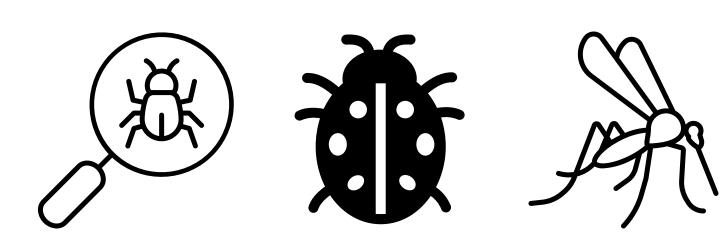


GOALS FOR THIS SESSION

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/languages (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

VCC 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:

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

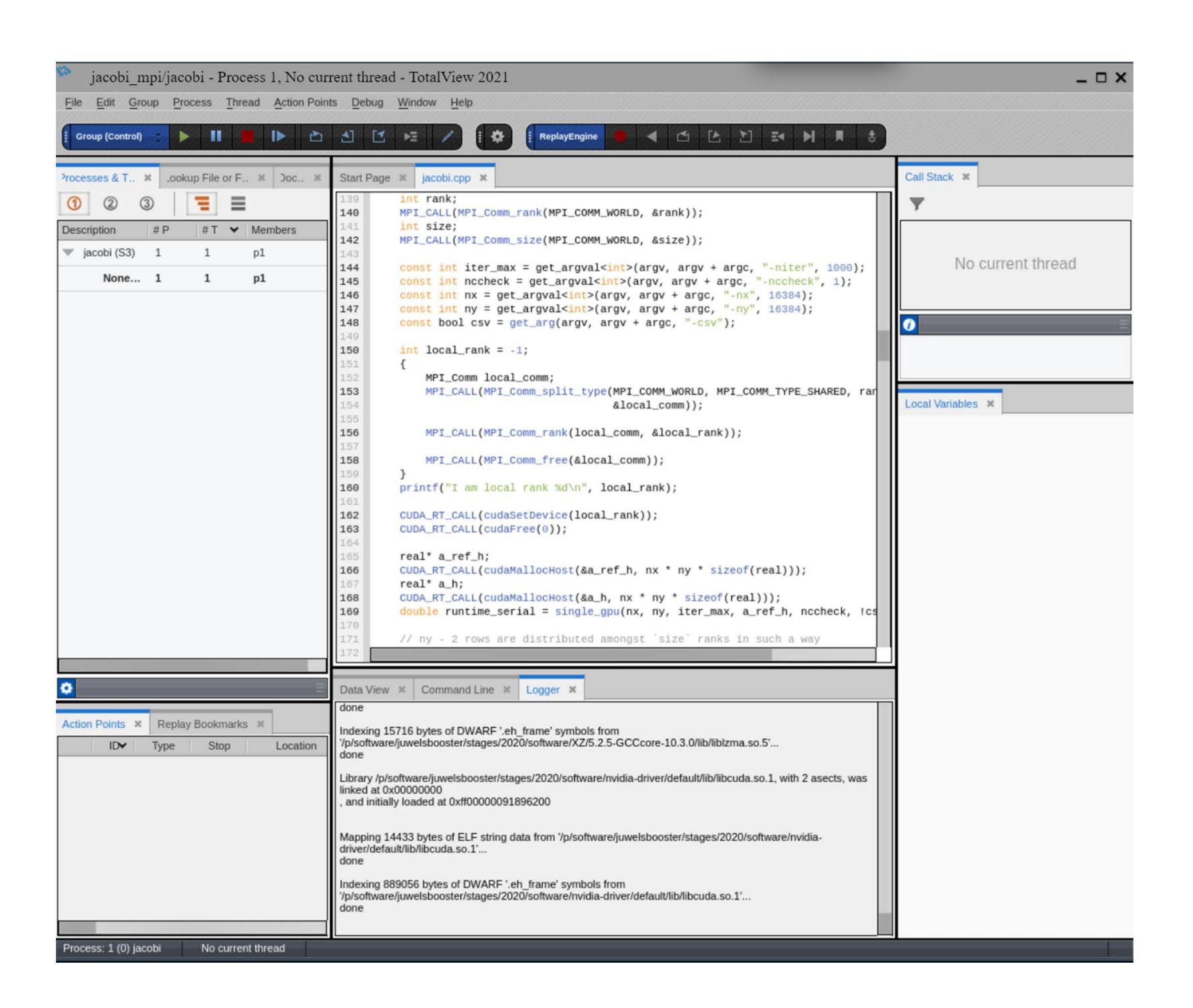
EXAMPLE: OPENING A CORE DUMP

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-gdb

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

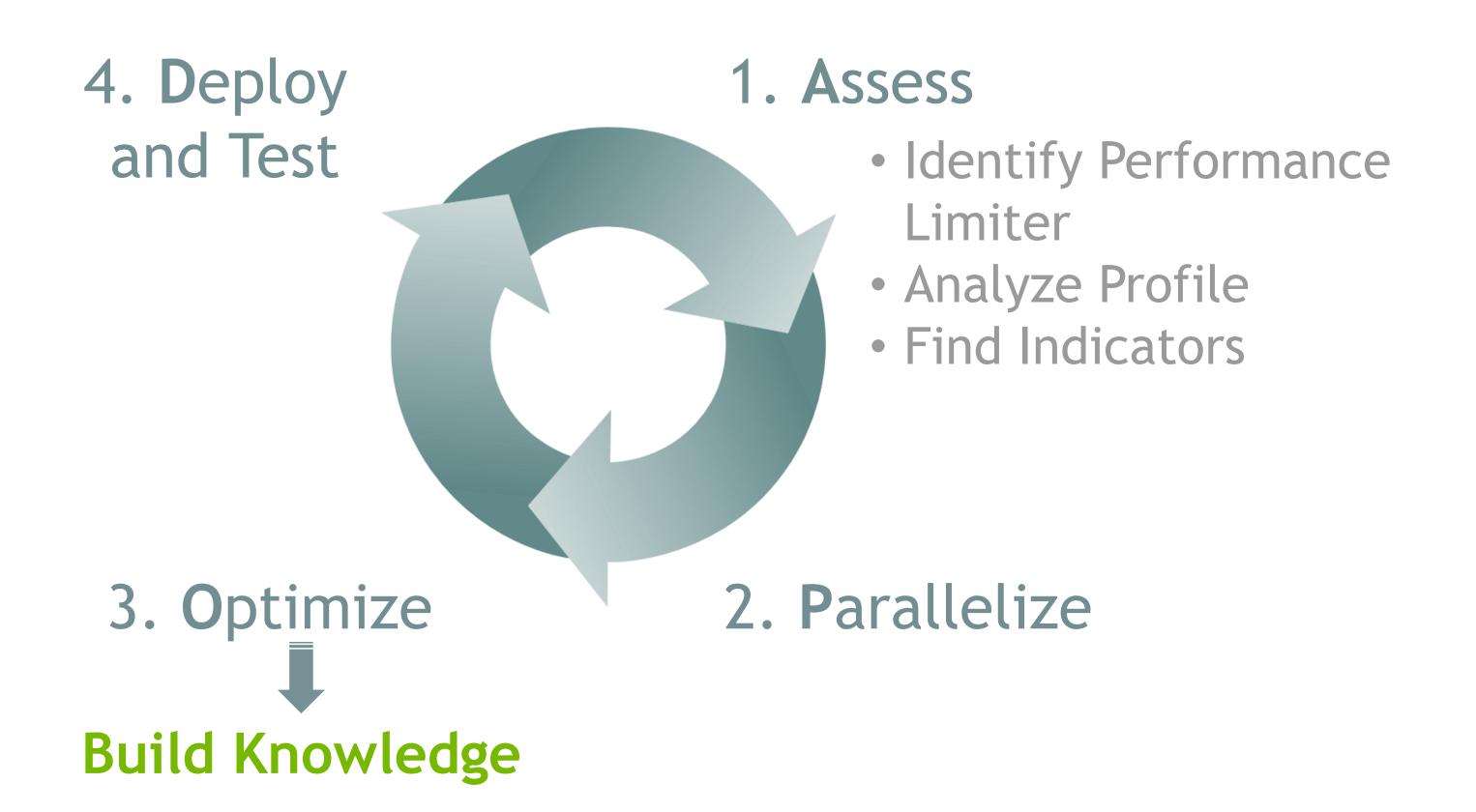
- Think about interfaces in your code: Which parts must depend on each other, etc.
 - Example: BLAS, linear algebra routines
- Think about structure and architecture ("the big picture")
- Don't go overboard: "I read this book, we need 100% test coverage", etc.
 - For many research codes that would be 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

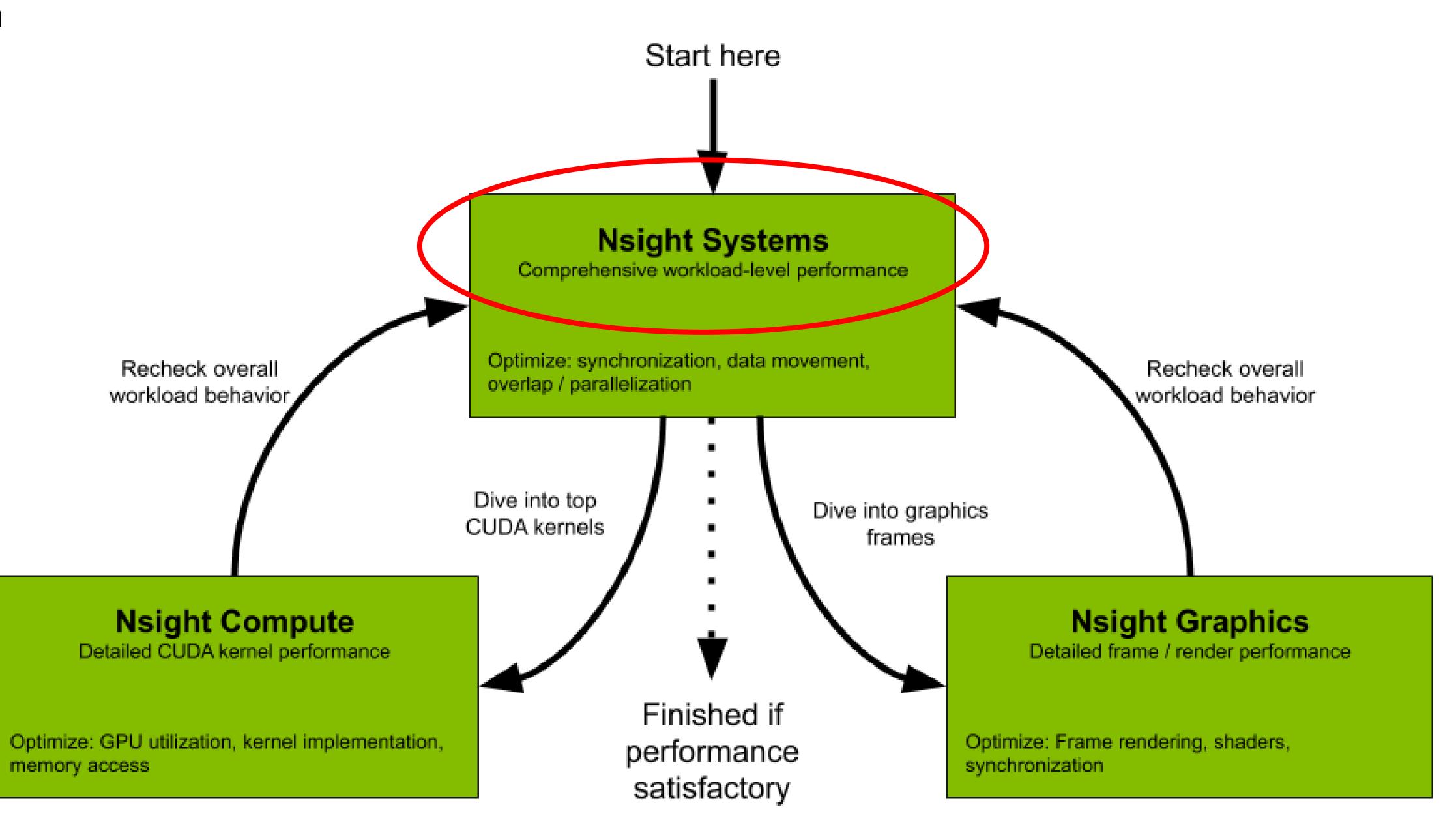


Nsight Systems: Coarse-grained, whole-application



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

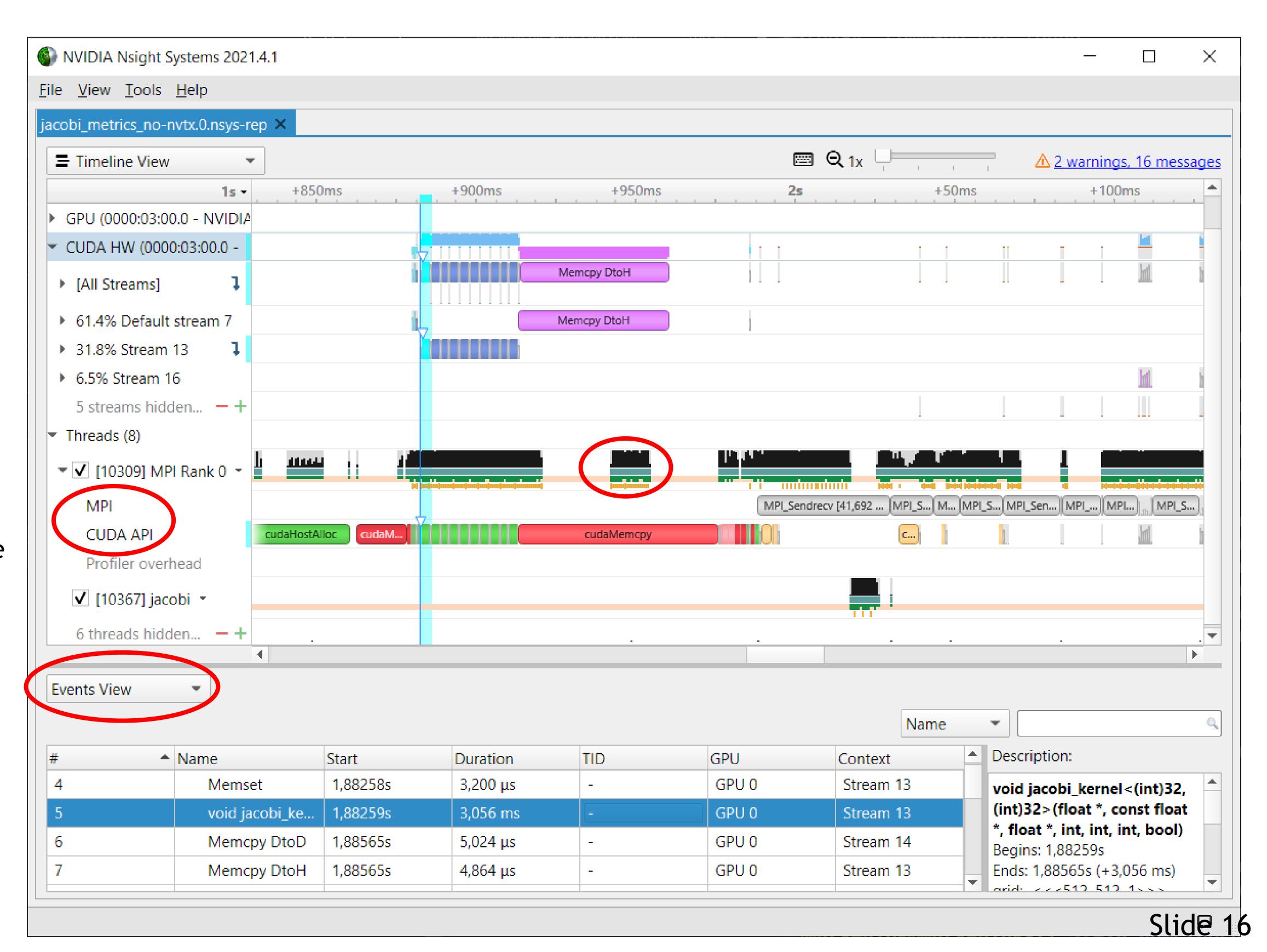
- Use the command line
 - srun nsys profile --trace=cuda, nvtx, mpi --output=my report.%q{SLURM PROCID} ./jacobi -niter 10
- Inspect results: Open the report file in the GUI
 - Also possible to get details on command line
 - Either add --stats to profile command line, or: nsys stats --help
- Runs set 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

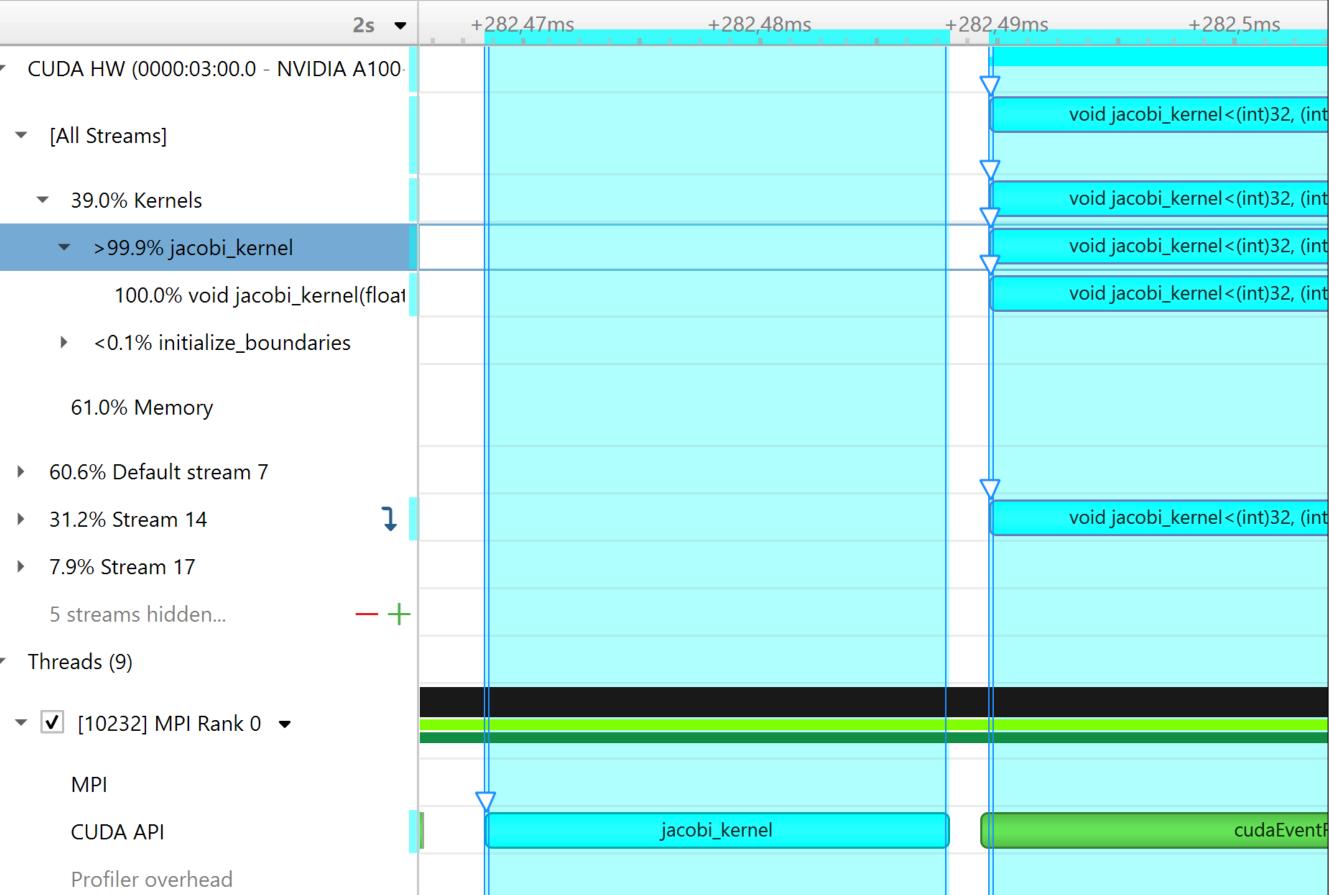
- Global timeline view
 - CUDA HW: streams, kernels, memory
- Different traces, e.g. CUDA, MPI
 - correlations API <-> HW
- Stack samples
 - bottom-up, top-down for CPU code
- GPU metrics
- Events View
 - Expert Systems
- looks at single process (tree)
 - correlate multi-process reports in single timeline



CORRELATING EVENTS ON THE TIMELINE

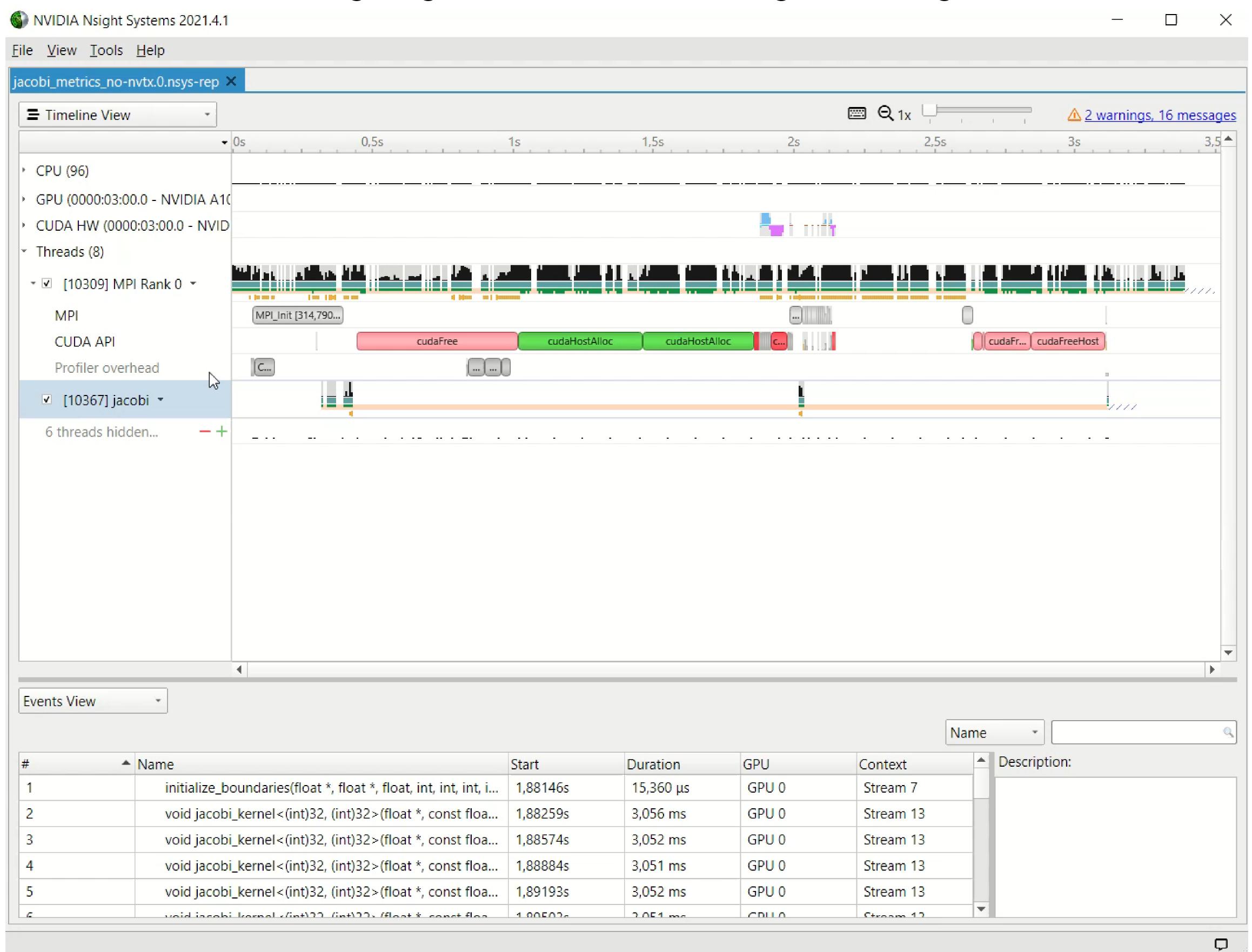
Selecting events in one row highlights related events





NSIGHT SYSTEMS BASIC WORKFLOW

Navigating the timeline and finding interesting areas



DISCOVERING OPTIMIZATION POTENTIAL

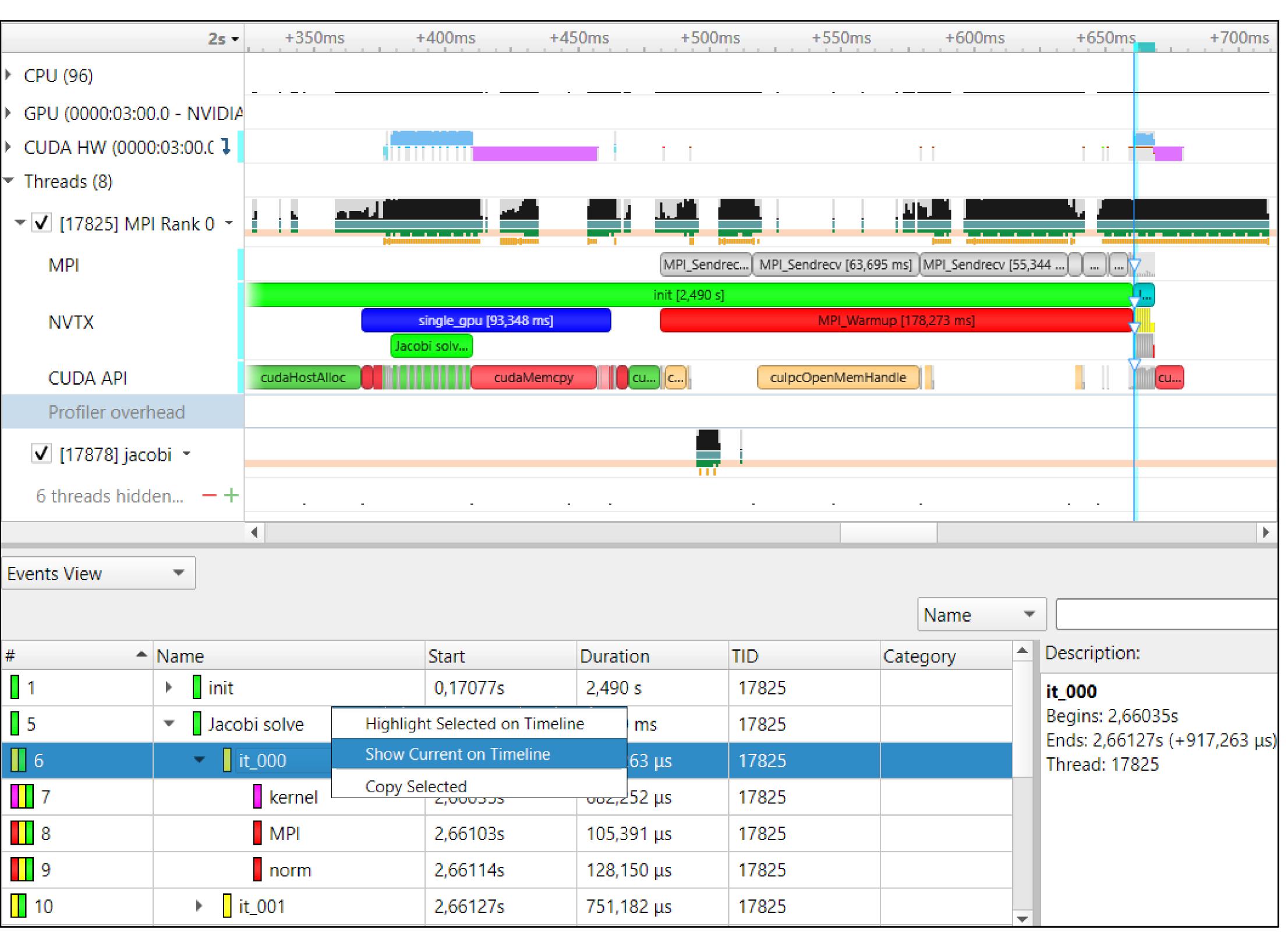
- Using our Jacobi example (see exercise)
- Spot kernels lots of whitespace
 - Which part is "bad"?
 - Enhance!
- MPI calls
 - Memory copies
 - We know: This is CUDA-aware MPI
- Even without knowing source, insight
- Too complicated for repeated/reliable usage
 - How to simplify navigating and comparing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

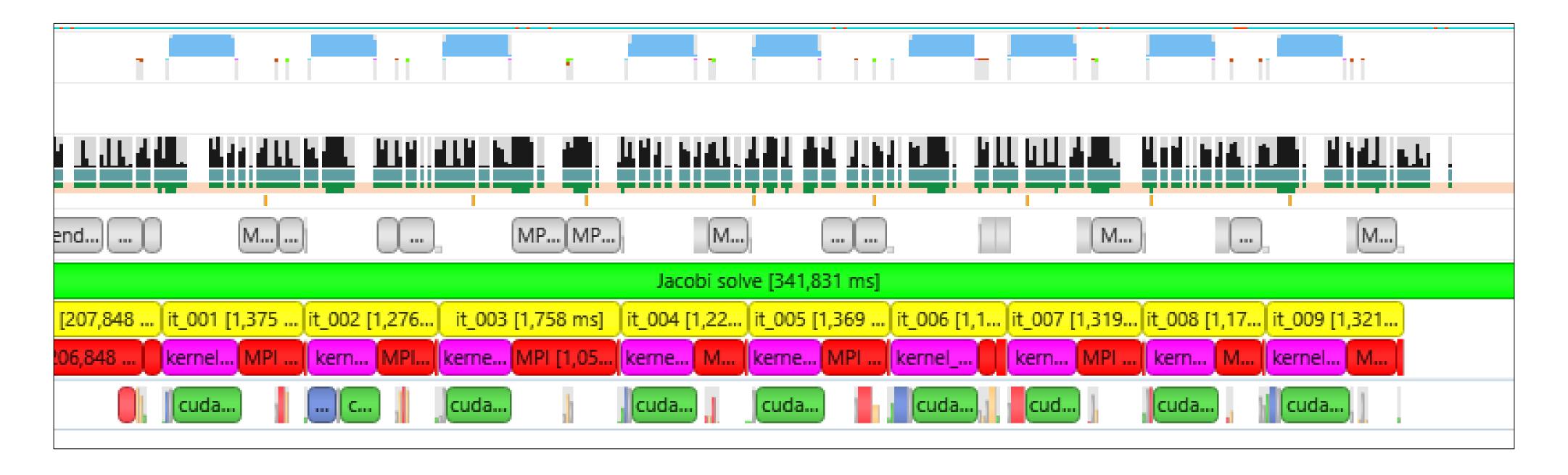
MINIMIZING PROFILE SIZE

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:

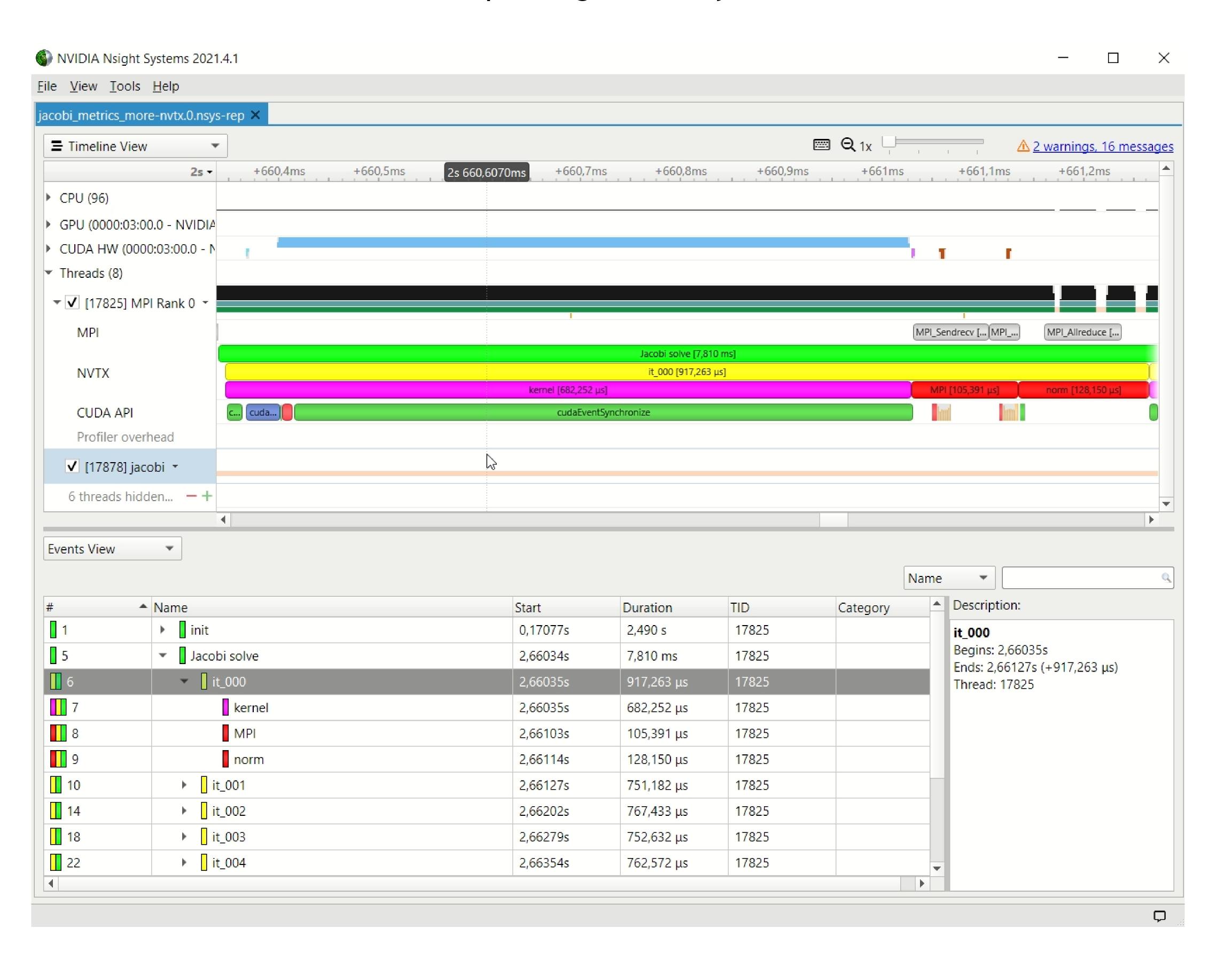
```
    --capture-range=nvtx --nvtx-capture=any_nvtx_marker_name \
        --env-var=NSYS_NVTX_PROFILER_REGISTER_ONLY=0 --kill none
```

- Use <u>NVTX registered strings</u> for best performance
- Alternatively: cudaProfilerStart() and -Stop()
 - --capture-range=cudaProfilerApi



NSIGHT SYSTEMS WORKFLOW WITH NVTX

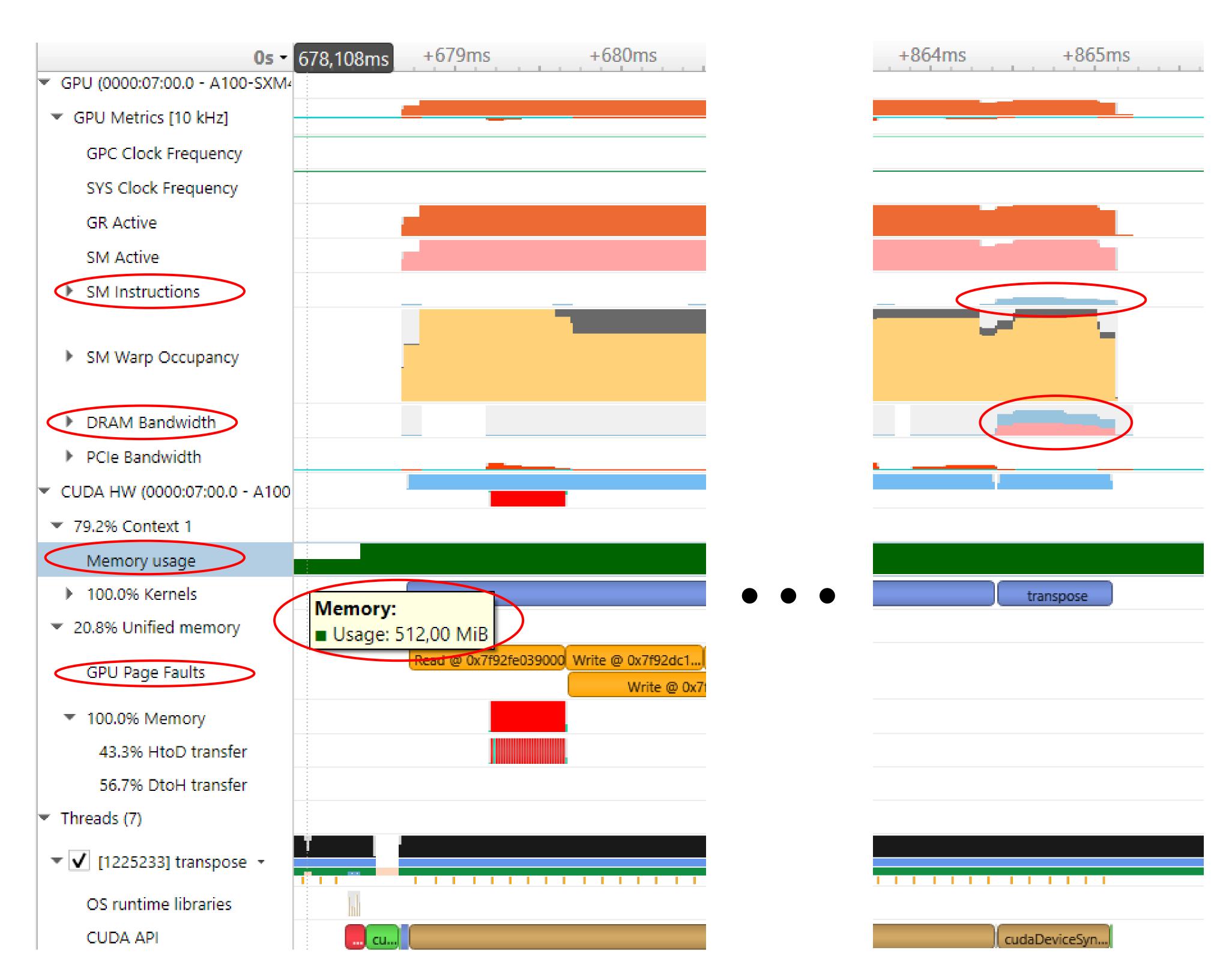
Repeating the analysis



GPU METRICS IN NSIGHT SYSTEMS

...and other traces you can activate

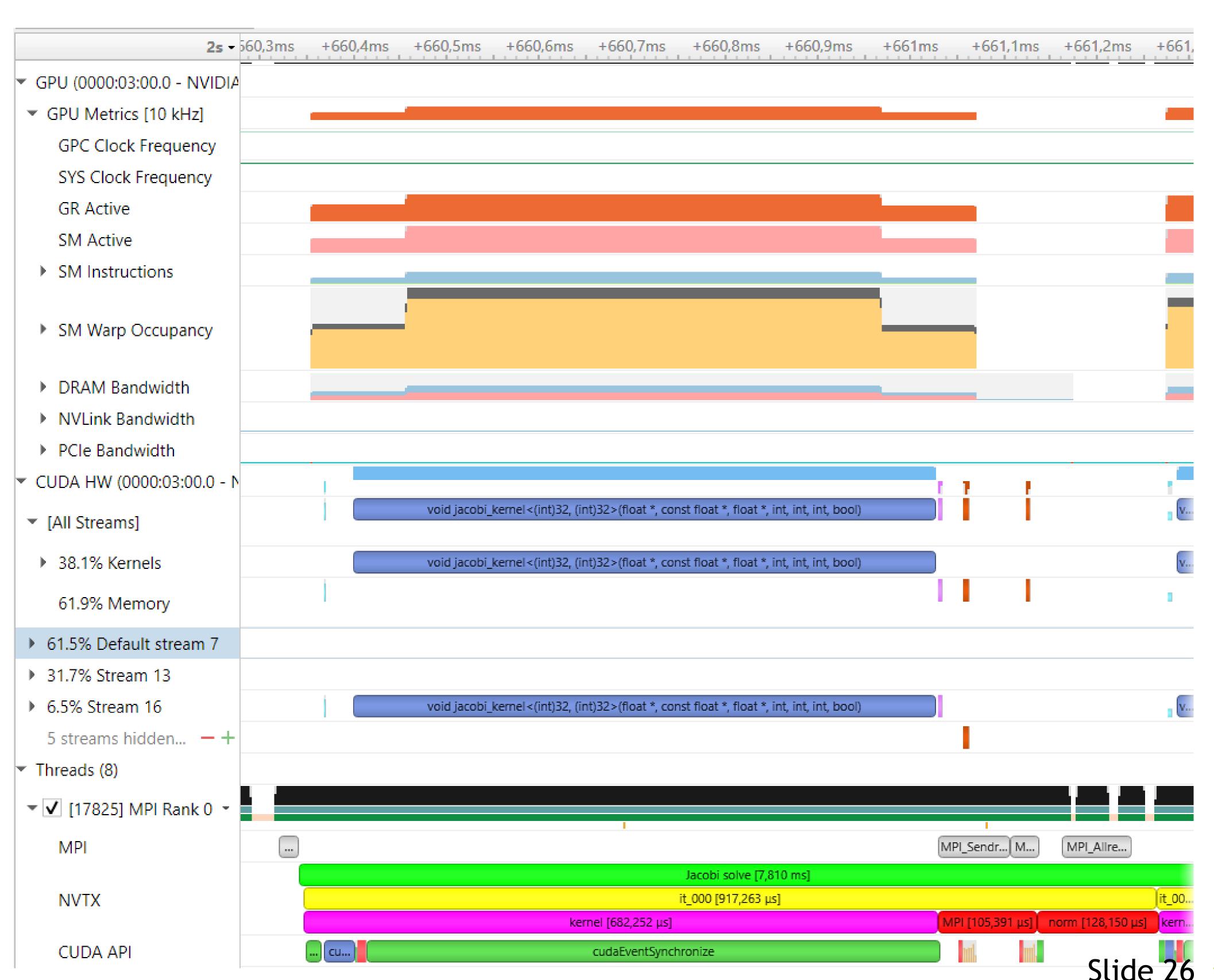
- Valuable low-overhead insight into HW usage:
 - SM instructions
 - DRAM Bandwidth, PCIe Bandwith (GPUDirect)
- Also: Memory usage, Page Faults (higher overhead)
 - CUDA Programming guide: <u>Unified Memory</u>
 <u>Programming</u>
- Can save kernel-level profiling effort!
- nsys profile
 - --gpu-metrics-device=0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



FOCUSING THE ANALYSIS

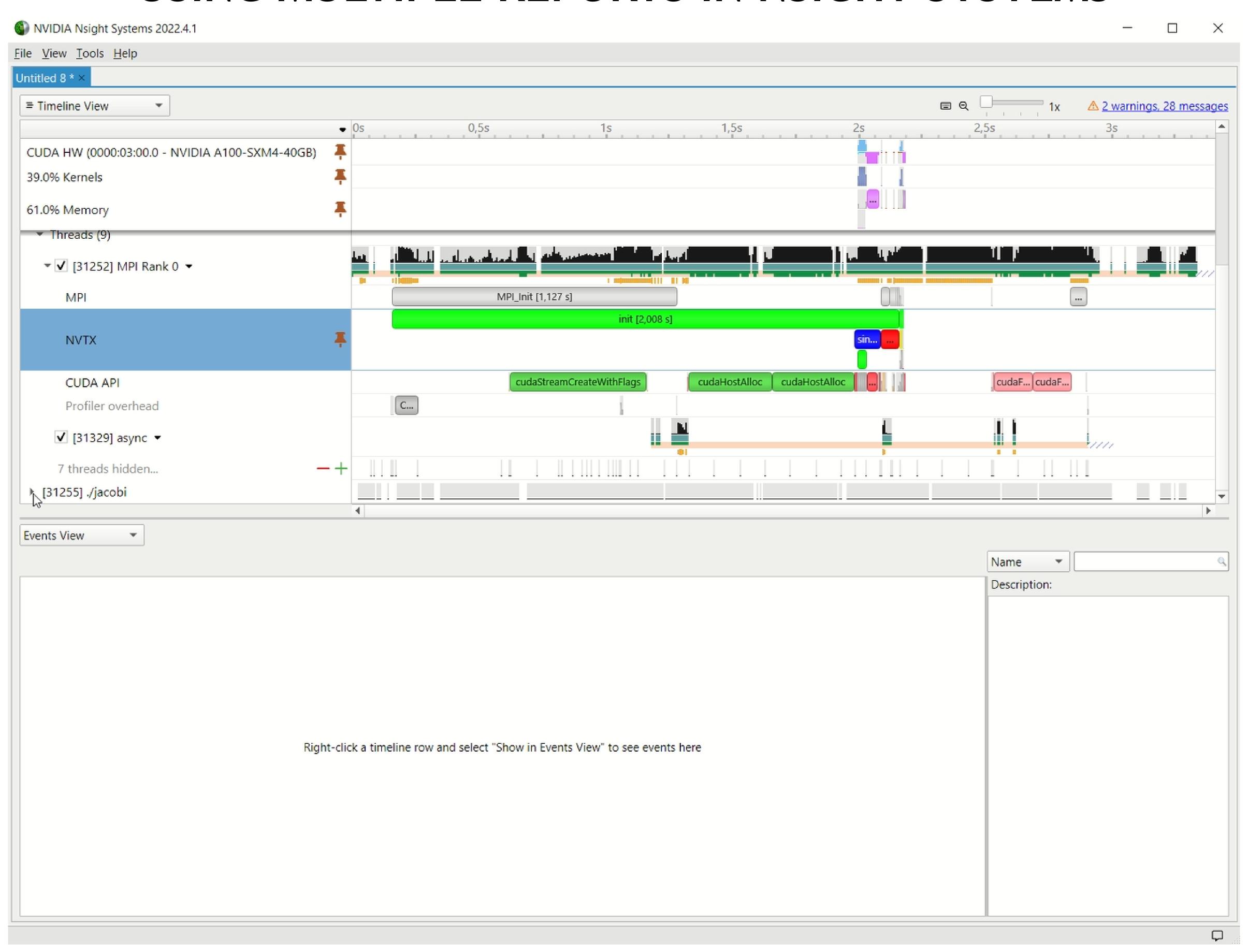
Introducing GPU metrics sampling

- 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



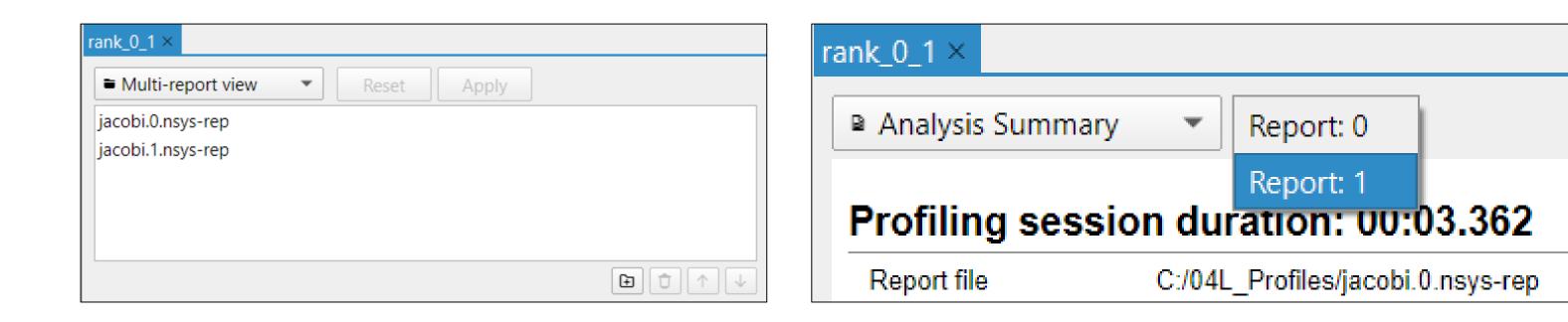


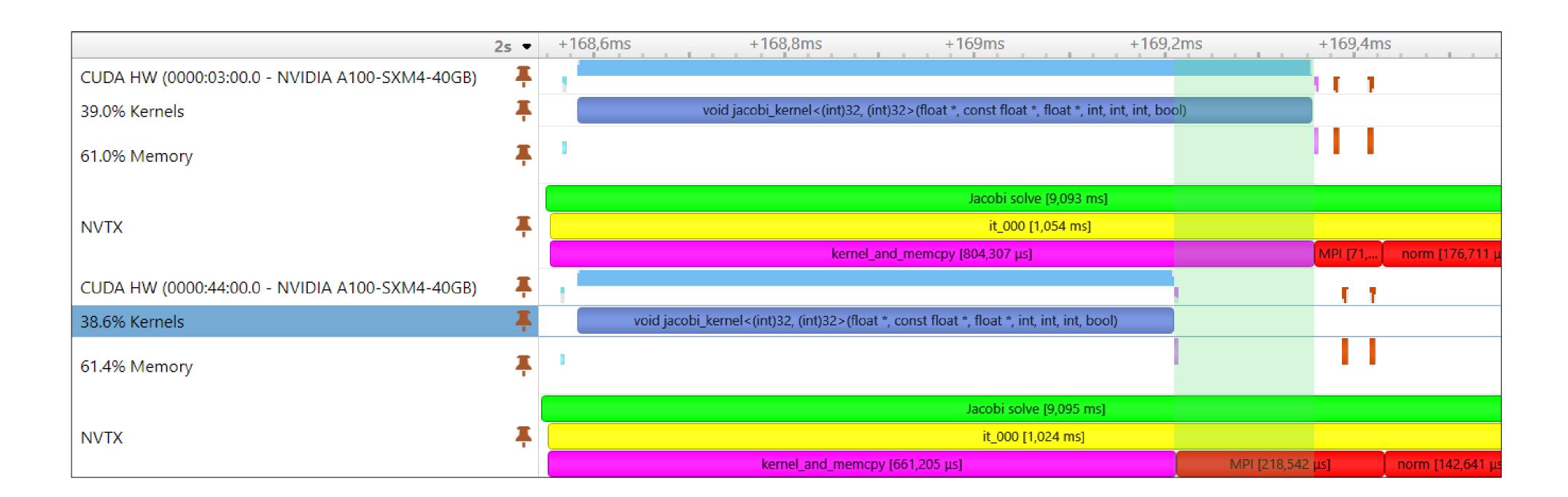
USING MULTIPLE REPORTS IN NSIGHT SYSTEMS



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

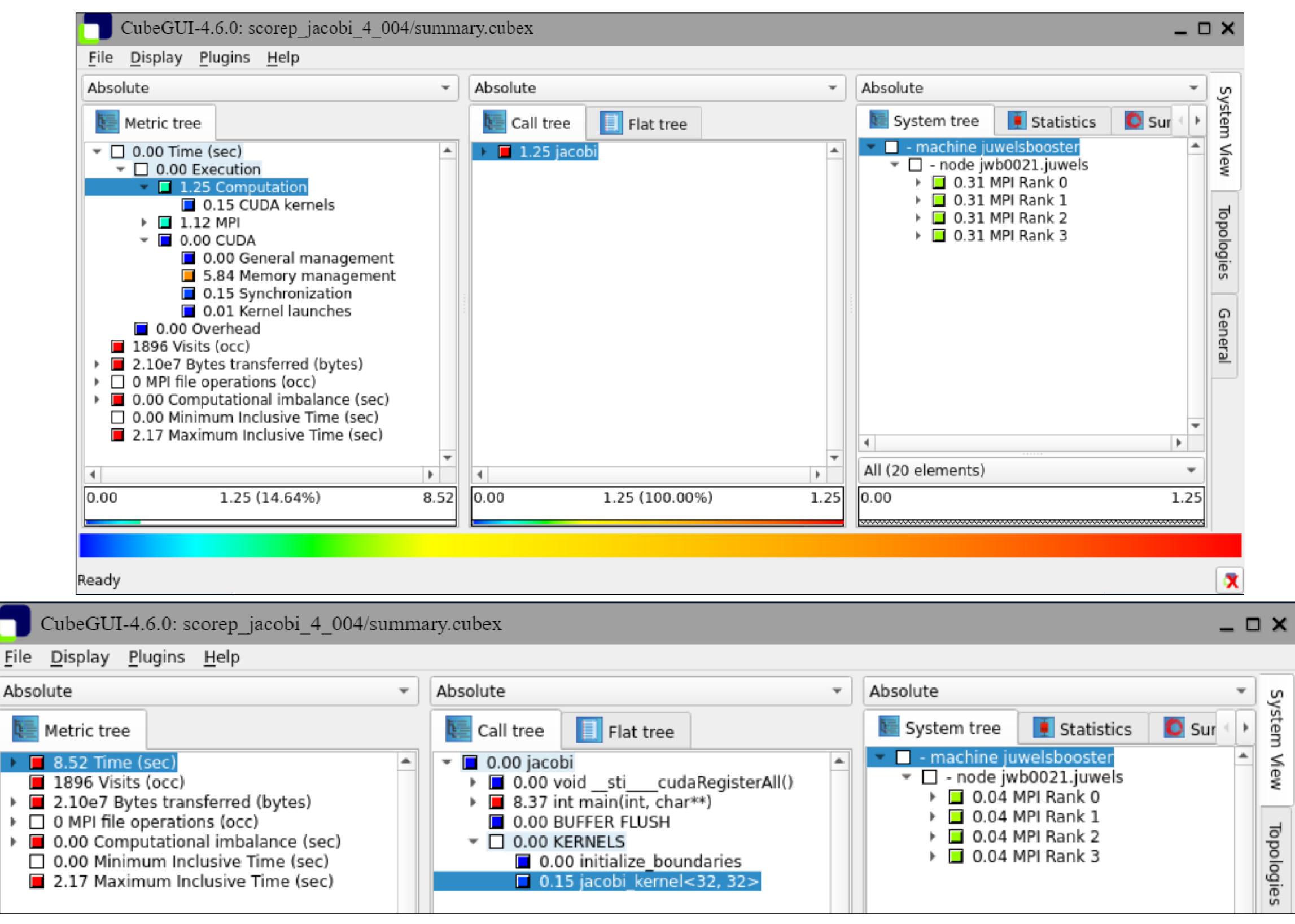
Specialized for large-scale distributed analysis

- 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)



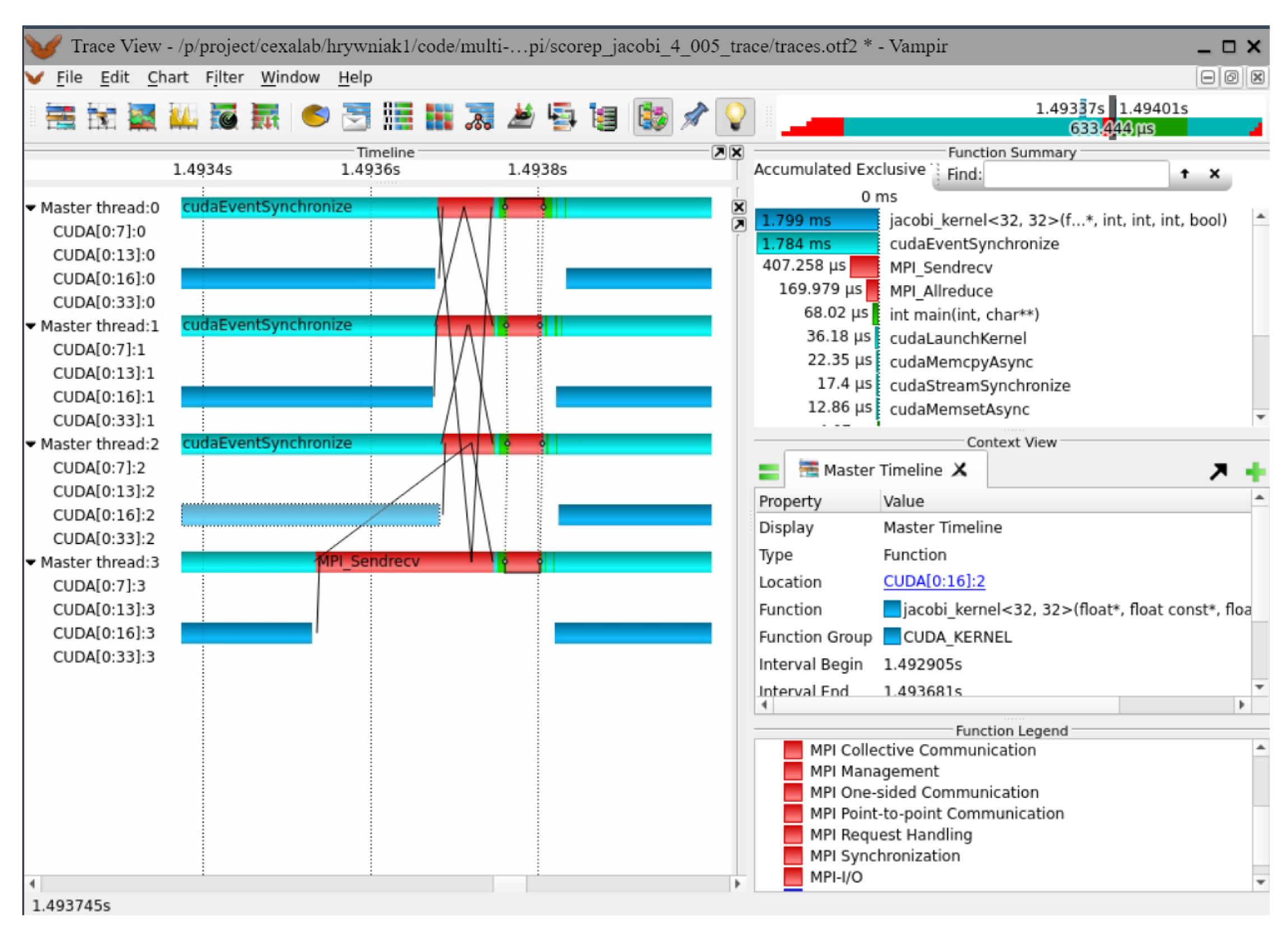
SCALASCA / CUBE

- 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

