

AMBER 11

Performance Benchmark and Profiling

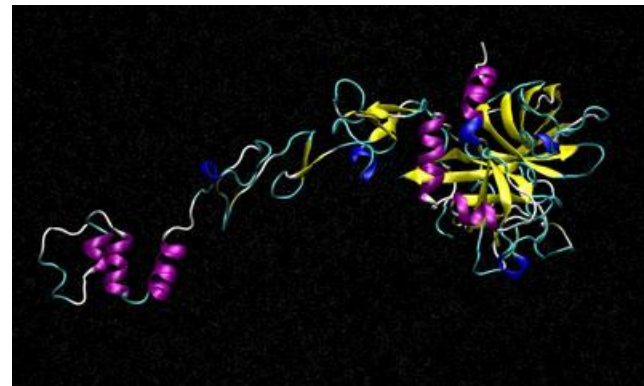
July 2011



- **The following research was performed under the HPC Advisory Council activities**
 - Participating vendors: AMD, Dell, Mellanox
 - Compute resource - HPC Advisory Council Cluster Center
- **For more info please refer to**
 - [http:// www.amd.com](http://www.amd.com)
 - [http:// www.dell.com/hpc](http://www.dell.com/hpc)
 - <http://www.mellanox.com>
 - <http://ambermd.org/>

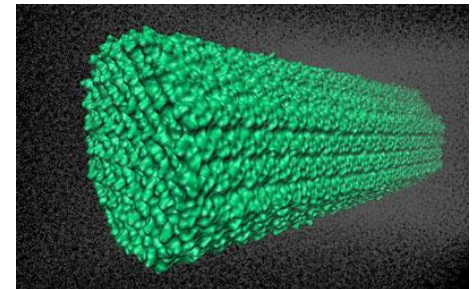
- **AMBER**

- Software for analyzing large-scale molecular dynamics (MD) simulation trajectory data
- Reads either CHARMM or AMBER style topology/trajectory files as input, and its analysis routines can scale up to thousands of compute cores or hundreds of GPU nodes with either parallel or UNIX file I/O
- AMBER has dynamic memory management, and each code execution can perform a variety of different structural, energetic, and file manipulation operations on a single MD trajectory at once
- The code is written in a combination of Fortran90 and C, and its GPU kernels are written with NVIDIA's CUDA API to achieve maximum GPU performance



- **The following was done to provide best practices**
 - AMBER performance benchmarking
 - Understanding AMBER communication patterns
 - Ways to increase AMBER productivity
 - Compilers and MPI libraries comparisons
- **The presented results will demonstrate**
 - The scalability of the compute environment
 - The capability of AMBER to achieve scalable productivity
 - Considerations for performance optimizations

- **Dell™ PowerEdge™ R815 11-node (528-core) cluster**
- **AMD Opteron™ 6174 Series processors (codenamed “Magny-Cour”) 12-cores @ 2.2 GHz**
- **4 CPU sockets per server node**
- **Mellanox ConnectX-2 VPI adapters for 40Gb/s QDR InfiniBand and 10Gb/s Ethernet**
- **Mellanox MTS3600Q 36-Port 40Gb/s QDR InfiniBand switch**
- **Memory: 128GB memory per node DDR3 1333MHz**
- **OS: RHEL 5.5, MLNX-OFED 1.5.2 InfiniBand SW stack**
- **MPI: Open MPI 1.5.3 with KNEM 0.9.6, Platform MPI 8.1.1**
- **Compilers: PGI 10.9, GNU Compilers 4.1.2**
- **Application: AMBER 11 (PMEMD), AmberTools 1.5**
- **Benchmark workload:**
 - Cellulose_production_NVE_256_128_128 (408,609 atoms)



- **HPC Advisory Council Test-bed System**
- **New 11-node 528 core cluster - featuring Dell PowerEdge™ R815 servers**
 - Replacement system for Dell PowerEdge SC1435 (192 cores) cluster system following 2 years of rigorous benchmarking and product EOL
 - System to be redirected to explore HPC in the Cloud applications
- **Workload profiling and benchmarking**
 - Characterization for HPC and compute intense environments
 - Optimization for scale, sizing and configuration and workload performance
 - Test-bed Benchmarks
 - RFPs
 - Customers/Prospects, etc
 - ISV & Industry standard application characterization
 - Best practices & usage analysis



About Dell PowerEdge™ Platform Advantages

Best of breed technologies and partners

Combination of AMD Opteron™ 6000 Series platform and Mellanox ConnectX InfiniBand on Dell HPC. Solutions provide the ultimate platform for speed and scale

- Dell PowerEdge R815 system delivers 4 socket performance in dense 2U form factor
- Up to 48 core/32DIMMs per server – 1008 core in 42U enclosure

Integrated stacks designed to deliver the best price/performance/watt

- 2x more memory and processing power in half of the space
- Energy optimized low flow fans, improved power supplies and dual SD modules

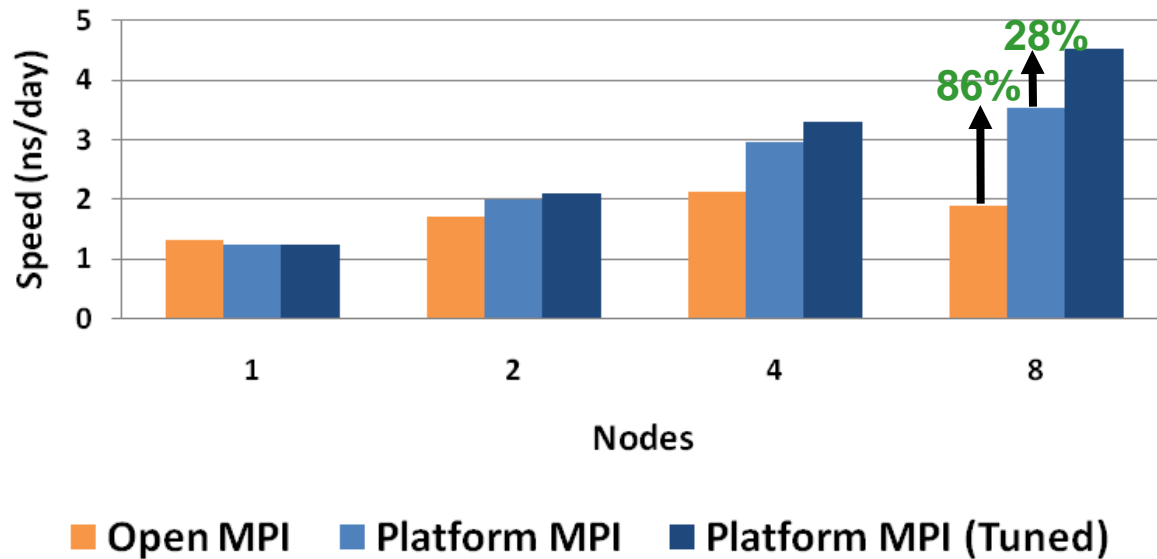
Optimized for long-term capital and operating investment protection

- System expansion
- Component upgrades and feature releases



- **Platform MPI enables better scalability than Open MPI**
 - Seen up to 86% better performance with Platform MPI
- **Tuned Platform MPI provides better performance for InfiniBand**
 - Seen an speed improvement of up to 28% at 8-node
 - Tuning parameters used: `-cpu_bind -e MPI_RDMA_MSGSIZE=65536,65536,4194304 -e MPI_RDMA_NSRQRECV=2048 -e MPI_RDMA_NFRAGMENT=128`

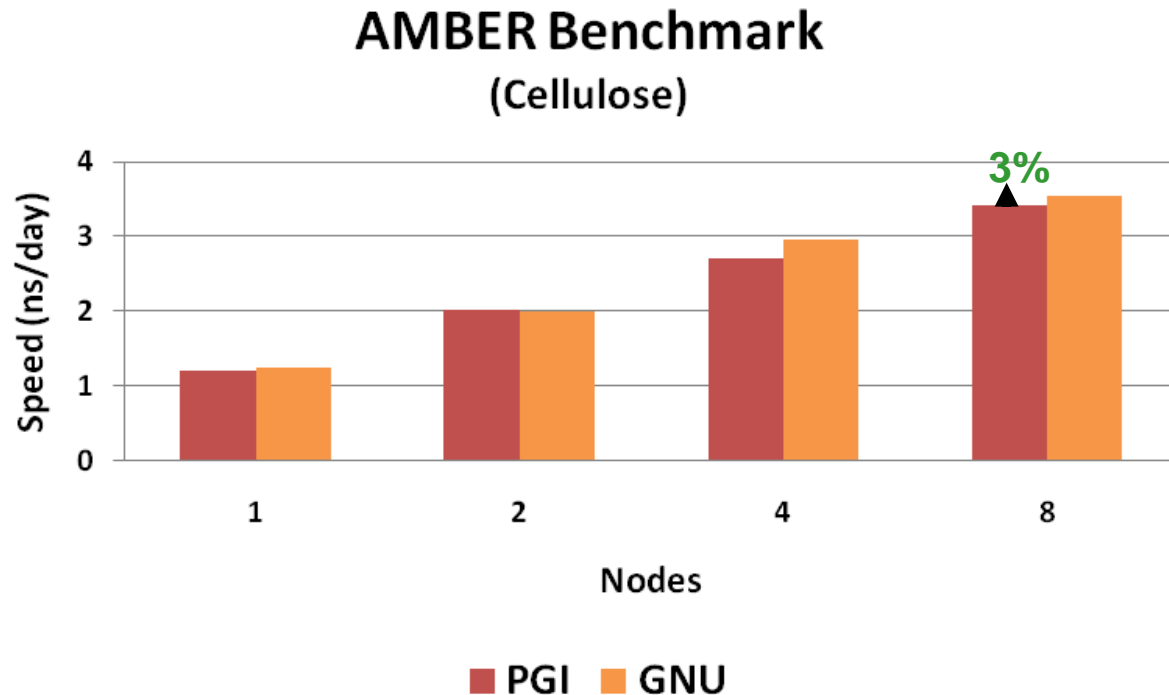
AMBER Benchmark (Cellulose)



Higher is better

48 Cores/Node

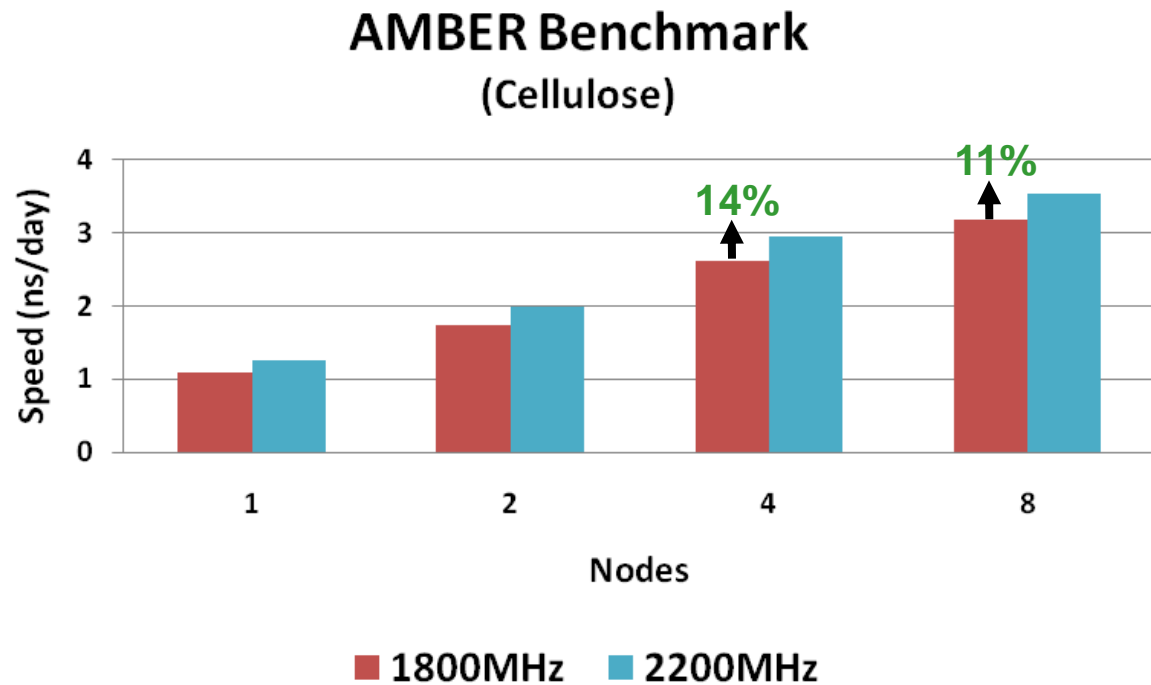
- Executable generated by the GNU compilers runs slightly faster
 - Up to 3% faster is seen than with PGI
- Using the default optimization and linker flags:
 - PGI: “-fast -O3 -fastsse”
 - GNU: “-O3”



Higher is better

*Platform MPI
48 Cores/Node*

- **Higher CPU core frequency enables higher job performance**
 - Up to 11-14% better job performance between 2200MHz vs 1800MHz
 - Shows the performance is somewhat affected by the change in the CPU frequency, and the percentage of Computation versus MPI Communication



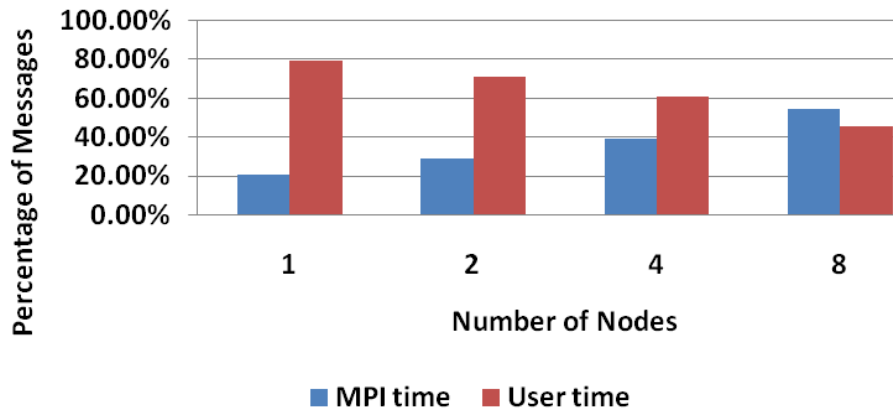
Higher is better

48 Cores/Node

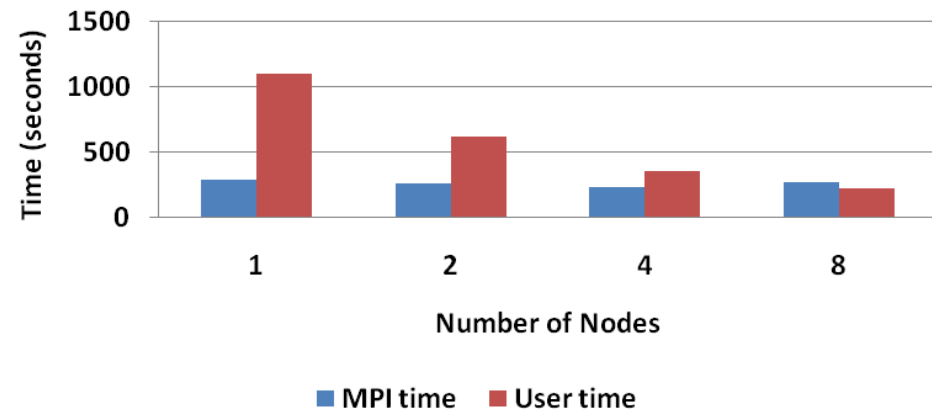
AMBER Profiling – MPI/User Time Ratio

- **Large data communications are seen**
 - Shows heavy data communications between parallel tasks
 - More communications than computation happens between 4 to 8 node
- **MPI time stays constant while CPU time reduces as more nodes in cluster**
 - Demonstrate the importance of the interconnect to handle addition network throughput

AMBER Profiling
(Cellulose)
MPI/User Time Ratio



AMBER Profiling
(Cellulose)
MPI/User Time



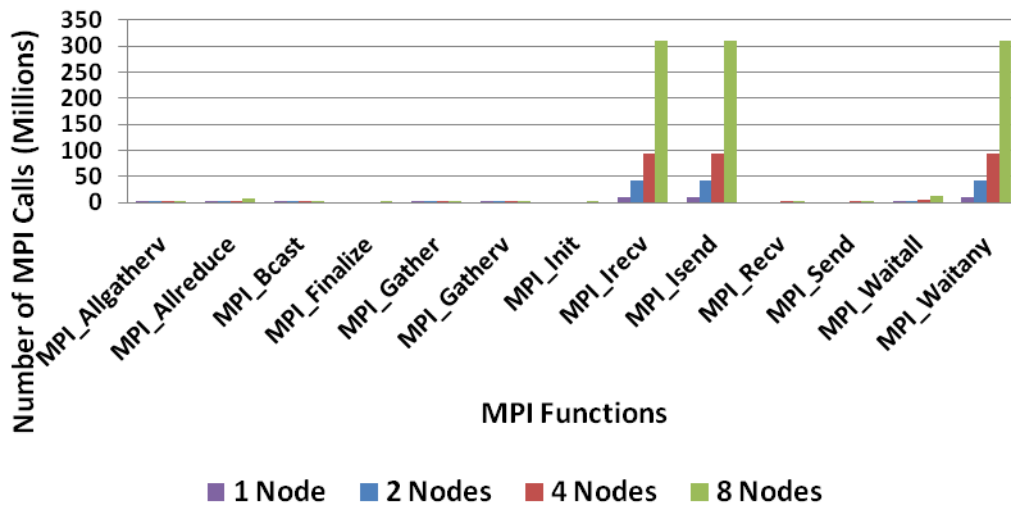
InfiniBand QDR

48 Cores/Node

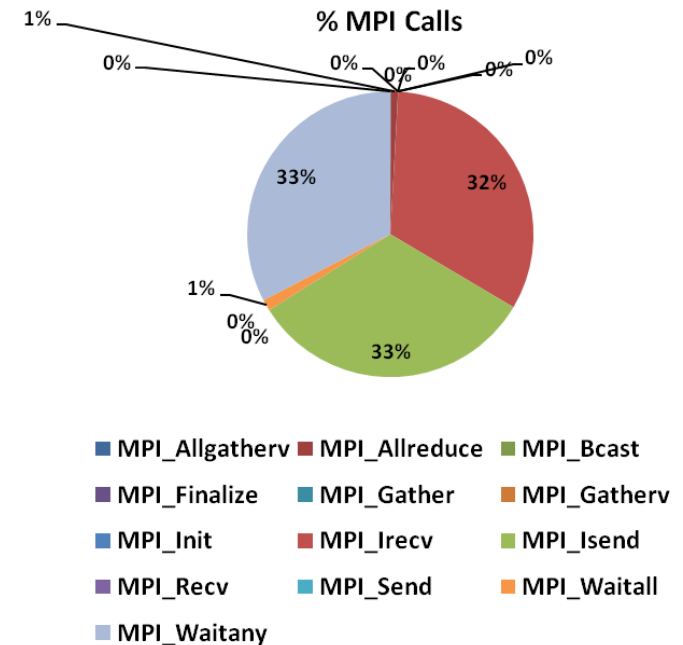
AMBER Profiling – Number of MPI Calls

- **The most used MPI functions are MPI_Isend, MPI_Irecv, MPI_Waitany**
 - These non-blocking MPI calls allows computation while communications take place
 - Each of these dominates as a third of MPI calls used on a 8-node job
- **The number of calls accelerates rapidly as the cluster scales**

AMBER Profiling
(Cellulose)
Number of MPI Calls

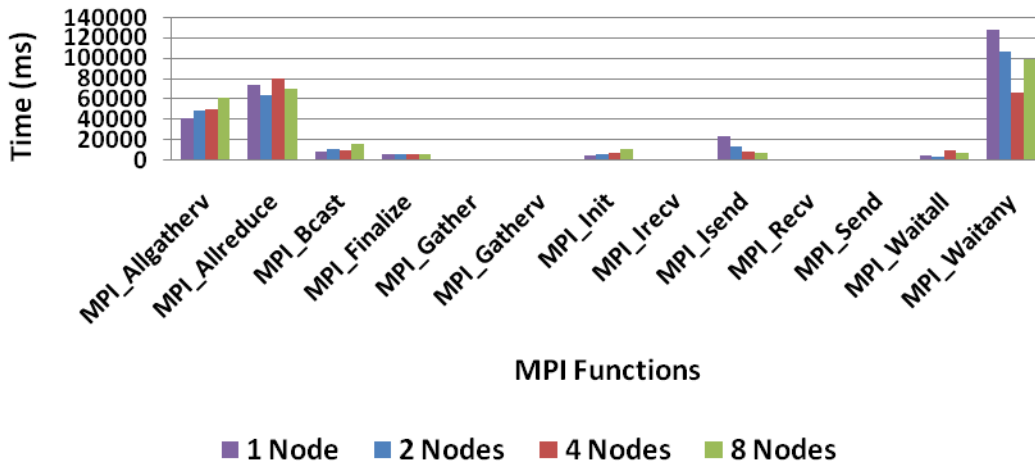


AMBER Profiling
(Cellulose, 8-node, InfiniBand)
% MPI Calls

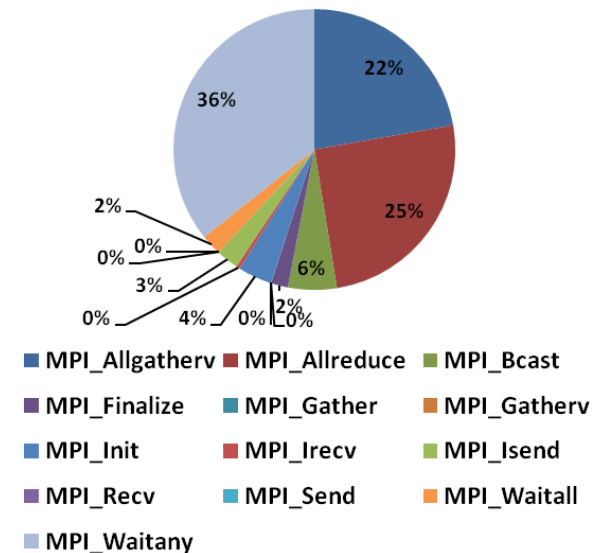


- The time in communications is taken place in the following MPI functions:
 - MPI_Waitany (36%)
 - MPI_Allreduce (25%)
 - MPI_Allgather (22%)
 - MPI_Bcast (6%)

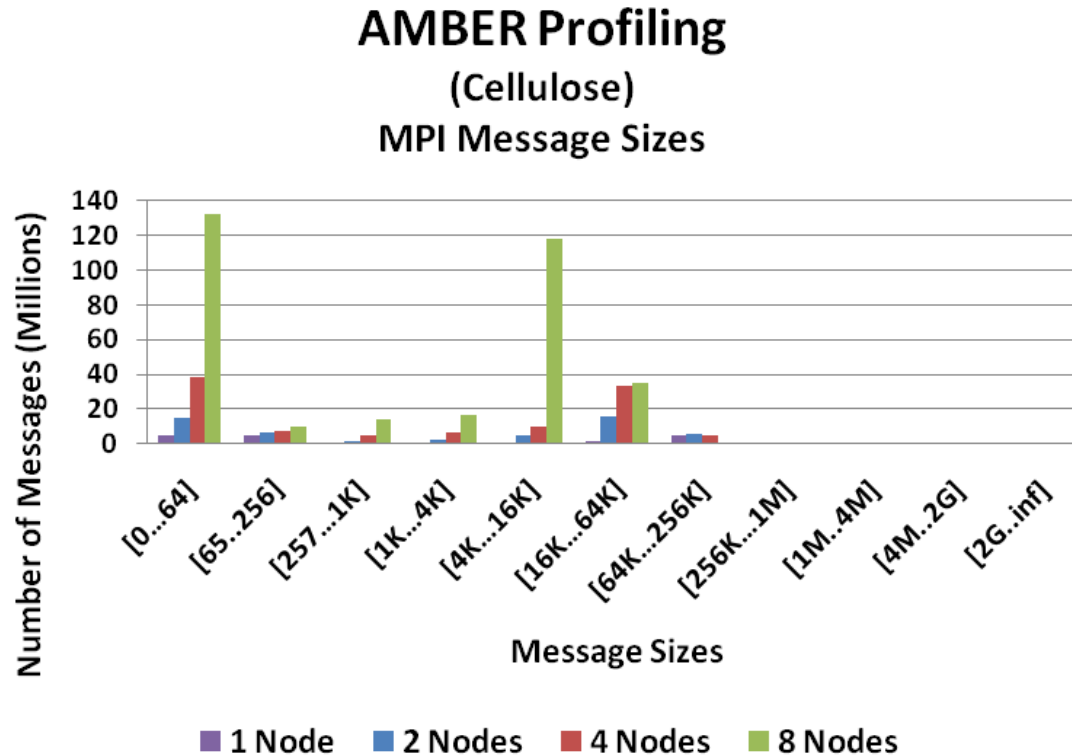
AMBER Profiling
(Cellulose)
Time Spent of MPI Calls



AMBER Profiling
(Cellulose, 8-node)
% Time Spent of MPI Calls

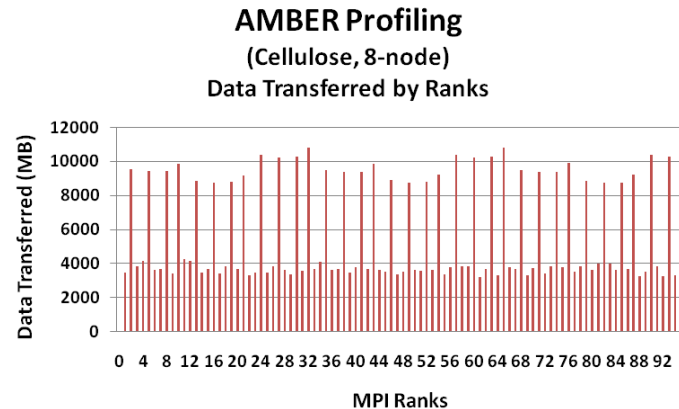
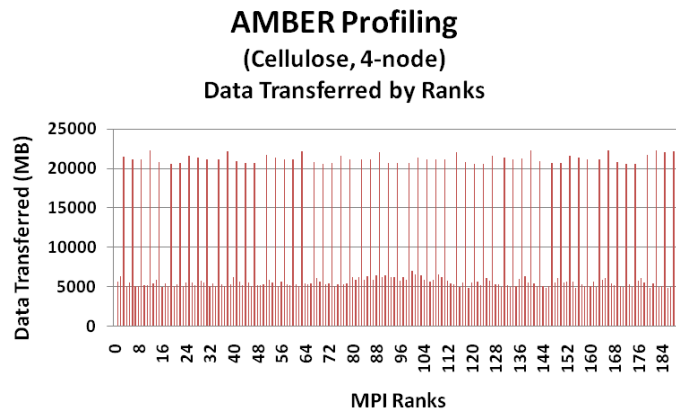
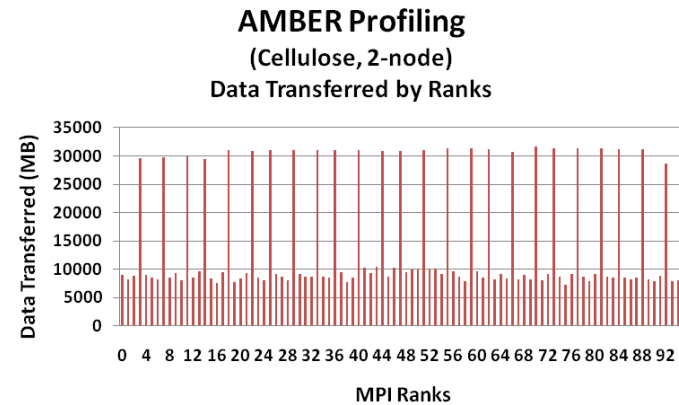
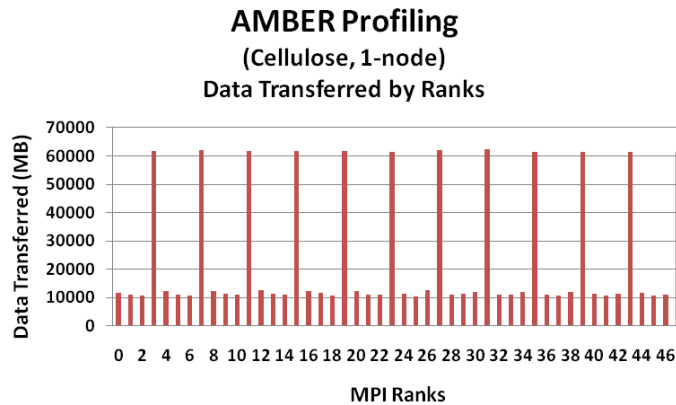


- **Majority of the MPI messages are small and median message sizes**
 - In the ranges of less than 64 bytes and between 4K and 16K

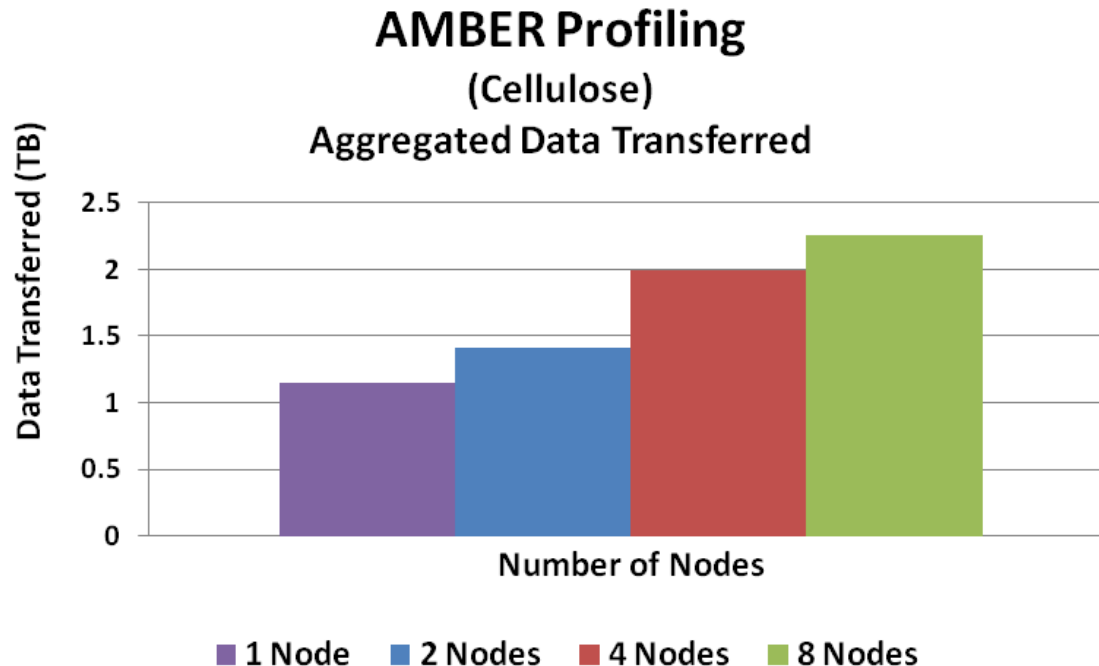


AMBER Profiling – Data Transfer By Process

- **Data transferred to each MPI rank is consistent for any number of processes**
 - Shows large amount of data transfers happened
 - Amount of data transfer to each rank is reduced as more nodes are in the job



- **Aggregated data transfer refers to:**
 - Total amount of data being transferred in the network between all MPI ranks collectively
- **The total data transfer increases steadily as the cluster scales**
 - Huge amount of data being sent and received across the network
 - As a compute node being added, more data communications will take place



InfiniBand QDR

- **AMBER is a compute and data communications intensive application**
 - Which has a high demand for CPU power and good network throughput
 - Using InfiniBand enables good scalability by spreading workload to compute nodes
- **MPI:**
 - Better scalability is seen with Platform MPI than with Open MPI
- **CPU:**
 - Shows higher job productivity when using CPU with higher core frequency
- **MPI Communications:**
 - Large data communications take place between MPI processes

Thank You

HPC Advisory Council



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