Tuning *HipGISAXS* on Multi and Many Core Supercomputers

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Lawrence Berkeley National Laboratory

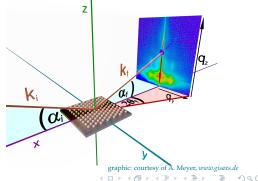


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Performance Modeling, Benchmarking and Simulation of High Performance Computer Systems Supercomputing 2013, Denver

Background: What is *HipGISAXS*?

- Massively parallel High-performance GISAXS simulation code.
- GISAXS: Grazing Incidence Small-Angle X-ray Scattering.
- X-ray scattering: a tool to measure structural properties of materials; characterize macromolecules and nano-particle systems at micro and nano-scales.
- Expose sample to high-energy X-rays, producing scattering patterns.
- Scattering patterns contain structural information about the sample.



Outline

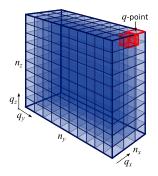
- 1 The main computational kernel.
- **2** Motivations and challenges.
- Contributions.
- ④ Platforms' overview.
- **6** Code optimization and tuning on clusters of:
 - Graphics processors (GPU);
 - Intel Phi coprocessors (MIC);
 - Intel Sandy Bridge and AMD Magny Cours CPUs.
- 6 Performance analysis and comparisons.
- Onclusions.

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HipGISAXS Kernel: Numerical Form Factor Calculations

- Q is grid of size $n_x \times n_y \times n_z$
- Described by 3 vectors,
 - $q_{\alpha} = \langle p_0 \cdots p_{n_{\alpha}-1} \rangle, \alpha \in \{x, y, z\}.$

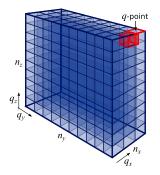


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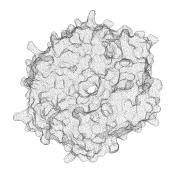
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HipGISAXS Kernel: Numerical Form Factor Calculations

- Q is grid of size $n_x \times n_y \times n_z$
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- *T* is set of *n*^{*t*} triangles describing a nano-structure model.
- $q_{\alpha} = \langle p_0 \cdots p_{n_{\alpha}-1} \rangle, \alpha \in \{x, y, z\}. \quad \bullet \quad \mathcal{T} = \{t_0 \cdots t_{n_{\ell}-1}\}, t_k = \langle \vec{r}_{t_{\ell}}, \vec{N}_{t_{\ell}}, s_{t_{\ell}} \rangle$



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HipGISAXS Kernel: Numerical Form Factor Calculations

• Form factor computed over model surface at each *q*-point:

$$\mathcal{F}: f(\vec{q}) = -rac{i}{|\vec{q}|^2} \sum_{t \in \mathcal{T}} e^{i \vec{q} \cdot \vec{r}_t} q_{N_t} s_t, \quad \forall \vec{q} \in \mathcal{Q}.$$

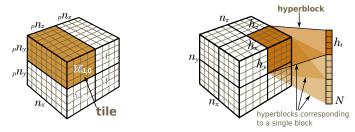
- Construct matrix \mathcal{F} of size $n_x \times n_y \times n_z$.
- Hence, computational complexity = $O(n_x n_y n_z n_t)$.
- Kernel is *compute-bound*.

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Generic Parallelization of the Form Factor Kernel

Covered in previous work ¹:

- "Embarrassingly parallel".
- *Tiles* across multiple nodes to decompose computations.
- Hyperblocks on a node to improve memory-based performance.
- Hyperblocks also helpful with accelerators.



1"Massively Parallel X-ray Scattering Simulations", Supercomputing, 2012. 🗇 😽 🖘 🖘 🖘 🖉 🗠 🔍

Motivations and Challenges

- This is a both *big data* and *big compute* problem.
- Data generation rate is continually increasing at light sources.
- Need for near real-time analysis.
- We utilize massive-parallelism to solve this.
- "Architecture-aware" algorithms and techniques necessary to extract raw computational power.
- Efficient mapping onto the processors' multi-level memory and parallelism hierarchies.
- Make effective use of state-of-the-art compute facilities, reducing cost (time and power).

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Contributions

- Optimizations on Nvidia Fermi and Kepler graphics processors.
- Parallelization and optimization on Intel MIC architecture.
- Optimization on the AMD Magny Cours and Intel Sandy Bridge processors.
- Implementation of auto-tuning of parallelization and optimization parameters.
- Detailed performance analysis and comparisons.

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Experimental Platforms

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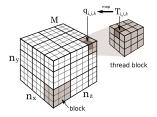
System	Titan (Cray XK7)	Stampede	Hopper (Cray XE6)	Edison-I (Cray XC30)
Architecture	GPU	MIC	Multicore	Multicore
Processor	Nvidia K20X	Intel Phi SE10P	AMD Magny Cours	Intel Sandy Bridge
Cores	14×192 CUDA	60 + 1	2×12	2×8
HW Threads/Core	1	4	1	2
Theoretical peak	3,950 GFLOP/s	2,021 GFLOP/s	403 GFLOP/s	664 GFLOP/s

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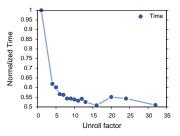
- Guided by performance profiling: Nvidia visual profiler, Intel VTune, PAPI.
- Expose more parallelism.
 - Loop interchange and loop collapsing.
 - New optimal CUDA thread-block decomposition scheme on GPUs.



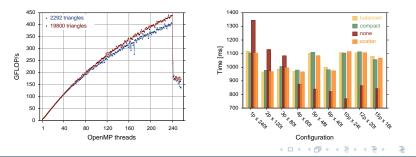
- Avoid memory transfers, read/writes.
 - Kernel fusion. E.g. 87% improvement on GPUs.
 - Use *persistent buffers* on MIC.
 - Data reuse. E.g. input data reuse through shared memory on GPUs.

- Overlap computations with memory transfers.
 - *k*-buffering. E.g. triple-buffering > 5% better than double-buffering.
- Optimize the unavoidable memory transfers.
 - Input triangles data (7 reals) padding (1 real) for better alignment (32B).
 - Avoid expensive operations, like vgatherdps on MIC, by input *data reorganization*. E.g. 33% improvement.

- Expose instruction level parallelism.
 - Shape triangles *loop unrolling*. 50% improvement on GPUs.



- Vectorize.
 - Reorganize data to facilitate better vectorization.
 - Hand vectorize and optimize complex number operations. E.g. 22% improvement on MIC over auto-vectorization.
 - E.g. use SSE2 on Magny Cours, AVX on Sandy Bridge.
 - Optimize exp and sincos. E.g. 25% improvement over SVML on MIC.
- Optimize execution environment configuration.
 - E.g. Thread configuration and thread affinities on MIC.



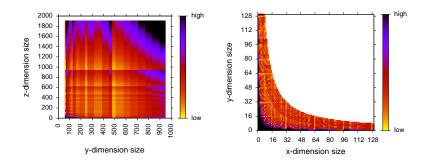


- Optimize parameter values. E.g. On GPUs,
- auto-tuning hyperblock size.

• auto-tuning CUDA thread block size.

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GPU Performance: Single Node

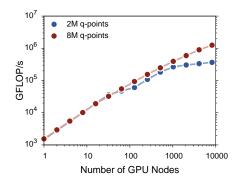
nt	$ \mathcal{Q} $	M2090	GFLOP/s	K20X	GFLOP/s
6,600	2M	0.68 s	813.6	0.25 s	2172.2
6,600	8M	2.73 s	813.2	1.02 s	2167.2
91,753	2M	9.47 s	813.4	3.56 s	2159.1
91,753	8M	37.9 s	813.4	14.25 s	2161.5

- Achieve 813 GFLOP/s on M2090, 61.2% of the theoretical peak.
- Achieve 2,172 GFLOP/s on K20X, 55% of the theoretical peak.

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GPU Performance: Scaling Across Multiple Nodes

Strong Scaling on Titan (7.5M input triangles)



• Achieve 1.25 PetaFLOP/s on 8,192 nodes.

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MIC Performance: Single Node

nt	$ \mathcal{Q} $	Xeon Phi	GFLOP/s
6,600	2M	2.27 s	453.58
6,600	8M	8.77 s	469.67
91,753	2M	30.77 s	465.22
91,753	8M	118.49 s	483.91
7,514,364	2M	2565.18 s	456.98

• Achieve 484 GFLOP/s, about 24% of the theoretical peak.

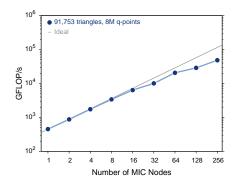
Tuning HipGISAXS on Multi and Many Core Supercomputers. PMBS @ SC13

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MIC Performance: Scaling Across Multiple Nodes

Strong Scaling on Stampede (91.7K input triangles)



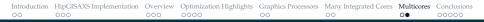
• Achieve 0.1 PetaFLOP/s on 1,024 nodes.

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Performance on Multicores (XE6/XC30): Single Node

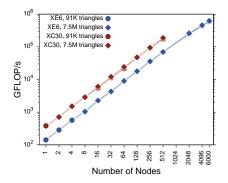
	nt	$ \mathcal{Q} $	XE6	GFLOP/s	XC30	GFLOP/s
6,6	00	2M	6.34 s	141.7	2.97 s	378.2
6,6	00	8M	25.28 s	142.1	$11.87 \mathrm{~s}$	378.3
91,75	53	2M	88.16 s	141.7	41.0 s	379.0
91 <i>,</i> 75	53	8M	352.0 s	142.0	164.1 s	380.0

- Achieve 142 GFLOP/s on XE6 node, 35.2% of the theoretical peak.
- Achieve 380 GFLOP/s on XC30 node, 57.2% of the theoretical peak.



Performance on Multicores (XE6/XC30): Scaling

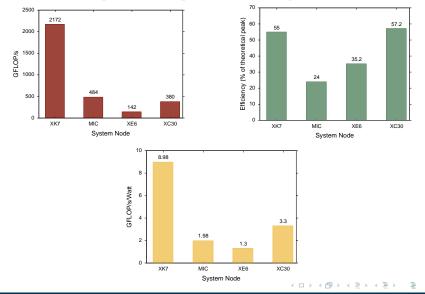
Strong Scaling on Hopper and Edison-I (91.7K and 7.5M input triangles)



- Achieve 0.63 PetaFLOP/s on 6,000 Hopper nodes (144,000 cores.)
- Achieve 0.19 PetaFLOP/s on 512 Edison-I nodes (8,192 cores.)

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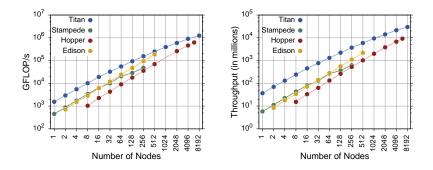
Performance Comparisons: HipGISAXS on Single Node



Tuning HipGISAXS on Multi and Many Core Supercomputers. PMBS @ SC13

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Performance Comparisons: HipGISAXS Strong Scaling



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Conclusions and Future Work

- Developing a basic code with acceptable performance is easier on multicores like the Magny Cours and Sandy Bridge.
- **But**, extraction of high-performance from them is no easier than implementations on GPUs or MIC.
- Architecture-aware implementations are inevitable for high-performance on these platforms.
- Graphics processors in general perform higher, and have high power-efficiency.
- Explore more architectures such as IBM BG/Q.
- Multi-level distributed memory parallel framework for simulations.
- GISAXS simulation is just one component of the under-development HipGISAXS suite.

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- Resource usage at *Texas Advanced Computing Center* (TACC) at the University of Texas at Austin.

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Thank you!

HipGISAXS Code and Documentation:

- http://portal.nersc.gov/project/m1285/hipgisaxs.html
- http://github.com/hipgisaxs/hipgisaxs/wiki

Additional Information:

- Massively Parallel X-ray Scattering Simulations, Supercomputing (SC12), no. 46, pp. 46:1–46:11, 2012.
- HipGISAXS: A High Performance Computing Code for Simulating Grazing Incidence X-Ray Scattering Data, Journal of Applied Crystallography, vol. 46 (6), pp. 1781–1795, 2013.

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