Parallel programming practice (1-2)

MPI example, environment: para.iit.bme.hu

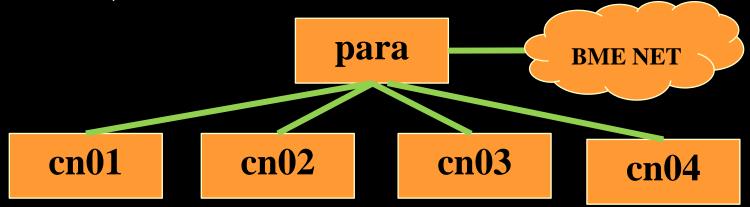
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Environment /1

- Cluster in the CIRCLE cloud
- para.iit.bme.hu main machine 4 vCPU
- cn01, cn02, cn03, cn04 workers 4 vCPU
- /users, /usr NFS



Environment /2

- module
 - Easy modification of the environment variables that different tools require.
- slurm
 - Sheduler and resource manager. This tool ensures an efficient and fair share of resources.
- compilers
- development tools

Login to the cluster



Using Linux based client (eg. from lab, win10) ssh neptun@para.iit.bme.hu

Using Windows based free client: (putty.org)

putty

host: para.iit.bme.hu

login: neptun

OR you can use VScode Remote ssh extension

module

• A bit forgotten tool for quick and maintainable change of environment variables to select alternative program variants.

Example:

```
module load mpi
printenv | grep MPI_LIB
MPI_LIB=/usr/local/openmpi/lib
module unload mpi
module load mvapich2
printenv | grep MPI_LIB
MPI_LIB=/usr/local/mvapich2/lib
http://modules.sourceforge.net
```



module example 2

module load mpi module load cuda module load dot module list



Currently Loaded Modulefiles:

1) mpi 2) cuda 3) dot

The dot inserts the current directory into PATH

Try the linda example

```
# Login to the cluster
module load linda
cd
cp -r ~szebi/para/linda . # Mind the dot!
cd linda
clc -w -linda compile_args -m32 -o chello chello.cl
./chello
# There is no resource allocation and scheduling
# The program runs on the para headnode
# It is not efficient, Solution: see later!
```

MPI example #1 (.c)

```
#include <stdio.h>
#include <stdlib.h>
#include <mpi.h>
int main(int argc, char **argv) {
 int rank, size, len;
  char hostname[MPI MAX PROCESSOR NAME];
 MPI Init(&argc, &argv);
 MPI Comm rank (MPI COMM WORLD, &rank);
 MPI Comm size(MPI COMM WORLD, &size);
 MPI Get processor name(hostname, &len);
 printf("I am %s %d of %d on host %s\n",
             arqv[0], rank, size, hostname);
 MPI Finalize();
  return 0;
```

MPI example #1 (.cc)

```
#include <iostream>
#include <mpi.h>
using namespace std;
int main(int argc, char *argv[]) {
 MPI::Init(argc, argv);
  int rank = MPI::COMM WORLD.Get rank();
  int size = MPI::COMM WORLD.Get size();
  int len;
  char hostname[MPI MAX PROCESSOR NAME];
  MPI::Get processor name(hostname, len);
  cout << "I am " << rank << " of " << size
       << " on host " << hostname << endl;
 MPI::Finalize();
  return 0;
```

MPI compile & run

module load mpi cd cp -r ~szebi/para/MPI . # Mind the dot! cd MPI mpicc -o mpihello mpihello.c mpirun -np 2 mpihello Hello, world! I am ./mpihello 1 of 2 on host para Hello, world! I am ./mpihello 0 of 2 on host para

There is no resource allocation and scheduling # The mpirun starts as many instances as we request.

MPI running on the cluster

The mpirun

- enters in to the nodes, or
- uses daemons (eg. LAM)

to launch the program instances, but does not do any resource allocation, nor scheduling.

In multi-user environment the resource allocation is essential. There are several tools for this (condor, pbs, sge, slurm, ...)

We use SLURM.

MPI example using SLURM

```
run_mpi.sh:
#!/bin/bash
#SBATCH -o std.out
mpirun $@
```

sbatch –n 3 run_mpi.sh ./mpihello more std.out



Hello, world! I am ./mpihello 0 of 3 on host cn01 Hello, world! I am ./mpihello 2 of 3 on host cn01 Hello, world! I am ./mpihello 1 of 3 on host cn01

SLURM

- Simple Linux Utility for Resource Management
- Resources are divided into partitions that can overlap. On para.iit.bme.hu machine we have 3 partitions: prod, debug, misi
- Major commands:
 - sinfo node and partition info
 - srun running a parallel job
 - sbatch start a batch script
 - squeue query the queue
 - salloc resource reservation for interactive usage
 - scancel delete job from the queue

SLURM example /1

sinfo

Information about the resources

srun -n 2 /bin/hostname

 Allocate 2 CPU's and start the command /bin/hostnae on each.

srun -N 2 /bin/hostname

Allocate 2 node's and start the command on each.

salloc -n 4 /bin/bash

 Allocate 4 CPU's and start an interactive bash command on the first one.



SLURM example /2

cd

cd slurm

cp -r ~szebi/para/slurm . # Mind the dot!



sbatch -n 3 simple.sh

Creates a job requesting3 CPUs and returns the command prompt.

simple.sh:
#!/bin/bash
printenv | grep SLURM

hostname

 When the resources are available, it allocates them and launches simple.sh on one. Pass the names of the reserved resources by environment variables.

Most important switches

- -n number of tasks to run
- --ntasks-per-node=n number of tasks per node
- -N number of nodes (N = min[-max])
- -o location of stdout
- -p partition requested
- --prolog=program run "program" before job step
- -i location of stdin
- -t time limit
- --mincpus=n minimum number of logical CPUs

Switches in the script

Switches can be integrated with simple syntax.Pl:

```
#!/bin/bash
#SBATCH -n 3
#SBATCH -o output_file
#SBATCH -D working_dir
#SBATCH --mail-type=end
#SBATCH --mail-user=xyz@xyz.de
#SBATCH -t 01:00:00
srun /bin/hostname # like srun -n 3
```

Running SPMD programs

File: run.sh

#!/bin/bash

mpirun \$@

Command:

sbatch –n 4 run.sh ./mpihello

File: runhello.sh

#!/bin/bash

#SBATCH -o hello.txt

#SBATCH -n 4

mpirun ./mpihello

Command:

sbatch runhello.sh

Running MPMD programs

File: multi.sh

#!/bin/bash

#SBATCH -N 2

#SBATCH -n 8

#SBATCH -o multi.out

srun --multi-prog ./multi.conf

File: multi.conf



0./master.sh

slaves

* ./slave.sh

Command:

sbatch multi.sh

Major environmental variables:

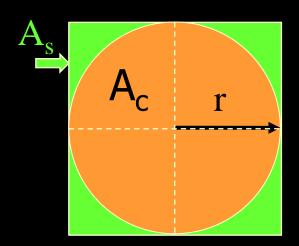
SLURM_PROCID

SLURM_NODEID

SLURM_LOCALID

PI example

```
A_{c} = \pi r^{2}
A_{s} = (2r)^{2}
\pi = 4 \frac{A_{c}}{A_{s}}
```



```
int isInside() {
   double x = ((double) rand()) / RAND_MAX;
   double y = ((double) rand()) / RAND_MAX;
   if ((x*x + y*y) <= 1.0) return(1);
   else return(0);
}</pre>
```

PI example (serial version)

cd ~/MPI

more pi.c gcc -o pi pi.c time ./pi 5000000



PI_MPI example

module load mpi cd ~/MPI mpicc -o pi_mpi pi_mpi.c



Measuring run times on vCPU's
sbatch -o pi.out run_mpi.sh ./pi_mpi 5000000
run times with 4 processors
sbatch -o pi4.out -n 4 run_mpi.sh ./pi_mpi 5000000

Problem: Primes in [1..n]

m

<u>Init.</u>	Pass 1	Pass 2	Pass 3
23456789011234567890122345	2 3← <i>m</i>	2 3	2 3
	5	5 <i>←m</i>	5
	7	7	7 ←
	9		×
	11	11	11
	13	13	13
	15		
	17	17	17
	19	19	19
	21		
	23	23	23
	25	25	

The sieve of Eratosthenes yielding a list of primes for n = 25. Multiples of marked element

erased from the list.

 $m^2 > n$, the computation stops

Serial implementation

```
m := 2
while m*m < n do
  if V[m] = 1 then
    i:= m*m
    while i <= n do
      v[i] = 0
      i := i + m
    od
  fi
  m := m+1
od
return all i such that V[i] = 1
```

Control-paralel implementation

```
shared
            V[2..n]:
memory
                  m:
j:= m, m:= m+1 // atomic
while j*j < n do
  if V[j] = 1 then
    i:= j*j
    while i <= n do
      v[i] = 0
      i := i + j
    od
  fi
  j:= m, m:= m+1 // atomic
od
wait for all and return all i such that V[i] = 1
```

Linda implementation

```
// init the shared "variables"
 in tuple space
 void init(int n) {
   int i;
    for (i = 2; i \le n; i++)
      out("vec", i);
    out("m", 2);
                    vec
                            vec
                                     vec
Tuple space
             vec
                   vec
                                     vec
                                     m
```

Linda implementation #2

```
int proc(int p, int n) {
  int i, j;
  in("m", ?j); // assume: j == 2
  out("m", j+1);
  while (j*j < n) {
    if (rdp("vec", j)) {
       i = j*j;
       while (i <= n) { inp("vec", i) i = i+j; }</pre>
    in("m", ?j);
    out("m", j+1); Vec
                                                vec
                              vec
                    vec 6 vec
  in("done", ?i);
  out("done", i+1);
  return 0;
                                vecl
           for coordination only
```

Linda implementation #3

```
int real main(int argc, char *argv[]) {
  int i, int N = 1000, NP = 4;
  init(N);
  out("done", 0);
  for (i = 1; i \le NP; i++)
                                    for coordination only
     eval(proc(i, N));
  in("done", NP);
 printf("\nAll (%d) processors done\n", NP);
  printf("Primes in [1..%d]:\n", N);
  for (i = 2; i \le N; i++) {
    if (inp("vec", i)) printf(" %d", i);
  return 0;
```

Data-paralel implementation

```
n/p
   m:
m := 2
while m <= n/p \&\&
              m*m < n do
  if V[m] = 1 then
     broadcast m →
    i := m*m
    while i <= n do
       V[i] = 0, i := i+m
    od
  fi
  m := m+1
od
broadcast 0 and wait for results
```

Assume that $p < \sqrt{n}$ $\rightarrow \sqrt{n} < n/p$ \rightarrow All the primes that have multiples are located on the P_1 processor.

 P_1 processor act as coordinator.

Data-paralel implementation #2

```
2n/p
         n/p+1
   m:
while \rightarrow m<>0 do
  i := m*m
  while i <= n/p do
    i := i+m
  od
  while i <= 2n/p do
    v[i] = 0
    i := i + m
  od
od
send back all i such that V[i] = 1
```

```
(p-1)n/p+1
                            n
    m:
while \rightarrow m<>0 do
  i:= m*m
  while i \le (p-1)n/p do
     i := i + m
  od
  while i <= n do
    v[i] = 0
     i := i+m
  od
od
send back all i such that V[i] = 1
```

Assignment

- 1. Learn / try the environment
- 2. Create a MPI porgram solving prime number generation. Use the data-parallel solution.

Theory:

Check the slides 22-29

Help: ~szebi/para/MPI/myprime_serial.cc

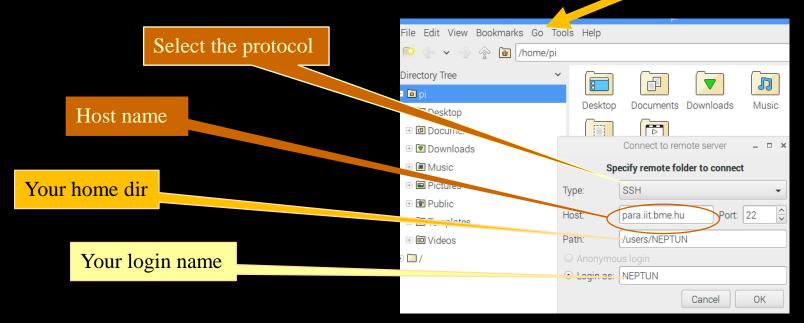
Tools/methods good to know

- General UNIX/LINUX commands
- File transfers between the local and remote machine.
- C/C++ development
- File transfer between tha local and remote sites
- MPI C/C++ API
 - MPI Python also available on the para cluster

Transferring files

Transferring files from para.iit.bme.hu to a Linux machine:

- Using Linux command line: (scp from to)
 - > scp NEPTUN@iit.bme.hu: . # mind the dot
- Using PCManFM file manager (menu: go)



Transferring files

Transferring files from para.iit.bme.hu to a Windows machine:

• Using WinSCP freeware utility (winscp.net):

