



Blue Gene/P

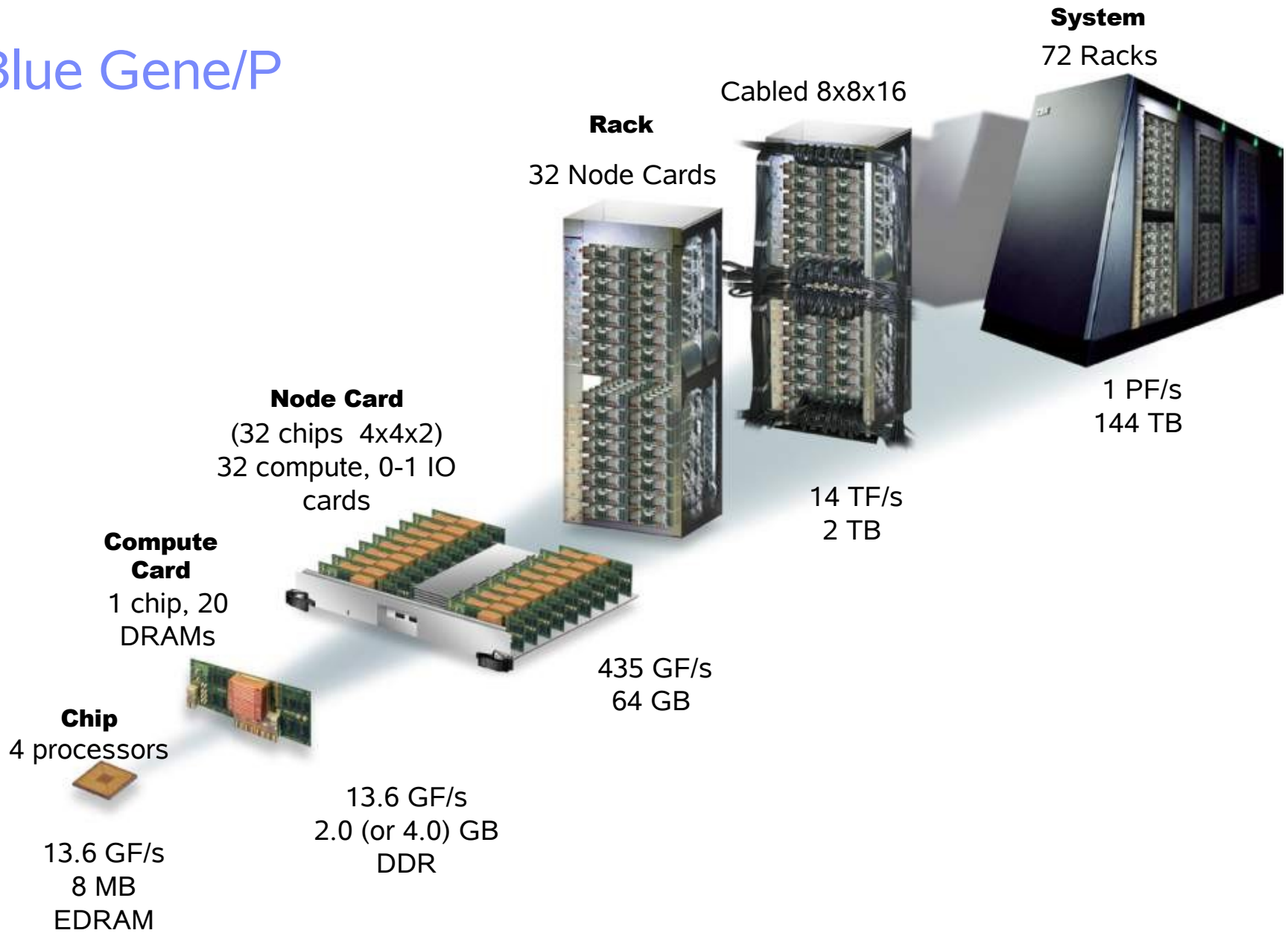
Porting Applications to Blue Gene/P

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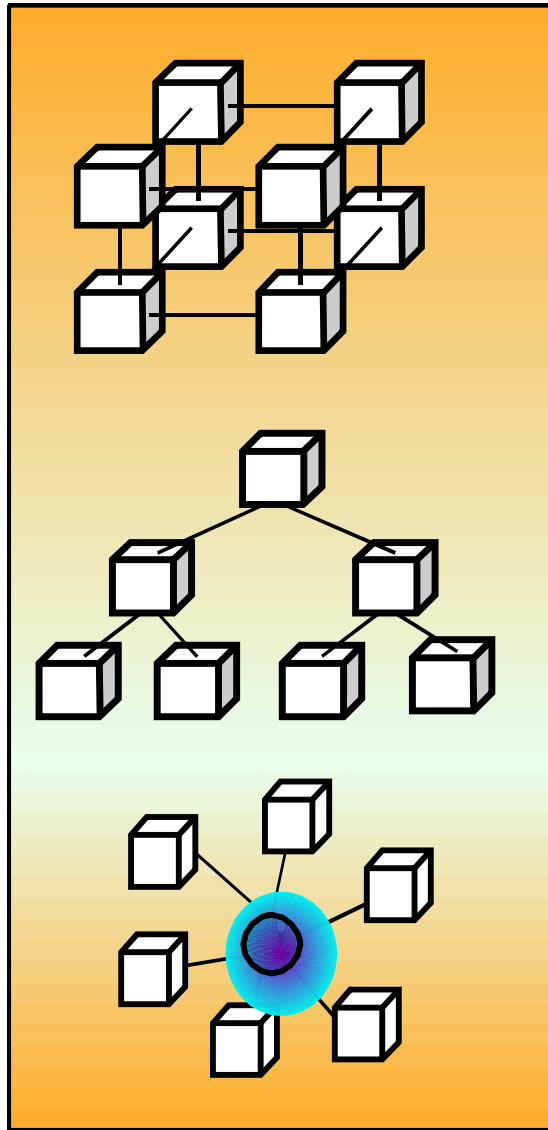
Agenda

- **What beast is this ?**
- Compile - link – go !
- MPI subtleties
- Help ! It doesn't work
(the way I want) !

Blue Gene/P



Blue Gene/P Spider Webs



3 Dimensional Torus

- Interconnects all compute nodes (73,728)
- Virtual cut-through hardware routing
- 3.4 Gb/s on all 12 node links (5.1 GB/s per node)
- 0.5 μ s latency between nearest neighbors, 5 μ s to the farthest
- MPI: 3 μ s latency for one hop, 10 μ s to the farthest
- Communications backbone for computations
- 1.7/3.9 TB/s bisection bandwidth, 188TB/s total bandwidth

Collective Network

- One-to-all broadcast functionality
- Reduction operations functionality
- 6.8 Gb/s of bandwidth per link
- Latency of one way tree traversal 1.3 μ s, MPI 5 μ s
- ~62TB/s total binary tree bandwidth (72k machine)
- Interconnects all compute and I/O nodes (1152)

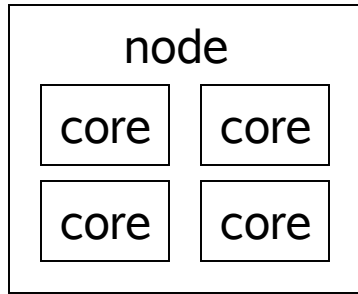
Low Latency Global Barrier and Interrupt

- Latency of one way to reach all 72K nodes 0.65 μ s, MPI 1.6 μ s

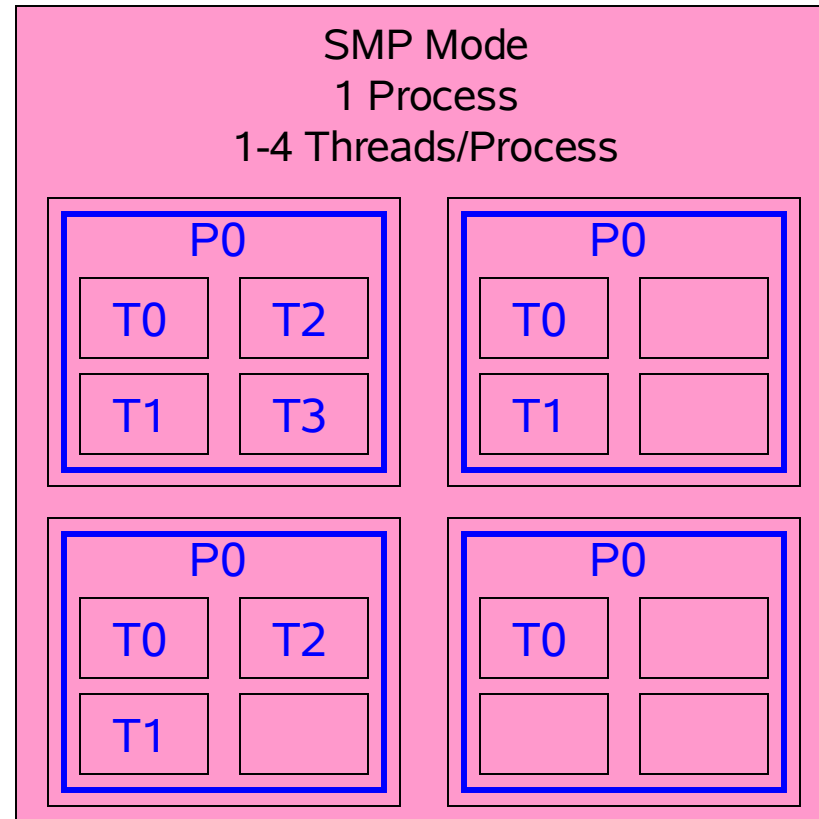
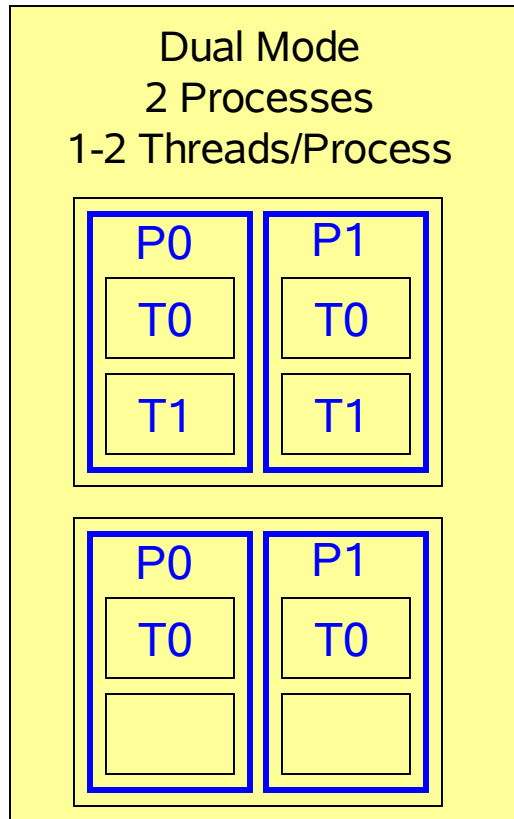
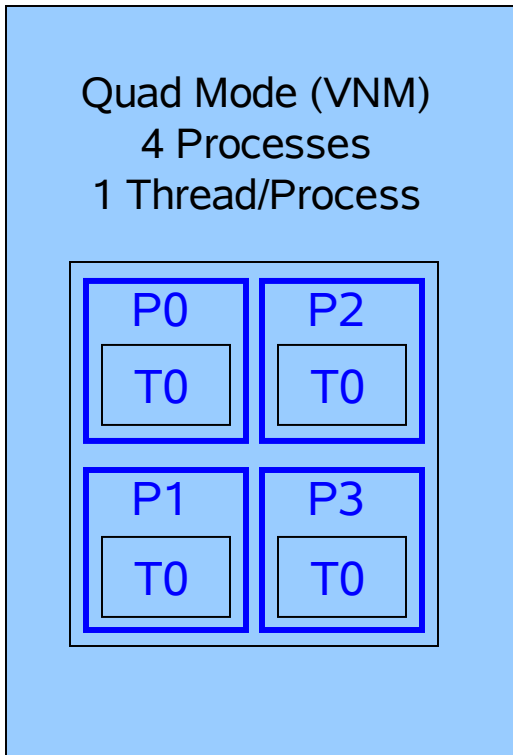
Other networks

- 10Gb Functional Ethernet
- I/O nodes only
- 1Gb Private Control Ethernet
- Provides JTAG access to hardware. Accessible only from Service Node system

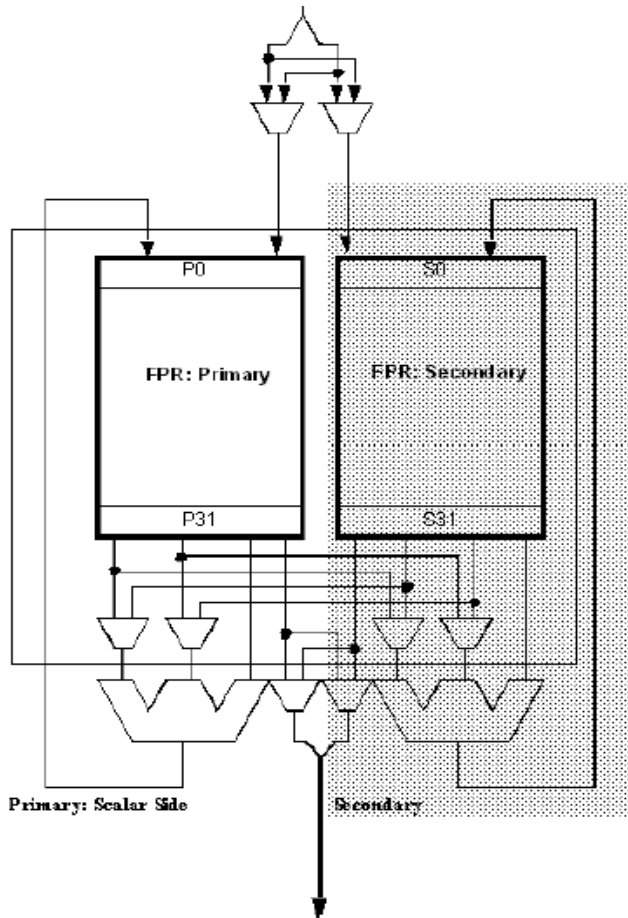
You have the choice !



Software Abstractions Blue



Dual FPU Architecture



- SIMD instructions over both register files
- FMA operations over double precision data
- Parallel (quadword) loads/stores
- Data needs to be 16-byte aligned

Caches

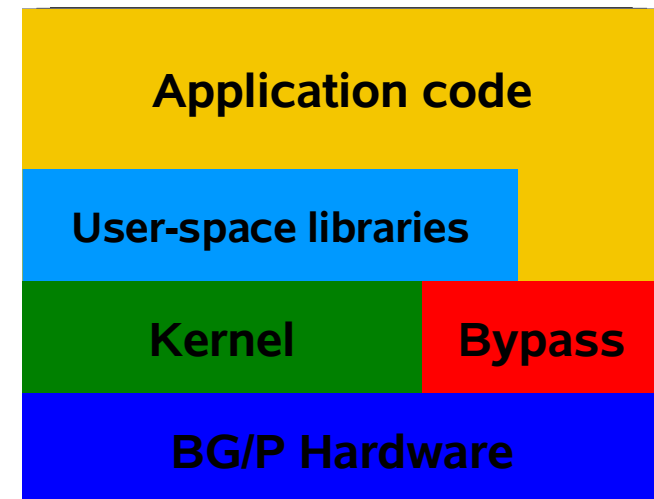
Cache	Total per node	Size	Replacement Policy	Associativity
L1 Instruction	4	32 KB	Round-Robin	64-way set-associative 16 sets 32B line size
L1 Data	4	32 KB	Round-Robin	64-way set-associative 16 sets 32B line size
L2 PreFetch	4	14x256 B	Round-Robin	Fully associative (15-way)128 B Line size
L3	2	2x4 MB	Least Recently Used	8way associative 2 Bank Interleaved 128 B Line

Jitter-free Execution

- **Compute node runs nothing but application**
- **I/O delegated to I/O nodes**
- **Cross-Compiling on the front end node**

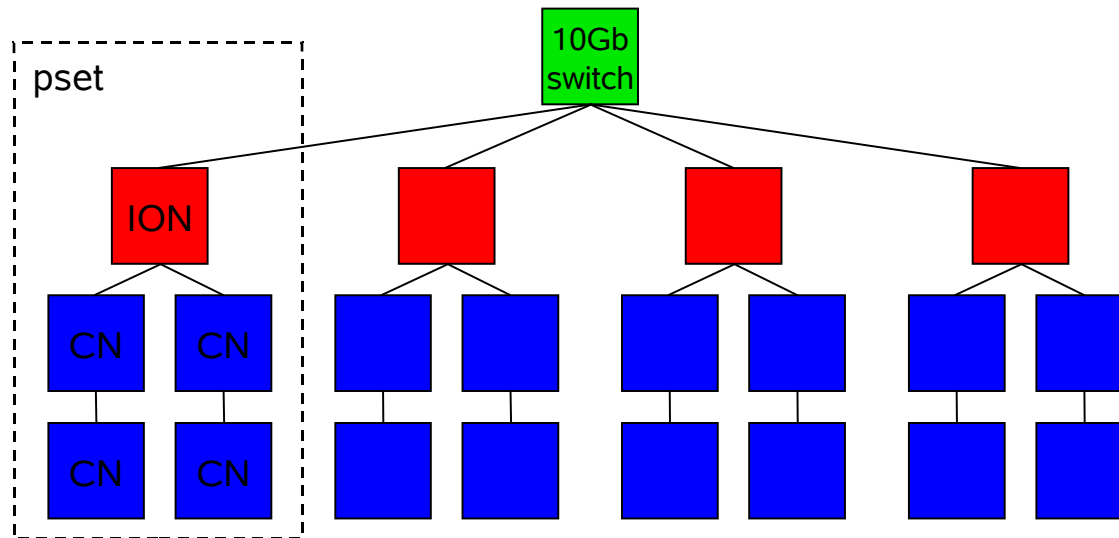
Software Stack in Blue Gene Compute Node

- Compute Node Kernel (CNK) controls access to hardware, and enables bypass for application use
- User-space libraries and applications can directly access torus and collective network through bypass
- Application code can use all processors in a compute node



Processing Sets (Psets)

- I/O node dedicated to a fixed group of compute nodes
- Compute to I/O ratio is fixed within a partition
 - 128:1, 64:1, 32:1, 16:1



I/O Node Kernel



- **SMP Linux**
- **No persistent store (network filesystems only; no swap)**
- **10Gb Ethernet interface**
- **Several CNK System calls are function shipped to here**
 - Linux compatibility by executing these syscalls on Linux
 - Function ship occurs over Collective network
 - The `ciod` daemon manages a fixed set of compute nodes in a `processing set` (pset)
 - Linux provides the portable filesystem and network layer interfaces

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Getting started is simple

- **...for simple cases**
- **BGP_SYS=/bgsys/drivers/ppcfloor**
- **make CC=\$BGP_SYS/comm/bin/mpixlc_r**
- **lrun -mode VN -np 512 ./hello_par**

IBM XL Compilers for Blue Gene

- **XLF 11.1/VACPP 9.0 will be the compiler releases**
 - /opt/ibmcmp/xlf/bg/11.1/bin
 - /opt/ibmcmp/vacpp/bg/9.0/bin
- **Differences in this release:**
 - xlf2003 (the 2003 Fortran standard) is available
 - BGP wrapper names are different
 - blrts_ is replaced by bg
 - bgxlf, bgxlc, bgcc, etc.
 - On BG/L for xlf 11.1/vacpp 9.0 both blrts_ and bg will be supported.
 - -qarch=450d/450 are accepted in addition to 440d/440

Some key options for IBM compilers

- **-qarch=440, 450** generates only instructions for one floating point (option minimal option with blrts_)
- **-qarch=440d, 450d** generates only instructions for 2 floating point pipes
- **-qtune=440**
- **-O3 (-qstrict)** minimal level for SIMDization
- **-O3 -qhot (=simd)**
- **-O4 (-qnoipa)**
- **-O5**
- **-qdebug=diagnostic** provide details about SIMDization, only with **-qhot**
- **-qreport -qlist -qsource** provide pseudo-assembler code .lst

What's new from BG/L ...

- **pthread and OpenMP support**
- **Dynamic linking**
- **Use of mmap for shared memory**
- **Protected readonly data and application code**
- **Protection for stack overflow**
- **Full socket support (client and server)**

ESSL for Blue Gene

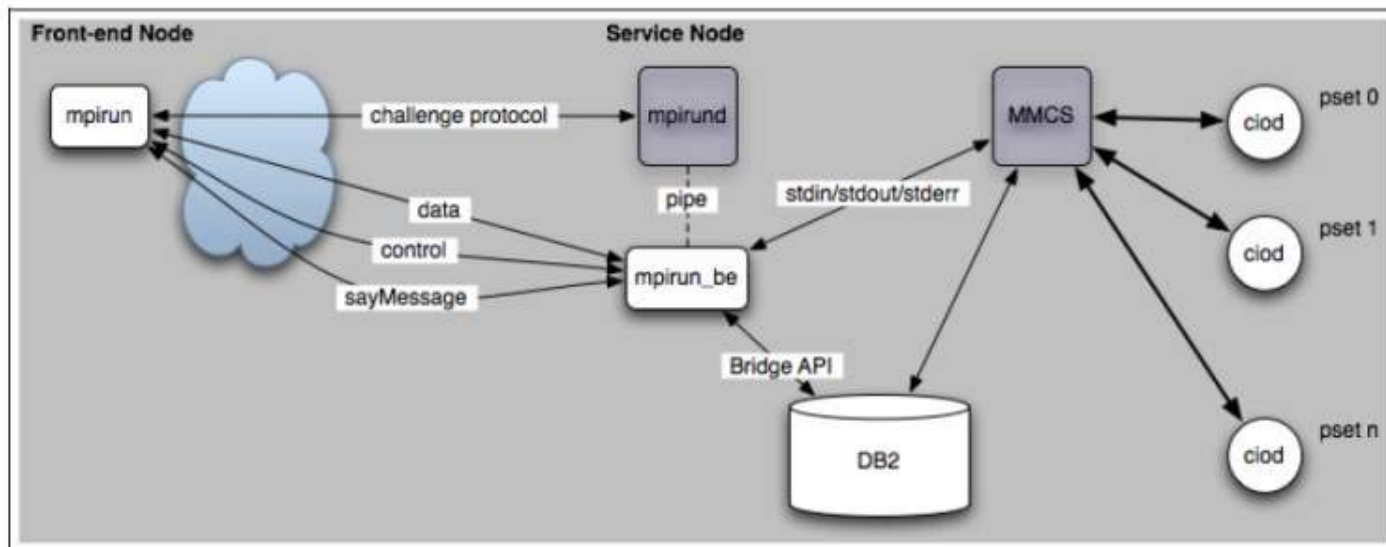
- **Engineering and Scientific Subroutine Library**
- **Optimization library and intrinsics for better application performance**
- **Serial Static Library supporting 32-bit applications**
- **Callable from FORTRAN, C, and C++**
- **SMP support and ppc450 tuning done for BG/P**
- **libesslbg.a (.so) and libesslsmpbg.a (.so)**

Lib Mass for Blue Gene

- **Mathematical Acceleration Subsystem (MASS) libraries) consists of libraries of tuned mathematical intrinsic functions**
- **Location:**
 - /opt/ibmcmp/xlmass/bg/4.4/bglib
 - libmass.a libmassv.a
 - /opt/ibmcmp/xlmass/bg/4.4/include

MPIRUN implementation on BGP

- no rsh/ssh mechanism
- Option -free
- STDIN handling



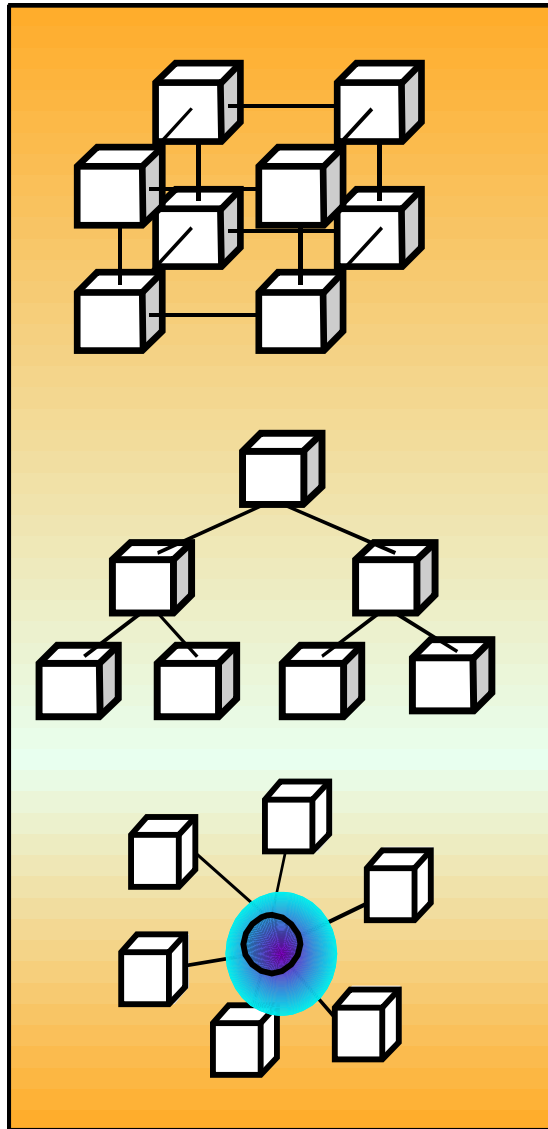
Partitioning

- **Subdivision of a single Blue Gene system**
- **Partitions are software defined**
- **Torus, Collective and Barrier networks are completely isolated from traffic from other partitions**
- **A single job runs on a partition**
 - i.e. jobs never share resources or interfere with each other
- **Custom kernels may be booted in a partition**

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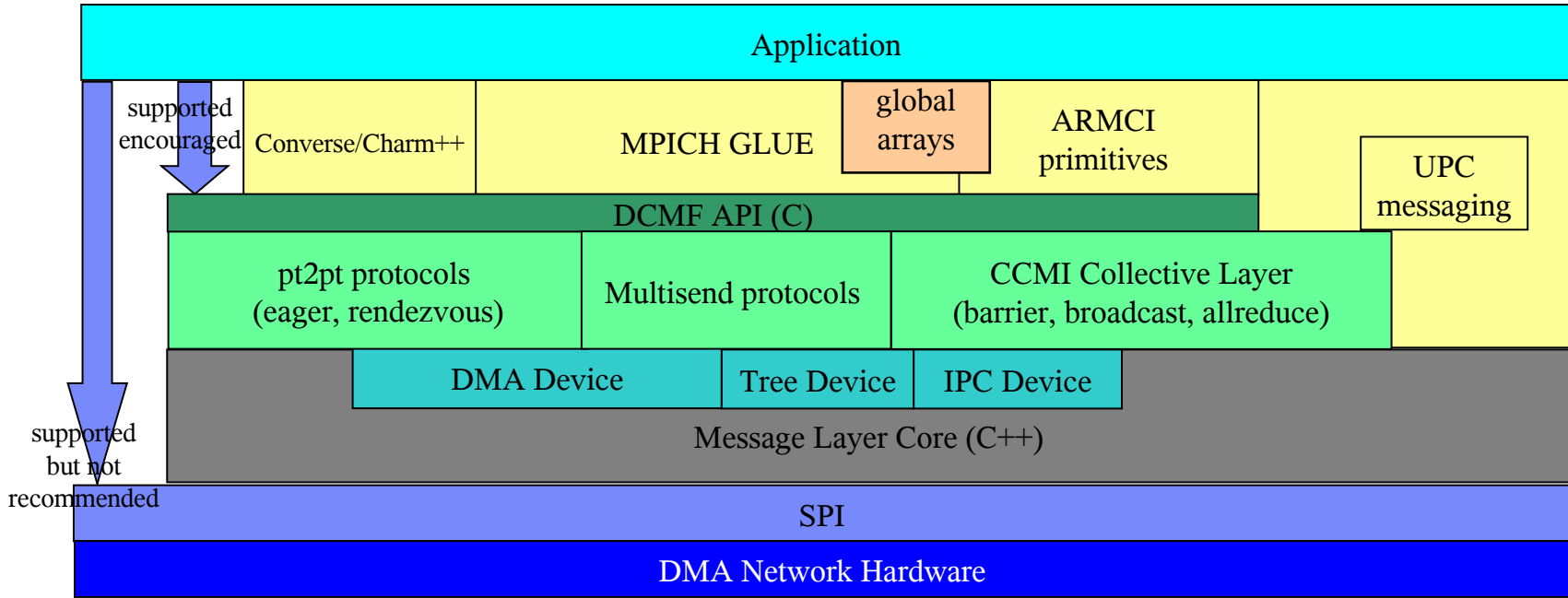
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Collectives

- **Use the hardware support in the collective network and global interrupt networks**
- **Supported operations**
 - Barrier
 - Broadcast
 - Allreduce
 - Alltoall
 - Allgather

Messaging Framework



Multiple programming paradigms supported

MPI, Charm++, ARMCI, GA, UPC (as a research initiative)

SPI : Low level systems programming interface

DCMF : Portable active-message API

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Debugger Interfaces

- **ptrace-like interfaces available via ciod**
 - non-parallel: gdbserver for direct use with gdb
 - parallel: Totalview, or other tools
- **lightweight core files**
 - each node writes a small file with regs, traceback, etc
 - superset of parallel tools consortium format
 - Use addr2line for translating HEX into source lines
- **Coreprocessor**

GNU Debugger

- Simple debug server call «gdbserver »
- Only one gdb instance for one compute node (to debug multiple CNs at the same time you need to launch multiple GDB clients)
- Limited subset of primitives (however enough to be useful)
- Standard Linux gdb client, not aware about Double FPU.
- Gdserver must start before the application; mpirun has a special option «-start_gdbserver »
- Compile and link with `-g (-O2)`
- Location: `/bgsys/drivers/ppcfloor/gnu-linux/bin`

Performance Tools

- **Low level SPI provided to configure, reset and read hardware perf counters**
- **PAPI interface to the perf counters**
- **HPC Toolkit**
- **Considering addition of per-job performance metrics recorded via job history**
- **Unix gprof command (compiler with `-g -pg`)**

IBM High Performance Computing (HPC) Toolkit

- **Message-passing performance**
 - MP_Profiler (MPI and SHMEM)
 - MP_Trace (MPI and SHMEM)
 - SHMEM Library (Cray API) – AIX only
- **CPU performance**
 - Xprofiler
 - HPM (Hardware counters)
- **Thread performance**
 - Pomp Profiler (OpenMP)
- **Memory performance**
 - SiGMA memory profiler
 - Prediction Assistant
- **Visualization and analysis**
 - PeekPerf

More Information

- **IBM Redbooks for Blue Gene**
 - Application Development Guide
 - System Administration Guide
 - Performance Tools
- **Open Source Communities (Argonne website, ...)**
- **BlueGene Rochester website**
 - <http://bgweb.rchland.ibm.com/~jratt/>
- ***Doxygen* documentations (DCMF, SPI, ...)**