Parallel Performance Analysis Tools

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Outline

- Motivations
- Discussion of CrayPAT and Scalasca
- Outline example code
- CrayPAT Usage
- Scalasca Usage





Motivations – What is Profiling?

- Examine the behaviour of the code
- Pick out any subroutines/functions that cause slowdown or have unusual behaviour
- Two types:
 - 1. Sampling (periodically queries running code to determine what function the code is in)
 - 2. Tracing (adds instructions into the code that report when entering/leaving functions, and various statistics)





Motivations – What is Profiling?







Picking an Example to Analyse

- Profiling generates a lot of extra data, and can cause your code to run more slowly
 - Need to choose a reasonably short example, but:
 - Program execution must be representative of a production run
 - Must be long enough to hide start-up and finalisation costs
 - Should include all the I/O of a normal job
- A good choice is something like a benchmark problem that takes a few minutes to run on a node/handful of nodes





Motivations - Why Profile?

- For developers:
 - Understand what the most time-consuming parts of the program are
 - Understand communication patterns and problems
 - E.g. load imbalance, synchronisation costs
 - Tool to help direct development efforts to give maximum benefits
- For users?
 - Understand why your program performs in a certain way
 - Help with choice of appropriate parameters, MPI processes...





Profilers: CrayPAT and Scalasca

- In this course we will consider two parallel performance analysis tools; CrayPAT and Scalasca
- With each tool you
 - 1. Instrument your code (typically during building)
 - 2. Run your code
 - 3. Analyse results







- + Various levels of detail
- + Extreme customisibility for expert users
- Only available on Cray Platforms
- GUI is not particularly useful





Scalasca

- + Open source
- + Portable

+ Allows you to determine early/late senders etc... + Useful GUI (Cube)

 Unable to trace CUDA, SHMEM events or OpenMP nested parallelism





- In this tutorial we will use a simple MPI code to demonstrate parallel performance analysis
- A computational fluid dynamics (CFD) code is employed, which calculates the flow of fluid within a cavity with an inlet in one side, and an outlet on another.
- The code can calculate the inviscid or viscid fluid flow.





Solves Poisson's Equation for the streamfunction:

$$\frac{\partial^2 \psi}{\partial x^2} + \frac{\partial^2 \psi}{\partial y^2} = -\zeta$$

$$u_x = \frac{\partial \psi}{\partial y}; \ u_y = -\frac{\partial \psi}{\partial x}; \ \zeta = \frac{\partial u_y}{\partial x} - \frac{\partial u_x}{\partial y}$$

Available in both C and Fortran





Iterate until convergence

$$\begin{split} \psi_{i,j}^{\text{new}} &= \frac{1}{4} \left(\psi_{i-1,j}^{\text{old}} + \psi_{i+1,j}^{\text{old}} + \psi_{i,j-1}^{\text{old}} + \psi_{i,j+1}^{\text{old}} \right) - \zeta_{i,j}^{\text{old}} \\ \zeta_{i,j}^{\text{new}} &= \frac{1}{4} \left(\zeta_{i-1,j}^{\text{old}} + \zeta_{i+1,j}^{\text{old}} + \zeta_{i,j-1}^{\text{old}} + \zeta_{i,j+1}^{\text{old}} \right) \\ &- \frac{R_e}{16} \left[\left(\psi_{i,j+1}^{\text{old}} - \psi_{i,j-1}^{\text{old}} \right) \left(\zeta_{i+1,j}^{\text{old}} - \zeta_{i-1,j}^{\text{old}} \right) \\ &- \left(\psi_{i+1,j}^{\text{old}} - \psi_{i-1,j}^{\text{old}} \right) \left(\zeta_{i,j+1}^{\text{old}} - \zeta_{i,j-1}^{\text{old}} \right) \right] \\ \hline \end{split}$$

Parallelised in the x (C) or y (Fortran) directions



Halos transferred via MPI_Sendrecv





- The code can be found on the course web pages
- To run it, use aprun –n [nprocs] ./cfd <scale> <numiter> <Re>

Where

- nprocs is the number of MPI processes
- scale scales the size of the box (32 x scale cells)
- numiter is the number of iterations
- Re (optional) is the Reynolds number ($0 \le \text{Re} < 3.7$)





 The output can be visualised using: \$ gnuplot –persist cfd.plt



$$R_e = 0$$





 $R_e = 3.0$





Examples of Performance Tools

- I will now go onto demonstrate CrayPAT and Scalasca on ARCHER using the CFD code.
- Afterwards you will get an opportunity to try using CrayPAT/Scalasca yourselves
- For best results, it is recommended that you to login to ARCHER with an X-windows connection, e.g. \$ ssh –X [username]@login.archer.ac.uk





- Load the CrayPAT modules:
 \$ module load perftools-base
 \$ module load perftools
- Build executable as normal \$ make clean; make
- Instrument the binary using pat_build \$ pat_build ./cfd





- Instrumentation creates a new binary cfd+pat
- Modify the job submission script to run this new binary, then submit the job
 \$ asub submit pbs
 - \$ qsub submit.pbs
- This will run the cfd code with sampling





- Once the job has completed, it will have created an additional file: cfd+pat+<number>.xf
- Generate a human-readable report using pat_report
 \$ pat_report cfd+pat+<number>.xf

(You can put this information into a file by using the argument '-o <file>')





Table 1: Profile by Function

Samp% | Samp | Imb. | Imb. |Group | |Samp | Samp% | Function | | | PE=HIDE

100.0% | 1,906.5 | -- | -- |Total

96.6% | 1,842.0 | -- | -- |USER

||------|| 74.9% | 1,427.2 | 15.8 | 1.5% |jacobistepvort

|| 21.0% | 401.0 | 8.0 | 2.6% |main

3.3% | 62.5 | -- | -- |MPI

3.1% | 58.5 | 25.5 | 40.5% |MPI_Sendrecv





Pat_report also produces two other files; an .ap2 file, and an .apa file:

- The ap2 file acts as an input to the Apprentice2 graphical interface for viewing performance statistics
 \$ app2 <file>.ap2
- The apa file contains suggested configuration options for a traced experiment





Using CrayPAT – Apprentice2









Using CrayPAT - Tracing

- Instrument the binary for tracing using the .apa file as an input to pat_build
 \$ pat_build -O cfd+pat+<number>.apa
- Modify the job submission script to use the new binary then submit the job
 \$ qsub submit.pbs
- View the results data using pat_report as before \$ pat_report cfd+apa+<number>.xf
- Then use Apprentice2 if desired \$ app2 cfd+apa+<number>.ap2





Using CrayPAT

- This process can be continued as necessary until the information you need has been obtained/you have gained the desired understanding of your code's performance
- More information on CrayPAT can be found using the commands
 - \$ pat_help
 - \$ man intro_pat
 - \$ man pat_build
 - \$ man pat_report





- Load the Scalasca module
 \$ module load scalasca
- Instrumentation must be carried out during compilation by prepending scorep to the compiler. For example \$ scorep cc -c foo.c or \$ scorep ftn –c foo.f90
- Modify the compiler line in Makefile to include scorep:
 CC = scorep cc
 FC = scorep ftn





- It is important to ensure that scorep is used during the linking of the object files.
- Functions/subroutines/files that you do not need/want to instrument do not need to be compiled with scorep
- Build the executable make clean; make





- Modify the submission script to launch the parallel job with scalasca –analyze, e.g.
 scalasca –analyze aprun –np 4 ./cfd <options>
- Submit the job
 - \$ qsub submit.pbs
- A measurement directory scorep_cfd_4_sum is created during the job's execution which contains all the log files





- To analyse the output data, first run \$ scalasca –examine scorep_cfd_4_sum
- This will open the cube browser, which allows you to examine the code's timings
- Using the –s option produces a file (scorep.score) that can be used to advise you about setting up a tracing experiment

\$scalasca -examine -s scorep_cfd_4_sum





Using Scalasca - Cube







Examining the scorep.score file in the measurement directory reveals information on the estimated final disk usage and memory usage of a trace

Estimated aggregate size of event trace: 128MB Estimated requirements for largest trace buffer (max_buf): 32MB Estimated memory requirements (SCOREP_TOTAL_MEMORY): 34MB (hint: When tracing set SCOREP_TOTAL_MEMORY=34MB to avoid intermediate flushes or reduce requirements using USR regions filters.)

type	max_buf[B]	visits tim	e[s] tim	ne[%] time	e/visit[us] r	egion
ALL	33,493,662	3,848,767	78.79	100.0	20.47	ALL
MPI	22,401,846	2,000,134	2.95	3.7	1.47 N	1PI
USR	7,491,672	1,248,609	57.90	73.5	46.37	USR
COM	3,600,144	600,024	17.95	22.8	29.91	COM





 To trace the code, alter your job submission script to contain:

scalasca –analyze –q –t aprun –np 4 ./cfd <options>

 Don't forget to also set SCOREP_TOTAL_MEMORY in the script as suggested in the .score file: export SCOREP_TOTAL_MEMORY=34MB





- A new directory scorep_cfd_4_trace is created, and the results can be examined using \$ scalasca -examine scorep_cfd_4_trace
- This time, more information is present, such as that on late senders/receivers.





 If the estimated disk/memory usage for tracing is too high, you may need to consider to avoid tracing certain functions by using a filter file:

SCOREP_REGION_NAMES_BEGIN EXCLUDE jacobistepvort MPI_Sendrecv SCOREP_REGION_NAMES_END

• Usage:

scalasca –examine –f filter.txt aprun ... scalasca –analyze –q –t –f filter.txt aprun ...





Using Scalsca

- More information can be found on the Scalasca website <u>http://www.scalasca.org</u>
- In particular their user's guide:

http://apps.fzjuelich.de/scalasca/releases/scalasca/2.3/docs/UserGuide.pdf





Practical: CFD

- Try out using CrayPAT and/or Scalasca to investigate the performance of the CFD code
- Options:
 - Try using different values for scale, and investigate turning viscosity on and off
 - How does the profile change when running on large numbers of processes?
 - Terminate calculation based on a tolerance value (see comments in code), investigate only computing this infrequently
 - Investigate using serialised Send / Receive functions (see alternative boundary source files) instead of Sendrecv



