



Performance Benchmarks and Tuning on QB3

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Objectives

- Show application performance benchmarks on QB3
- Show methods that may improve the performance









Disclaimers

- Target audience: users who run (and sometimes compile) applications developed by others
 - Not an in-depth guide for programmers and developers









Outline

- Summary of QB3 architecture
- Single node performance
- Multi-node performance









Summary of QB3 architecture



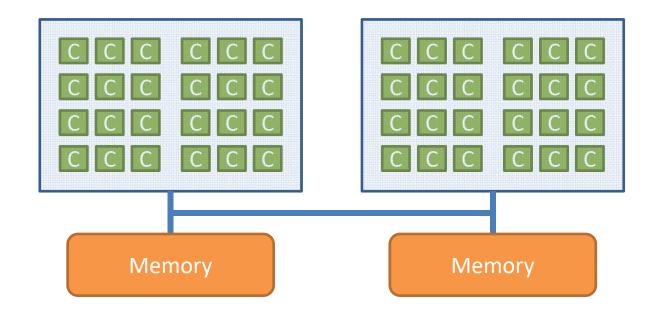






QB3 Architecture

Node level view





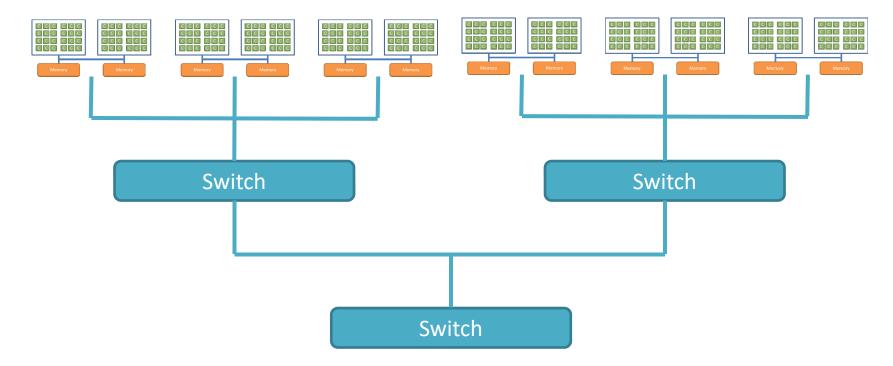






QB3 Architecture

Cluster level view











Parallel Paradigms

Con Pro Low latency, high bandwidth Shared memory system Implicit only (Limited to one Intranode communication node) Fine granularity Dynamic load balancing High latency, low Scalability beyond bandwidth Internode 1 node **Explicit communication** Hard to load balancing Switch









Parallel Paradigms

Pro Con Low latency, high bandwidth Shared memory system Implicit 1 to one **OpenMP (Multi-thread)** Intranode Fine granularity Dynamic load balancing High latency, low Scalability beyond **MPI (Multi-process)** Internode munication Hard to load balancing Switch

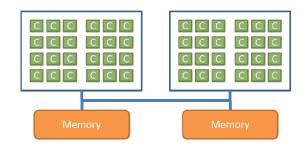


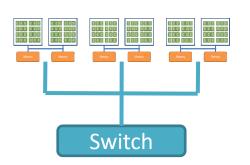






What About MPI+OpenMP Hybrid?





- Getting the benefits from both worlds?
- In theory, yes
- But adding OpenMP to (wellwritten) MPI programs might hurt the performance
- Hybrid helps to
 - Reduce memory footprint
 - Extend scalability

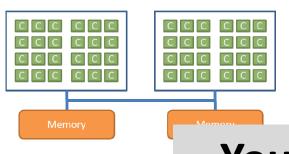








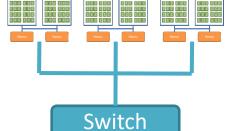
What About MPI+OpenMP Hybrid?



- Getting the benefits from both worlds?
- In theory, yes

Your mileage may vary!

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- Hybrid helps to
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Single Node Performance









QB3 Specification

CPU	Intel Cascade Lake (Xeon Platinum 8260) (2 sockets *24 cores/socket)	
CPU frequency	2.4 G Hz	
Floating operation per clock cycle (double precision)	512	
Memory	192 GB DDR4	
Interconnect	Mellanox 100 Gbps Infiniband	









QB3 vs QB2 (Node over Node)

	QB3	QB2
CPU frequency	2.4 x 10 ⁹	2.8 x 10 ⁹
CPU cores	48	20
Operation per cycle	32	8
Memory bandwidth	~115 GB/s	~48 GB/s
Interconnect	100 Gbps	56 Gbps

Node peak performance

QB3: $48 \text{ cores/node} * 2.4 \times 10^9 \text{ cycles/second} * 32 \text{ flop/cycle} = 3.69 \times 10^{12} \text{ flops}$

QB2: 20 cores/node * 2.8×10^9 cycles/second * $8 \text{ flop/cycle} = 0.45 \times 10^{12}$ flops



Theoretical speedup = 8.2







STREAM Benchmark

 The de facto industry standard benchmark in HPC domain for the measurement of sustainable memory bandwidth (in GB/s).

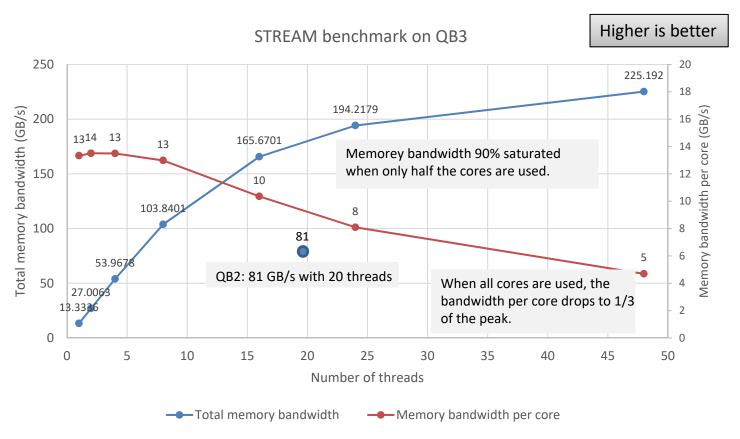








STREAM Benchmark - Results





STREAM 5.10 Array size = 4000 MB Intel 19.0.5 with "-O3 -xCORE-AVX512 -qopt-zmm-usage=high" Default thread affinity







Compiler Flags (Intel)

- -02, -03
 - Generic, aggregated optimization flag
- -xCORE-AVX512
 - Turns on optimization for the Skylake/Cascade Lake processor
- -qopt-zmm-usage=high
 - Improves performance for some code

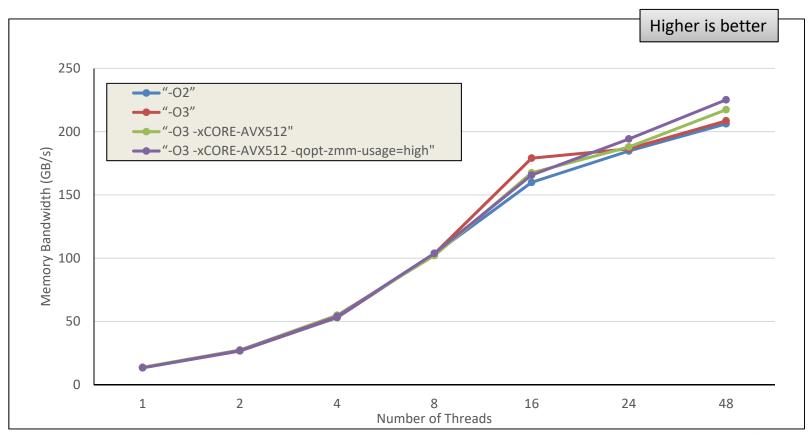








STREAM – Compiler Flags





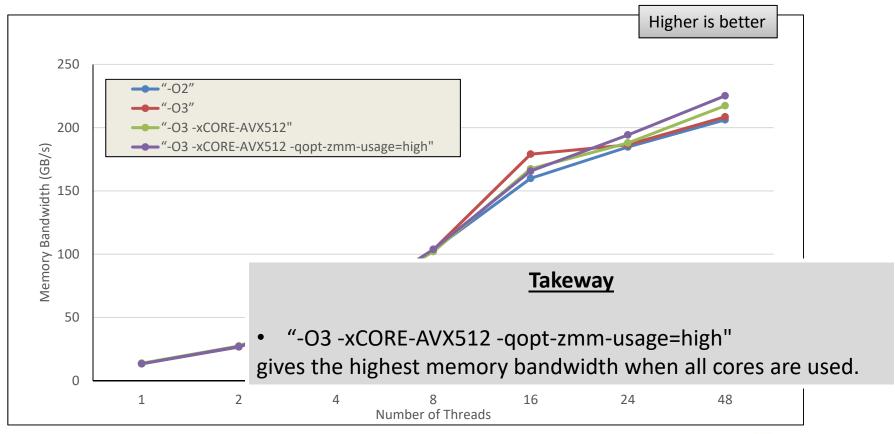
STREAM 5.10 Array size = 4000 MB Intel 19.0.5 Default thread affinity







STREAM – Compiler Flags





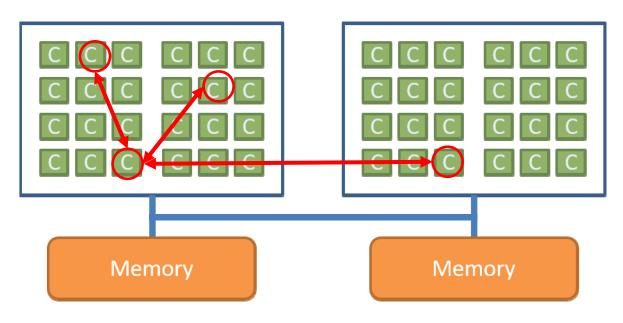
STREAM 5.10 Array size = 4000 MB Intel 19.0.5 Default thread affinity







Thread Affinity



- The 48 cores on a node are grouped into 4 sets.
- The data exchange cost is not homogeneous.
- Depending on the data exchange pattern, how the threads are arranged could affect performance significantly.

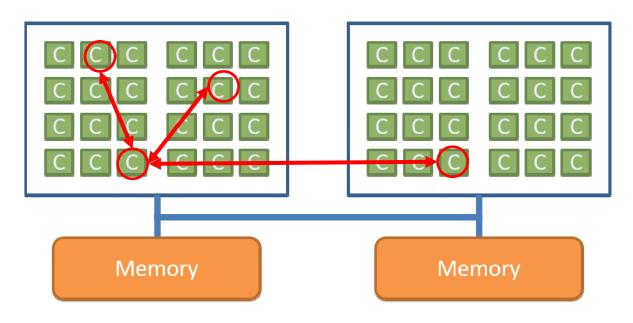








Thread Affinity



- For Intel compiler, use the KMP_AFFINITY environment variable
- The options are "none", "disabled", "balanced", "compact", and "scatter".

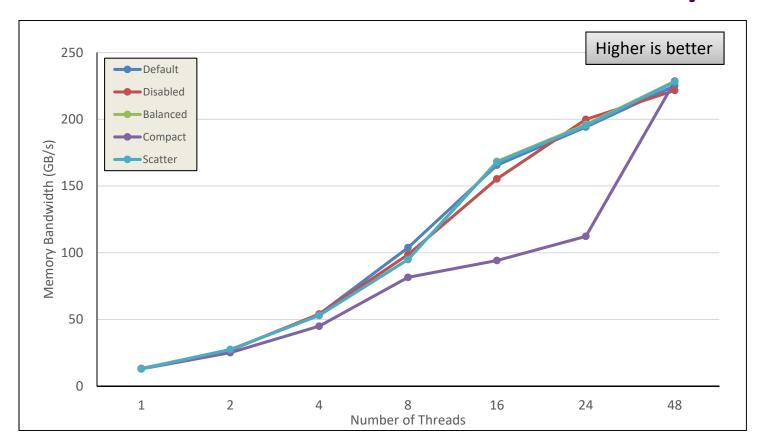








STREAM – Thread Affinity





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HPL Benchmark

- High Performance Linpack
 - Standard benchmark for CPU-bound HPC applications
- Results
 - QB3: 2540 GFLOPS per node
 - QB2: 424 GFLOPS per node
 - Speedup = 6 (compared to 8.2 theoretical)



HPL 2.3 (Intel MKL version)

Problem size = 90% installed memory

MVAPICH2 2.3.3

Intel 19.0.5 with "-03 -xCORE-AVX512"







HPCG Benchmark

- High Performance Conjugate Gradient
 - Standard benchmark for memory-bound HPC applications
- Results
 - QB3: 32.4 GFLOPS per node
 - QB2: 14.5 GFLOPS per node
 - Speedup: 2.2 (compared to 8.2 theoretical)



HPCG 3.0 (Intel MKL version)
Problem size = 336 336 336
MVAPICH2 2.3.3
Intel 19.0.5 with "-O3 -xCORE-AVX512 -qopt-zmm-usage=high"







NPB Benchmark Suite

- NAS Parallel Benchmarks
 - a small set of programs derived from computational fluid dynamics applications
- Five kernels and three pseudo-applications
 - IS Integer Sort, random memory access
 - EP Embarrassingly Parallel
 - CG Conjugate Gradient, irregular memory access and communication
 - MG Multi-Grid on a sequence of meshes, long- and short-distance communication, memory intensive
 - FT discrete 3D fast Fourier Transform, all-to-all communication
 - BT Block Tri-diagonal solver
 - SP Scalar Penta-diagonal solver
 - LU Lower-Upper Gauss-Seidel solver

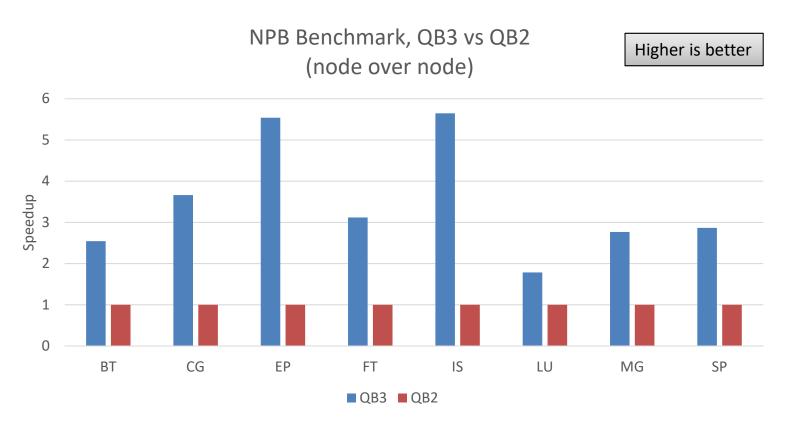








NPB Benchmarks – Node over Node





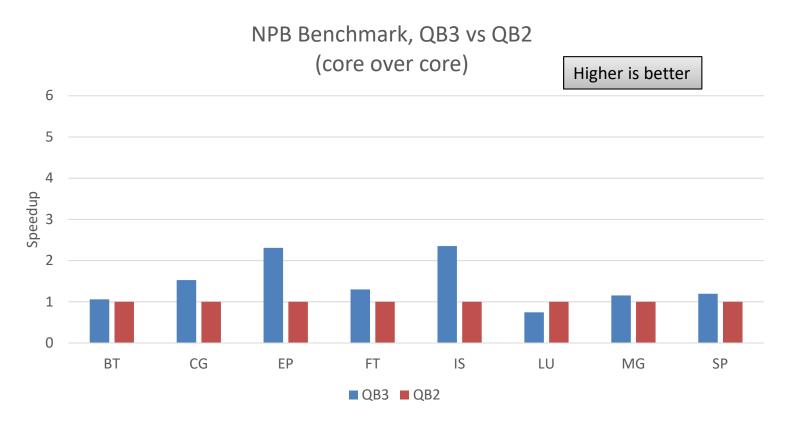
NPB 3.4.1, Class C Intel 19.0.5 with "-O3 –xCORE-AVX512" Default thread affinity







NPB Benchmarks – Core over Core





NPB 3.4.1, Class C Intel 19.0.5 with "-O3 –xCORE-AVX512" Default thread affinity







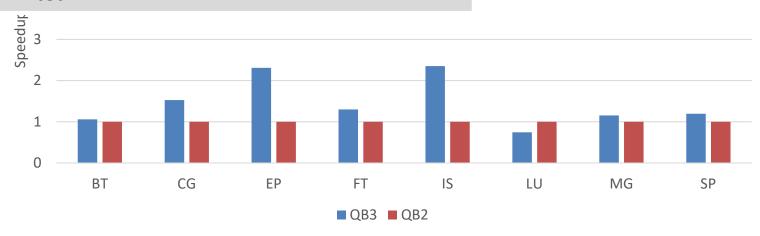
NPB Benchmarks – Core over Core

Takeway

- Performance gain on QB3 varies a lot from application to application.
- Core-over-core performance gain could be very limited.

QB2

Higher is better





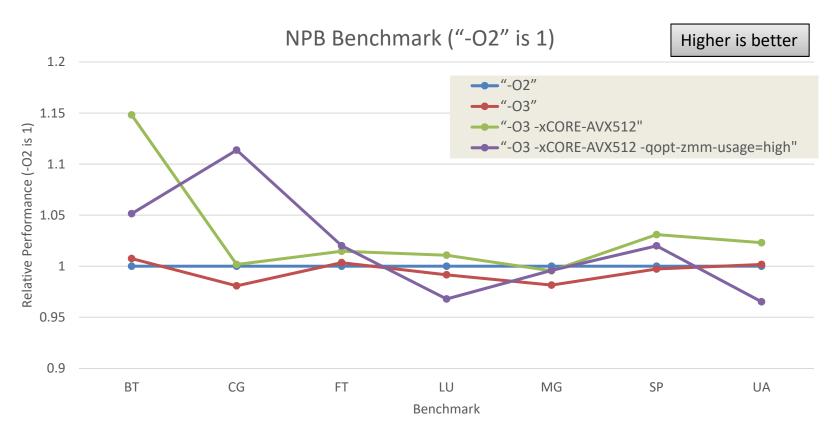
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NPB Results – Compiler Flags





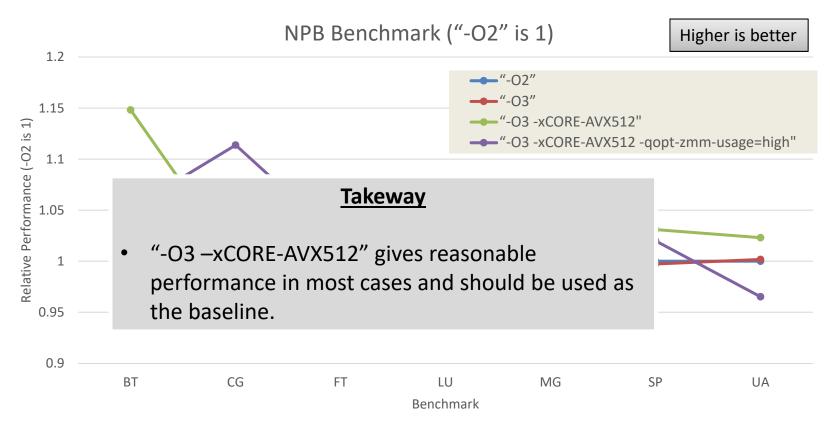
NPB 3.4.1, Class D Intel 19.0.5 Default thread affinity







NPB Results – Compiler Flags





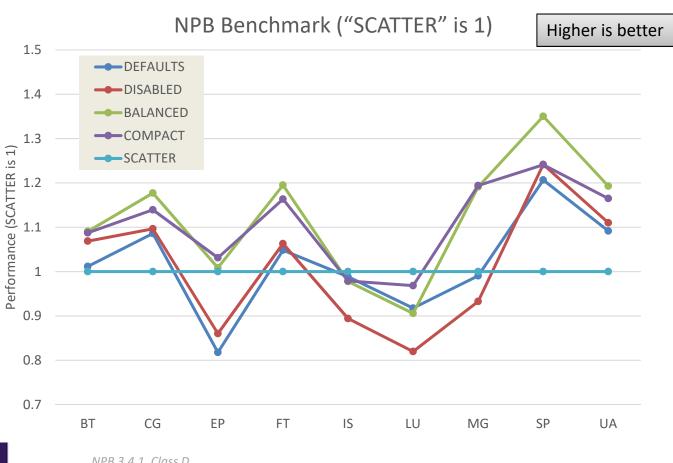
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NPB Results – Thread Affinity





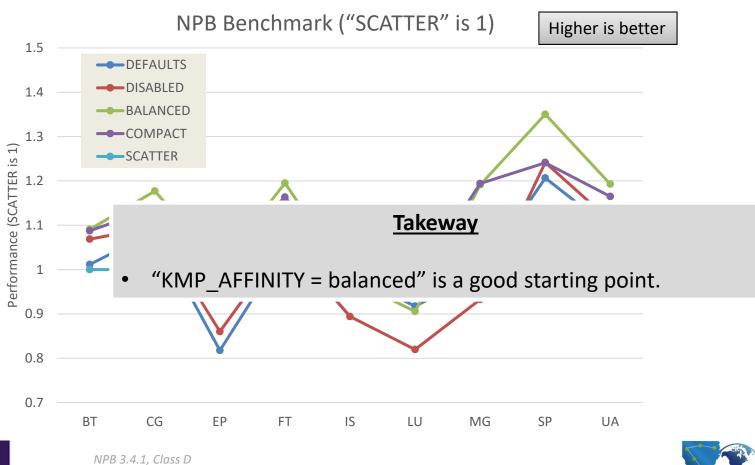
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NPB Results – Thread Affinity





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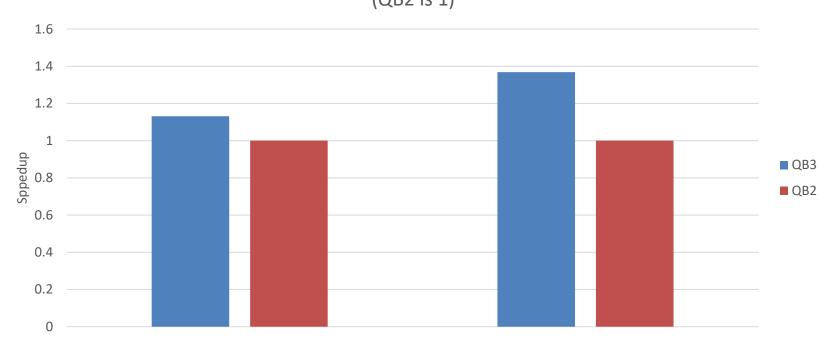
LSU



Bioinformatics

QB3 Performance Compared to QB2 (QB2 is 1)

Higher is better



CellRanger (single cell RNAseq)

Supernova (de novo assembly)



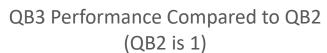
Both are binary distribution from 10x Genomics



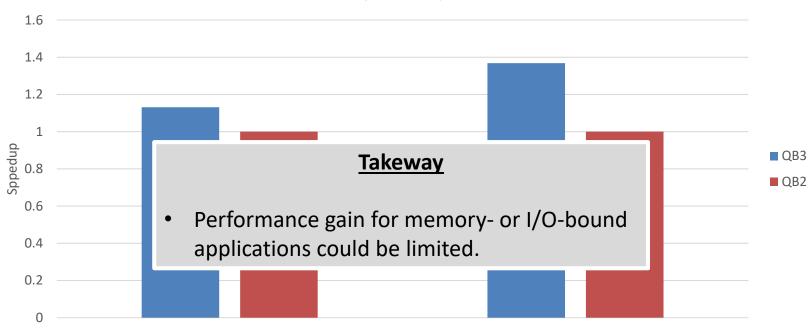




Bioinformatics



Higher is better



CellRanger (single cell RNAseq)

Supernova (de novo assembly)



Both are binary distribution from 10x Genomics Cellranger: mouse intestine cell, data size = 134 GB Supernova: fruit fly genome, data size = 46 GB







Multi-node Performance

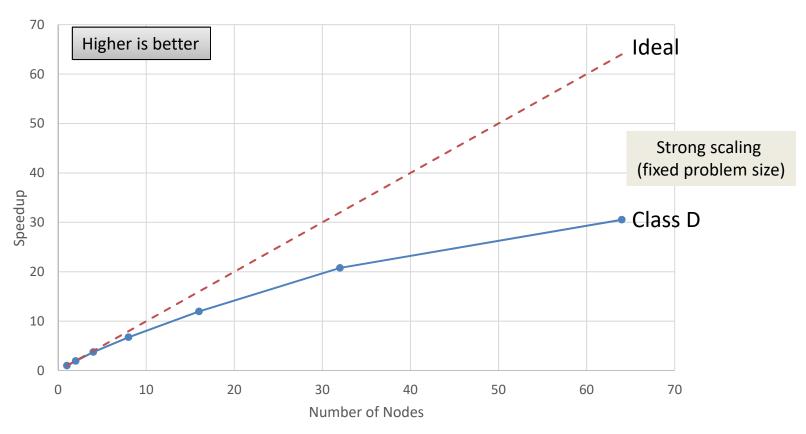








Pure MPI - NPB LU Benchmark





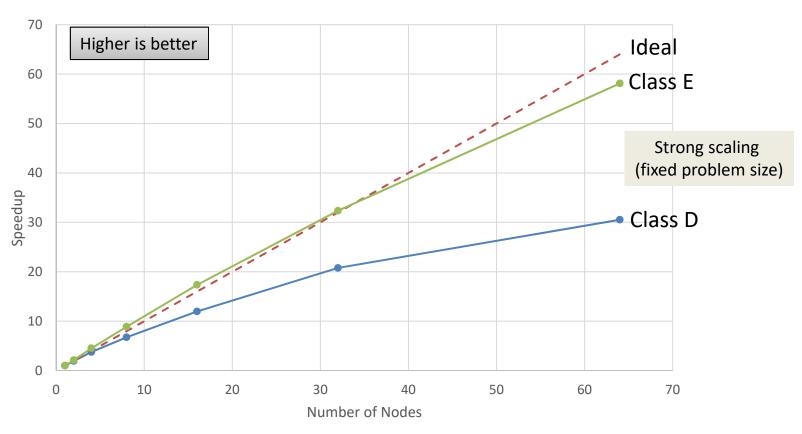
NPB 3.4.1, Class D and E MVAPICH2 2.3.3 Intel 19.0.5 with "-O3 -xCORE-AVX512"







Pure MPI - NPB LU Benchmark





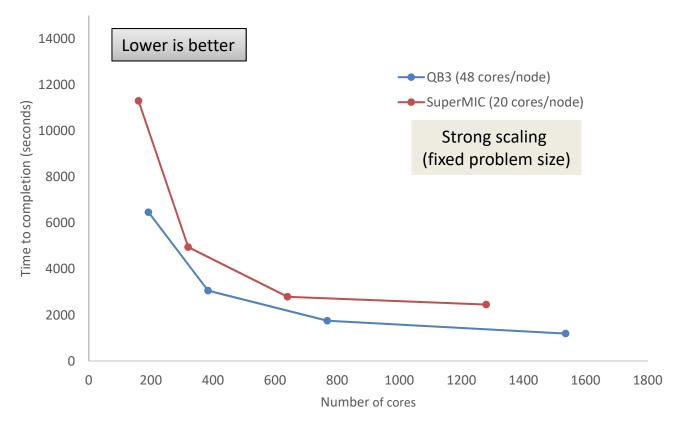
NPB 3.4.1, Class D and E MVAPICH2 2.3.3 Intel 19.0.5 with "-O3 -xCORE-AVX512"







Pure MPI - PADCIRC





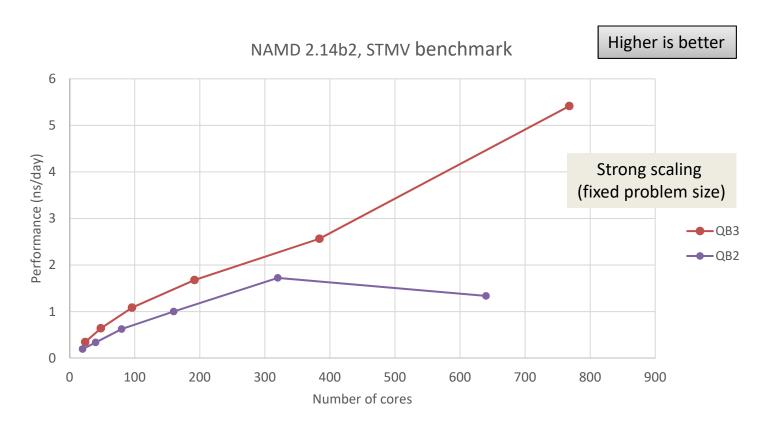
PADCIRC version 55 MVAPICH2 2.3.3 Intel 19.0.5







NAMD





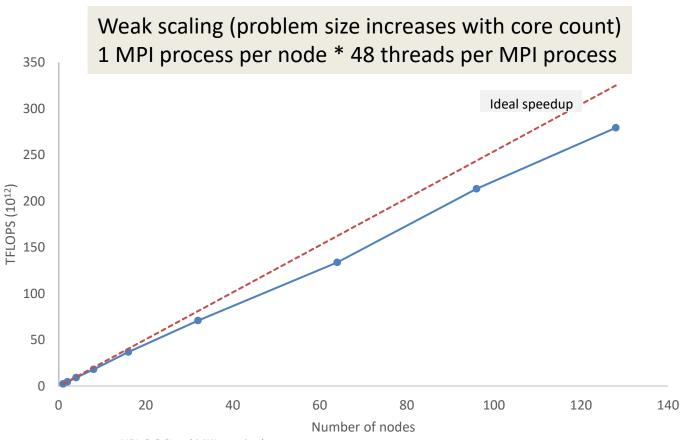
NAMD 2.14b2 Iverbs version of charmrun Intel 19.0.5







Hybrid – HPL





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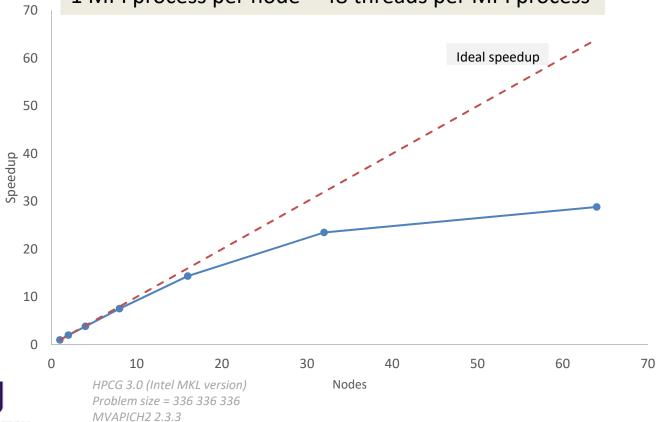




Hybrid - HPCG

Weak scaling (problem size increases with core count)

1 MPI process per node * 48 threads per MPI process





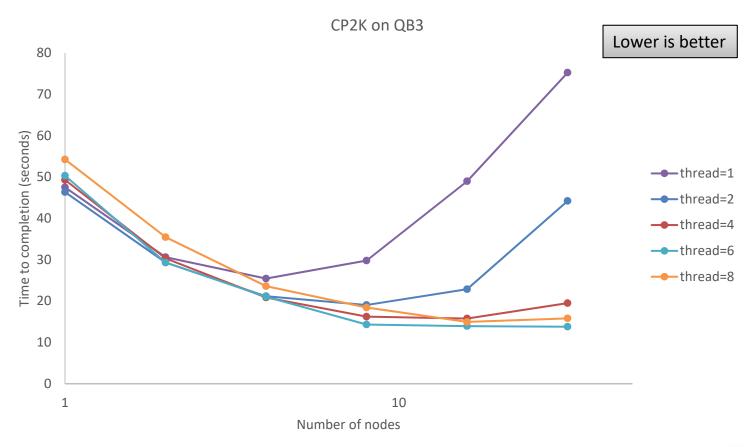


Intel 19.0.5 with "-O3 -xCORE-AVX512 -qopt-zmm-usage=high"





Hybrid - CP2K





CP2K 7.1 Intel MPI 2019u5 Intel 19.0.5 with "-O3 –xCORE-AVX512 -qopt-zmm-usage=high"







I/O Consideration

- You want to avoid letting your program accesses to disk excessively
 - Reading/writing hundreds of GBs to checkpoint or output files frequently
 - Running with thousands of MPI tasks all reading/writing individual files
- What you should and should not do
 - Avoid writing intermediate/checkpoint files unless necessary
 - Look for and use options that allow one or a few big files instead of files per process
 - Reduce the frequency of writing output files
 - Do not use /home for productive jobs use /work instead









Takeaways

- In the majority cases, your application will run faster and scale better on QB3 (compared to QB2)
- That being said, how much faster depends on a lot of factors
- You need to figure that out before making a (educated) decision whether or not switch to QB3
- You need to run your own experiments









Takeways

- Baseline
 - Use "-O3 -xCORE-AVX512" to compile
 - The default settings work reasonably well in most cases
- Serial programs
 - The performance gain will be limited
- OpenMP programs
 - Try different KMP_AFFINITY settings
- MPI programs
 - Remember the scaling behavior depends on the problem size
- Hybrid programs
 - It does not always help
 - Finding the optimal number of threads could be tricky and certainly varies from application to application









Takeaways

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