOK);
    if (something.is.word) sum = sum + 1;
}

Close (BOOK);

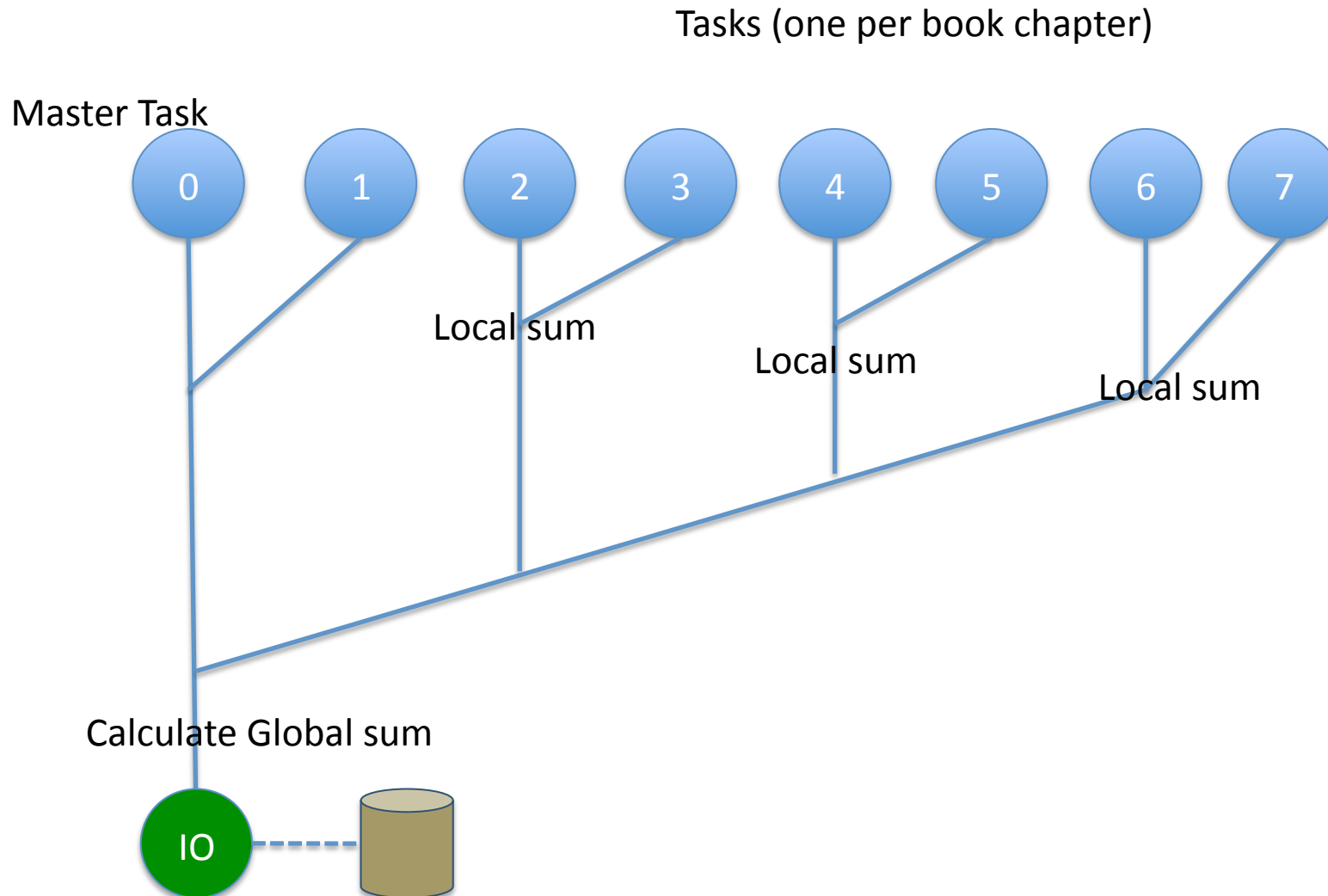
print (sum);
```



Problem: We need to count how many words are in a book



Problem: We need to count how many words are in a book



## Parallelism Considerations: **Communications**

Splitting a large problem into a large number of tasks:

decreases the execution time attributable to **computation**, but  
increases the execution time attributable to **communication**.

**(1) Communication Latency**

**(2) Communication Bandwidth**

**(3) Communication Pattern** (nearest-neighbor, broadcast, one-to-all, etc.)

*NetPerf*, *Iperf* (Open Source tools) provide accurate measurements of network latency and bandwidth.

The Unix *ping* command:

Used commonly to check network connectivity, some basic network characteristics, find out the IP address of a remote server, etc.

Try on your laptop's terminal:

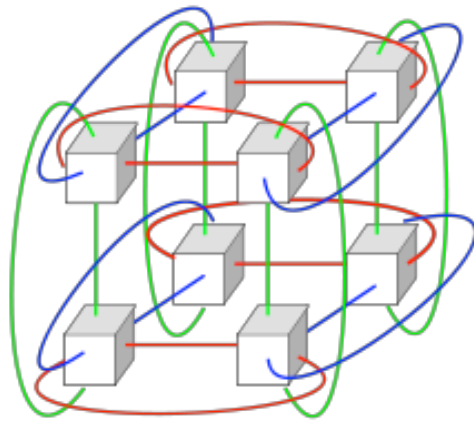
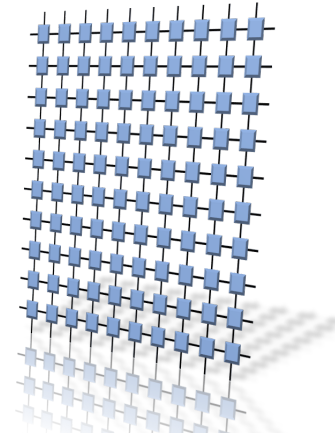
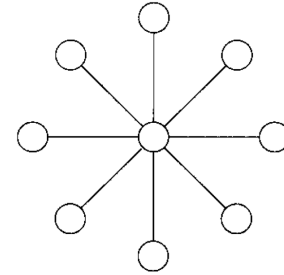
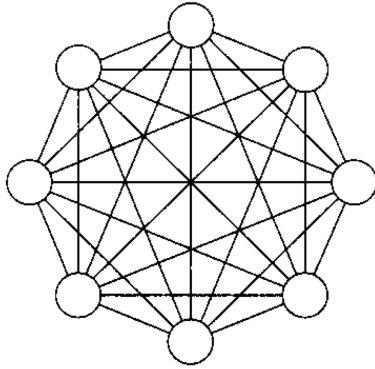
➤ *ping phoenix.med.myu.edu*

To Exit, press Control-C

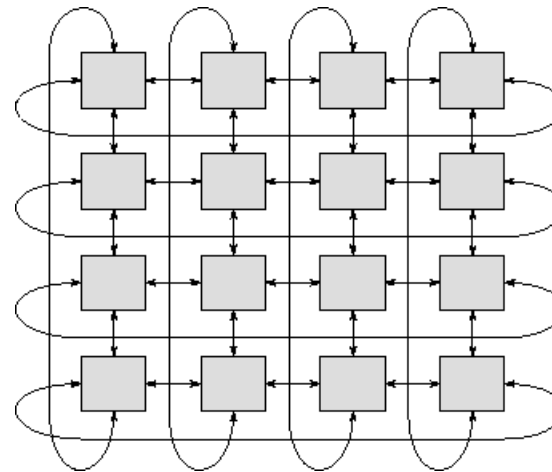
➤ *ping -c 5 google.com*

➤ *man ping*

## Parallelism Considerations:



2 x 2 x 2 Torus



2 x 2 Torus

## Communications/ Node Interconnects

## Parallelism Considerations: *I/O Operations*

Many tasks making up a parallel job may need to read/write to the same file system, at the same time, over a conventional network. This can become a serious bottleneck.

- (1) **Reduce I/O** when possible.
- (2) **Parallel file systems** (GPFS, Lustre, etc.) can be helpful.
- (3) **Stage-in, Stage-out.** On a cluster environment, consider writing/reading from local (on the node) file system, rather than a shared file system over the network.
- (4) **Create unique filenames** using filename extensions that correspond to a task Id is a good practice.
- (5) **Separate IO from message-passing:** The IO network is usually separate from the inter-node network.
- (6) **IO-Nodes:** Only a small number of nodes have access to the external file system. The rest of the nodes (compute nodes) will need to send IO operations to an IO node. **P-set ratio.**  
What is the p-set ratio of the HPC cluster?



# High Performance Computing in Biomedical Informatics

## Class2 Homework

(1) Read cloud-related docs at :

[https://genome.nyumc.org/hpcf/wiki/HPC\\_course/spring\\_2014#Session\\_3](https://genome.nyumc.org/hpcf/wiki/HPC_course/spring_2014#Session_3)

(2) Create a directory, under your home directory on phoenix, called *HPCCache*

- Put a test file, named *dataFile* , in the *HPCCache* directory so that *hpcclass* group members can view the contents of the *dataFile* and also copy the file.
- Allow members of the *hpcclass* group to write files in the *HPCCache* directory.
- Create a directory under your home directory, called *HPCTest* where members of the *hpcci* group can write files.

(3) Write a Shell script that runs the *uptime* command on the first 10 nodes of the HPC cluster and sorts the output based on the *1-min* average load on each node.

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# High Performance Computing in Biomedical Informatics

Class 3: Thursday February 20<sup>th</sup>, 2014, 2:30PM

## AGENDA

- Cloud Computing (presented by Dr C. Krampis, JCVI)