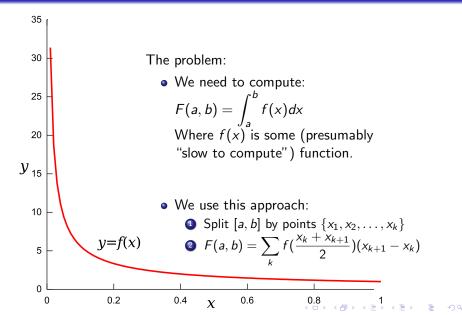
MPI profiling with Allinea MAP

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Problem: Calculation of a definite integral.



Problem size behavior: how to measure.

Do we even need to parallelize?

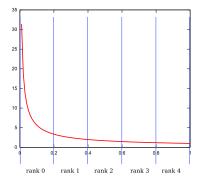
```
1 | # Compilation:
   $ gcc -03 -o integral_seq.x integral_seq.cxx\
         -L./mylib -lmymath
4 | # Timed runs:
   $ time -p ./integral_seq.x 1000000
   Result=3.775045 Exact=3.775063 Difference=-0.000019
   real 2.12
   $ time -p ./integral_seq.x 2000000
   Result=3.775058 Exact=3.775063 Difference=-0.000005
   real 4.28
10
   $ time -p ./integral_seq.x 8000000
11
   Result=3.775062 Exact=3.775063 Difference=-0.000001
12
   real 17.61
13
```

The time grows linearly with the problems size. Acceptable accuracy at 8M points. Can we speed it up?

Parallellization: domain decomposition.

Approach:

- Split [a, b] into several domains;
- Compute integrals independently.



$$F(a,b) = \int_a^b f(x) dx$$

- Assign a process to each domain $[x_k, x_{k+1}]$
- 2 Let each process compute $F(x_k, x_{k+1})$

3
$$F(a,b) = \sum_{k} F(x_{k}, x_{k+1})$$

A sketch of the parallel code

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```
MPI_Init(&argc, &argv);
int rank, nprocs:
MPI_Comm_rank(MPI_COMM_WORLD, &rank);
MPI_Comm_size(MPI_COMM_WORLD, &nprocs);
// ... Get total number of steps, broadcast...
// Each rank figures out its integration limits and number of steps
unsigned long my_stepbase, my_nsteps;
get_steps(nsteps_all, nprocs, rank, &my_stepbase, &my_nsteps);
const double per_step=(global_b-global_a)/nsteps_all;
const double x1=global_a + my_stepbase*per_step;
const double x2=x1 + mv nsteps*per step;
// Compute my own part of the integral
double my_y=integral(integrand, my_nsteps, x1, x2);
// Sum all numbers on master
double y=0;
MPI_Reduce(&mv_v, &y, 1, MPI_DOUBLE, MPI_SUM, 0, MPI_COMM_WORLD);
// ... print results ...
MPI Barrier(MPI COMM WORLD):
// Here we could start another computation.
MPI_Finalize();
```

Parallel performance: how to measure.

Now let's see how much we achieved...

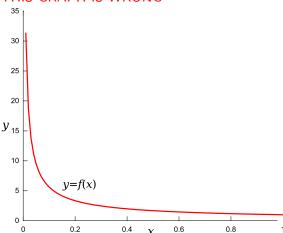
- Strong scaling: as we add processes, how do we fare?
- Weak scaling: as we add both processes and work?

```
$ mpicc -03 -o integral_par.x integral_par.cxx \
           -L./mylib -lmymath
2
   $ time -p mpirun -np 1 ./integral_par.x 8000000
   Result=3.775062 Exact=3.775063 Difference=-0.000001
   real 17.23
   user 17.08
   sys 0.02
   $ time -p mpirun -np 2 ./integral_par.x 8000000
   Result=3.775062 Exact=3.775063 Difference=-0.000001
   real 17.24
10
   user 31.98
11
   sys 0.05
12
```

Parallel performance: results

Is there a performance problem?

THIS GRAPH IS WRONG



How does performance analysis work?

How to collect data?

- Instrumentation:
 - Insert timers & counters in the code
 - Requires source or binary processing
- Sampling:
 - Interrupt & check the program at regular intervals
 - Introduces statistical error

What kind of data?

- Profile:
 - Summary information only
 - Relatively small file
- Trace:
 - Detailed recording during the run
 - Potentially huge file
 - Profile can be restored

Allinea MAP does tracing by sampling.

Prepare for profiling

2

To prepare for profiling, one needs:

- Compile with full optimization
- Generate debugging symbols
- Link with system libs dynamically
 - Usually the default
 - Notable exception: Cray
- On Flux: load ddt module

Running Map: simple way (demo)

- Get interactive access to a compute node
- Change to your working directory
- Optionally, set sampling interval
- Run as you would, prefixed by map

Caution:

- Too small interval: large overhead!
- Too large interval: not enough samples!

Running Map: other options

What if you can not or would not run GUI?

- Have slow or non-existing X connection to compute nodes
- Do not want to run interactively

Use -profile option.

```
#PBS -V
#PBS -q flux -l qos=flux -A account_flux
#PBS -l nproc=12,walltime=10:0:0

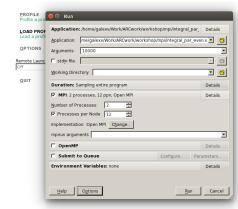
cd $PBS_0_WORKDIR
export ALLINEA_SAMPLER_INTERVAL=5
map -profile mpirun -np 12 ./integral_par.x 10000
```

This will create a *.map file. Then run from the login node:

```
$ map integral_par_even_12p_*.map
```

If you are submitting to a Flux queue...

- Run map from the login node:
- \$ map ./integral_par.x 8000000
- Set number of processes
- Check Submit to queue
- Click Configure...
- Load a proper submission template file (see next page)
- Click OK
- Click Run



Submission template for Flux

```
#PBS -V
#PBS -1 walltime=WALL_CLOCK_LIMIT_TAG
#PBS -1 nodes=NUM_NODES_TAG:ppn=PROCS_PER_NODE_TAG
#PBS -q QUEUE_TAG -1 qos=flux -A account_flux
#PBS -o PROGRAM_TAG-allinea.stdout
#PBS -e PROGRAM_TAG-allinea.stderr

cd $PBS_O_WORKDIR
AUTO_LAUNCH_TAG
```

Time for a live demo!

SCREENSHOT HERE

- Most of the time is spent in MPI
- As the run progresses, even more time is spent in MPI
- Problem: some processes spend more time in calculating the integrand f(x)!
- It's called "Load Imbalance"

Possible solutions:

- Distribute work unevenly (but how?)
- Implement dynamic load balancing

Dynamic load balancing

Manager-Workers approach:

Manager

- Listen to all workers
- Worker sent READY ? send GO with a job chunk
- Worker sent DONE? add result to the sum
- No more job chunks? send STOP to the worker
- No more workers? we are done

Worker

- Send READY to the Manager
- Listen to the Manager
- Manager sent GO ?
 - Get job chunk
 - Do the calculation
 - Send DONE with result to the Manager
 - Go to (1)
- Manager sent STOP? exit.



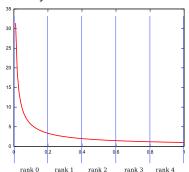
Strong scaling graph.

Map demo and screenshot.

The problem and the first approach

$$I(a,b) = \int_a^b x^{-3/4} dx$$

Analytic solution to cross-check: $I(a,b) = 4(b^{1/4} - a^{1/4})$

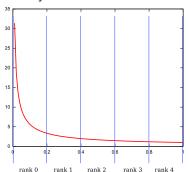


- Split [a, b] equally between workers
- $I(a,b) = \sum_{k} I(x_k, x_{k+1})$

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- Split [a, b] equally between workers
- $I(a,b) = \sum_{k} I(x_k, x_{k+1})$
- Caveat: integrand takes more time to compute at low x!

Dynamic load balancing

Try various chunk sizes, various number of processes. Where are the bottlenecks?

(Alternative: give smaller chunks to low X. Scalability. Profile.)