

# A brief introduction to profiling with the Intel toolchain

Mark Turner and Mark Dixon Advanced Research Computing

July 7th, 2022

#### Welcome and icebreaker

Thank you for joining!

Meet the team

- 1 hour course (12pm-1pm)
- Please do post in the chat or speak up when questions arise

 Feel free to post something in the chat about where in the university you come from and something about your research interests





#### **Course structure**

- What is performance analysis?
- Automatic tracing of MPI codes using ITAC
- Leveraging the Intel Traceanalyzer
- User defined instrumentation using ITAC

An exercise...





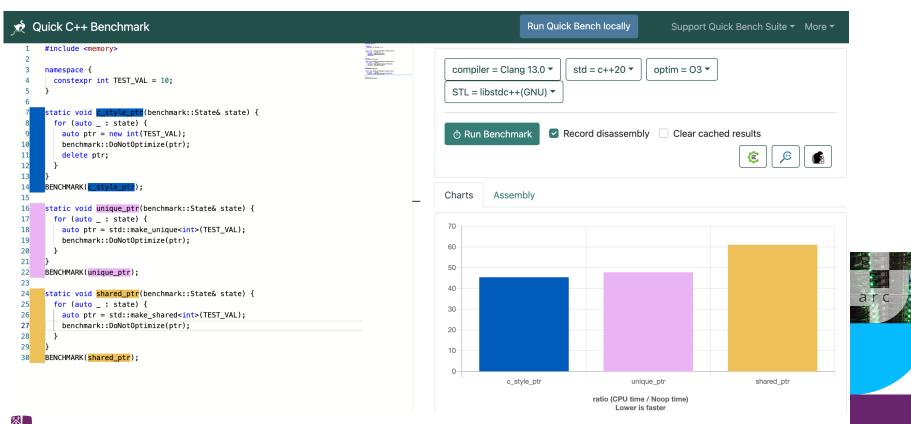
### What is performance analysis?

Strongly depends on what you are trying to investigate / achieve



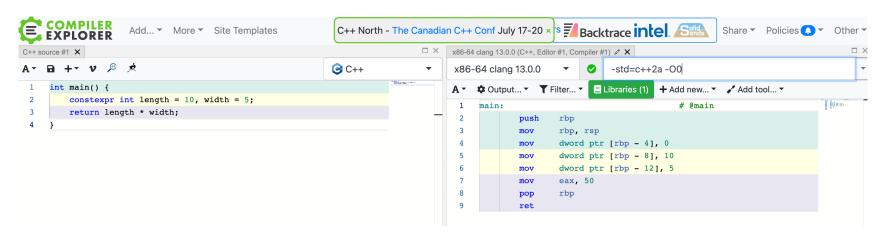


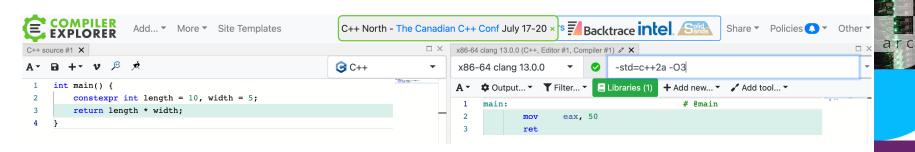
#### quick-bench.com





#### godbolt.org







Measuring like this is great, but...

how do we study performance of entire codes – in particular, those that run across supercomputers?

What if we want to understand the behaviour of an entire simulation?





### A flavour of performance analysis in HPC

- Identifying code flaws:
  - Hotspots (always a good place to focus dev time)
  - Load imbalance
  - Inefficient communication patterns
  - Poor scaling
  - Inefficient use of hardware (cache misses, excessive I/O etc.)

- Establishing the obstacles to being compute bound
  - Memory bound; MPI bound







#### How can tools understand our codes?

#### Sampling

- Regular reading of typical averages, patterns etc.
- Recording runtime characteristics (e.g., hardware counters)
- likwid-perfctr

#### Tracing

- Chronologically ordered events/timelines
- Logging events (e.g. function calls) in code





# For today, we're interested in tracing





### **Intel Trace Analyzer and Collector**





Presents the chronology of your runtime in a visual way

- Easier to reason about behaviour of parallel codes and identify:
  - temporal dependencies and bottlenecks
  - load imbalance
  - Hotspots
  - And more...







# So, on the surface, what does ITAC give us?





#### A high-level overview page:

#### Summary: peano4.stf

Total time: 1.07e+04 sec. Resources: 2 processes, 4 threads, 1 node. Continue > Top MPI functions Ratio This section represents a ratio of all MPI calls to the rest of your code in the application. This section lists the most active MPI functions from all MPI calls in the application. MPI Iprobe 244 sec (4.58 %) 75.7 sec (1.42 %) MPI Test MPI Isend 0.0473 sec (0.000887 %) Serial Code - 1.03e+04 sec 96.9 % MPI Comm dup 0.0466 sec (0.000873 %) OpenMP - 0 sec MPI Irecv 0.0178 sec (0.000335 %) MPI calls - 320 sec 3 %

#### Where to start with analysis

For deep analysis of the MPI-bound application click "Continue >" to open the tracefile View and leverage the Intel® Trace Analyzer functionality:

- Performance Assistant to identify possible performance problems
- Imbalance Diagram for detailed imbalance overview
- Tagging/Filtering for thorough customizable analysis

To optimize node-level performance use: Intel® VTune™ Profiler for:

- algorithmic level tuning with hpc-performance and threading efficiency analysis;
- microarchitecture level tuning with general exploration and bandwidth analysis; Intel® Advisor for:
- vectorization optimization and thread prototyping.

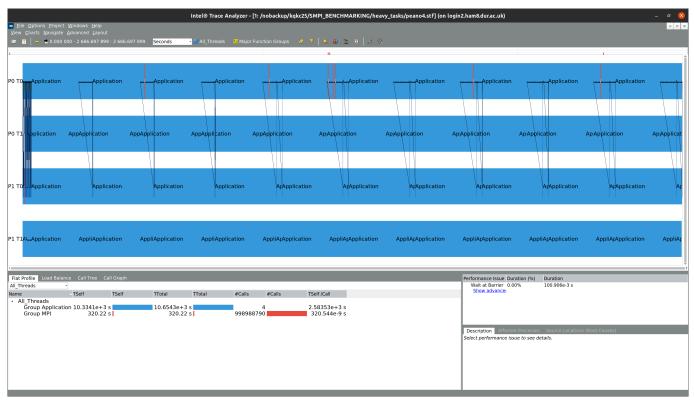
For more information, see documentation for the respective tool:

Analyzing MPI applications with Intel® VTune™ Profiler

Analyzing MPI applications with Intel® Advisor

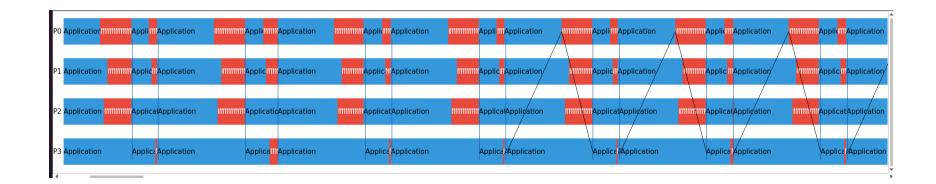


#### An event timeline:



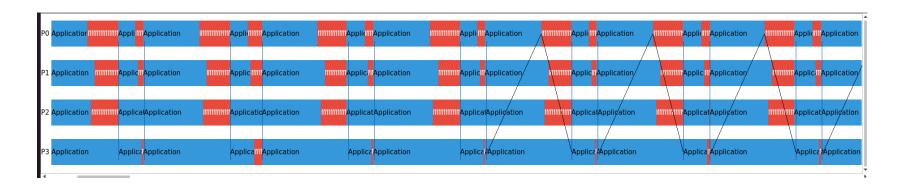


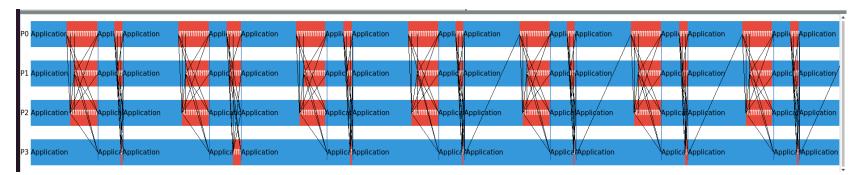
### A more problematic snapshot:





#### **Ability to filter MPI communications:**







#### A demo



How to navigate the traceanalyzer and explore an event timeline





# And how can I generate these profiles for my code on Hamilton?







- Compile with Intel MPI (Open MPI is not supported...)
- Load the ITAC module:
  - module load itac
- The trace files can be **BIG**. It is worth Asking SLURM for more temporary disk space on your nodes when doing big runs:
  - #SBATCH --gres=tmp:400G

- Then pass the -trace flag through to mpirun:
  - mpirun -trace -np <N> <executable>







#### **Exercise**

## Ask us questions!





# So far, we have the tools to better understand MPI behaviour

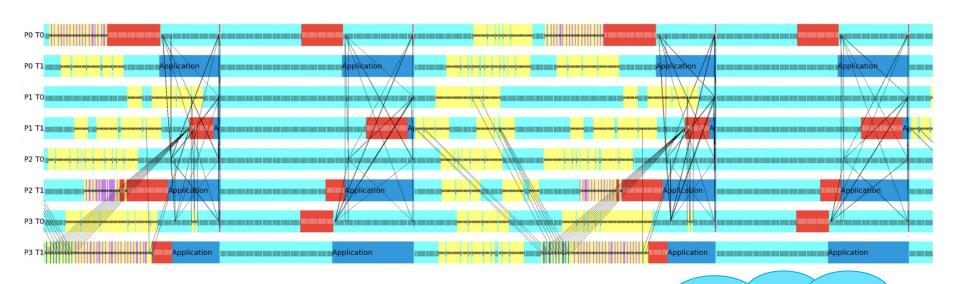
... But how can we visualize application behaviour?







#### **User defined instrumentation:**



- Red = MPI
- Light Blue = AMR framework
- Yellow = Physics
- Dark blue = Open MP

Zoom in to see more detail. Right click a cell and select "ungroup" to view function call.



# And how can I generate these profiles for my code on Hamilton?







1. Add instrumentation boilerplate

```
namespace {
  std::map<std::string, const int> itac handles;
void trace_in(const std::string& trace) {
  if (itac handles.count(trace) == 0) {
    int new handle;
    VT funcdef( trace.c str() , VT NOCLASS, &new handle );
    itac handles.insert({trace, new handle});
  VT_begin(itac_handles[trace]);
void trace out(const std::string& trace) {
  VT_end(itac_handles[trace]);
```







- 1. Add instrumentation boilerplate
- 2. Add your code (whatever that might be ©)

```
void do_fancy_physics(const int rank) {
  float aggregator = 0;
  for (int i = 0; i < 100'000'000; ++i) {
    aggregator += sin(i);
  }
  std::cout << "Result on " << rank << ": " << aggregator << std::endl;
}</pre>
```







- 1. Add instrumentation boilerplate
- 2. Add your code (whatever that might be ©)
- 3. Add tracing around your code

```
int main(int argc, char* argv[]) {
  MPI Init(&argc, &argv);
  int rank;
 MPI Comm rank(MPI COMM WORLD, &rank);
  const std::array<std::string, 3> phases {"alpha", "beta", "gamma"};
  for (const auto &phase : phases) {
    trace in(phase);
    do_fancy_physics(rank);
    trace out(phase);
  MPI_Finalize();
  return 0;
```







#### **Building/executing the code**

- 1. Load modules:
  - module load oneapi intelmpi itac
- 2. Launch the build:
  - mpiicpc -I\$ITAC\_HOME/itac/latest/include -L\$ITAC\_HOME/itac /latest/slib -lVTnull itac\_test.cpp -o itac\_test
- 3. Run with ITAC tracing enabled:
  - mpirun -trace -np 2 ./itac\_test







### **Ungrouped output:**

traceanalyzer itac\_test.stf

```
Po alpha beta gamma

Pl alpha beta gamma

pl alpha beta gamma
```



# Code for reference:



```
#include <VT.h>
#include <cmath>
#include <iostream>
#include <map>
#include <mpi.h>
#include <string>
namespace {
  std::map<std::string, const int> itac handles;
void trace in(const std::string &trace) {
 if (itac_handles.count(trace) == 0) {
    int new handle;
    VT_funcdef(trace.c_str(), VT_NOCLASS, &new_handle);
    itac handles.insert({trace, new handle});
  VT begin(itac handles[trace]);
void trace_out(const std::string &trace) {
  VT end(itac handles[trace]);
void do fancy_physics(const int rank) {
  float aggregator = 0;
  for (int i = 0; i < 100'000'000; ++i) {
    aggregator += sin(i);
  std::cout << "Result on " << rank << ": " << aggregator << std::endl;</pre>
int main(int argc, char *argv[]) {
  MPI_Init(&argc, &argv);
  MPI_Comm_rank(MPI_COMM_WORLD, &rank);
  const std::array<std::string, 3> phases{"alpha", "beta", "gamma"};
  for (const auto &phase : phases) {
    trace_in(phase);
    do_fancy_physics(rank);
    trace_out(phase);
  MPI_Finalize();
  return 0;
```



# Thank you!

- Feedback would really be appreciated: https://forms.office.com/r/YeFwCZ2kQi
- Other training courses at ARC
- RSE support
- Join the Society of Research Software Engineering: https://society-rse.org/join-us/