DVIDIA.

Throughput Computing The Quest for Efficiency and Programmability

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The Exascale Challenge

Sustain 1EFLOPs on a "real" application

Power less than 20MW



The Cellphone Challenge

Deliver 50GFLOPs on mobile applications

Power < 1W





From cell phones to supercomputers we are *Power Limited*

Perf/W is Performance



And we need to make it easy to program both these devices



TITAN

18,688 NVIDIA Tesla K20X GPUs27 Petaflops Peak: 90% of Performance from GPUs17.59 Petaflops Sustained Performance on Linpack



Tsubame KFC 4.5GFLOPS/W #1 on Green500 List





Its not about the FLOPs

DFMA 0.05mm² 10pJ/OP – 2GFLOPs

A chip with 10⁴ FPUs: 500mm² 200W 20TFLOPS

Pack 50,000 of these in racks 1EFLOPS 10MW

16nm chip, 25mm on a side, 200W



Overhead

Locality



How is Power Spent in a CPU?



Dally [2008] (Embedded in-order CPU)

Natarajan [2003] (Alpha 21264)











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Payload Arithmetic 20pJ

Overhead 20pJ



The Locality Challenge Data Movement Energy 77pJ/F





Energy Shopping List

Processor Technology	40 nm	10nm
Vdd (nominal)	0.9 V	0.7 V
DFMA energy	50 pJ	7.6 pJ
64b 8 KB SRAM Rd	14 pJ	2.1 pJ
Wire energy (256 bits, 10mm)	310 pJ	174 pJ

Memory Technology	45 nm	16nm
DRAM interface pin bandwidth	4 Gbps	50 Gbps
DRAM interface energy	20-30 pJ/bit	2 pJ/bit
DRAM access energy	8-15 pJ/bit	2.5 pJ/bit



Keckler [Micro 2011], Vogelsang [Micro 2010]



Minimize Data Movement

Move Data More Efficiently







GRS Test Chips













Test Chip #2 fabricated on production GPU



Optimized Circuits 77pJ/F -> 18pJ/F



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While Improving Locality

Simplify Programming

Parallel programming is not inherently any more difficult than serial programming

However, we can make it a lot more difficult



A simple parallel program

```
forall molecule in set { // launch a thread array
  forall neighbor in molecule.neighbors { // nested
     forall force in forces { // doubly nested
     molecule.force =
     reduce_sum(force(molecule, neighbor))
  }
```



Why is this easy?

```
forall molecule in set { // launch a thread array
   forall neighbor in molecule.neighbors { // nested
      forall force in forces { // doubly nested
      molecule.force =
      reduce_sum(force(molecule, neighbor))
   }
```

No machine details All parallelism is expressed Synchronization is semantic (in reduction)



```
pid = fork() ; // explicitly managing threads
```

```
lock(struct.lock) ; // complicated, error-prone synchronization
// manipulate struct
unlock(struct.lock) ;
```

code = send(pid, tag, &msg) ; // partition across nodes

We could make it hard



Programmers, tools, and architecture Need to play their positions



Map molecules across memories Stage data up/down hierarchy Select mechanisms







Autotuned Software 18pJ/F -> 9pJ/F



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Conclusion

Power-limited: from data centers to cell phones

Perf/W is Perf

Throughput cores

Reduce overhead



Data movement

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- **Circuits:** 200 -> 20
- Optimized software

Parallel programming is simple - we can make it hard

Target-independent programming - mapping via tools







"Super" Computing From Super Computers to Super Phones