Goal

Discover where differences arise in scientific codes between CPU and GPU implementations and evaluate the effects on performance and accuracy.

Myths about Debugging Scientific Code

Many scientists do not thoroughly debug their code or compare results between GPU and CPU due to the assumption that results will differ on different platforms. Our research shows that comparing the results helps to debug both CPU code and GPU code. By understanding the sources of differences, errors can be corrected. In the medical reconstruction code we studied, all differences were removed and a bug was fixed in the CPU version of the code.

[1] Devon Yablonski, "Numerical Accuracy Differences in CPU and GPGPU Codes" MS Thesis Northeastern Uni. Sept. 2011.

Compiler Induced Differences

NVIDIA's CUDA compiler aggressively combines double precision addition and multiplication into **floating point**multiply-add (FMAD) instructions [3] on GT200 GPU hardware. The addition is truncated to be faster on the GPU which makes the CPU more accurate.

Example Equation	Device	Result	
1151 * 2.12221 + 221.99993	CPU	2220.663818	
	GPU	2220.663574	
Pseudo-Code float result = test1 * test2 + test3 GPU PTX CPU x86			
%f1 = test2 %f2 = test3 %f3 = t	-12(%rbp) = test	1 $-8(\%rbp) = test2 -4(\%rbp) = test3$	
	addss	-8(%rbp), -4(%rbp)	

Fig. 2: Instruction level (IL) MAD code. PTX is GPU IL code.

On Fermi, a **fused-multiply-add (FMA)** is used. The addition and multiplication only encounter a single rounding. The GPU is more accurate.

fma.f32 %**f4**, %**f2**, %**f3**, %**f1**; Rounded 1x

mulss -12(%rbp), -4(%rbp) addss -8(%rbp), -4(%rbp)

Fig. 3: Instruction level (IL) FMA code. PTX is GPU IL code.

Test Example - Tomosynthesis

Digital Breast Tomosynthesis (DBT) is a mammography algorithm that creates a 3D image from the data of 15 Xray scans to aid in the search for cancerous tissues. • Original code was developed by MGH and ported to the GPU at Northeastern by Schaa et al [4].





Rounded 2x!

Fig. 8: 15 X-Rays scan the subject

Fig. 9: Result from a test dataset used for this work

 Table 2: Code Performance* for 8 iterations of Tomosynthesis

Implementation	Time	Speedup
CPU Intel XEON W3580 @ 3.33GHz	29min 47s	-
GPU NVIDIA C1060	19s	97x

* These results are based on our test setup and may differ from published results of Schaa et al.

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[1] "IEEE Std 754-2008," *IEEE Std 754-2008,* pp. 1-58, 29 August 2008. [2] N. Whitehead, A. Fit-Florea. "Precision & Performance: Floating Point and IEEE 754 Compliance for NVIDIA GPUs". NVIDIA, 2011. [3] NVIDIA, "NVIDIA CUDA Programming Guide," http://developer.download.nvidia.com/compute/cuda/3_2/toolkit/docs/CUDA_C_Programming_Guide.pdf [4] D. Schaa, B. Jang, P. Mistry, R. Dominguez, D. Kaeli, R. Moore, D. B. Kopans, "GPU Acceleration of Iterative Digital Breast Tomosynthesis," GPU Gems 4