



From Live Multi-Camera 3D Modeling to In Situ Processing for HPC

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What this talk is about

Data flow programming for data intensive high performance applications

Tool: FlowVR

Evolved from applications to interactive applications (virtual reality, 3D modeling) to computational steering and in situ processing for large scale numerical simulations







Get/put messages from/to ports: no explicit origin/destination

A simple API to limit code intrusion Direct access to memory: no hidden copies Adapted to develop domain specific layers

http://flowvr.sf.net





Python scripting:

- 1. Assemble components (data flow graph)
- 2. Instantiate parameters: component mapping,...











N-to-1 Communication Pattern (tree with arity k - unkown) (N unknown – get value from incoming component)





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Run one daemon per node

- Components (applications) attach to the daemon when starting
- Daemons in charge of routing messages and triggering component executions:
 - Pointer exchange inside a node (shared memory)
 - MPI or TCP between nodes





Interactive Applications









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Interactive Applications





Equipex Kinovis: 70 cameras



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Itermediate Summary

- Data Flow Programming well adapted to this type of interactive data intensive applications
- Avoid moving data, process them as closely as possible to their source
- No intermediate storage to disk, process data as soon as available (some buffering possible)

Actually an approach that is today relevant for more traditional large scale numerical simulations !



The Data Challenge

More compute capabilities -> larger simulations -> more data

Usability Challenge:

- How to extract meaningful information from this huge amount of data in a reasonable time
- Analysis tools have not been considered as first class citizen so far. They did not receive the same attention as simulation codes. Today analysis codes are either:
 - In the simulation codes
 - Scripts (with limited parallelism)
 - Rely on on scientific visualization tools like Paraview/VTK or Visit (reasonable parallelism support)

Performance Challenge:

- Moving data becomes the bottleneck for simulation as well as data analytics
- Compute capabilities increase faster than data transfer ones
- Data movements and storage consume 50%-70% of total energy (ScidacReview 1001)



A Data Challenge Already Present

Scientists already spend a significant part of their efforts in the data analysis

A simple but classical strategy to limit the impact of the data challenge: **Reduce output frequency**

Need for more advanced strategies to better manage the available I/O and storage budget

In Situ Processing!!



Traditional Workflow



WorkFlow with In situ Analytics





In Situ Processing: What for ?

Data compression (Isabela [Lehmann & al. LDAV'14])
Indexing (FastBit, Dirac [Lakshminarasimhan & al. HPDC'13])
Analytics (1D, 2D, 3D descriptors)





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In Simulation Processing





In Transit Processing







Asynchronous In Situ + In Transit Processing





Molecular Dynamics

Collaboration with Marc Baaden (IBPC):

- Realistic molecular systems
- Realistic analytics scenarios



Gromacs:

- Standard open source molecular dynamics simulation code
- Hybrid MPI+openMP+GPU parallelism, with integrated dynamic load balancing strategies
- Heterogeneous: long range interactions handled through dedicated processes (PME nodes)
- Very high iteration frequency (500 Hz)
- Often a benchmark in exascale related publications



Exemple: Benefit of the In Situ Helper Core Strategy



Aggregate the results in situ on helper-core (1 per node) to the master node



Exemple: Benefit of the In Situ Helper Core Strategy



Aggregate local results in situ on helper-core (1 per node) and write to disk





In Sim vs. In Situ I/O [Dreher,CCGRID'14]



Gromacs without I/O: 15 cores/node 3% slower than 16 cores/node (- 6% if scalability would have been perfect)



Parallel In Situ Quicksurf





Classical way to visualize a molecule surface (isosurface based on atom density)



Parallel In Situ Isosurface Extraction [Dreher, CCGRID'14]



Gromacs Merge Gromacs Merge Gromacs Gromacs Merge Merge Gromacs Gromacs Gromacs Gromacs Morton Morton Morton Morton Gromacs Gromacs Gromacs Gromacs Compute-3 Compute-2 Compute-1 Rooter Rooter Rooter Compute-4 Rooter Merger Merger Staging-2 Staging-1 Grid Grid Density Density lsosurface lsosurface

Compute a molecule surface based on atom density

Tested different distributions of processing steps to in situ and in transit nodes.



Densities Redistributed In Situ

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Densities Redistributed In Transit



Atom Positions Redistributed In Transit



Performance [Dreher,CCGRID'14]



- In transit: 1 staging node every 64 compute nodes
- Density-intransit:
 costs 7% comp. to
 gromacs 15 cores
- Density-insitu costs 8% but use 1.5% less nodes than density-intransit
- Atoms-intransit costs 8.6% but enables other in transit analytics (3x more data to move on stagging nodes than Density-intransit)



From In Situ Analytics to Computational Steering **Enterobactin Iron** Lipid Complex bilayer **FepA Channel** Ínría

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Conclusion

DataFlow Programming:

- FlowVR
- Google TensorfFlow (Machine learning)
- Apache Flink (Big Data stream + batch processing)

In-situ analytics:

- A paradigm shift that will very likely influence the HPC ecosystem (SW and HW)
- Not standard framework yet
- An opportunity to rethink the use of the I/O budget
- Bring some interactivity into the HPC world. Put the "user in the loop"

HPC and Big Data Convergence ?



Infratructures

- Dedicated platforms combining compute (Cluster + GPUs), data acquisition (Cameras) and visualization (display wall, VR headset) resources
 - Grimage (2002-2013)
 - Equipex Kinovis (since 2014)
 - Led to experiment sharing compute resources for batch and interactive computing through Grid'5000.
- Grid'5000 multi-site experiments (Bordeaux, Grenoble, Orléans)
- Grid'5000 transcontinental experiments (Japan, France)
- Mésocentre Ciment
- EDF & GENCI machines



References

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