

Active Learning: Lunarc/Iridium, batch systems

COMPUTE RESEARCH SCHOOL COURSE NTF004F



Basic concepts of parallelism



Outline

- Multi-core computing in a nutshell
 - motivation, terminology, difficulties
- Handy tools for remote sessions
 - ssh, screen
- Riding on the clusters
 - Batch system basics
- Task farming
 - Scaling experiment





What is parallel computing?

- <u>Traditional computing</u>: serial execution of a single stream of instructions on a single processing element
- Parallel computing: simultaneous execution of stream(s) of instructions on multiple processing elements
 - Non-sequential execution of a computational task
 - (part of) the problem solved by simultaneous subtasks (processes)
 - Relies on the assumption that problems can be divided (decomposed) into smaller ideally independent ones that can be solved **parallel**





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What is parallel computing (cont.)?

- **Parallelism levels** ("distance" among the processing elements):
 - Bit and Instruction level: inside the processors (e.g. 64 bits processor can execute 2 32 bits operations)
 - Multicore/multi cpu level: inside the same chip/computer. The processing elements share the memory, system bus and OS.
 - Network-connected computers: clusters, distributed computing. Each processing element has its own memory space, OS, application software and data

» Huge difference depending on the interconnects: e.g. High Performance Computing (supercomputers) vs. High Throughput Computing (seti@home)



Some classifications

SMP vs. MPP (or the shared memory vs. distributed memory debate):

- SMP: Symmetric Multi Processors system: shared memory approach
 - "single box" machines, OpenMP programming family
- MPP: Massively Parallel Processors system: distributed memory, network-connected CPUs
 - "clusters", MPI programming family (message passing)
- SMPs are easier to program but scale worse than the MPPs







Why parallel computing?

- It is cool
- Sometimes the problem does not fit into a single box: you need more resources than you can get from a single computer
- To obtain at least 10 times more power than is available on your desktop
- To get exceptional performance from computers
- To be couple of years ahead of what is possible by the current (hardware) technology
- The frequency scaling approach to increase performance does not work any longer (power consumption issues):
 - The new approach is to stuff more and more processing units into machines, introducing <u>parallelism everywhere</u>



Measuring performance gain: the Speedup

- In an ideal scenario a program running on P processing elements would execute P times faster..., giving us a linear speadup
- Speedup S(n,P): ratio of execution time of the program on a single processor (T₁) and execution time of the parallel version of the program on P processors (T_P):
 - » In practice, the performance gain depends on the way the problem was divided among the processing elements and the system characteristics.
- Amdahl's law: gives an upper estimate for maximum theoretical speedup and states that it is limited by the non-parallelized part of the code:

$$S(n, P) \leq \frac{1}{\alpha + (1 - \alpha)/P} \leq \frac{1}{\alpha}$$

- alpha is the sequential fraction of the program
- e.g. if 10% of the code is non-parallizable, then the maximum speedup is limited by 10, independent of the number of used processors (!)

$$S(n, P) = \frac{T(n, 1)}{T(n, P)}$$

n denotes the problem size.

T denotes the execution time.





The dark side

"the bearing of a child takes nine months, no matter how many women are assigned"

- Not everything is suitable for parallelization
- Complexity increases as more and more communication is involved:
 - embarrasingly paralell -> coarse-grained -> fine-grained problem domains
- Parallel computing opens up new set of problems:
 - Communication overheads
 - Concurrency problems
 - Synchronization delays
 - Race conditions and dead locks
- Nobody wants to debug a parallel code...
- Developing & deploying a parallel code usually consume more time than the expected speedup
- A practical advice for parallelization:
 - Unless you have an embarrasingly parallel problem, forget it
 - If you are stubborn, then at least use an available parallel (numerical) library and start with the profiling (understanding) of your program
 - Wait for the holy grail of computational science: automatic parallelization by compilers





Now comes active learning ③



Accessing remote computers







Task 1: Howto avoid loosing all your work on a remote computer

IMAGINE that:

- You are being logged on a remote computer
- In the middle of a long task (e.g. compilation, download, etc..)
- Then, suddenly the network connection dies
- or you'd like to go home and continue the same work from your home

Is there a way to avoid loosing all your work? How can one disconnect & "session" without the need to restart everything from scratch?

TODO: try to use the **screen** utility to keep your remote session alive. Launch screen and start working in several screen session. Then, imitate a network failure and with the help of screen resume your work on the remote machine.

screen allows you to

- Keep a session active even through network disruptions
- Disconnect and re-connect to a sessions from multiple locations (computers)
- Run a long remote running process without maintaining an active remote login session





Task 1: helpdesk

Use the Linux **screen** utility to manage remote screen sessions, connect, reconnect to active session, survive a network failure ☺

- · Screen is started from the command line just like any other command
 - [iridium ~]\$: screen
 - You can create new "windows" inside screen, ctr+a c then rotate, switch between windows with ctrl+a n
- Listing your screens:
 - [iridium ~]\$: screen -list
- Disconnecting from your active session, screen (your task keeps running!):
 - [iridium ~]\$: screen -d or ctrl+a d
- Re-connecting to an active screen session (re-attach to screen):
 - » [iridium ~]\$: screen -r
- Terminating, logging out of screen
 - » type exit from inside all your active screen sessions
- Using screen to log your activity:
 - » [iridium ~]\$: screen -L or ctrl+a H turns on/off logging during a screen session



Task 2: Understand the layout of a cluster: Head node vs. Worker node

- Head (login) node: pptest-iridium.lunarc.lu.se
- Worker nodes: n3, n4, ..., n12

TODO: try to use various linux commands to discover the structure of the iridium cluster:

Hint: you can "login" to a node by using the interactive command

- is there shared file system?
- are the nodes identical (cpus, memory)?
- how about network connectivity?
- how about user accounts?
- how about the operating system, available software?
- how about running processes, cpu-load?





Task 3: Understand the basic concept of ,,resource schedulling" by using SLURM

- There is no infinite size cluster $\ensuremath{\textcircled{\odot}}$
 - Always limited number of cores, memory, walltime
 - Workload Management System, Scheduller, Batch System: SLURM

TODO: get greedy and eat all the cake! Here is the "menu":

- slurm.schedmd.com/quickstart.html
- lunarc-documentation.readthedocs.io/en/latest/batch_system/#firstexample-for-a-job-submission

Hint: grab a piece of cake by creating a so-called job on the cluster

- Find out howto create (submit) a job. Hint: use the sbatch command
- Find out the state of the cake. Hint: use sinfo, squeue
- Find out how to control the size of a piece of cake? (number of nodes, cores, memory, etc..)
- Find out how to put back a piece of cake (cancel job)
- Find out how to obtain status of the job, receive notification







Task 3: helpdesk

- List SLURM queues (partitions)
 - > sinfo
- Create file **myscript** (use provided examples)
- Submit simple jobs and check their status:
 - > sbatch myscript
 - > cat slurm-<jobid>.out
 - > squeue
 - > scontrol show job <jobid>
- Repeat with multi core/node jobs
 - sbatch -N4 myscript
 - sbatch -n6 myscript
 - In a multi-core advanced example, pay attention how jobs are distributed across nodes and cores
- Receive notification on state changes using the directives within the scripts (not enabled on iridium):
 - #SBATCH --mail-user=fred@institute.se
 - #SBATCH --mail-type=END (BEGIN, END, FAIL, REQUEUE, ALL)

Simple myscript:

```
#!/bin/sh
#SBATCH -J "simple job"
#SBATCH --time=1
echo "we are on the node"
hostname
who
sleep 2m
```

Multicore/node myscript:

```
#!/bin/sh
#SBATCH -J "multi job"
#SBATCH --time=1
srun hostname |sort
sleep 5m
```



Task 4: Understand the concept of work directory, input & output of batch jobs

Unless it is a trivial "hello world" exercise, most of the real-life jobs process some input data and produces output data. It is very important to understand where all that data is located relative to the worker node.

TODO: Execute a non-trivial task as a batch job submitted to SLURM that either reads in some data from a file and/or generates some output. As a second example, use the sysbench toolkit submitted as a batch job to measure IO performance on various filesystems.

- Find out what happens to the standard output and standard error of the "executable"
- Find out what is the execution directory on the Worker Node
- · Find out where the output files are placed.
- Use special variables understood by SLURM to place your job into a specific directory (SLURM_SUBMIT_DIR, TMPDIR)
- Execute the sysbench --test=fileio testkit as part of a batch job to measure disk performance
 - Info and manual about sysbench: wiki.gentoo.org/wiki/Sysbench





Task4: helpdesk

Stdout/Stderr script:

#!/bin/sh
#SBATCH -J "ioperf"
#SBATCH --time=4
#SBATCH -o ioperf_%j.out
#SBATCH -e ioperf_%j.err

echo "we are on the node and testing io" hostname cd \$TMPDIR mkdir testdir cd testdir pwd

sysbench --test=fileio --file-total-size=4G prepare sysbench --test=fileio --file-total-size=4G --file-test-mode=rndrw --max-time=120 --max-requests=0 run sysbench --test=fileio --file-total-size=4G cleanup

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Sysbench myscript:

#!/bin/sh
#SBATCH -J ``ioperf"
#SBATCH --time=10

echo "we are on the node" hostname who sleep 2m

Task farming



Implementing trivial parallelism with a master –worker system

- With a help of a master script you are going to execute X number of subtasks on Y number of processing units
- The master script (master.sh) takes care of launching (new) subtasks as soon as a processing element becomes available
- The worker.sh script imitates a payload execution that corresponds to a subtask

<u>Steps:</u>

- 1. Copy the scripts to a new directory on pp-test-iridium
- 2. Set the problem size (*NB_of_subtasks*) and the number of processing elements (#SBATCH -*n*) in the *master.sh*, the payload size (i.e. How long a subtask runs, PAYLOAD) in the *worker.sh*
- 3. Launch the taskfarm (*sbatch master.sh*), monitor the execution of the subtasks (*squeue j* <*jobid> -s*) and finally check how much time the taskfarm processing required (check the output files of the subtasks and the slurm job)
- 4. Repeat the taskfarming with modified parameters, What is the speedup?

