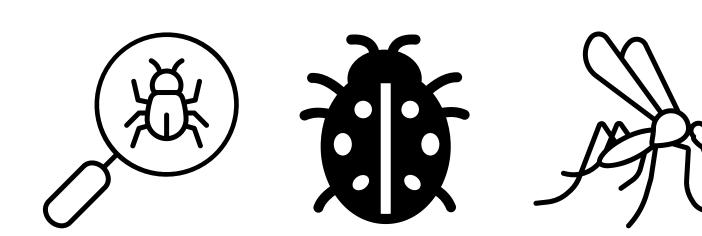


GOALS FOR THIS SESSION

Why you should use tools, and what they do

- All code has bugs. The more code, the more bugs.
- Different classes of "bugs"
 - Crashes, undefined behavior, deadlocks, correctness issues, ...
 - Time to solution, parallel efficiency, energy efficiency
- Treat performance issues as bugs, especially at scale



- Manual debugging: Mimic program execution, print state, time code regions
- Tool-assisted debugging: Automate "tedious" part
 - More information, less effort
 - Otherwise unavailable information (hardware counters!)
- Fixing bugs not (yet ⊕) automated: Tools simplify and enable analysis
- Introduce workflow and representative tools, focus on distributed GPU applications

DEBUGGING CORRECTNESS: BEST PRACTICES

Before you start

- Crashes are "nice" the stacktrace often points to the bug
- Prerequisite: Compile flags
 - While developing, always use -g -lineinfo
 - Use -g -G for manual debugging
 - Specific flags for compilers/lanugages (e.g. gfortran): -fcheck=bounds
- Memory corruption: Out-of-bounds accesses may or may not crash
 - compute-sanitizer: Automate finding these errrors
- Other issues: Manual debugging
 - cuda-gdb: Command-line debugger, GPU extensions

IVCC compile flags for debugging							
-g	Embed symbol info for <i>host</i> code						
-lineinfo	Generate line correlation info for <i>device</i> code						
-G	Device debug - slow						

COMPUTE-SANITIZER

Functional correctness checking suite for GPU

- compute-sanitizer is a collection of tools
- memcheck (default) tool comparable to Valgrind's memcheck.
- Other tools include
 - racecheck: shared memory data access hazard detector
 - initcheck: uninitialized device global memory access detector
 - synccheck: identify whether a CUDA application is correctly using synchronization primitives

Example run:

```
srun -n 4 compute-sanitizer \
--log-file jacobi.%q{SLURM_PROCID}.log \
--save jacobi.%q{SLURM_PROCID}.compute-sanitizer \
./jacobi -niter 10
```

- Stores (potentially very long) text output in *.log file, raw data separately, once per process.
- One file per MPI rank more on %q{} later

COMPUTE-SANITIZER

Anatomy of an error

- Look into log file, or use compute-sanitizer --read <save file>
- Actual output can be very long, if many GPU threads produce (similar) errors.

• We introduced an off-by-one error in line 60 ourselves:

```
a_{new}[iy * nx + (nx - 1) + 1] = y0;
```

APPROACHES FOR MULTI-PROCESS TOOLS

- Tools usually run on a single process adapt for highly distributed applications?
 - Bugs in parallel programs are often serial bugs in disguise
- Common MPI paradigm: Workload distributed; bug classes/performance similar for all processes
 - Not: Load imbalance, parallel race conditions; require parallel tools
- Ergo: Run tool N times in parallel, have N output files, only look at 1 (or 2, ...)
- %q{ENV_VAR} supported by all the NVIDIA tools discussed here, embed environment variable in file name
 - ENV_VAR should be one set by the process launcher, unique ID
 - Evaluated only once tool starts running (on compute node) not when launching job
- Other tools: Use a launcher script, for late evaluation

OpenMPI:

OMPI_COMM_WORLD_RANK
OMPI_COMM_WORLD_LOCAL_RANK

MVAPICH2:

MV2_COMM_WORLD_RANK
MV2_COMM_WORLD_LOCAL_RANK

Slurm:

SLURM_PROCID
SLURM LOCALID

https://www.open-mpi.org/faq/?category=running#mpi-environmental-variables
http://mvapich.cse.ohio-state.edu/static/media/mvapich/mvapich2-2.2-userguide.html#x1-32100013
https://slurm.schedmd.com/srun.html#SECTION_OUTPUT-ENVIRONMENT-VARIABLES

USING CUDA-GDB WITH MPI

- Launcher (mpirun/srun/...) complicates starting process inside debugger
- Workaround: Attach later

```
#include <unistd.h>
if (rank == 0) {
   char name[255]; gethostname(name, sizeof(name)); bool attached;
   printf("rank %d: pid %d on %s ready to attach\n", rank, getpid(), name);
   while (!attached) { sleep(5); }
}
```

Launch process, sleep on particular rank

```
$ srun -n 4 ./jacobi -niter 10 rank 0: pid 28920 on jwb0001.juwels ready to attach
```

Then attach from another terminal (may need more flags)

```
[jwlogin]$ JOBID=$(squeue -ho %i --me) # obtain job ID of user's first job
[jwlogin]$ srun -n 1 --jobid ${JOBID} --pty bash -i # launch interactive shell on job's node
[jwb0001]$ cuda-gdb --attach 28920
```

Wake up sleeping process and continue debugging normally

```
(cuda-gdb) set var attached=true
```

JSC system shortcut: sgoto --help

USING CUDA-GDB WITH MPI

Environment variables for easier debugging

• Automatically wait for attach on exception without code changes:

```
$ CUDA_DEVICE_WAITS_ON_EXCEPTION=1 srun ./jacobi -niter 10 Single GPU jacobi relaxation: 10 iterations on 16384 x 16384 mesh with norm check every 1 iterations jwb0129.juwels: The application encountered a device error and CUDA_DEVICE_WAITS_ON_EXCEPTION is set. You can now attach a debugger to the application (PID 31562) for inspection.
```

Same as before, go to node (see previous slide), then attach cuda-gdb:

```
$ cuda-gdb --pid 31562
CUDA Exception: Warp Illegal Address
The exception was triggered at PC 0x508ca70 (jacobi_kernels.cu:88)

Thread 1 "jacobi" received signal CUDA_EXCEPTION_14, Warp Illegal Address.
[Switching focus to CUDA kernel 0, grid 4, block (0,0,0), thread (0,20,0), device 0, sm 0, warp 21, lane 0]
0x000000000508ca80 in jacobi_kernel<32, 32><<<(512,512,1),(32,32,1)>>> (/*...*/) at jacobi_kernels.c
u:88

**Real foo = *((real*)nullptr);**
```

DEBUGGING MPI+CUDA APPLICATIONS

More environment variables for offline debugging

- With CUDA ENABLE COREDUMP ON EXCEPTION=1 core dumps are generated in case of an exception
 - CUDA_ENABLE_LIGHTWEIGHT_COREDUMP=1 does not dump application memory faster
 - Can be used for post-mortem debugging
 - Helpful if live debugging is not possible
- Enable/Disable CPU part of core dump (enabled by default)
 - CUDA_ENABLE_CPU_COREDUMP_ON_EXCEPTION
- Specify name of core dump file with CUDA_COREDUMP_FILE
- Open GPU
 - (cuda-gdb) target cudacore core.cuda
- Open CPU+GPU
 - (cuda-gdb) target core core.cpu core.cuda

https://docs.nvidia.com/cuda/cuda-gdb/index.html#gpu-coredump

EXAMPLE: OPENING A CORE DUMP

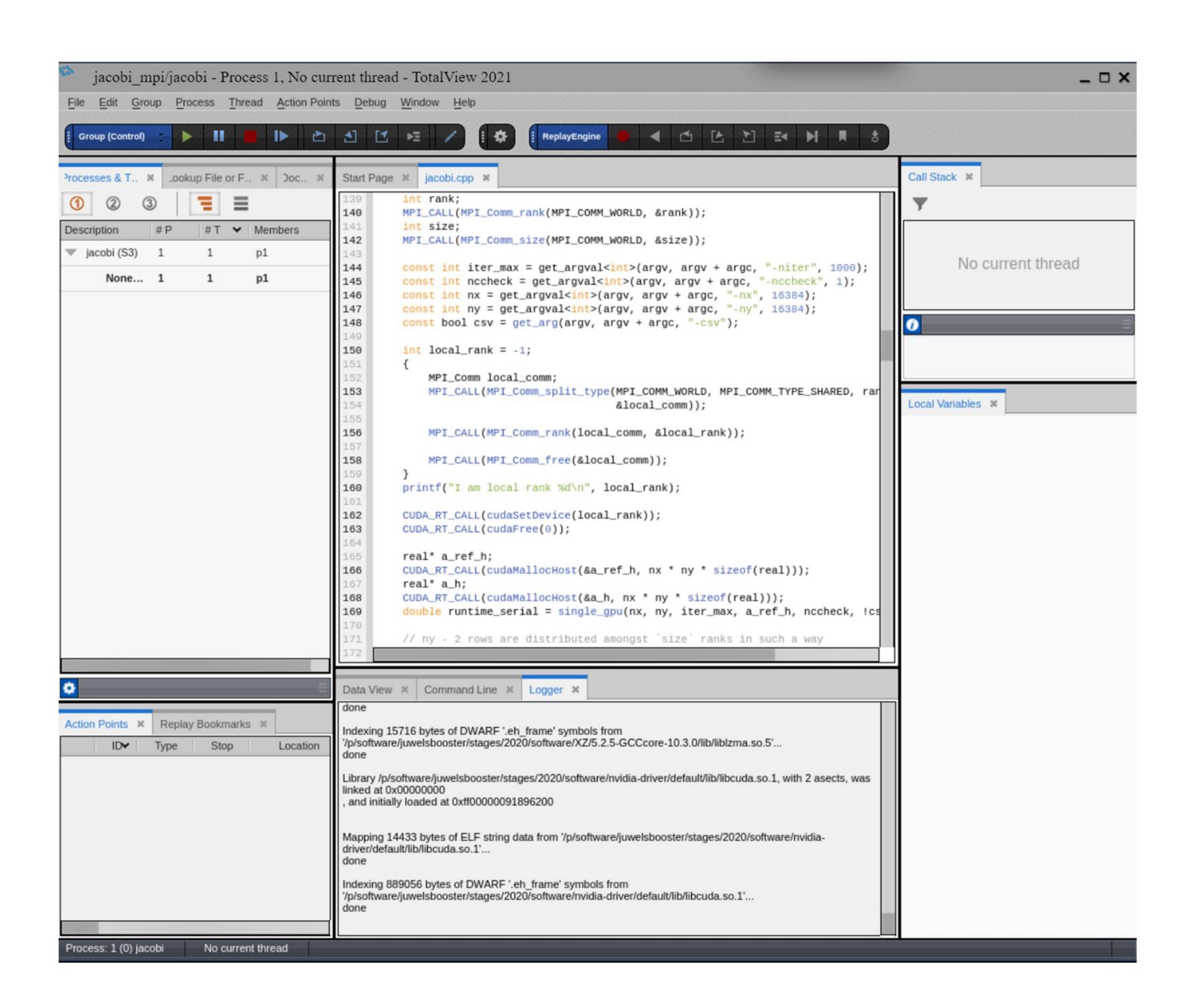
Running and generating the core file

```
$ CUDA_ENABLE_COREDUMP_ON_EXCEPTION=1 CUDA_ENABLE_LIGHTWEIGHT_COREDUMP=1 srun ./jacobi -niter 10 Single GPU jacobi relaxation: 10 iterations on 16384 x 16384 mesh with norm check every 1 iterations srun: error: jwb0021: tasks 0-3: Aborted (core dumped)

$ ls core*
core.jwb0021.juwels.23959 core_1633801834_jwb0021.juwels_23959.nvcudmp ...
```

And opening the core dump in cuda-qdb

SPECIALIZED PARALLEL DEBUGGERS



- cuda-gdb can debug multiple processes (add-inferior), although...
- For truly parallel bugs (e.g. multi-node, multi-process race conditions), third-party tools offer more convenience
 - Or enable "live" analysis in the first place
- ARM DDT
- Perforce TotalView (screenshot)
 - Available as module on JSC systems

WRITE DEBUGGABLE SOFTWARE

A case for modularity, and proper test cases

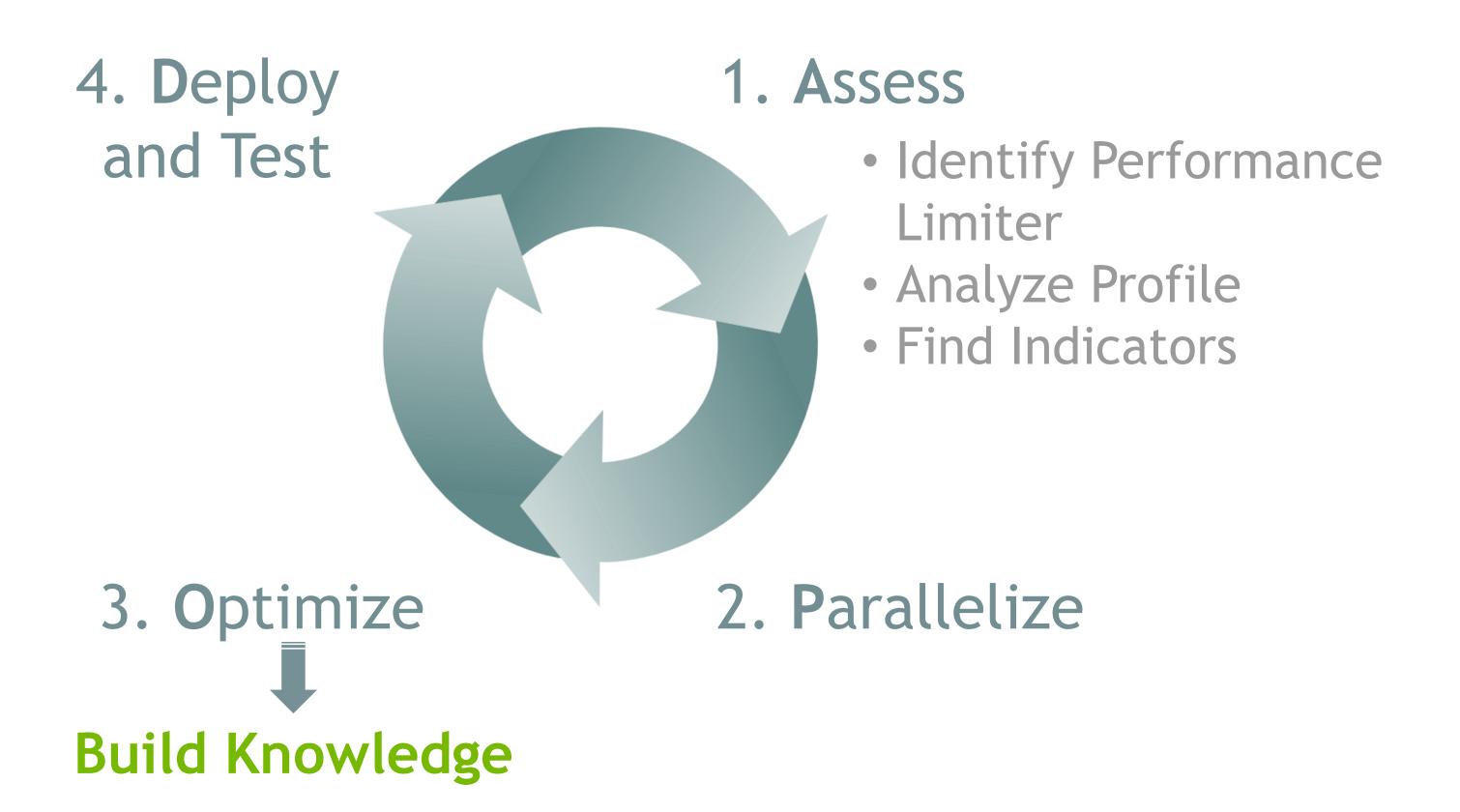
- Think about interfaces in your code: Which parts must depend on each other, etc.
 - Example: BLAS, linear algebra routines
- Think about structure and architecture ("the big picture")
- Don't go overboard: "I read this book, we need 100% test coverage", etc.
 - For many research codes that would be overkill
- "Everything should be made as simple as possible, but no simpler."
- Badly structured legacy code slows you down as well, as it resists change
 - Today's code is tomorrow's legacy
 - Strike a balance, avoid full rewrites. Code encapsulates hard-earned bug fixes and knowledge

- Representative test cases
 - Contain the correct science, walk the code paths
 - But run quickly, best on a single process, should run on a single node
 - Some (but not all) tests at full scale

DEBUGGING PERFORMANCE

Why you *must* use profilers

- Paraphrasing Donald Knuth:
 - Don't overoptimize, but meta-optimize your own time by using tools to focus on relevant parts
- Do not trust your gut instinct very often very misleading
 - Easy to waste a lot of time chasing the "perceived" issue
- Getting the same information, you end up reimplementing your own profiler
- Iterative workflow
- Different kinds of measurement tools, different tradeoffs
 - Instrumenting/Sampling
 - Profiling/Tracing
 - multi-process, single-process, kernel-level
- Here: Focus on GPU and system-level: Nsight Systems



THE NSIGHT SUITE COMPONENTS

How the pieces fit together

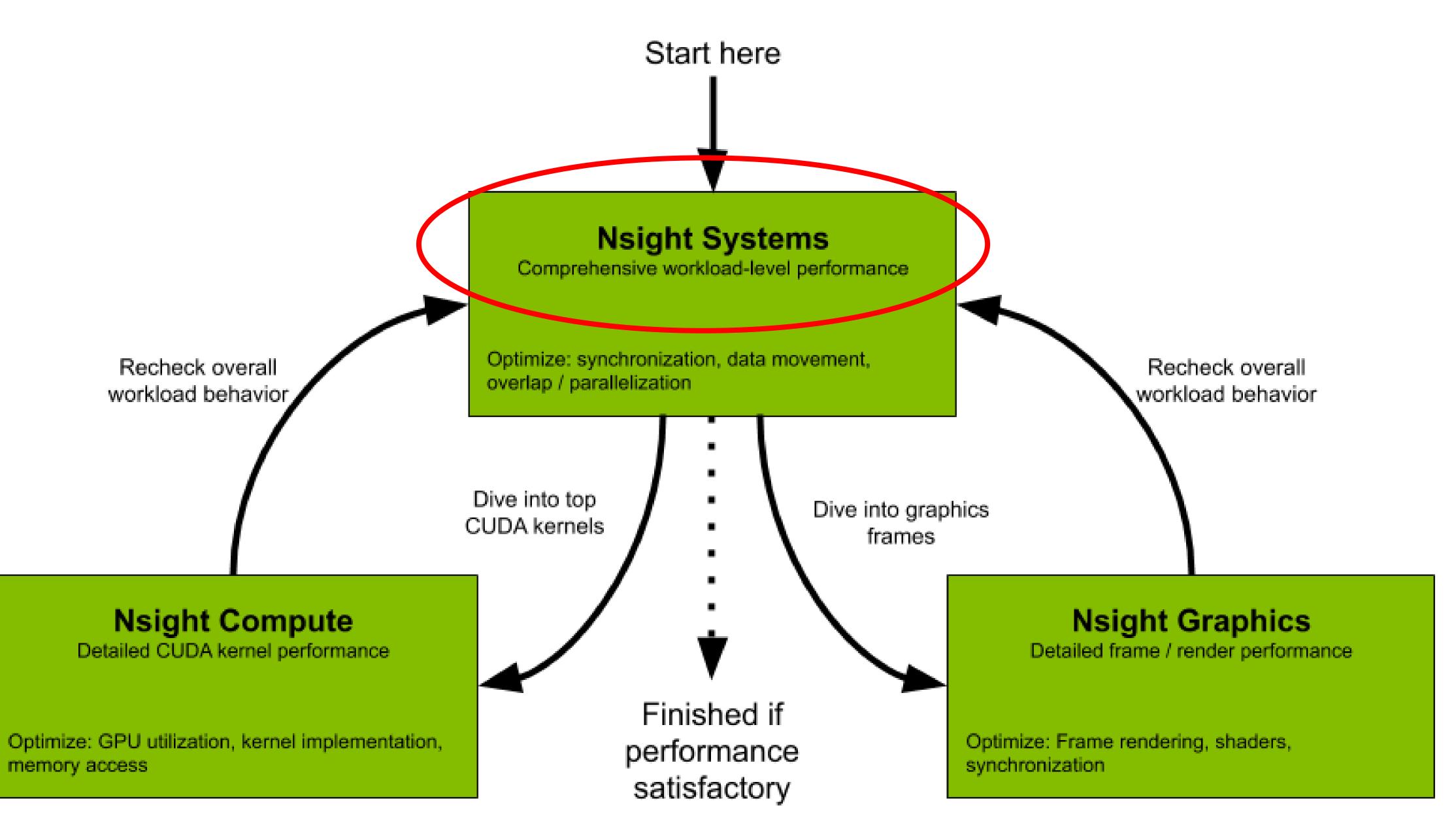


Nsight Systems: Coarse-grained, whole-application



Nsight Compute: Fine-grained, kernel-level

- NVTX: Support and structure across tools
- Main purpose: Performance optimization
 - But at their core, advanced measurement tools



A FIRST (I)NSIGHT

Recording with the CLI

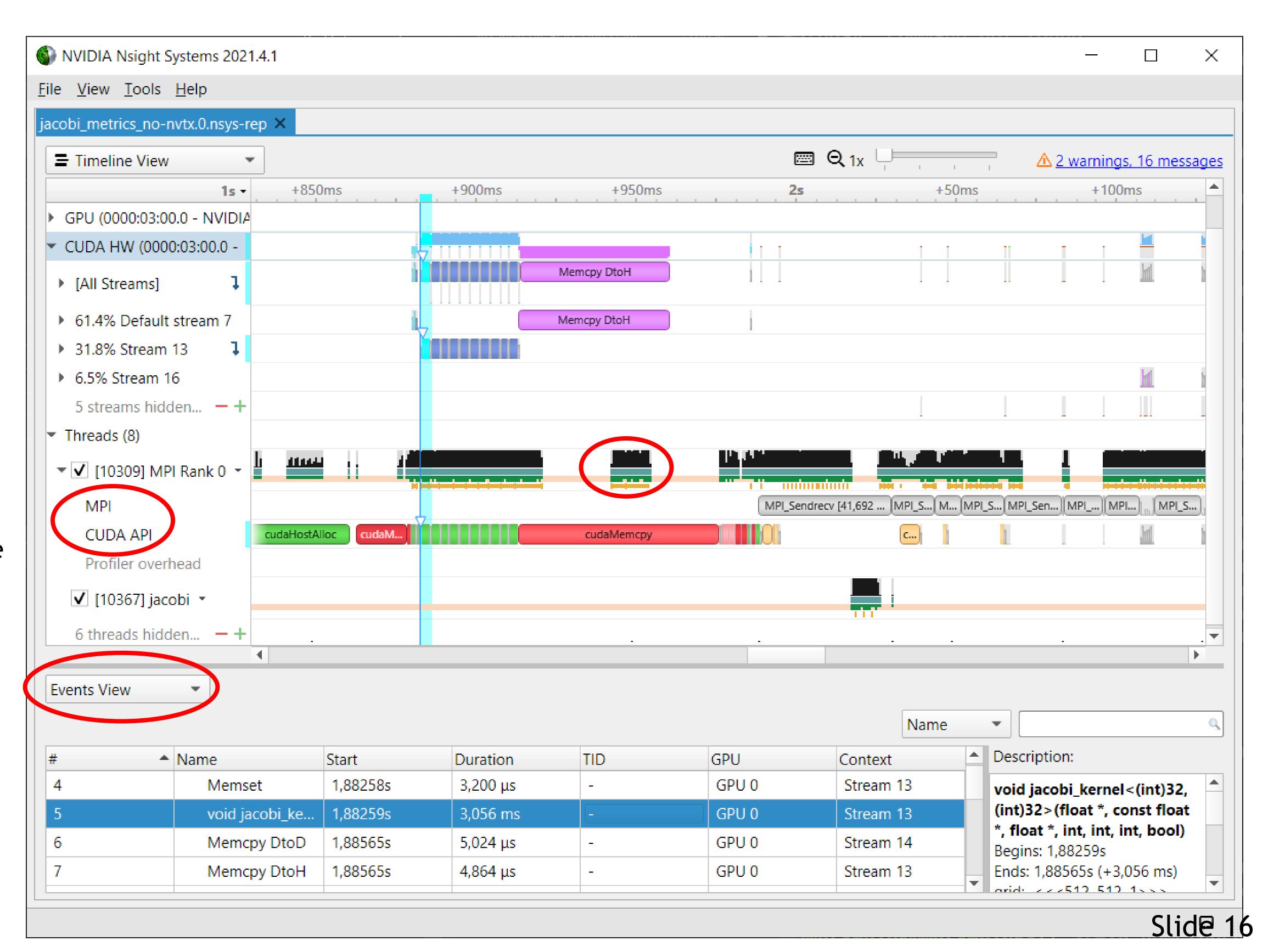
- Use the command line
 - srun nsys profile --trace=cuda, nvtx, mpi --output=my report.%q{SLURM PROCID} ./jacobi -niter 10
- Inspect results: Open the report file in the GUI
 - Also possible to get details on command line
 - Either add --stats to profile command line, or: nsys stats --help
- Runs set of reports on command line, customizable (sqlite + Python):
 - Useful to check validity of profile, identify important kernels

Running [.../reports/gpukernsum.py jacobi_metrics_more-nvtx.0.sqlite]...

Time(%)	Total Time (ns)	Instances	Avg (ns)	Med (ns)	Min (ns)	Max (ns)	StdDev (ns)	Name
99.9	36750359	20	1837518.0	1838466.5	622945	3055044	1245121.7	void jacobi kernel
0.1	22816	2	11408.0	11408.0	7520	15296	5498.5	initialize_boundaries

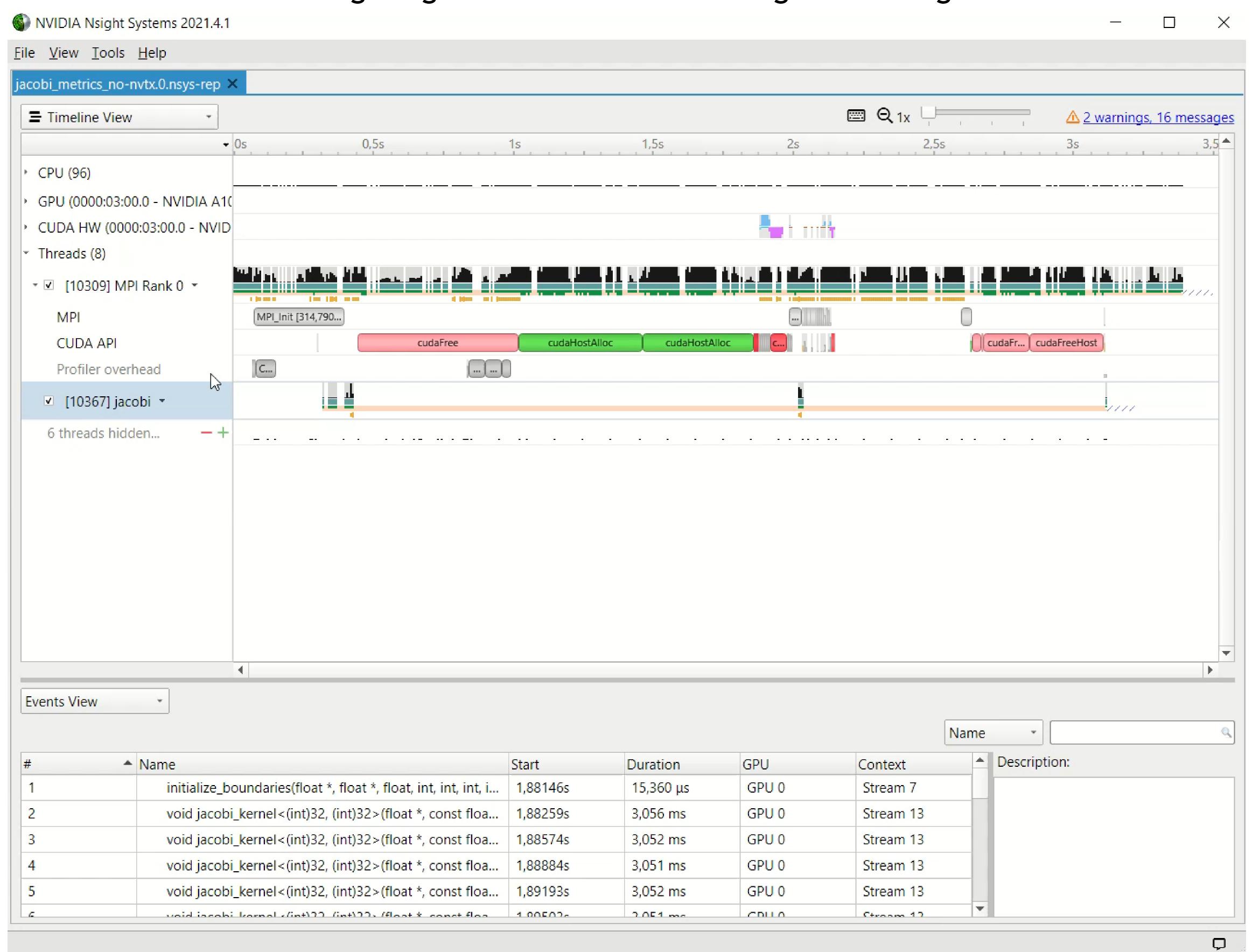
SYSTEM-LEVEL PROFILING WITH NSIGHT SYSTEMS

- Global timeline view
 - CUDA HW: streams, kernels, memory
- Different traces, e.g. CUDA, MPI
 - correlations API <-> HW
- Stack samples
 - bottom-up, top-down for CPU code
- GPU metrics
- Events View
 - Expert Systems
- looks at single process (tree)
 - correlate multi-process reports in single timeline



NSIGHT SYSTEMS BASIC WORKFLOW

Navigating the timeline and finding interesting areas



DISCOVERING OPTIMIZATION POTENTIAL

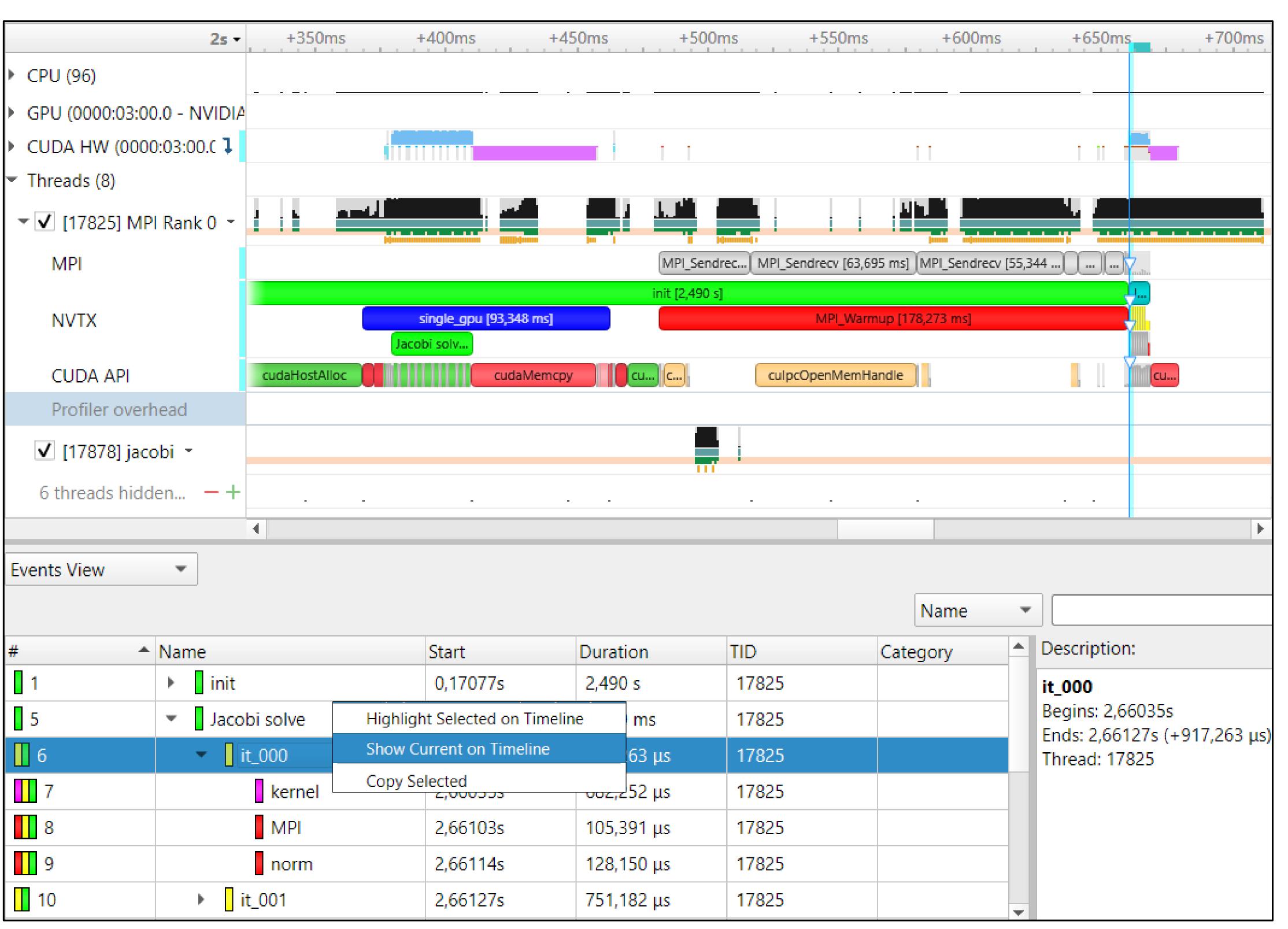
- Using our Jacobi example (see exercise)
- Spot kernels lots of whitespace
 - Which part is "bad"?
 - Enhance!
- MPI calls
 - Memory copies
 - We know: This is CUDA-aware MPI
- Even without knowing source, insight
- Too complicated for repeated/reliable usage
 - How to simplify navigating and comparing reports?



ADDING SOME COLOR

Code annotation with NVTX

- Same section of timeline as before
 - Events view: Quick navigation
- Like manual timing, only less work
- Nesting
- Correlation, filtering



ADDING NVTX

Simple range-based API

- #include "nvtx3/nvToolsExt.h"
 - NVTX v3 is header-only, needs just -ldl
 - C++ and Python APIs
- Fortran: NVHPC compilers include module
 - Just use nvtx and -lnvhpcwrapnvtx
 - Other compilers: See blog posts linked below
- Definitely: Include PUSH/POP macros (see links below)

```
PUSH RANGE (name, color idx)
```

- Sprinkle them strategically through code
 - Use hierarchically: Nest ranges
- Not shown: Advanced usage (domains, ...)
- Similar range-based annotations exist for other tools
 - e.g. <u>SCOREP_USER_REGION_BEGIN</u>

```
int main(int argc, char** argv) {
   PUSH_RANGE("main", 0)
   PUSH_RANGE("init", 1)
   do_initialization();
   POP_RANGE
   /* ... */
   PUSH_RANGE("computation", 2)
    jacobi_kernel<<</* ... */, compute_stream>>>(...);
   cudaStreamSynchronize(compute_stream);
   POP_RANGE
   /* ... */
   POP_RANGE
}
```

https://github.com/NVIDIA/NVTX and https://nvidia.github.io/NVTX/#how-do-i-use-nvtx-in-my-code

https://developer.nvidia.com/blog/cuda-pro-tip-generate-custom-application-profile-timelines-nvtx/https://developer.nvidia.com/blog/customize-cuda-fortran-profiling-nvtx/

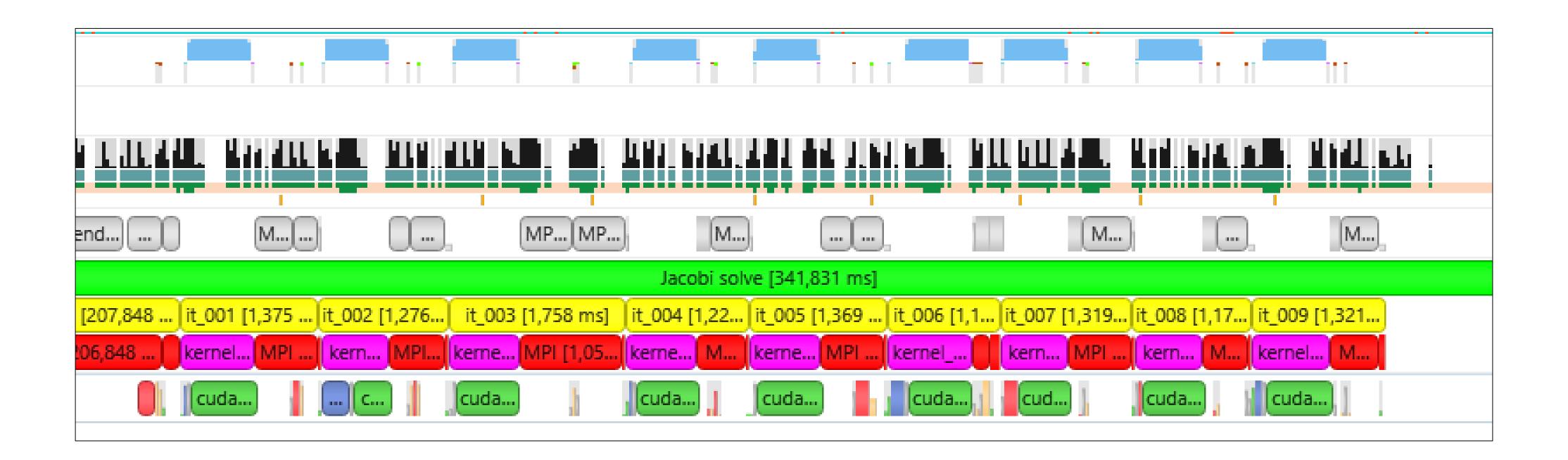
MINIMIZING PROFILE SIZE

Shorter time, smaller files = quicker progress

- Only profile what you need all profilers have some overhead
 - Example: Event that occurs after long-running setup phase
- Bonus: lower number of events leads to smaller file size
- Add to nsys command line:

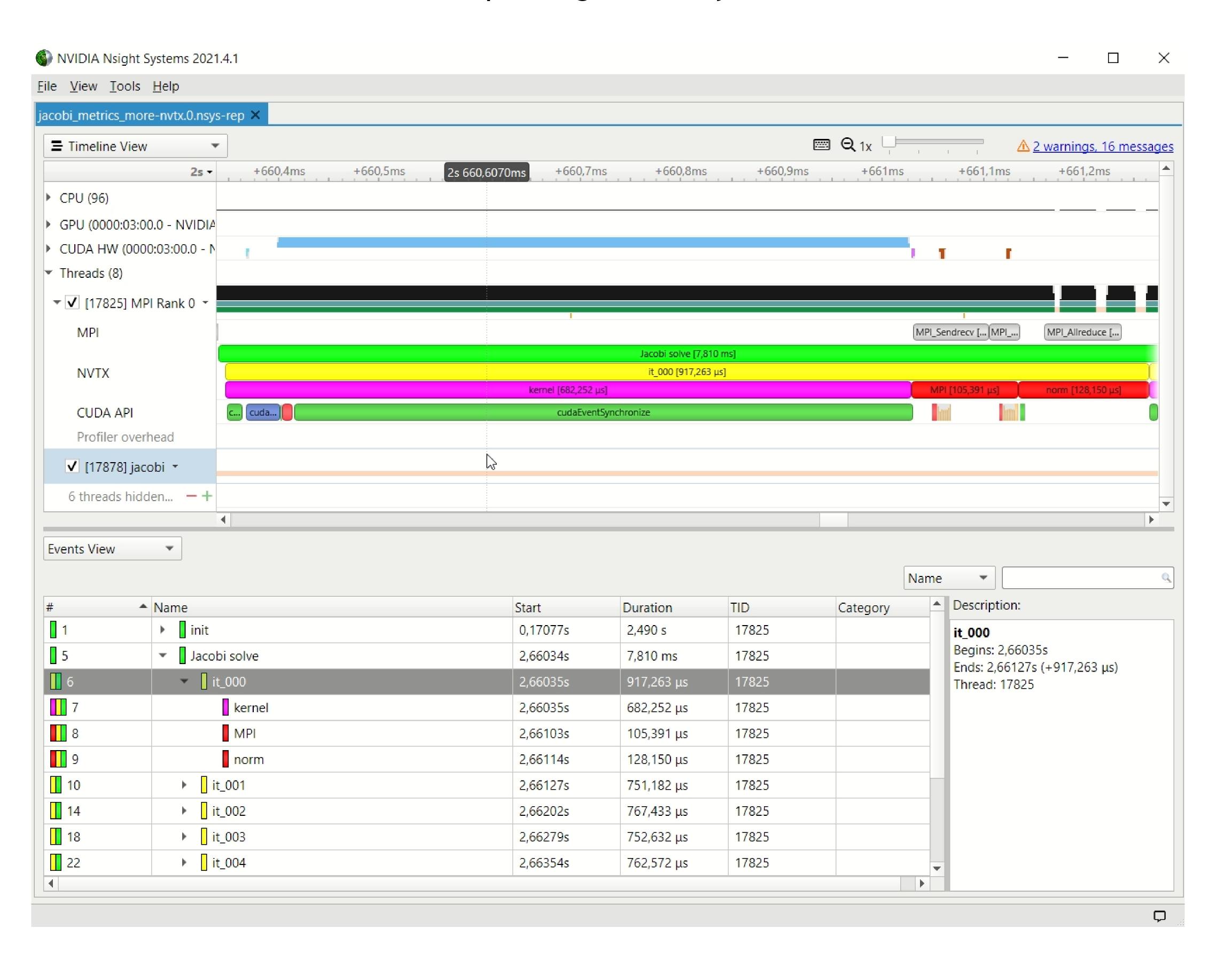
```
    --capture-range=nvtx --nvtx-capture=any_nvtx_marker_name \
        --env-var=NSYS_NVTX_PROFILER_REGISTER_ONLY=0 --kill none
```

- Alternatively: cudaProfilerStart() and -Stop()
 - --capture-range=cudaProfilerApi



NSIGHT SYSTEMS WORKFLOW WITH NVTX

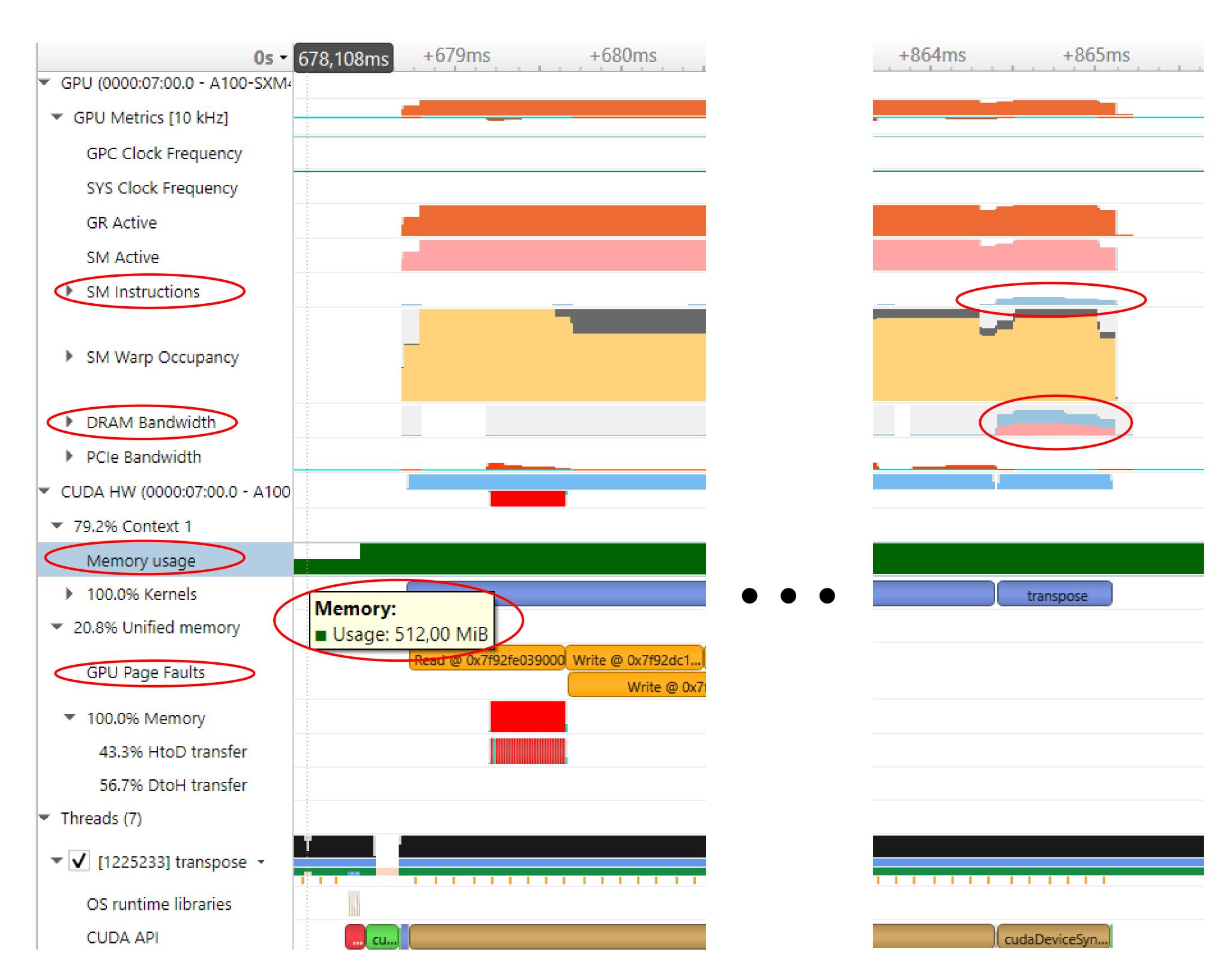
Repeating the analysis



GPU METRICS IN NSIGHT SYSTEMS

...and other traces you can activate

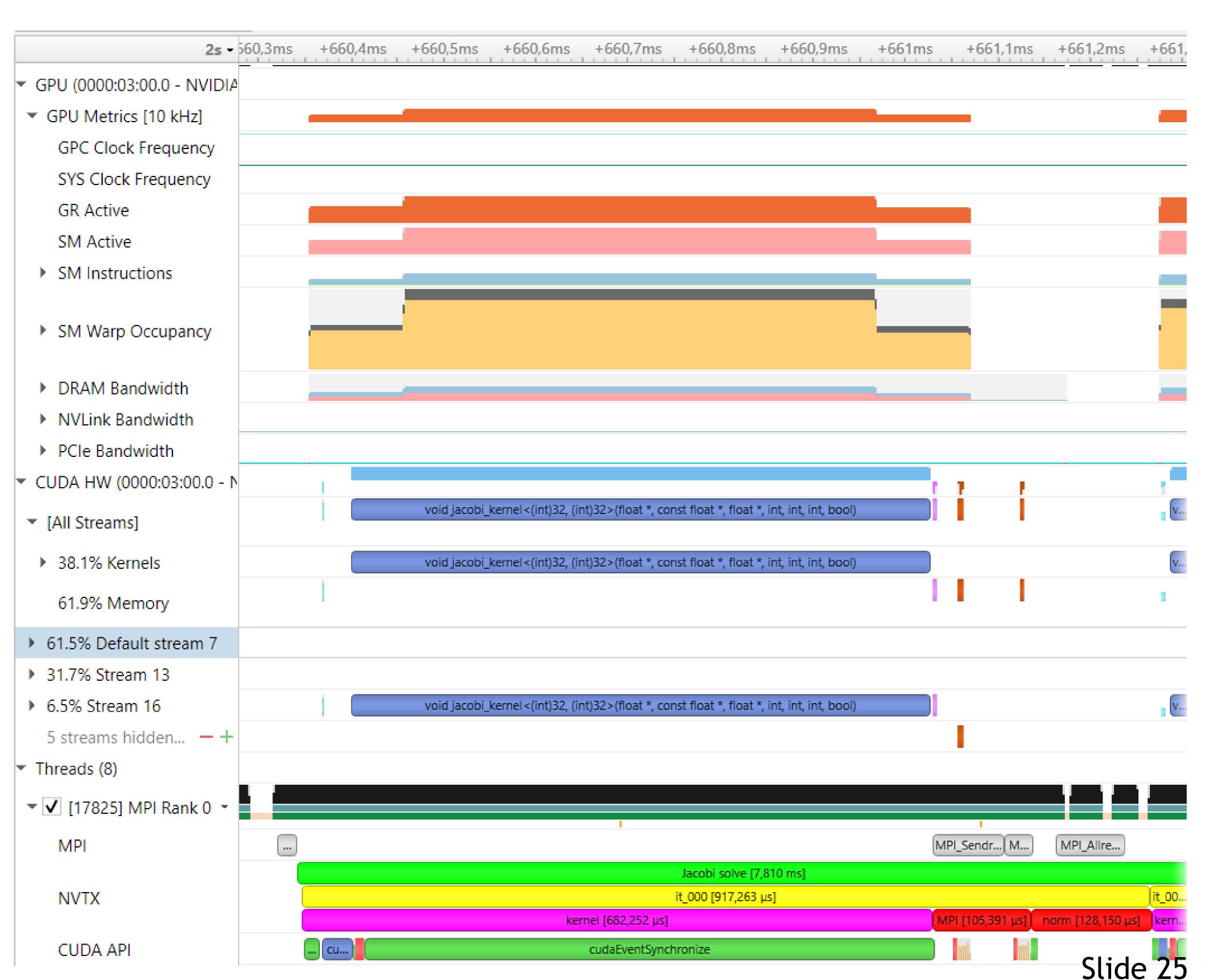
- Valuable low-overhead insight into HW usage:
 - SM instructions
 - DRAM Bandwidth, PCIe Bandwith (GPUDirect)
- Also: Memory usage, Page Faults (higher overhead)
 - CUDA Programming guide: <u>Unified Memory</u>
 <u>Programming</u>
- Can save kernel-level profiling effort!
- nsys profile
 - --gpu-metrics-device=0
 - --cuda-memory-usage=true
 - --cuda-um-cpu-page-faults=true
 - --cuda-um-gpu-page-faults=true ./app



FOCUSING THE ANALYSIS

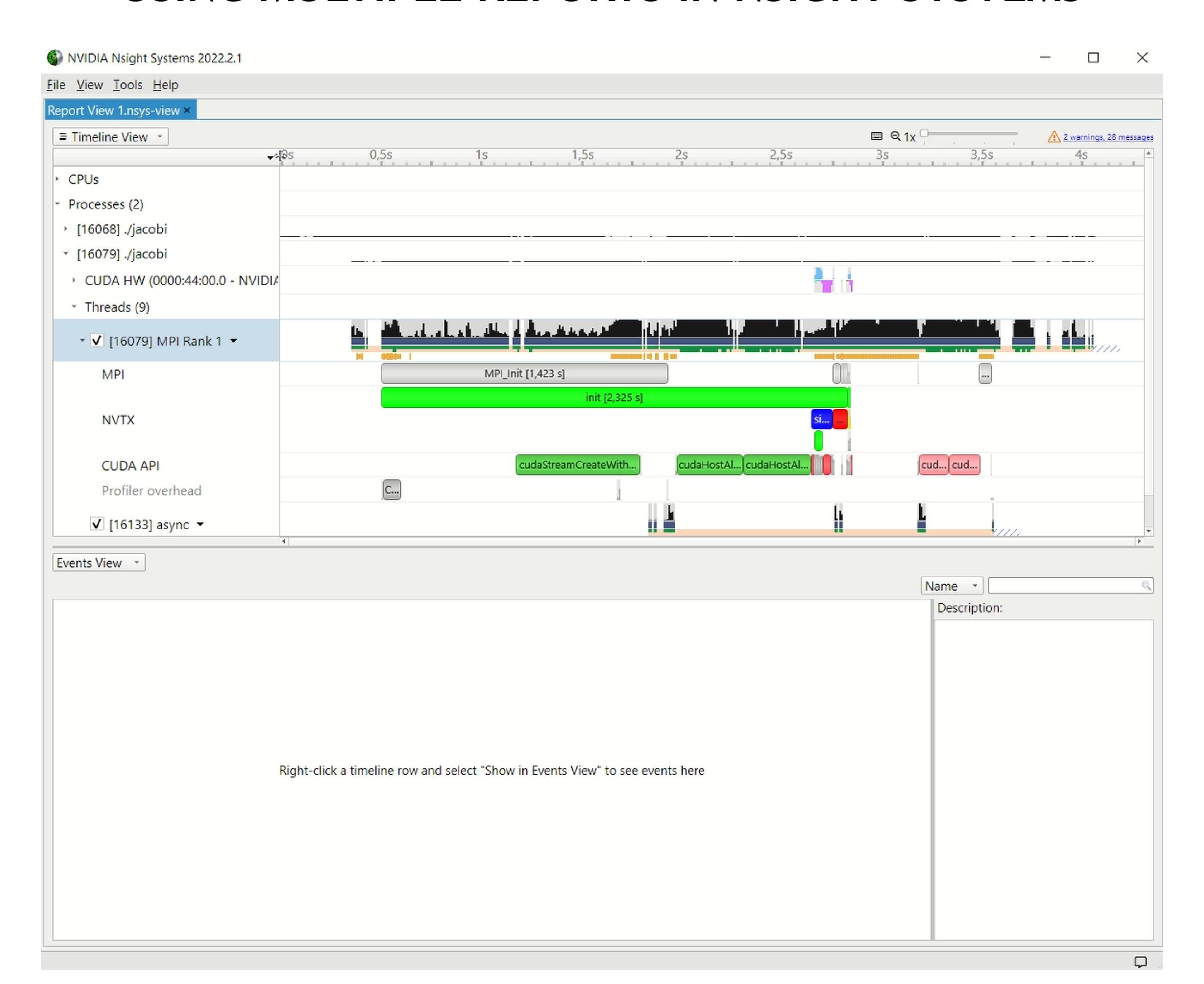
Introducing GPU metrics sampling

- Discover the "unit cell" of performance
 - in our case: single iteration
- Other blank spots during setup can be ignored (amortized, many more iterations)
- Maybe: Too small for proper comms profiling
- Kernel itself adequately using GPU
 - Remaning blank spots?
- Norm calculation
 - Can be turned off
- But still: Overlap potential? Can we run kernel during MPI?
 - later lectures



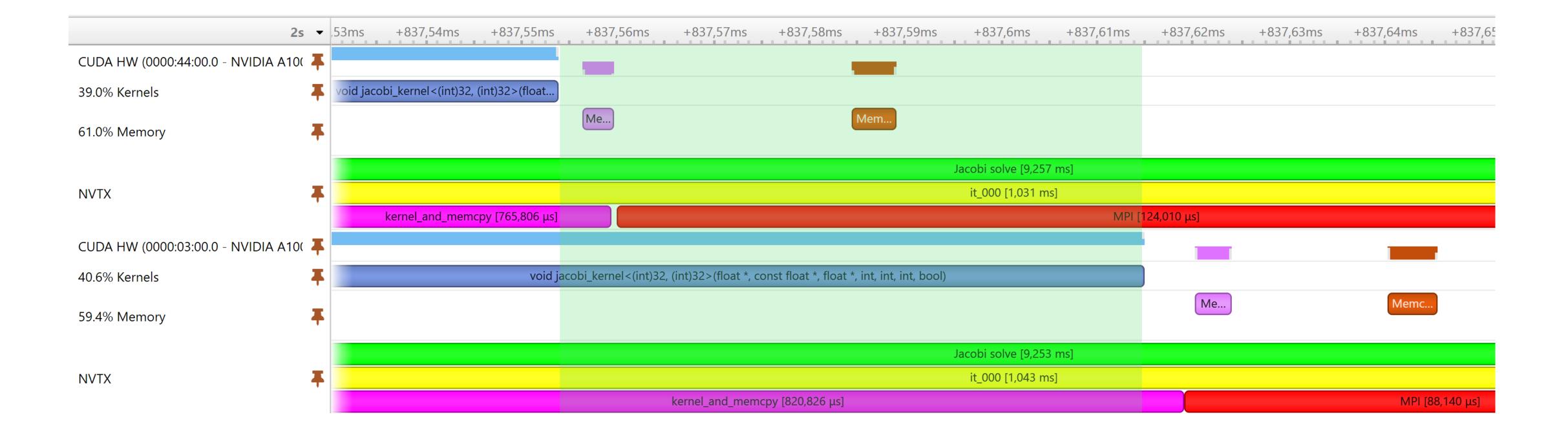


USING MULTIPLE REPORTS IN NSIGHT SYSTEMS



MULTI-PROCESS GPU ANALYSIS

- Load multiple reports into timeline
 - analyze differences in execution, GPU utilization
- Pin rows for comparison
- Example: End time of kernel execution



COMMUNITY PROFILING TOOLS

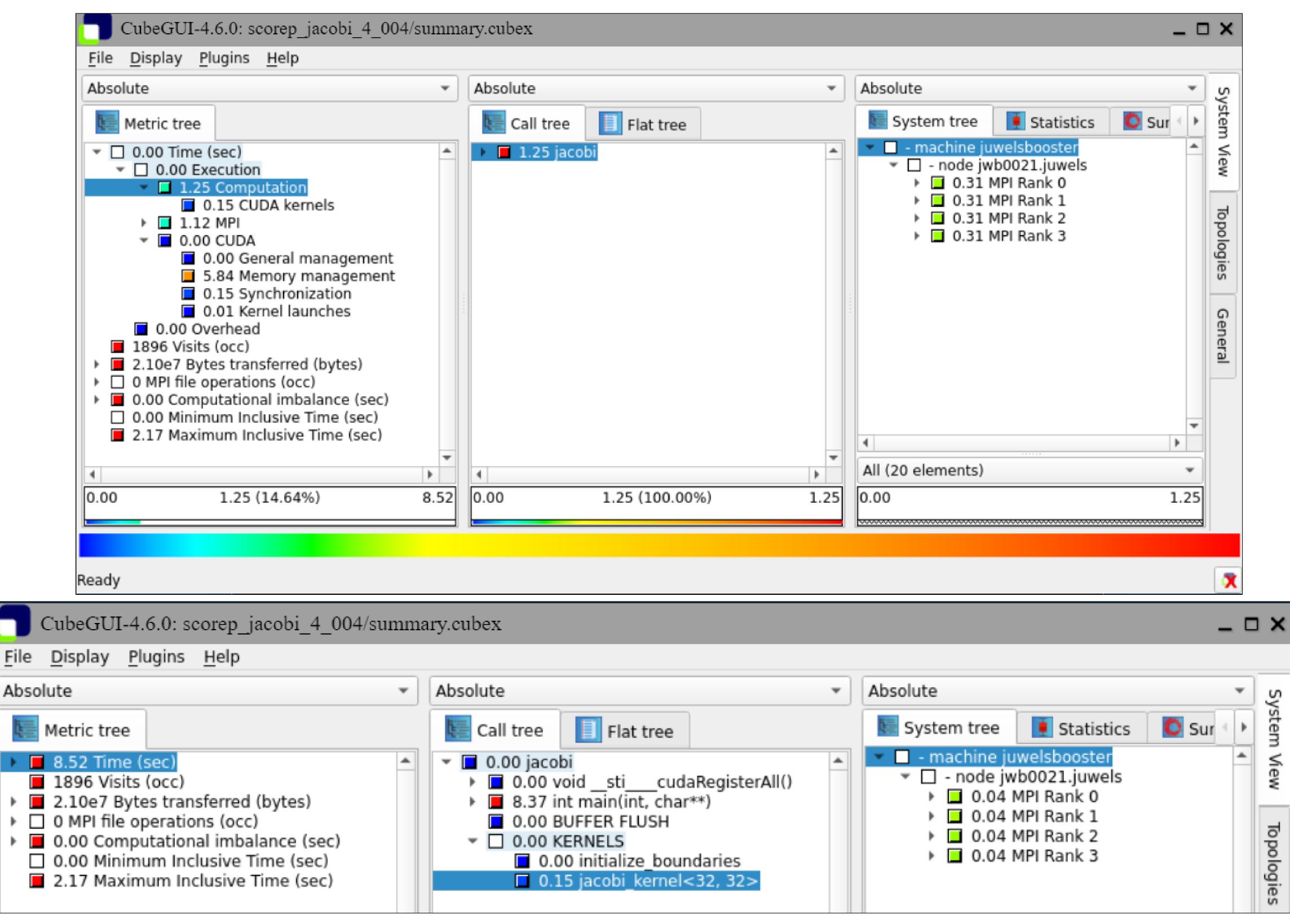
Specialized for large-scale distributed analysis

- Detecting issues at scale of thousands of GPUs (and processes)
 - Need to slice and dice data, too much to make sense of raw data
- Common measurement/instrumentation infrastructure: Score-P
 - Prefix all compilation/linker commands with scorep -cuda
- GPU data integration
 - CUDA profiling tools interface (CUPTI)
- Run the application to collect...
 - profiling data, for Scalasca
 - tracing data, for Vampir
 - (selection of tools not exhaustive)
- Tracing in particular: Careful tuning to keep overhead low (filtering)



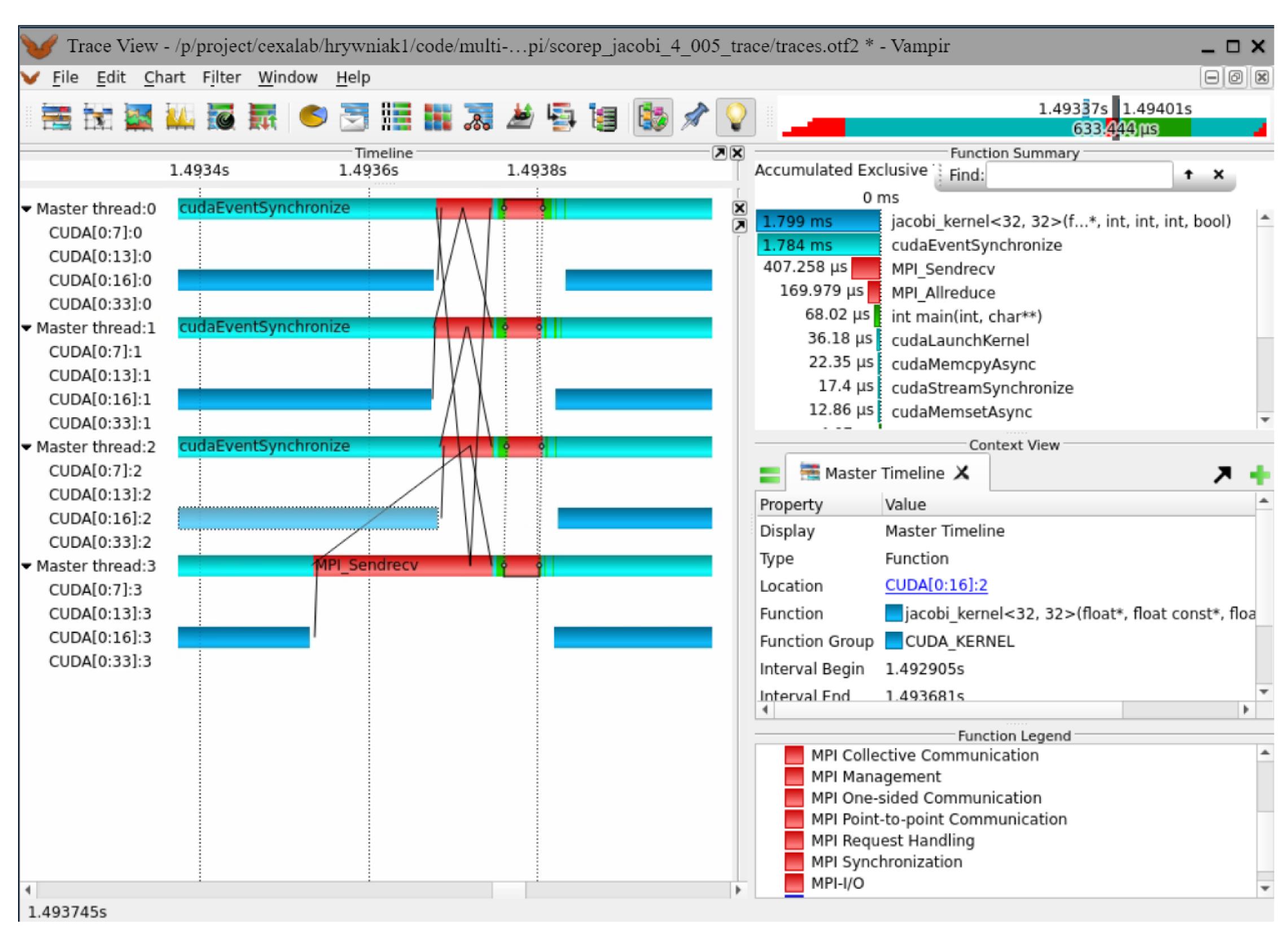
SCALASCA / CUBE

- Breakdown of different metrics across functions and processes
- Left-to-right: Selection influences breakdown
- Expanding changes inclusive/exclusive
- Example analysis:
 - Detect computational imbalance
- https://scalasca.org/



VAMPIR TRACE

- Analyze multi-process patterns
- What you can see in screenshot
 - Main timeline
 - Function summary
- Example analysis: Pinpoint MPI message relationships
 - e.g. late sender issues
- https://vampir.eu/



SUMMARY

- Looked at a wide selection of different tools
 - compute-sanitizer
 - cuda-gdb
 - Nsight-Systems
 - Score-P: Scalasca, Vampir, ...
 - and don't forget compiler flags and checks
- Correctness is paramount, but so is optimal resource usage
- Pick right tool for the job and take the time to learn it thoroughly
 - Do not trust your gut when analyzing performance, easy to be misled
 - How to adapt serial (or small-scale) tooling to highly distributed applications
- Meant as guideline, not gospel
 - Especially performance issues often require creativity to solve
- Workflow is equally important

FURTHER MATERIAL

- GTC on-demand talks
 - What, Where, and Why? Use CUDA Developer Tools to Detect, Locate, and Explain Bugs and Bottlenecks (s41493, GTC 2022)
 - Tuning GPU Network and Memory Usage in Apache Spark (s31566, GTC 2022)
- Documentation for <u>cuda-gdb</u>, <u>compute-sanitizer</u> and <u>Nsight Systems</u>
- GTC labs from Nsight teams: https://github.com/NVIDIA/nsight-training
- GPU bootcamp material, e.g., https://github.com/gpuhackathons-org/gpubootcamp/tree/master/hpc/multi_gpu_nways

