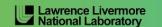
Analysis report examination with Cube

Brian Wylie
Jülich Supercomputing Centre



























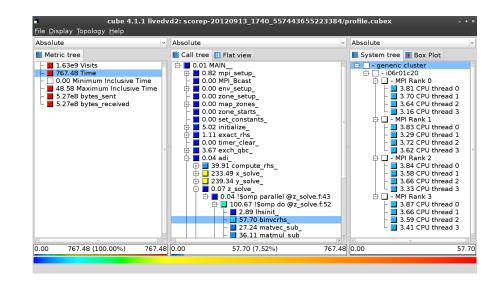






Cube

- Parallel program analysis report exploration tools
 - Libraries for XML+binary report reading & writing
 - Algebra utilities for report processing
 - GUI for interactive analysis exploration
 - Requires Qt4 ≥4.6 or Qt 5
- Originally developed as part of the Scalasca toolset
- Now available as a separate component
 - Can be installed independently of Score-P, e.g., on desktop or laptop
 - Latest release: Cube v4.5 (May 2020)

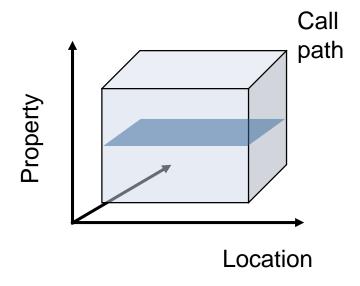


Note: source distribution tarballs for Linux, as well as binary packages provided for Windows & MacOS, from www.scalasca.org website in software/Cube-4x



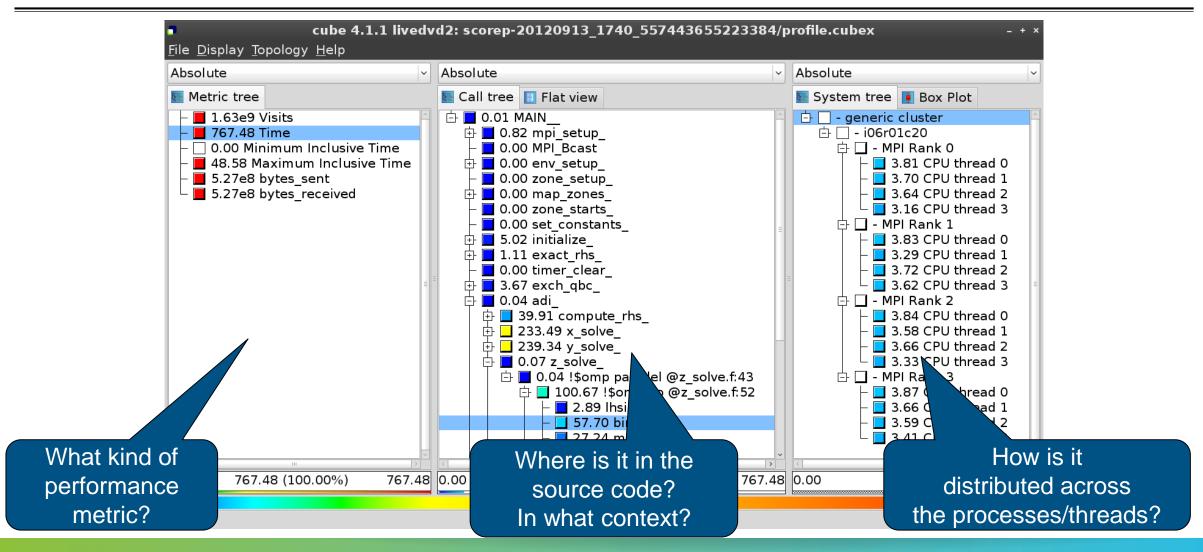
Analysis presentation and exploration

- Representation of values (severity matrix)
 on three hierarchical axes
 - Performance property (metric)
 - Call path (program location)
 - System location (process/thread)
- Three coupled tree browsers
- Cube displays severities
 - As value: for precise comparison
 - As colour: for easy identification of hotspots
 - *Inclusive* value when closed & *exclusive* value when expanded
 - Customizable via display modes





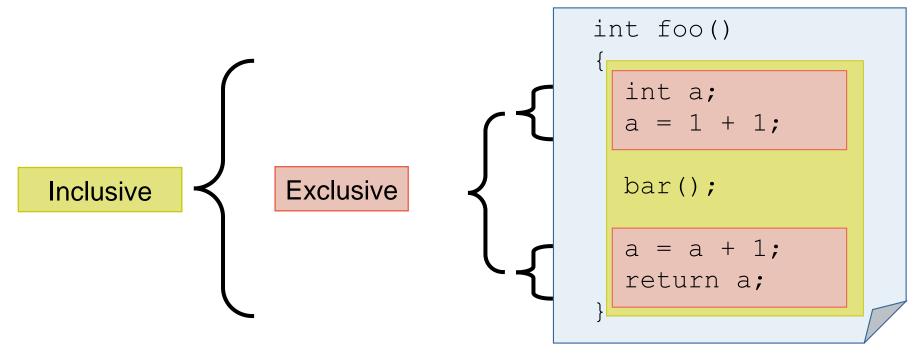
Analysis presentation



Inclusive vs. exclusive values

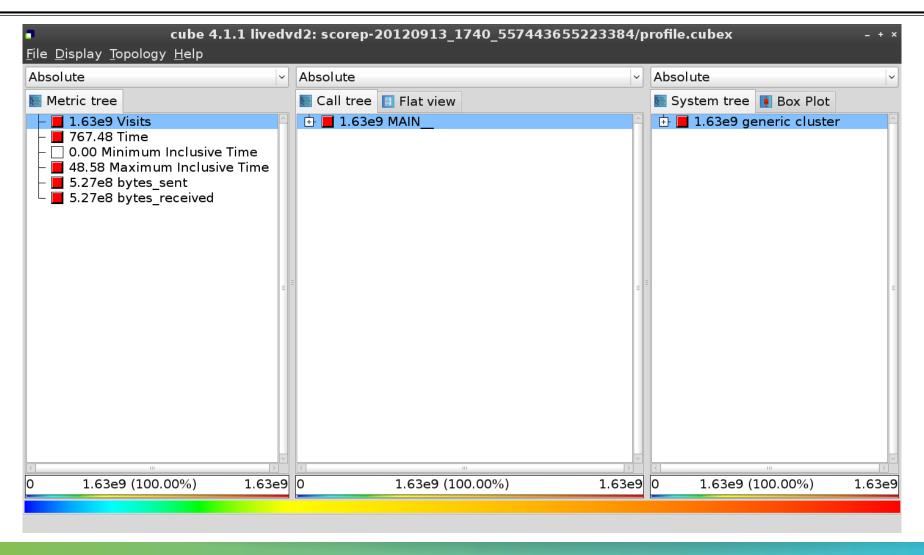


- Inclusive
 - Information of all sub-elements aggregated into single value
- Exclusive
 - Information cannot be subdivided further



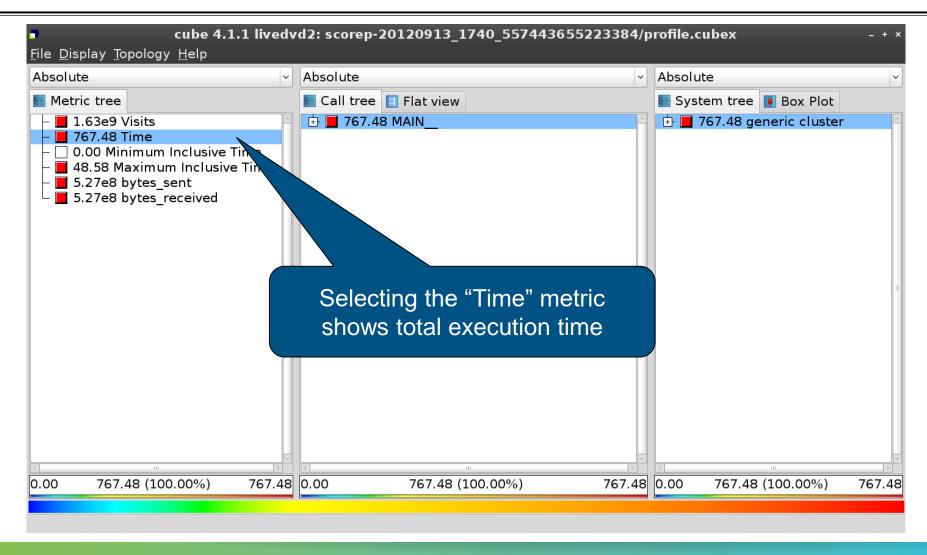
Score-P analysis report exploration (opening view)





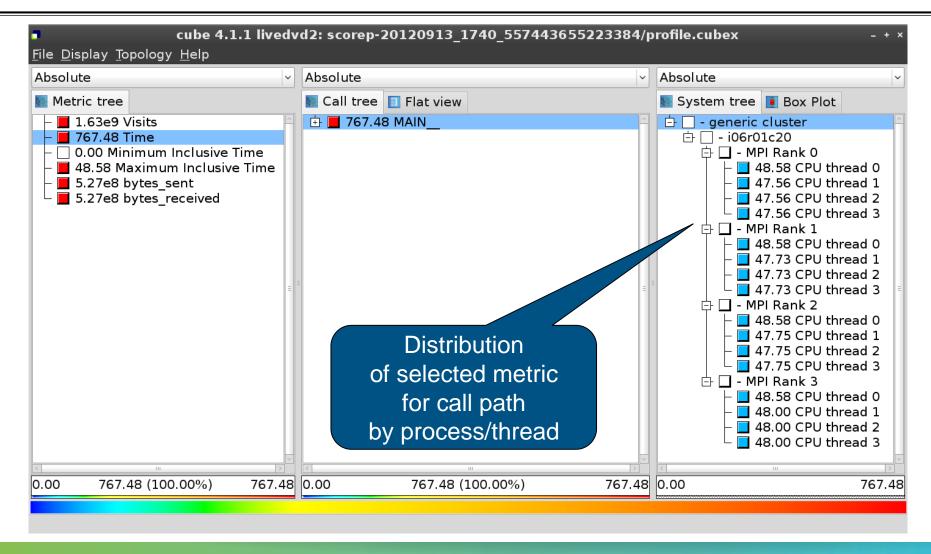
Metric selection





Expanding the system tree

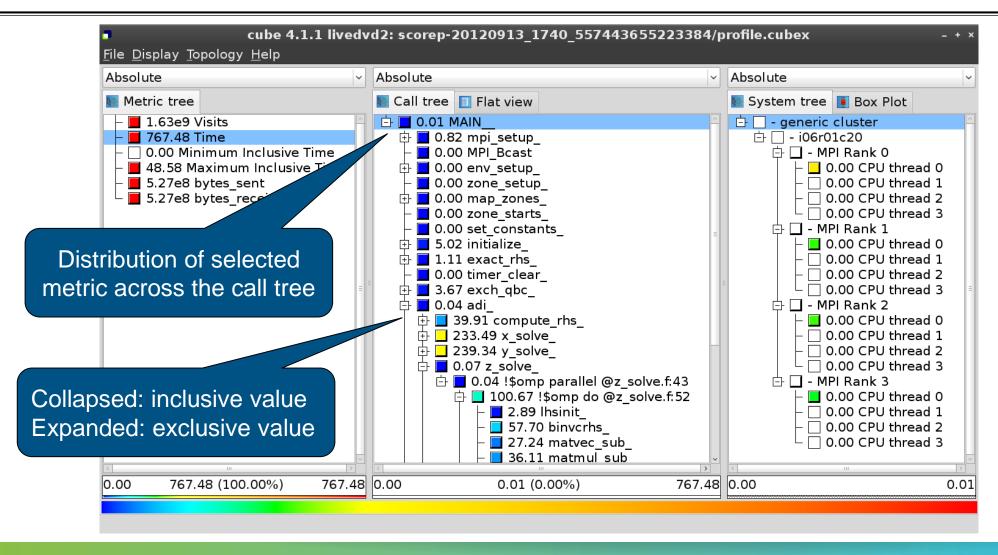






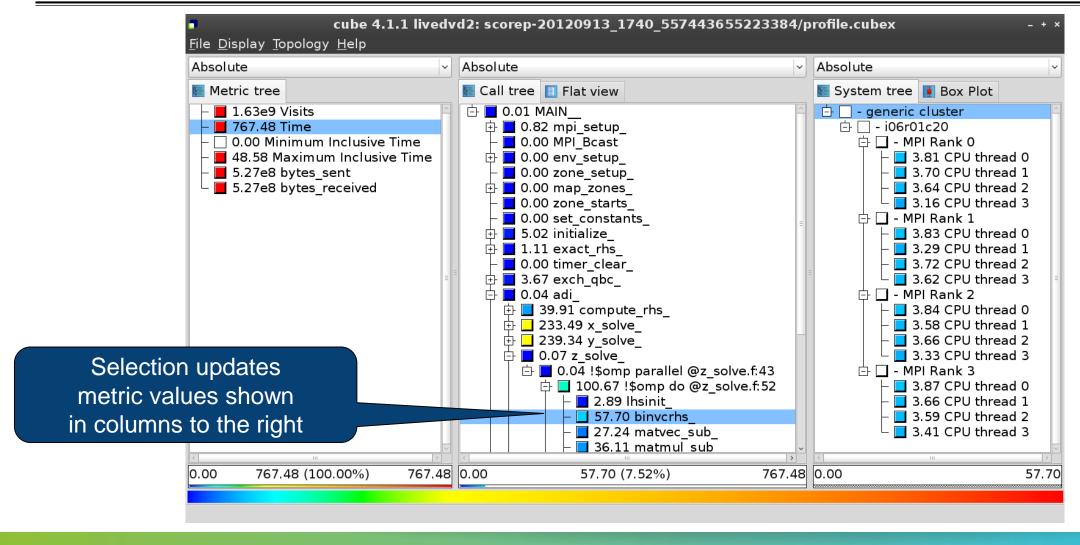
Expanding the call tree





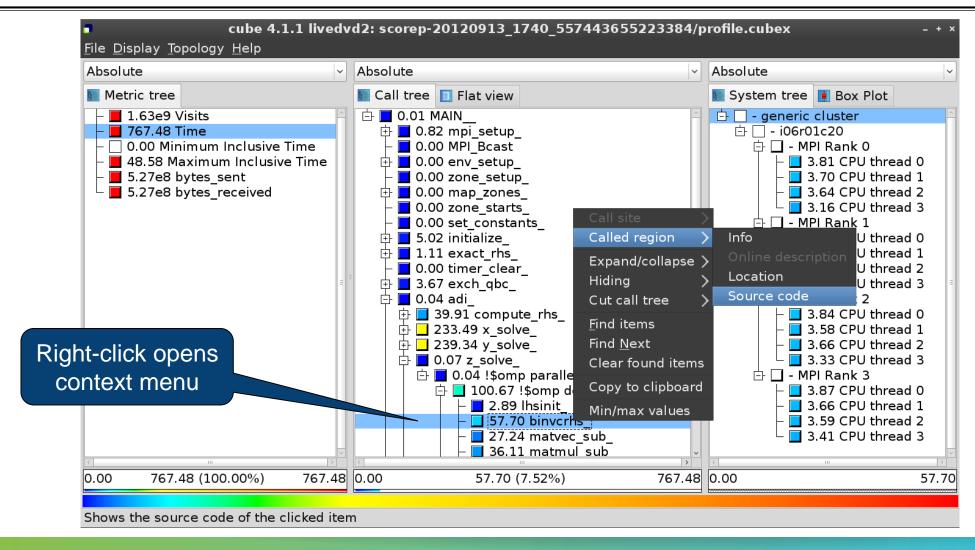
Selecting a call path





Source-code view via context menu

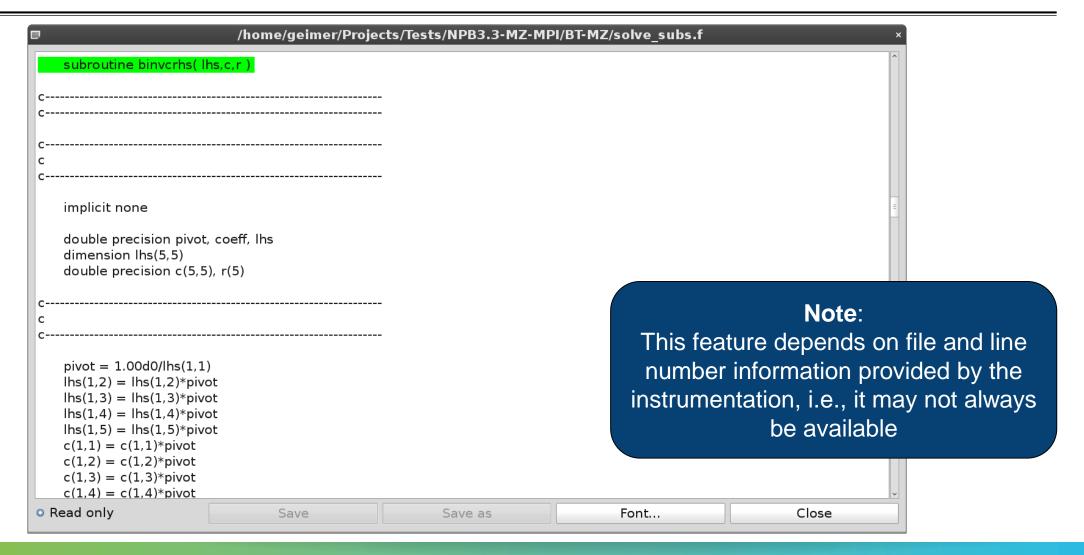






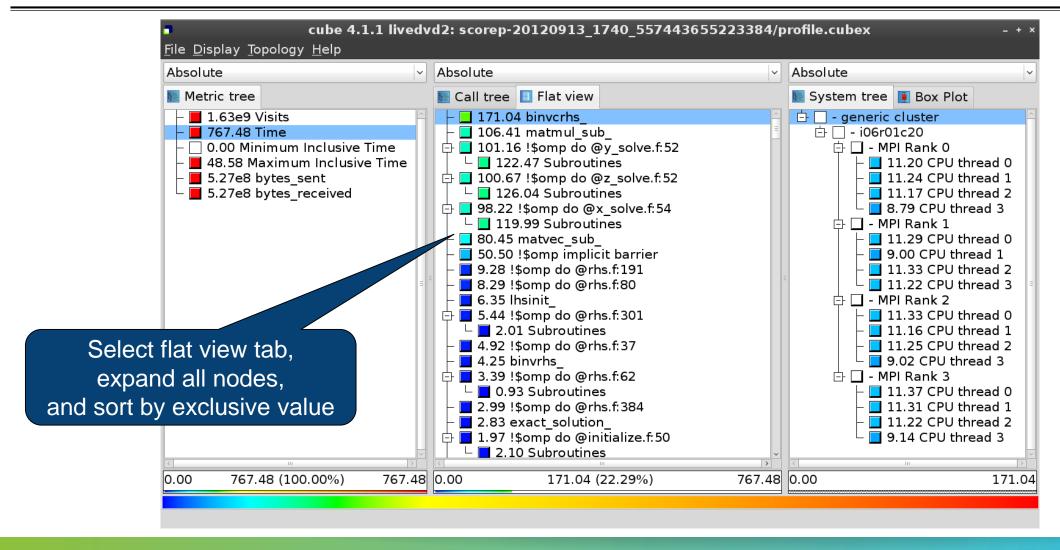
Source-code view





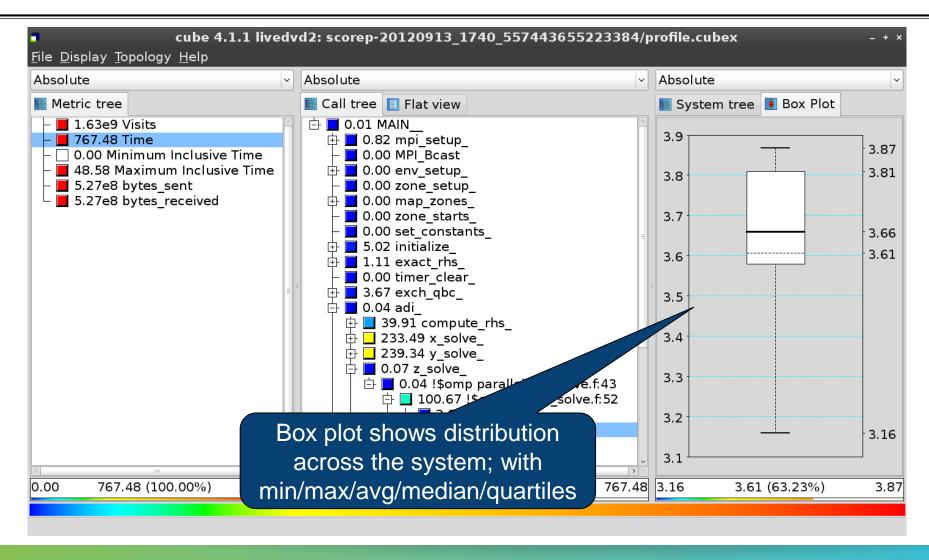
Flat profile view





Box plot view

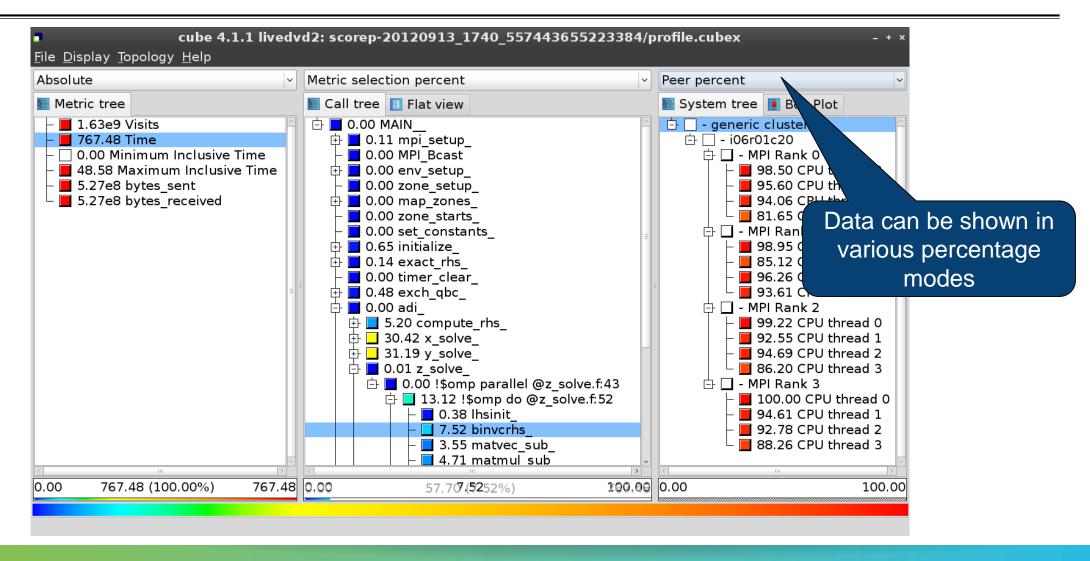






Alternative display modes





Important display modes

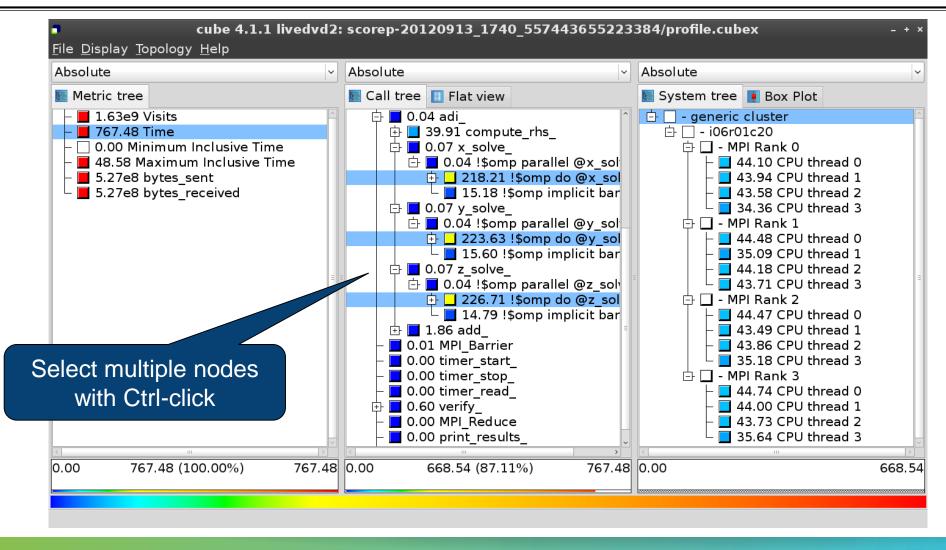


- Absolute
 - Absolute value shown in seconds/bytes/counts
- Selection percent
 - Value shown as percentage w.r.t. the selected node "on the left" (metric/call path)
- Peer percent (system tree only)
 - Value shown as percentage relative to the maximum peer value



Multiple selection

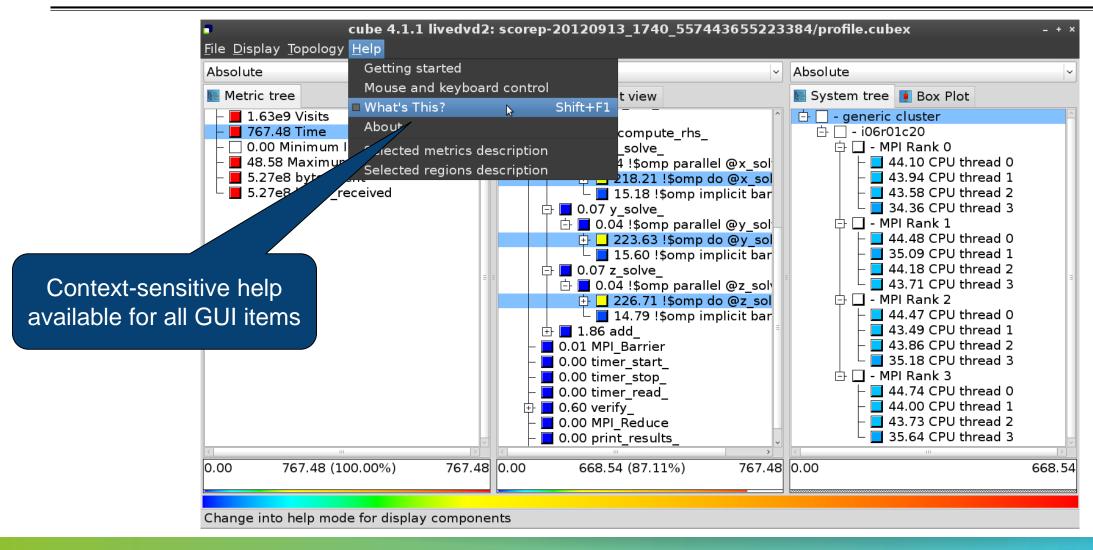






Context-sensitive help





Derived metrics



Derived metrics are defined using CubePL expressions, e.g.:

metric::time(i)/metric::visits(e)

- Values of derived metrics are not stored, but calculated on-the-fly
- Types of derived metrics:
 - Prederived: evaluation of the CubePL expression is performed before aggregation
 - Postderived: evaluation of the CubePL expression is performed after aggregation
- Examples:
 - "Average execution time": Postderived metric with expression

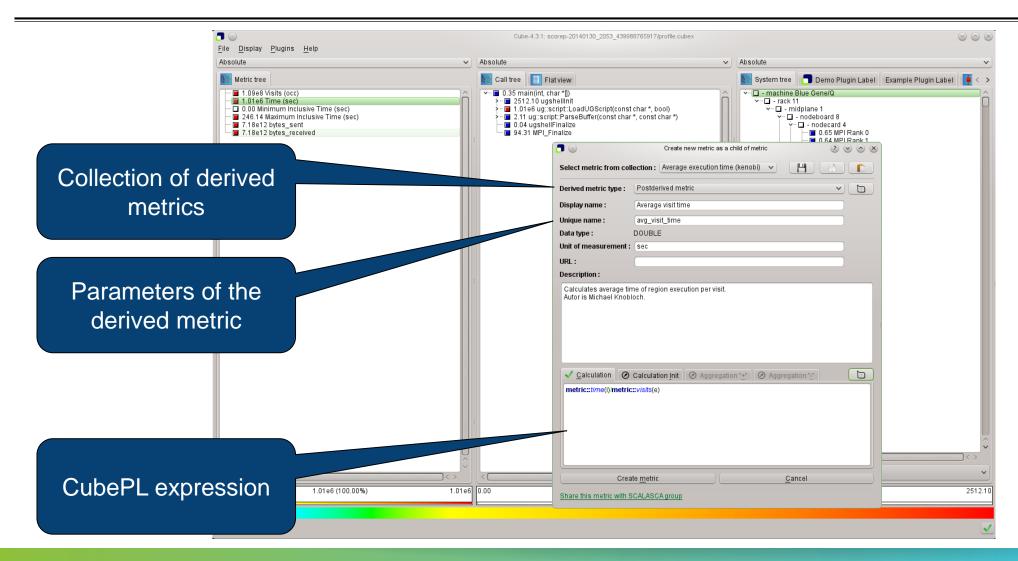
metric::time(i)/metric::visits(e)

"Number of FLOP per second": Postderived metric with expression

metric::FLOP()/metric::time()

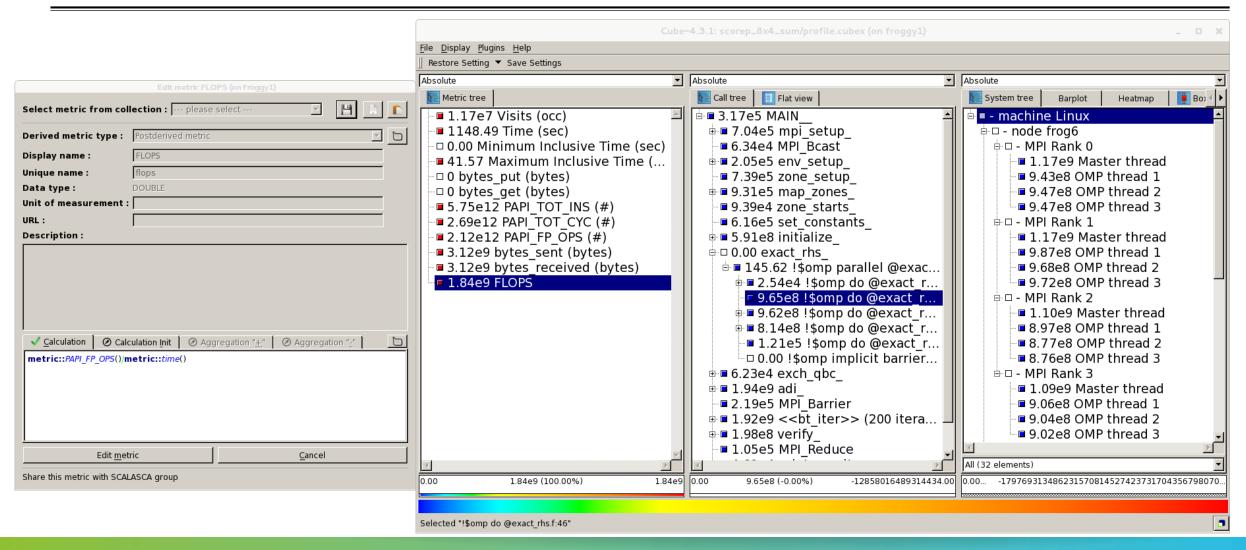
Derived metrics in Cube GUI





Example: FLOPS based on PAPI_FP_OPS and time





CUBE algebra utilities



Extracting solver sub-tree from analysis report

```
% cube_cut -r '<<ITERATION>>' scorep_bt-mz_C_32x4_sum/profile.cubex
Writing cut.cubex... done.
```

Calculating difference of two reports

```
% cube_diff scorep_bt-mz_C_32x4_sum/profile.cubex cut.cubex Writing diff.cubex... done.
```

- Additional utilities for merging, calculating mean, etc.
- Default output of cube_utility is a new report utility.cubex
- Further utilities for report scoring & statistics
- Run utility with `-h' (or no arguments) for brief usage info

Iteration profiling



- Show time dependent behavior by "unrolling" iterations
- Preparations:
 - Mark loop body by using Score-P instrumentation API in your source code

```
SCOREP_USER_REGION_DEFINE( scorep_bt_loop )
SCOREP_USER_REGION_BEGIN( scorep_bt_loop, "<<bt_iter>>", SCOREP_USER_REGION_END( scorep_bt_loop )
```

- Result in the Cube profile:
 - Iterations shown as separate call trees
 - > Useful for checking results for specific iterations

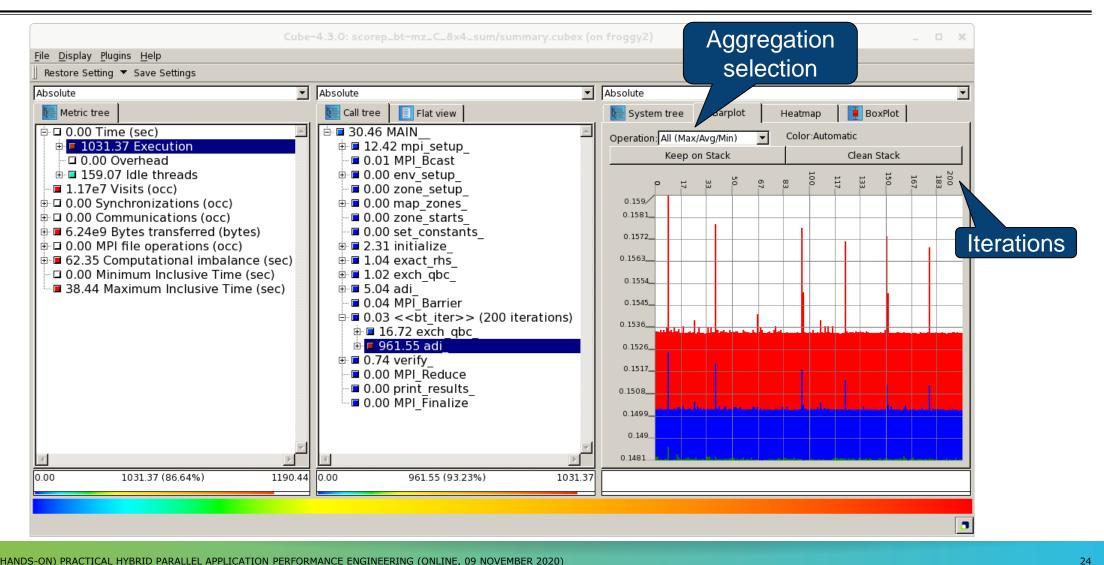
or

- Select your user-instrumented region and mark it as loop
- Choose "Hide iterations"
- ➤ View the Barplot statistics or the (thread x iterations) Heatmap



Iteration profiling: Barplot

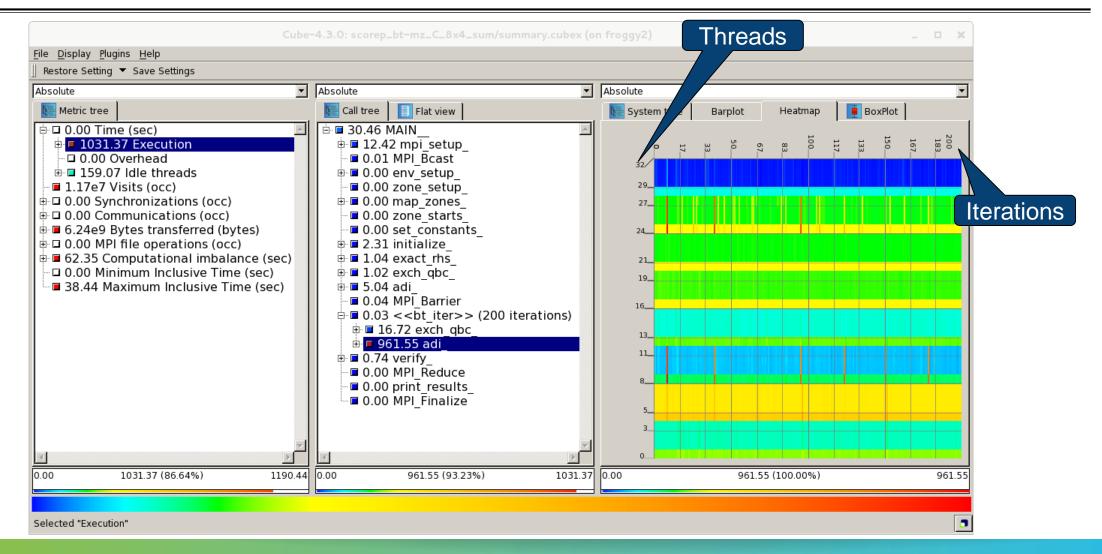






Iteration profiling: Heatmap

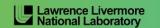




Case study HemeLB





























HemeLB (SuperMUC-NG: no GPUs)

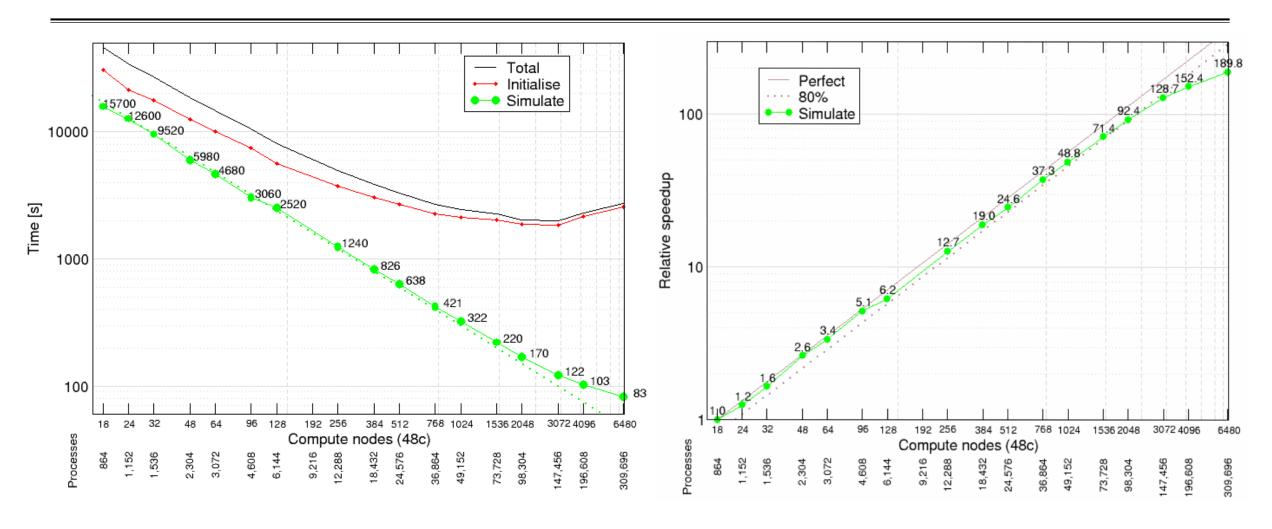


- 3D macroscopic blood flow in human arterial system developed by UC London (UK)
 - lattice-Boltzmann method tracking fluid particles on a lattice grid with complex boundary conditions
 - exascale flagship application of EU H2020 HPC Centre of Excellence for Computational Biomedicine
- Hemelb open-source code and test case: www.hemelb.org
 - C++ parallelized with MPI [+ CUDA unused]
 - Intel Studio 2019u4 compiler and MPI library (v19.0.4.243)
 - configured with 2 'reader' processes (intermediate MPI file writing disabled)
 - MPI-3 shared-memory model employed within compute nodes to reduce memory requirements when distributing lattice blocks from reader processes



- Focus of analysis 5,000 time-step (500μs) simulation of cerebrovascular "circle of Willis" geometry
 6.4μm lattice resolution (21.15 GiB): 10,154,448,502 lattice sites
- Executed on *SuperMUC-NG* Lenovo ThinkSystem SD650 (LRZ):
 - 2x 24-core Intel Xeon Platinum 8174 ('Skylake') @ 3.1GHz
 - 48 MPI processes/node, 6452 (of 6480) compute nodes: 309,696 MPI processes
 - 190x speed-up from 864 cores: 80% scaling efficiency to over 100,000 cores
- ⇒ Identification & quantification of impact of load balance and its variation

HemeLB@SNG strong scaling of FOA RunSimulation



[Execution of 9,216 processes on 192 compute nodes not possible due to insufficient compute nodes with adequate memory in 'fat' partition (768 GiB vs. regular 96 GiB node memory)



HemeLB@SNG strong scaling efficiency of FOA RunSimulation

| Compute nodes | 24 | 32 | 48 | 64 | 96 | 128 | 192 | 256 | 384 | 512 | 768 | 1024 | 1536 | 2048 | 3072 | 4096 | 6452 |
|---------------------------|-------|-------|-------|-------|-------|-------|------|-------|-------|-------|-------|-------|-------|-------|--------|--------|--------|
| Processes | 1152 | 1536 | 2304 | 3072 | 4608 | 6144 | 9216 | 12288 | 18432 | 24576 | 36864 | 49152 | 73728 | 98304 | 147456 | 196608 | 309696 |
| | | | | | | | | | | | | | | | | | |
| Global scaling efficiency | 0.79 | 0.79 | 0.84 | 0.80 | 0.82 | 0.75 | | 0.73 | 0.72 | 0.73 | 0.74 | 0.68 | 0.68 | 0.65 | 0.62 | 0.57 | 0.45 |
| - Parallel efficiency | 0.79 | 0.80 | 0.87 | 0.83 | 0.86 | 0.80 | | 0.75 | 0.74 | 0.74 | 0.77 | 0.71 | 0.72 | 0.70 | 0.72 | 0.70 | 0.73 |
| Load balance efficiency | 0.79 | 0.80 | 0.88 | 0.84 | 0.86 | 0.80 | | 0.75 | 0.74 | 0.75 | 0.78 | 0.72 | 0.74 | 0.72 | 0.74 | 0.73 | 0.80 |
| Communication efficiency | 1.00 | 1.00 | 1.00 | 1.00 | 1.00 | 1.00 | | 1.00 | 1.00 | 0.99 | 0.99 | 0.99 | 0.98 | 0.98 | 0.97 | 0.96 | 0.92 |
| - Computation scaling | 1.00 | 0.99 | 0.96 | 0.96 | 0.95 | 0.93 | | 0.98 | 0.98 | 0.98 | 0.96 | 0.96 | 0.94 | 0.93 | 0.87 | 0.81 | 0.61 |
| Instructions scaling | 1.00 | 1.00 | 1.00 | 1.00 | 1.00 | 1.00 | | 1.00 | 1.00 | 1.00 | 0.99 | 0.97 | 0.94 | 0.89 | 0.79 | 0.67 | 0.45 |
| IPC scaling | 1.00 | 0.99 | 0.96 | 0.96 | 0.95 | 0.93 | | 0.98 | 0.98 | 0.99 | 0.98 | 0.99 | 1.00 | 1.04 | 1.11 | 1.21 | 1.36 |
| IPC | 1.411 | 1.395 | 1.353 | 1.355 | 1.342 | 1.316 | | 1.377 | 1.387 | 1.396 | 1.383 | 1.390 | 1.417 | 1.473 | 1.566 | 1.704 | 1.919 |
| | | | | | | | | | | | Key: | <0.65 | <0.75 | <0.85 | <0.95 | <1.00 | >1.00 |

Global scaling efficiency fairly good around 80%, before degrading at larger scales

- Parallel efficiency deteriorating following Load balance efficiency
 - Communication efficiency excellent throughout
- Computation scaling (relative to 1152 processes) very good except at largest scale
- Degradation of Instructions scaling partially compensated by improving IPC scaling
 [POP CoE scaling efficiency model: www.pop-coe.eu]



HemeLB@SNG instrumentation & measurement



- Score-P/5.0 configured with MPIFLAGS=-DSCOREP_MPI_NO_MINI
 - don't generate wrappers for MPI_Comm_rank & MPI_Comm_size
- HemeLB instrumentation
 - application dependencies built without modification (or instrumentation)
 - src/main.cc annotated with SCOREP_RECORDING_OFF/SCOREP_RECORDING_ON macros
 - to pause measurement recording during initialization
 - configured using CXX=scorep-icpc with
 - SCOREP WRAPPER INSTRUMENTER FLAGS="--user --mpp=mpi --thread=none --nomemory"
 - SCOREP_WRAPPER_COMPILER_FLAGS="-tcollect-filter=<path_to_file>"
 - disable Intel compiler instrumentation apart from main routine and key classes such as SimulationMaster
- HemeLB execution configuration
 - SLURM --ear=off to enable 'performance' profile of processor and access to hardware counters
 - SCOREP_DEVELOPMENT_MEMORY_STATS=aggregated # report Score-P memory usage
 - SCOREP_METRIC_PAPI=PAPI_TOT_INS,PAPI_TOT_CYC,PAPI_REF_CYC,PAPI_RES_STL
 - SCOREP_TIMER=gettimeofday # use globally synchronized clock
 - SCOREP_TOTAL_MEMORY=200MB # as determined from Score-P memory usage stats



HemeLB (JUWELS-Volta)

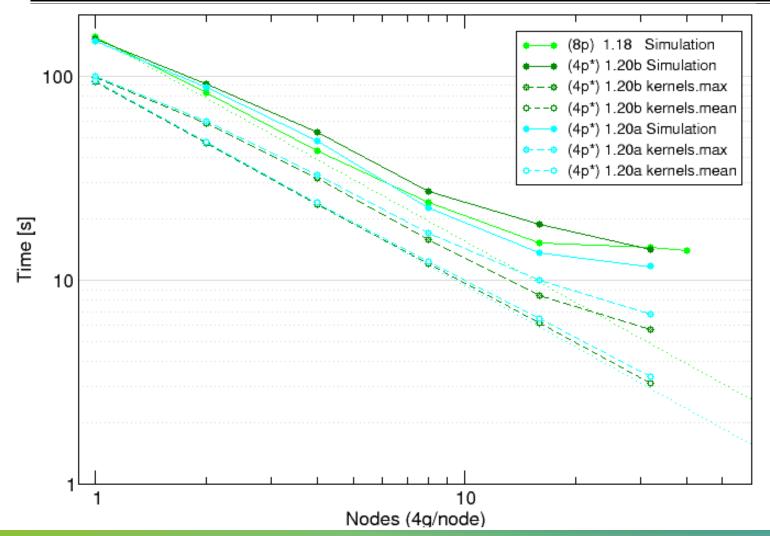


- 3D macroscopic blood flow in human arterial system developed by UC London (UK)
 - lattice-Boltzmann method tracking fluid particles on a lattice grid with complex boundary conditions
 - exascale flagship application of EU H2020 HPC Centre of Excellence for Computational Biomedicine
- Hemelb open-source code and test case: www.hemelb.org
 - C++ parallelized with MPI + CUDA (in development)
 - GCC/8.3.0 compiler, CUDA/10.1.105 and ParaStationMPI/5.4 library
 - configured with 2 'reader' processes and intermediate MPI file writing
 - rank 0 'monitor' process doesn't participate in simulation



- Focus of analysis 2,000 time-step (each 100µs) simulation of CBM2019_Arteries_patched geometry
 - 1.78 GiB: 66,401,494 lattice sites, 1+38 iolets
- Executed on JUWELS-Volta (@JSC):
 - 2x 20-core Intel Xeon Platinum 8168 ('Skylake') CPUs + 4 Nvidia V100 'Volta' GPUs
 - 4* MPI processes/node (one per GPU), 32 (of 56) compute nodes: 129 MPI processes
- ⇒ Identification & quantification of impact of load balance and its variation

HemeLB@JUWELS-Volta strong scaling of FOA RunSimulation



- Reference execution with 8ppn
 - multiple processes offloading GPU kernels generally unproductive
- Comparison of versions (4ppn)
 - v1.20a generally better
- Synchronous MPI file writing is the primary bottleneck
- CUDA kernels on GPUs
 - less than half of Simulation time (therefore GPUs mostly idle)
 - total kernel time scales very well (0.93 scaling efficiency)
 - load balance deteriorates
 (0.95 for single node,
 0.50 for 32 nodes)

HemeLB@JUWELS/Volta strong scaling efficiency of RunSimulation

| Simulation time [s] | 1n 5p 147.87 | 2n 9p 88.38 | 4n 17p 48.13 | 8n 33p 22.66 | 16n 65p 13.68 | 32n 129p 11.67 |
|--|------------------------|-----------------------|------------------------|------------------------|----------------------|--------------------------|
| Global scaling efficiency | 0.64 | 0.53 | 0.49 | 0.52 | 0.43 | 0.25 |
| – Parallel efficiency | 0.64 | 0.53 | 0.50 | 0.54 | 0.47 | 0.29 |
| – – Load balance efficiency (GPU) | 0.95 | 0.78 | 0.73 | 0.73 | 0.65 | 0.50 |
| – Communication efficiency (GPU) | 0.67 | 0.68 | 0.68 | 0.75 | 0.73 | 0.58 |
| Computation scaling (GPU) | 1.00 | 1.00 | 0.99 | 0.96 | 0.92 | 0.87 |

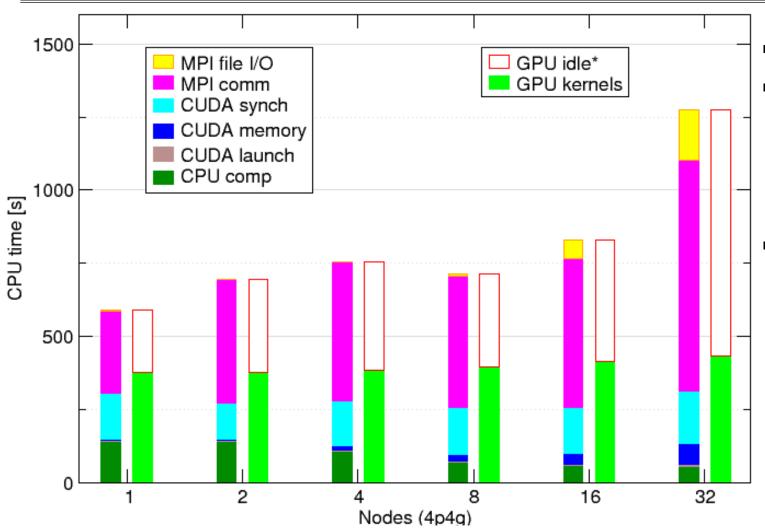
Only considering GPUs (ignoring all CPU cores, 90% of which are completely unused)

- Single (quad-GPU) node already suffers significant communication inefficiency
 - includes MPI file writing, but doesn't degrade much as additional nodes are included
- Load balance of GPUs deteriorates progressively
- GPU computation scaling remains reasonably good
 [POP CoE scaling efficiency model: www.pop-coe.eu]





HemeLB@JUWELS-Volta strong scaling of FOA RunSimulation



- CPU+GPU time breakdown
- CUDA kernels on GPUs
 - less than half of Simulation time (therefore GPUs mostly idle)
 - total kernel time scales very well (0.87 scaling efficiency)
- MPI processes on CPUs
 - computation time decreases
 - CUDA synchronization time fairly constant, but time for memory management increases somewhat
 - MPI communication time dominates, with much more time for file writing with 16+ nodes



HemeLB@JUWELS-Volta instrumentation & measurement



- Score-P/6.0 instrumentation
 - CMake: CUDACXX=scorep-nvcc (no instrumentation of CXX)
 - SCOREP_WRAPPER_COMPILER_FLAGS="--relocatable-device=true -c"
 - SCOREP_WRAPPER_INSTRUMENTER_FLAGS=
 - "--cuda --mpp=mpi --thread=none --instrument-filter=hemelb.filt"
 - hemelb.filt specifying patterns to exclude most source modules and include only routines of particular interest
- Scalasca/2.5 runtime measurement configuration
 - SCAN_TRACE_ANALYZER=none # CUDA streams not supported
 - SCOREP_CUDA_ENABLE=runtime,memcpy,kernel,sync,flushatexit
 - SCOREP_CUDA_BUFFER=10MB
 - SCOREP_MPI_ENABLE_GROUPS=coll,env,io,p2p,rma,topo,xnonblock # no 'cg' group
 - SCOREP_TIMER=gettimeofday # use globally synchronized clock
 - SCOREP_TOTAL_MEMORY=64MB # as determined from Score-P memory usage stats

Cube: Further information

- Parallel program analysis report exploration tools
 - Libraries for Cube report reading & writing
 - Algebra utilities for report processing
 - GUI for interactive analysis exploration
- Available under 3-clause BSD open-source license
- Documentation & sources:
 - http://www.scalasca.org
- User guide also part of installation:
 - prefix>/share/doc/CubeGuide.pdf
- Contact:
 - mailto: scalasca@fz-juelich.de

