



Accelerated Computing from Mobile Devices to Supercomputers

Dale Southard, NVIDIA



Power of 600 Petaflop
CPU-only Supercomputer



=



Power for the city
of San Francisco



HPC's Biggest Challenge: Power

GPUs Power World's 10 Greenest Supercomputers



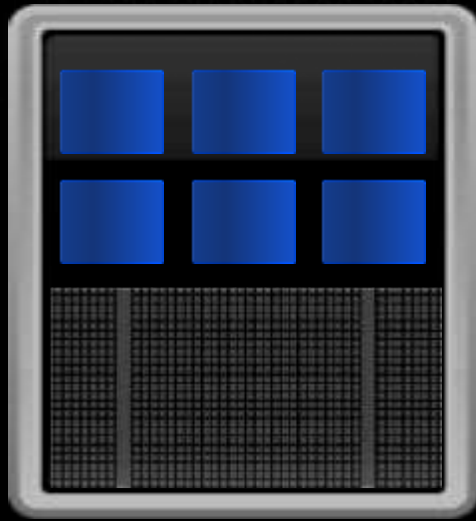
Green500 Rank	MFLOPS/W	Site
1	4,503.17	GSIC Center, Tokyo Tech
2	3,631.86	Cambridge University
3	3,517.84	University of Tsukuba
4	3,185.91	Swiss National Supercomputing (CSCS)
5	3,130.95	ROMEO HPC Center
6	3,068.71	GSIC Center, Tokyo Tech
7	2,702.16	University of Arizona
8	2,629.10	Max-Planck
9	2,629.10	(Financial Institution)
10	2,358.69	CSIRO
37	1959.90	Intel Endeavor (top Xeon Phi cluster)
49	1247.57	Météo France (top CPU cluster)

Accelerated Computing

10x Performance & 5x Energy Efficiency for HPC

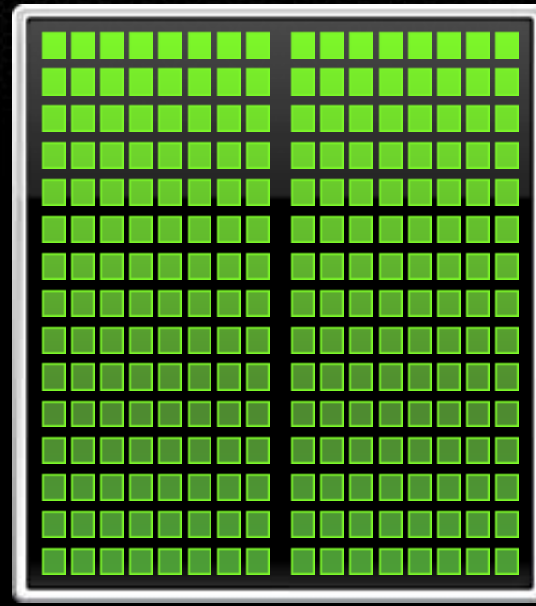
CPU

Optimized for
Serial Tasks

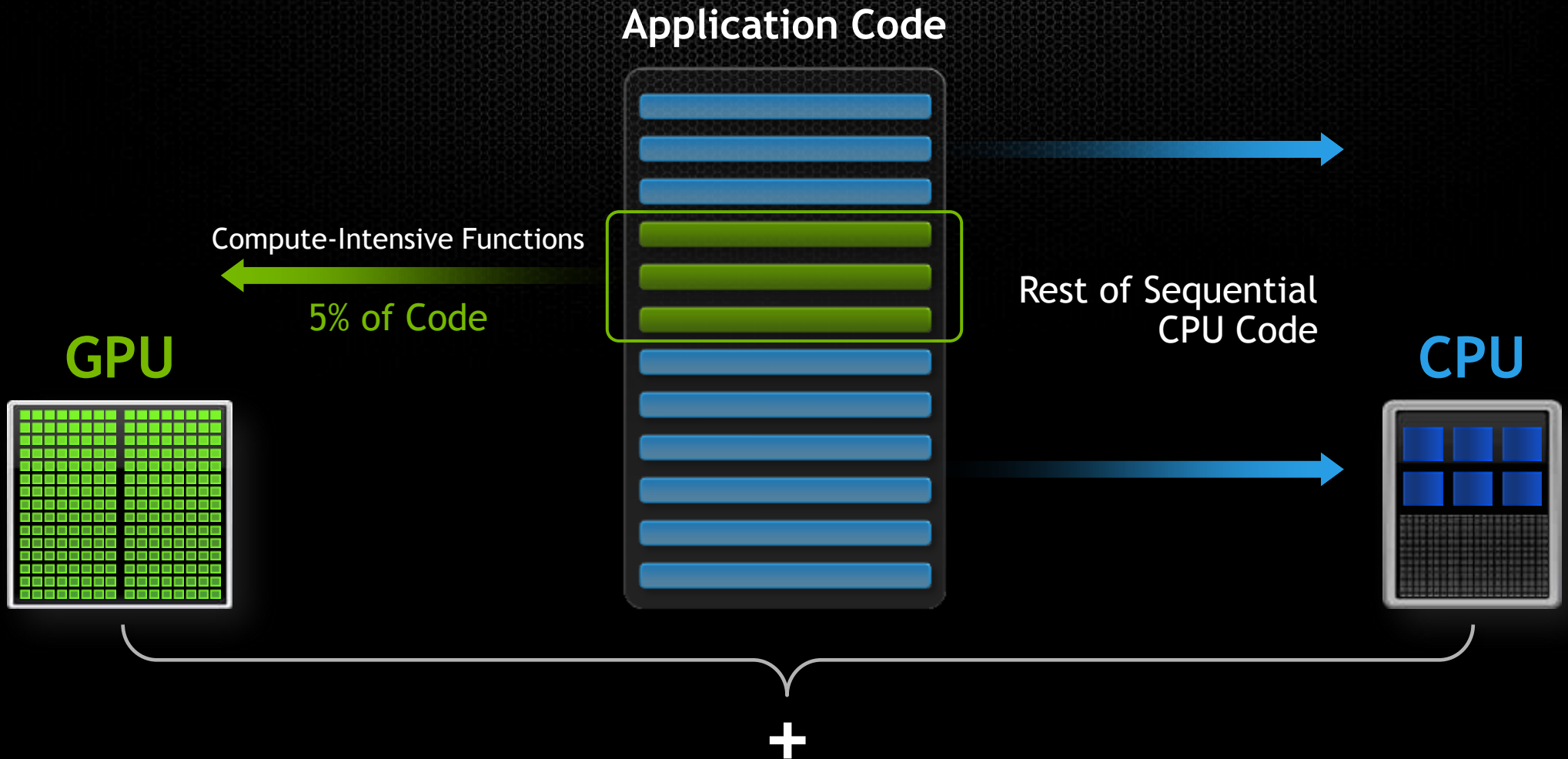


GPU Accelerator

Optimized for
Parallel Tasks

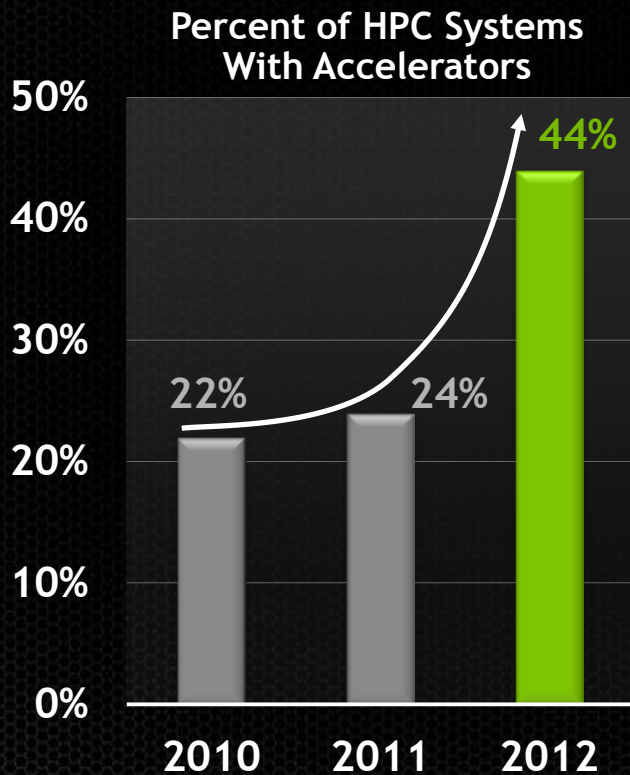


How GPU Acceleration Works



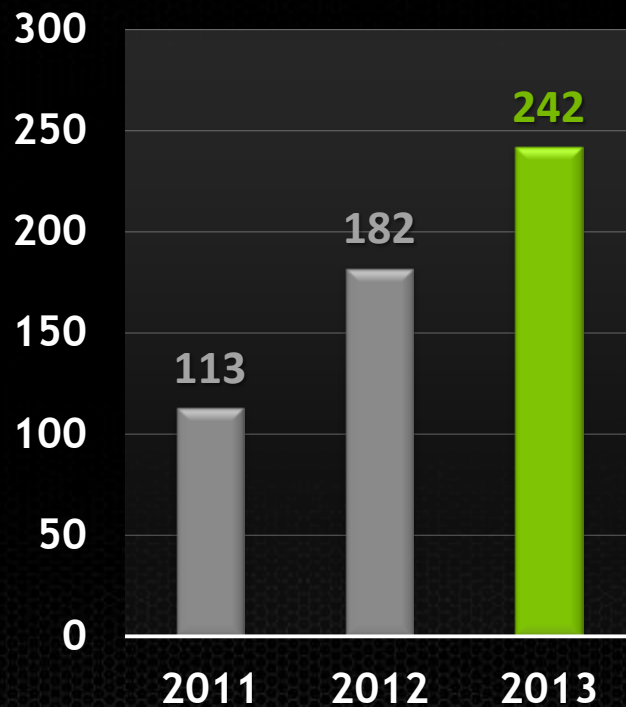
Accelerated Computing Growing Fast

2x Growth in One Year

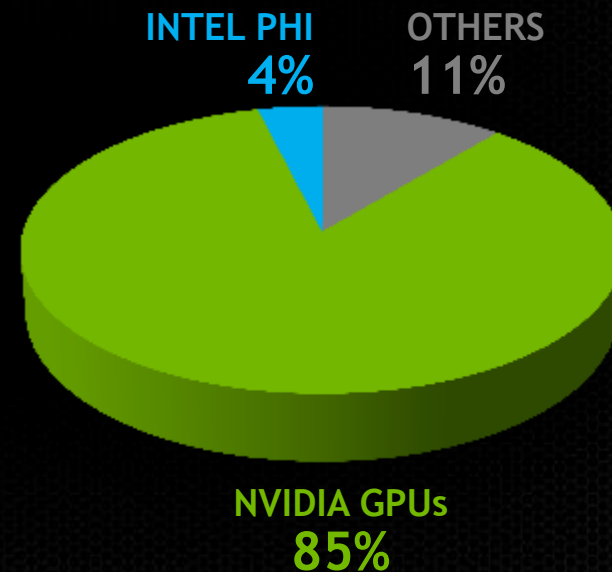


Intersect360 Research
HPC User Site Census: Systems, July 2013

Hundreds of GPU Accelerated Apps



NVIDIA GPU is Accelerator of Choice



Intersect360 Research
HPC User Site Census: Systems, July 2013



POPULAR GPU-ACCELERATED APPLICATIONS

02 Research: Higher Education and Supercomputing

COMPUTATIONAL CHEMISTRY AND BIOLOGY
BIOMEDICAL ANALYTICS
PATHS
WEATHER AND CLIMATE FORECASTING

06 Defense and Intelligence

COMPUTATIONAL FINANCE

MANUFACTURING: CAD AND CAE

COMPUTATIONAL FLUID DYNAMICS
COMPUTATIONAL STRUCTURAL MECHANICS
ELECTROMECHANICAL SIMULATION

10 Media and Entertainment

ANIMATION, RENDERING AND RENDERING
COLOR CORRECTION AND GRAFIC WORKFLOW
COMPOSITING, FINISHING AND EFFECTS
EDITING
ENCODING AND DIGITAL DISTRIBUTION
ON-SET, RENDER AND EDITING WORK
SIMULATION
WEATHER FORECASTS

14 Oil and Gas

Research: Higher Education and Supercomputing

COMPUTATIONAL CHEMISTRY AND BIOLOGY

Bioinformatics

Application	Description	Hardware Support	Operating System	Architecture	GPU Support	Availability
BerryCUDA	Sequence mapping software	Alignment of short sequencing reads	4-16x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 3.0.1
CUDA-MH	Open source software for Smith-Waterman protein database searches on GPUs	Parallel search of Smith-Waterman database	16-32x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 2.0.8
CUSAMM	Parallelized short read aligner	Parallel, accurate long read aligner - gapped alignments to large genomes	16x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 1.0.0E
GPU-BLAST	Local search with fast k-mer heuristic	Protein alignment according to k-mer, multi-epitope	3-4x	7.20T, 20P, 41G, 42G, 43G	Single only	Available now Version 2.2.24
GPU-HMMER	Parallelized local and global search with profile Hidden Markov models	Parallel local and global search of hidden Markov Models	48-1024x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 2.0.3
mDUSA-NEMO	Ultra-fast scalable motif discovery algorithm based on MEME	Scalable motif discovery algorithm based on MEME	4-16x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 3.0.1E
SeqFlux	A GPU Accelerated Sequence Analysis Toolkit	Reference assembly, BLAST, motif enrichment, term, de-novo assembly	400x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now
USENET	Open-source Smith-aligner for SGE/CUDA. Supports array based repeats filter and output	Fast short read aligner	9-16x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 1.1E
WidLM	Fits numerous linear models to a fixed design and response	Parallel linear regression on multiple similarly-shaped models	700x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 1.1-1

Molecular Dynamics

Application	Description	Hardware Support	Operating System	Architecture	GPU Support	Availability
Ashara	Models molecular dynamics of biopolymers for simulations of proteins, DNA and ligands	Simulation on 100k GPUs	4-256x	7.20T, 20P, 41G, 42G, 43G	Single Only	Available now Version 1.8.4B
ACEMD	GPU simulation of molecular mechanics force fields, implicit and explicit solvent	Written for use on GPUs	143 NVIDIA GPUs version only	7.20T, 20P, 41G, 42G, 43G	Yes	Available now
AMBER	Suite of programs to simulate molecular dynamics on biomolecules	PME, MD, explicit and implicit solvent	89-44 NVIDIA and 1 NVX	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 12.0 + cuda9
DL-POLY	Simulate macromolecules, polymers, ionic systems, etc on a distributed memory parallel computer	Two-body forces, Link-cell pairs, Rapid SPME forces, SHAKE-W	4x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 0 Tensor only
CHARMM	MD package to simulate molecular dynamics on biomolecules	Implicit Sol, Explicit Sol, Solvent for QM/MM	700x	7.20T, 20P, 41G, 42G, 43G	Yes	In Development 04/12
CHARMM-Blue	Simulation of biochemical molecules with complicated bath interactions	Implicit Sol, Explicit Sol, Solvent	145 NVIDIA GPUs	7.20T, 20P, 41G, 42G, 43G	Single only	Available now Version 8.8 in 04/12
HOOMD-blue	Particle dynamics package written primarily for GPUs	Written for GPUs	3x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now
LAMMPS	Classical molecular dynamics package	Lennard-Jones, Morse, Buckingham, CHARMM, tabulated, Colvars, coarse grain, SCX, Anisotropic dipole, BE-resp, hybrid combinatorial	3-18x	7.20T, 20P, 41G, 42G, 43G	Yes	Available now
MOE	Designed for high performance simulation of large molecular systems	MM/GBSA, MM/BSA, MM/BSA	4-64 NVIDIA GPUs	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 2.9
MOE-M	Library and application for molecular dynamics for xPC with GPUs	Implicit and explicit solvent, coarse force	Implicit: 127-243 NVIDIA GPUs Explicit: 16-32 NVIDIA GPUs	7.20T, 20P, 41G, 42G, 43G	Yes	Available now Version 6.1.1


272 GPU-Accelerated Applications
www.nvidia.com/appscatalog

Artificial Neural Network at a Fraction of the Cost with GPUs

“Now You Can Build Google’s \$1M Artificial Brain on the Cheap”

-Wired


GOOGLE BRAIN



1,000 CPU Servers
2,000 CPUs • 16,000 cores

600 kWatts
\$5,000,000

STANFORD AI LAB



3 GPU-Accelerated Servers
12 GPUs • 18,432 cores

4 kWatts
\$33,000

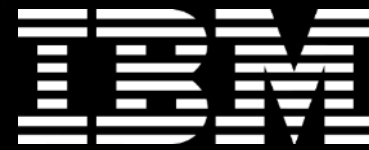
GPUs Accelerate Machine Learning & Data Analytics



Auto Tagging in Creative Cloud



Speech/Image Recognition



Hadoop-based Clustering



Recommendation Engine

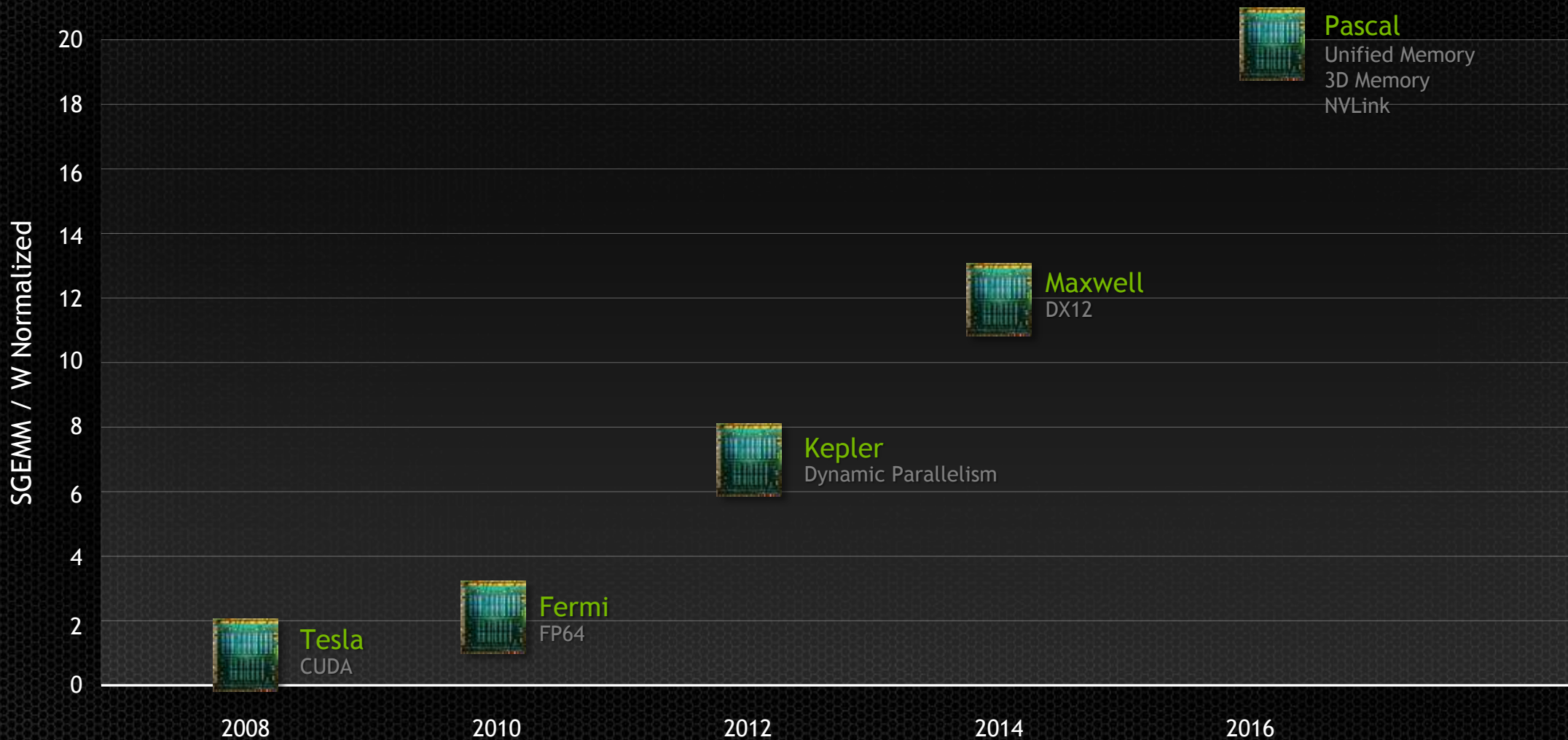


Database Queries

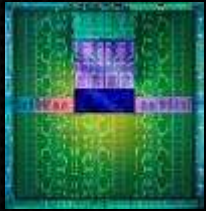


Search Ranking

Strong CUDA GPU Roadmap



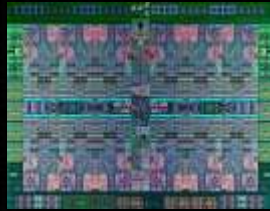
IBM Partners with NVIDIA to Build Next-Generation Supercomputers



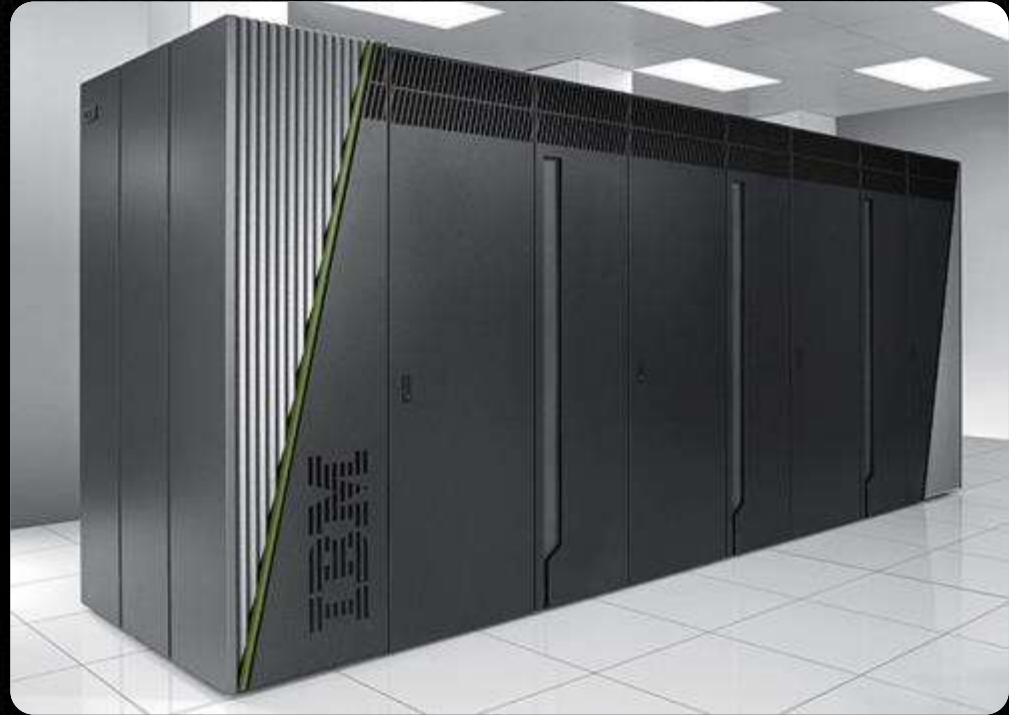
Tesla
GPU



+



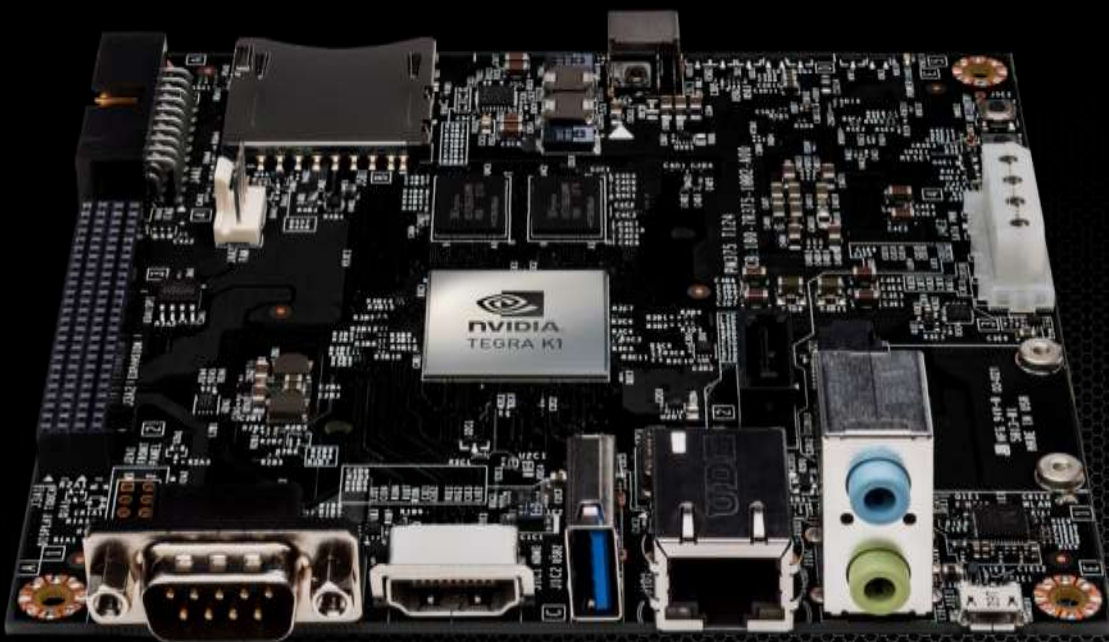
POWER 8
CPU



GPU-Accelerated POWER-Based Systems Available in 2014

JETSON TK1

THE WORLD'S 1st EMBEDDED SUPERCOMPUTER



Development Platform for Embedded
Computer Vision, Robotics, Medical

192 Cores · 326 GFLOPS FP32

CUDA Enabled

Available Now

GPU Accelerated Libraries

“Drop-in” Acceleration for your Applications

Linear Algebra

FFT, BLAS,
SPARSE, Matrix



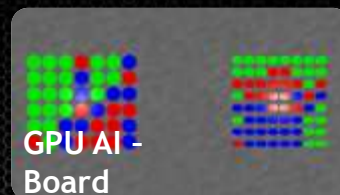
Numerical & Math

RAND, Statistics



Data Struct. & AI

Sort, Scan, Zero Sum

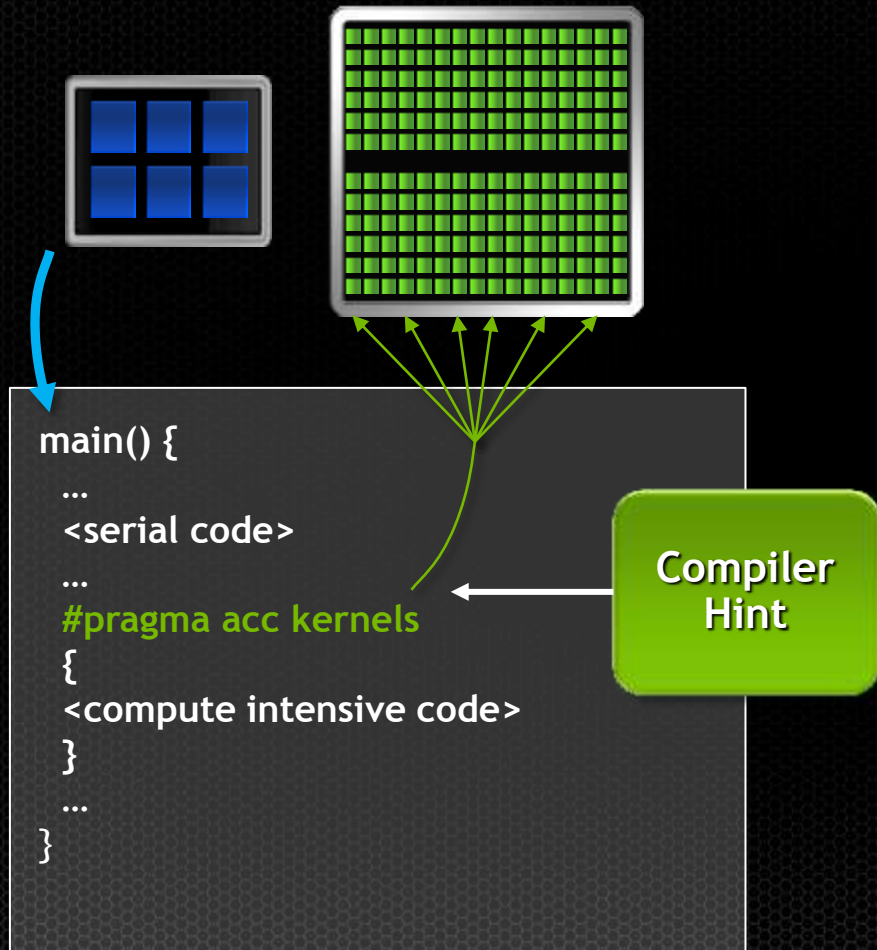


Visual Processing

Image & Video



OpenACC: Open, Simple, Portable



- Open Standard
- Easy, Compiler-Driven Approach
- Portable on GPUs and Xeon Phi

CAM-SE Climate
6x Faster on GPU
Top Kernel: 50% of Runtime



Linux GCC Compiler to Support GPU Accelerators

Open Source

GCC Efforts by Samsung & Mentor Graphics

Pervasive Impact

Free to all Linux users

Mainstream

Most Widely Used HPC Compiler



“ *Incorporating OpenACC into GCC is an excellent example of open source and open standards working together to make accelerated computing broadly accessible to all Linux developers.* **”**

Oscar Hernandez
Oak Ridge National Laboratories



ANNOUNCING CUDA 6

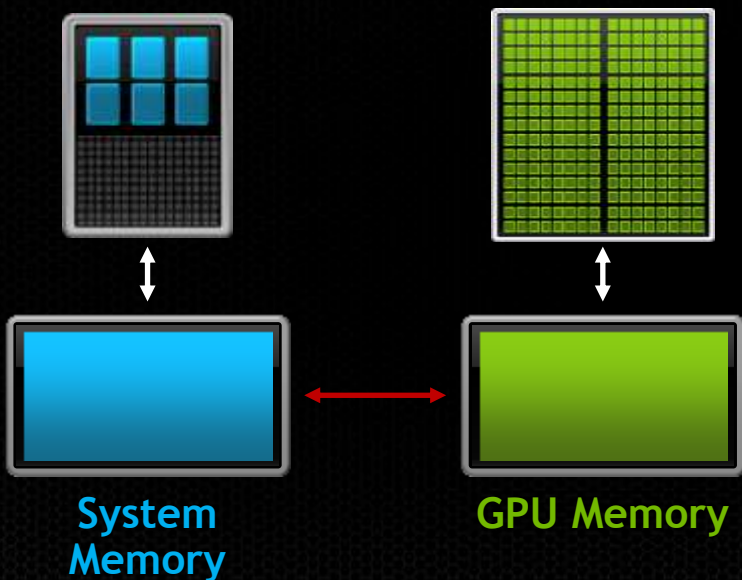


Dramatically Simplifies
Parallel Programming with
Unified Memory

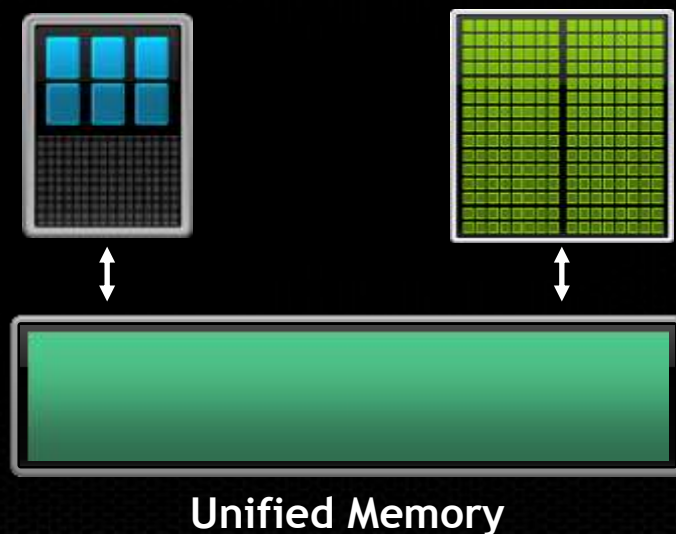
Unified Memory

Dramatically Lower Developer Effort

Developer View Today



Developer View With Unified Memory



CUDA: World's Most Pervasive Parallel Programming Model

14,000

Institutions with
CUDA Developers

2,000,000

CUDA Downloads

487,000,000

CUDA GPUs Shipped

700+ University Courses
In 62 Countries





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