

Advancing Atmospheric Science with Blue Gene/L

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Summary of remarks...

- Blue Gene/L seems to us to be an important architecture for many reasons.
 - Power/space & “fuel” efficiency
 - \$/Tflops sustained
 - Fast reduction network
- 80 km, 20 level explicit HOMME model
 - We see 587 Gflops on 1944 nodes of BG/L.
 - 13.7 simulated years/day (useful climate rate)
- 1/10 degree “eddy permitting” POP ocean model would be the next interesting thing to look at. (barotropic CG needs fast global sums)



BlueGene/L Architecture



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Why Blue Gene/L is Attractive

•Pros

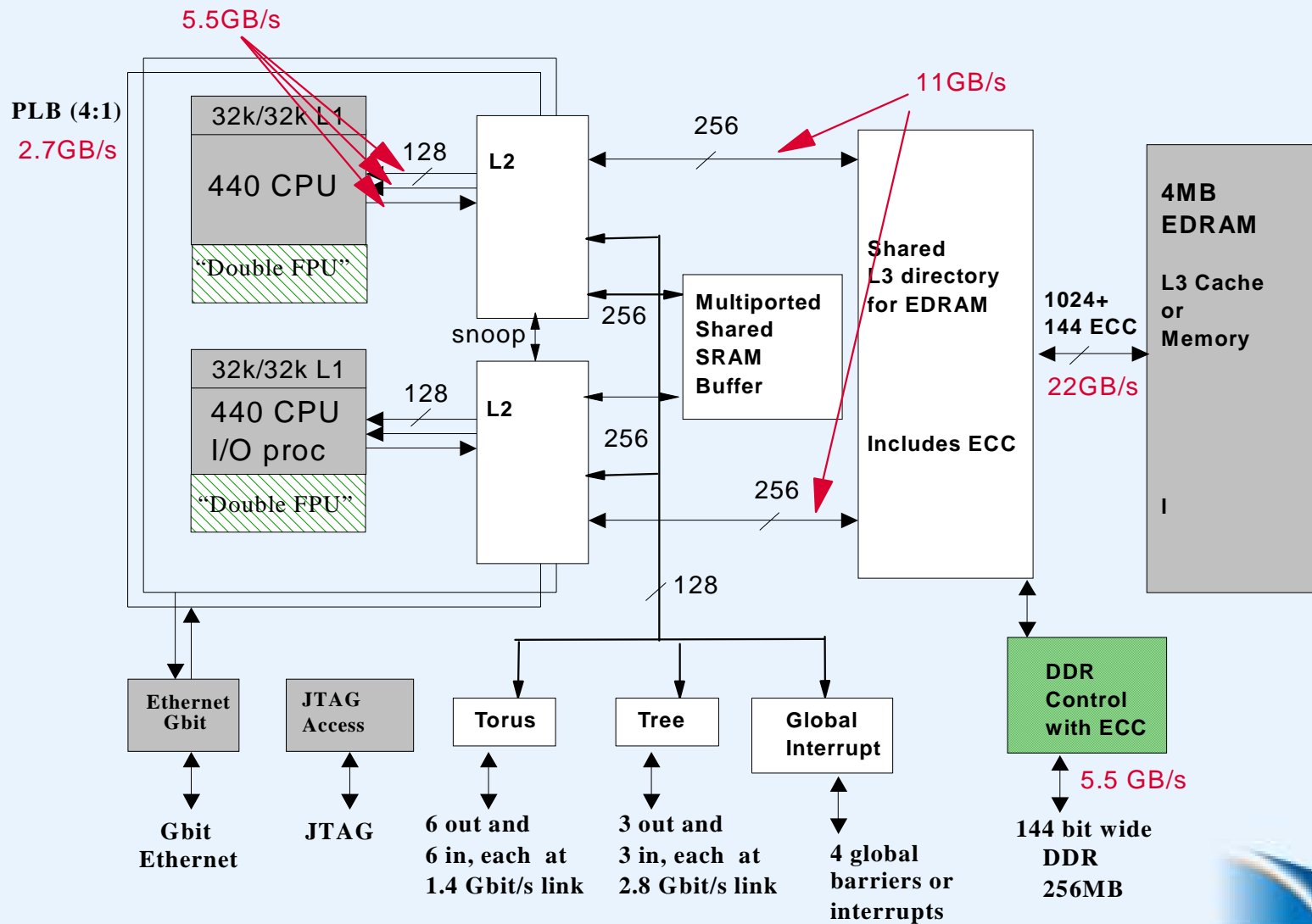
- Achieves high packaging density.
- Lower power density.
- Dedicated reduction network.
- Puts network interfaces on chip.
- Low cost per sustained FLOPs.

•Questions

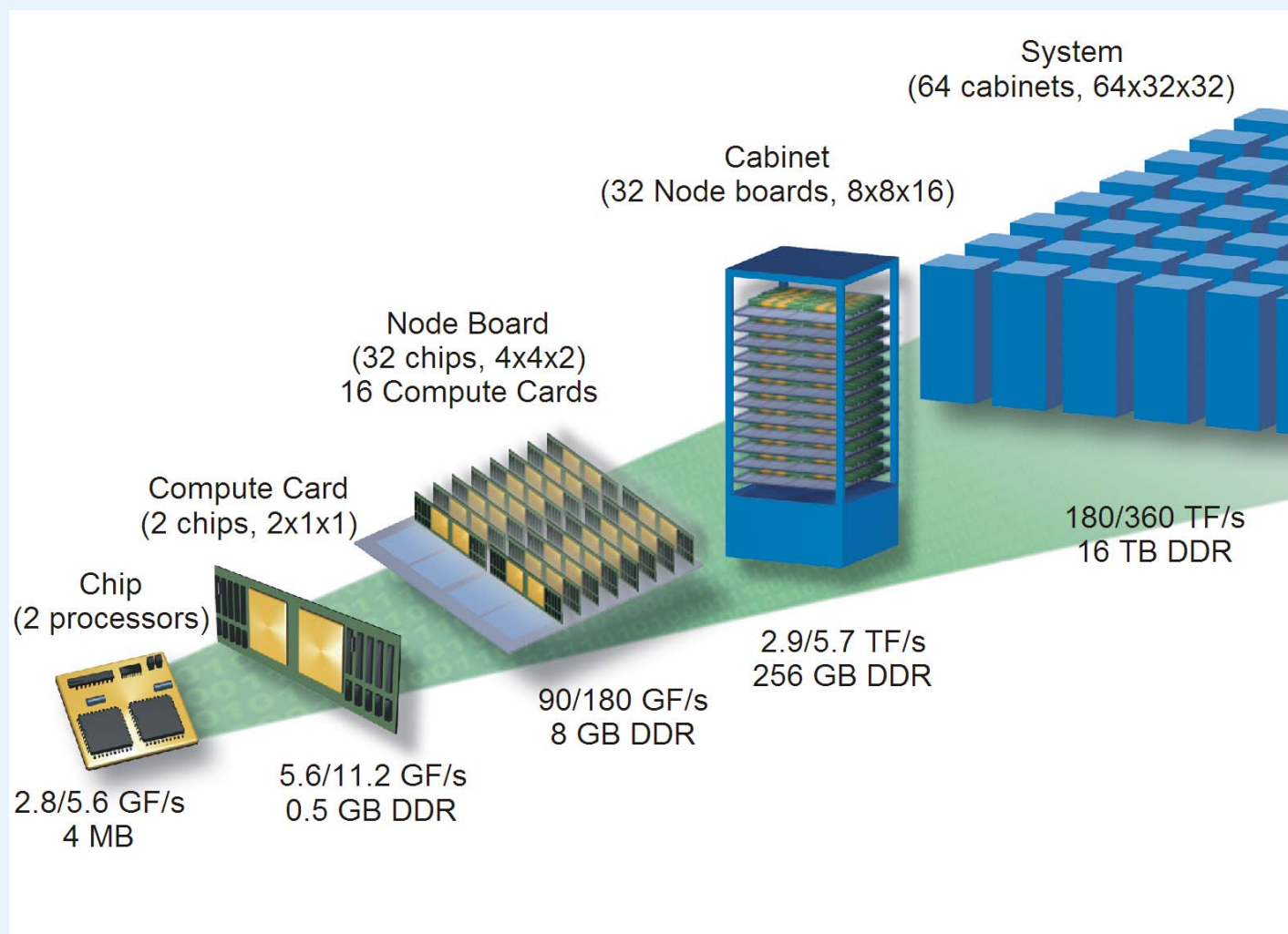
- High reliability?
- Applications for 100k processors?
- System robustness: I/O, scheduling flexibility.



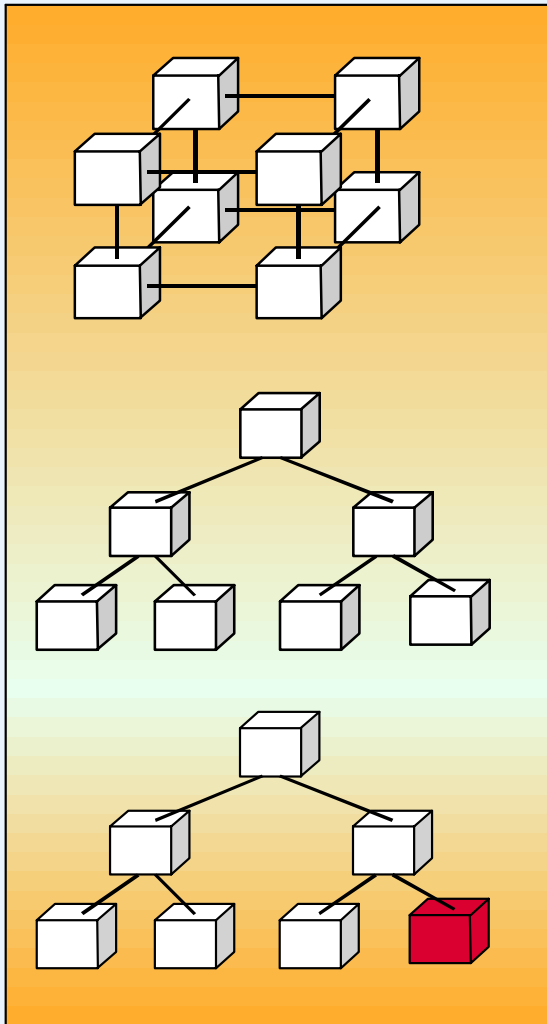
BlueGene/L ASIC



The Blue Gene/L Architecture



BlueGene/L Has Five Networks



3-Dimensional Torus

- interconnects all compute nodes (65,536)

Global Tree

- point-to-point, one-to-all broadcast, reduction functionality

Global Interrupts

- AND/OR operations for global barriers
- 1.5 microseconds latency (64K system)

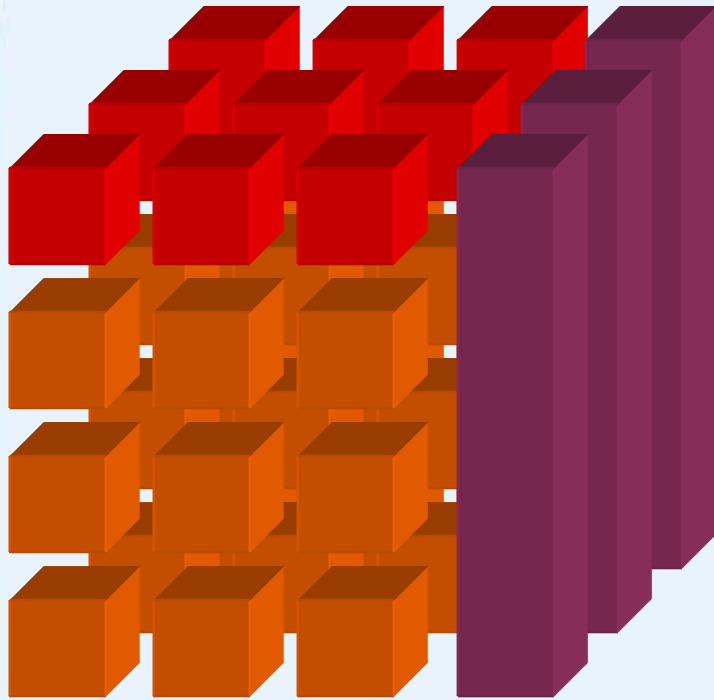
Ethernet

- incorporated into every node ASIC
- active in the I/O nodes (1:64 in LLNL configuration)
 - 1K 1Gbit links
- all external comm. (file I/O, control, user interaction, etc.)

JTAG (Control)



BlueGene/L System Software Architecture



- User applications execute exclusively in the **compute nodes**
 - avoid asynchronous events (e.g., daemons, interrupts)
- The outside world interacts only with the **I/O nodes**, an offload engine
 - standard solution: Linux
- Machine monitoring and control also offloaded to **service nodes**: large SP system or Linux cluster.



Blue Gene/L system overview

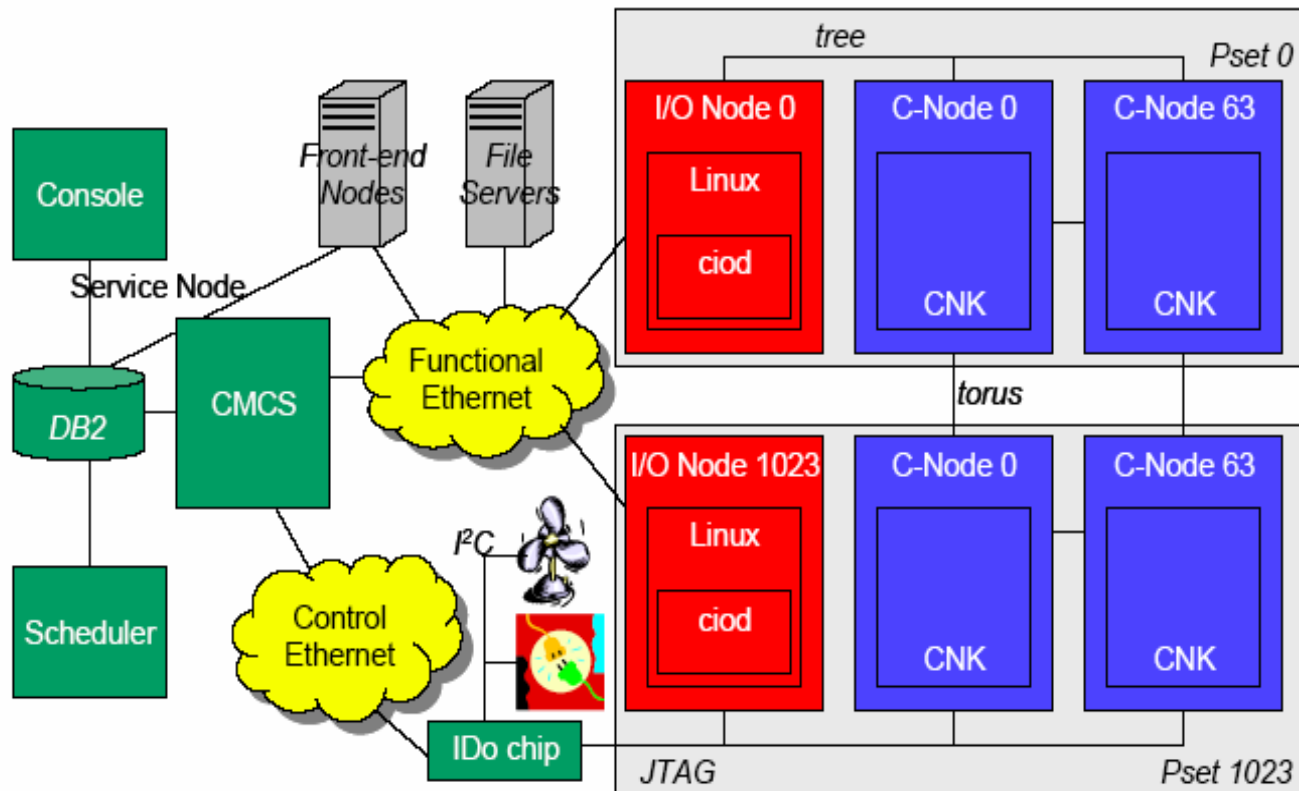


Figure 1: Overview of a complete system with BlueGene/L as the computational core.

Scalable atmospheric dynamics...



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Description of HOMME

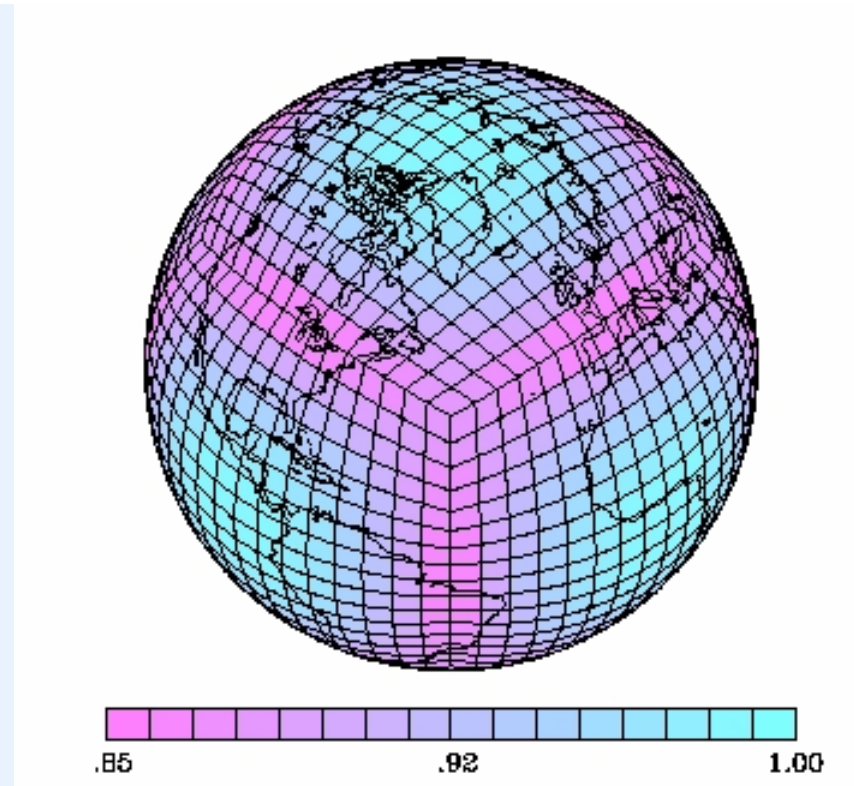
Atmospheric Dynamical Core

- **High Order Method Modeling Environment (HOMME)**
 - Solving the moist “primitive” equations on **cube-sphere**
 - CAM finite difference vertical discretization
 - Held-Suarez simplified test physics package
- **Algorithmic Advantages of High Order Methods**
 - h-p finite element method on quadrilaterals ($N_e \times N$)
 - Exponential convergence in p (N)
- **Computational Advantages of High Order Methods**
 - Naturally cache-blocked $N \times N$ computations
 - Nearest-neighbor communication between elements
 - Well suited to parallel μ processor systems
- **Semi-Implicit Time Step to eliminate fast gravity wave modes**
 - Crank-Nicholson time integration scheme
 - Use iterative CG solver for the resultant elliptic equation
 - Simple metric term/mass matrix preconditioner.
 - Acceleration 2.5x over explicit integration



The Cube-Sphere

- Equal angular grid
– Rancic et al (1996)
- Avoids pole problems
- Quasi-uniform
- Curvilinear coordinates: metric terms
- $N_e=16$ shown above

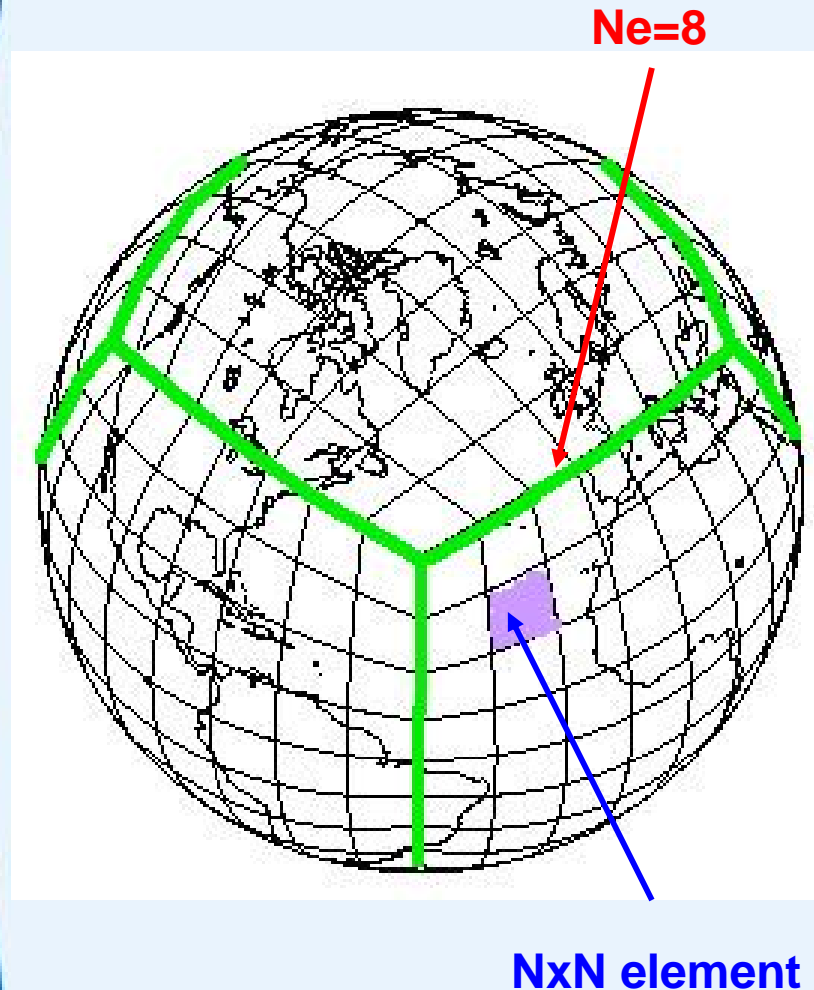


**$N_e=16$ Degree of
on-uniformity**

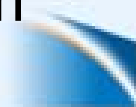


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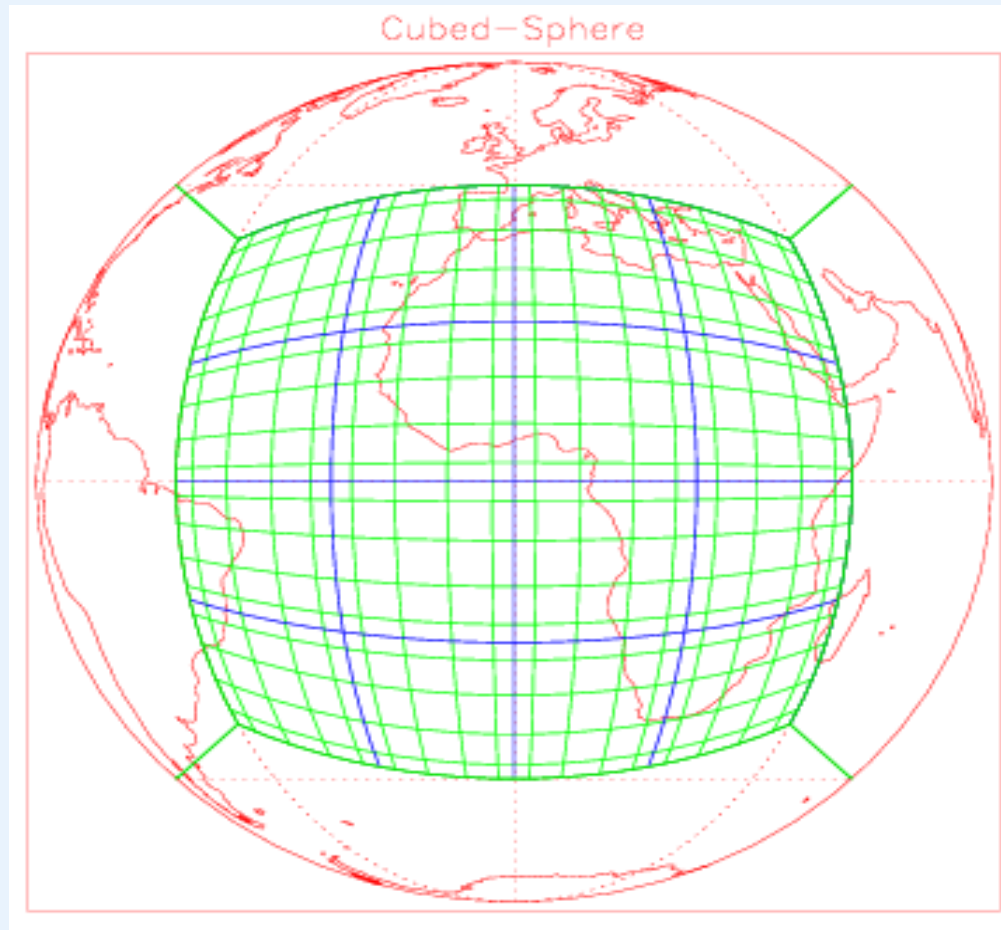
The Spectral Element Computational Mesh: the “Cube-Sphere”



- Spectral Elements:
 - A quadrilateral “patch” of gridpoints $N \times N$
 - Gauss-Lobatto Grid
 - $N=8$ is optimal (Taylor)
- Cube
 - N_e = Elements on an edge
 - $6 \cdot N_e \cdot N_e$ elements total
- Cube Partitioning
 - Metis
 - Space filling curve partitioning algorithm
- $N_e=8$ shown ~ 180 km

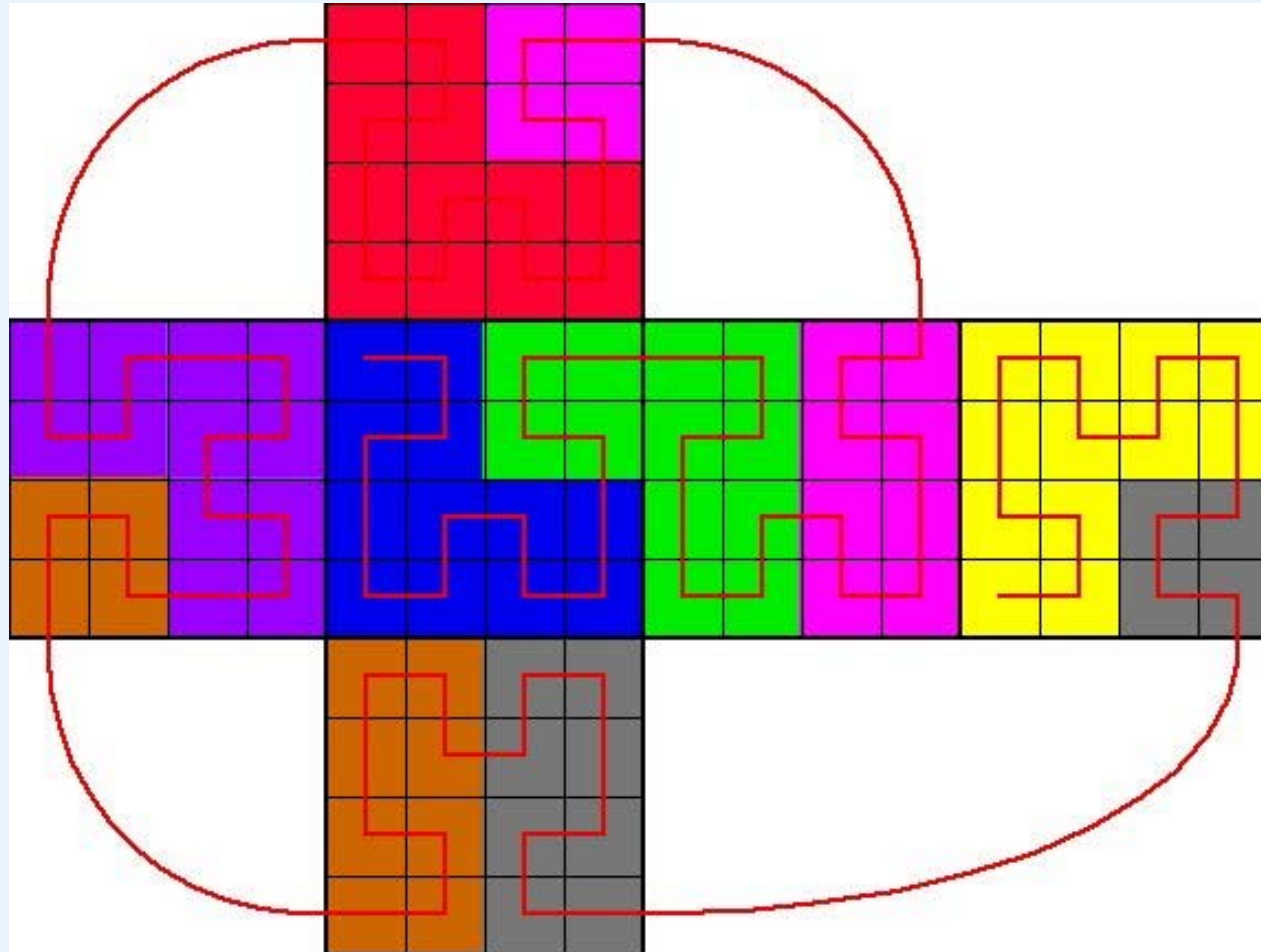


6th Order Spectral Elements on the Ne=4 Cube Sphere



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Partitioning a cubed-sphere on 8 processors

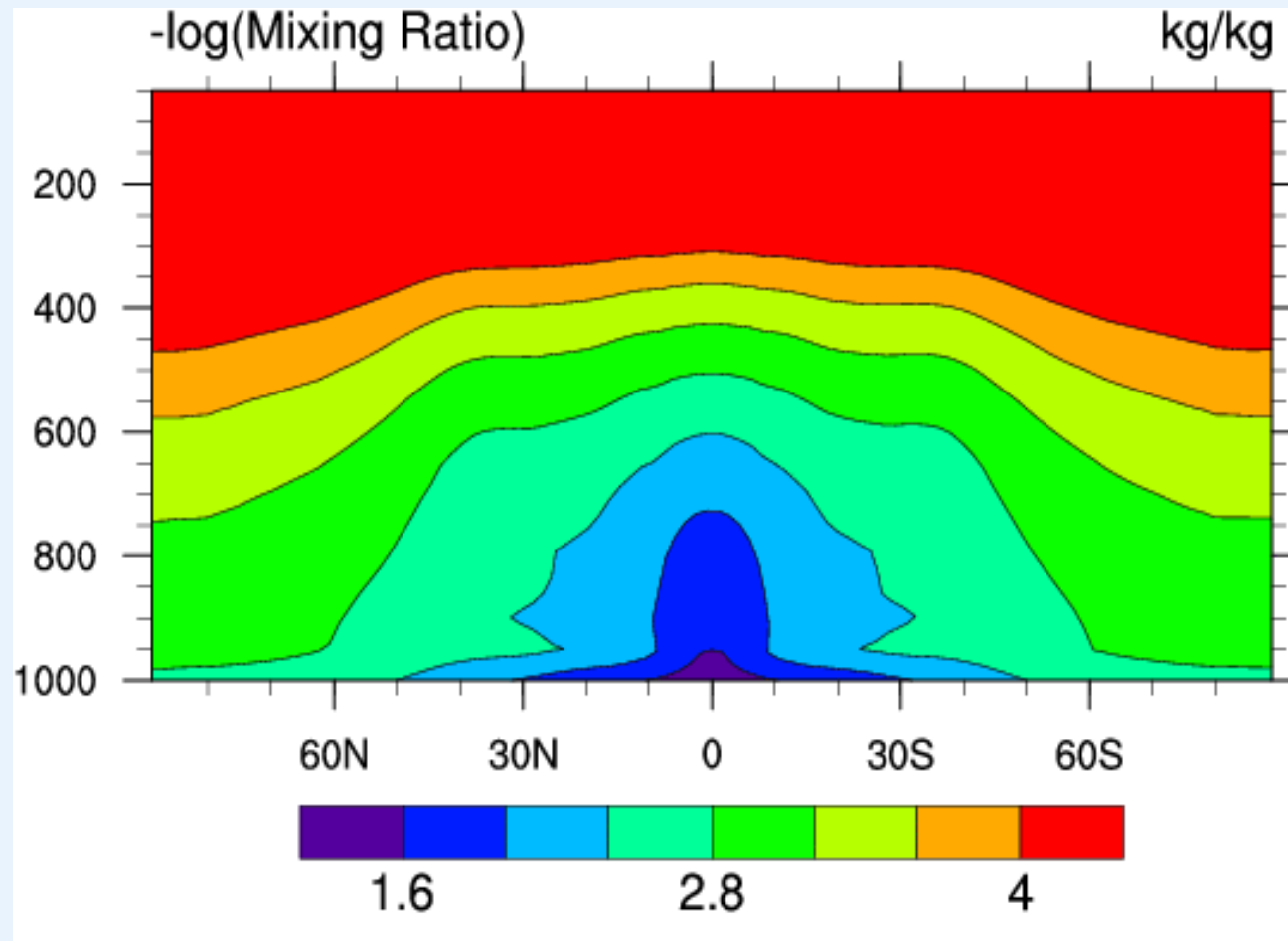


HOMME on Blue Gene/L...



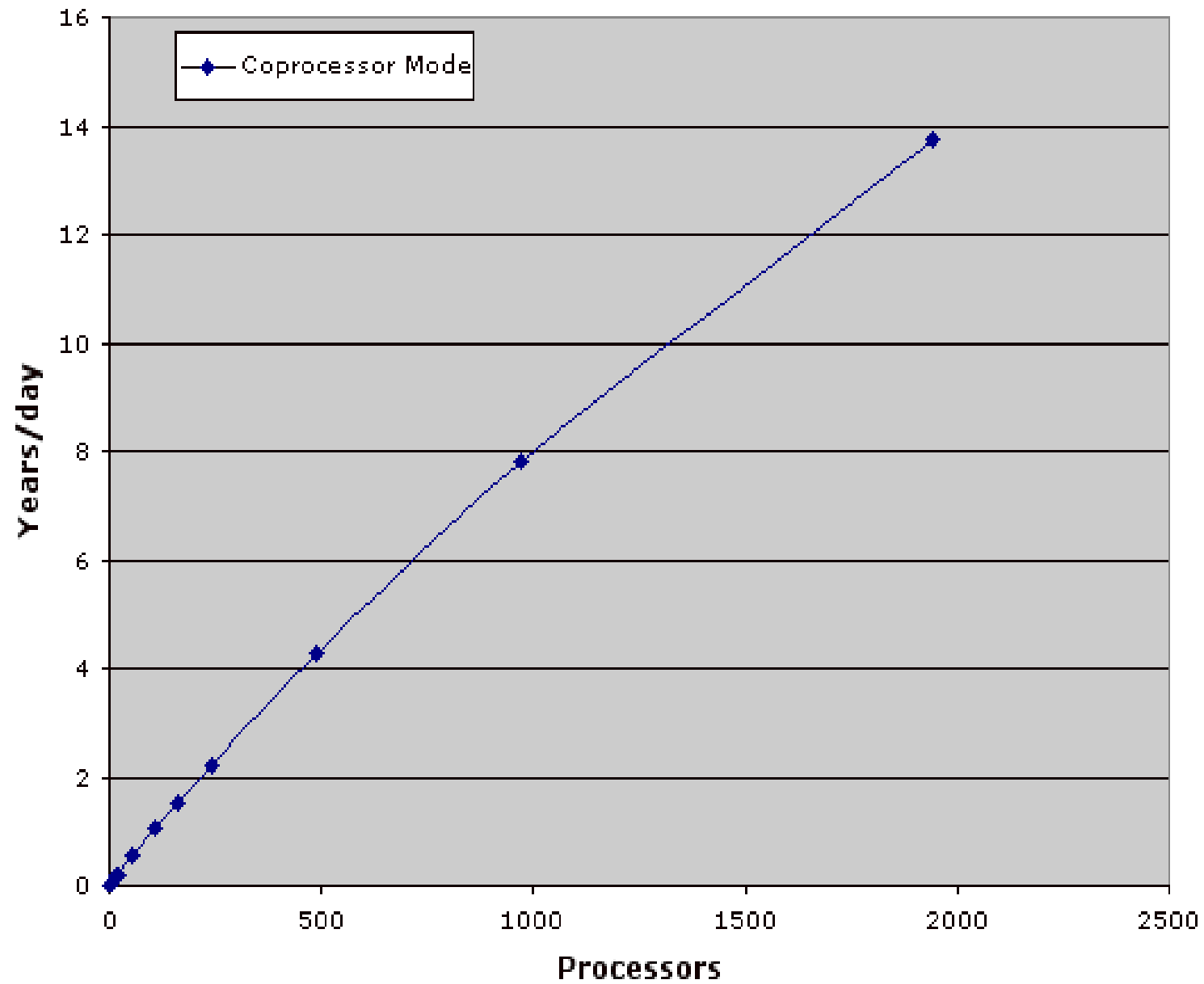
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Moist Held-Suarez Test Case: Temporally and Zonally Averaged Mixing Ratio

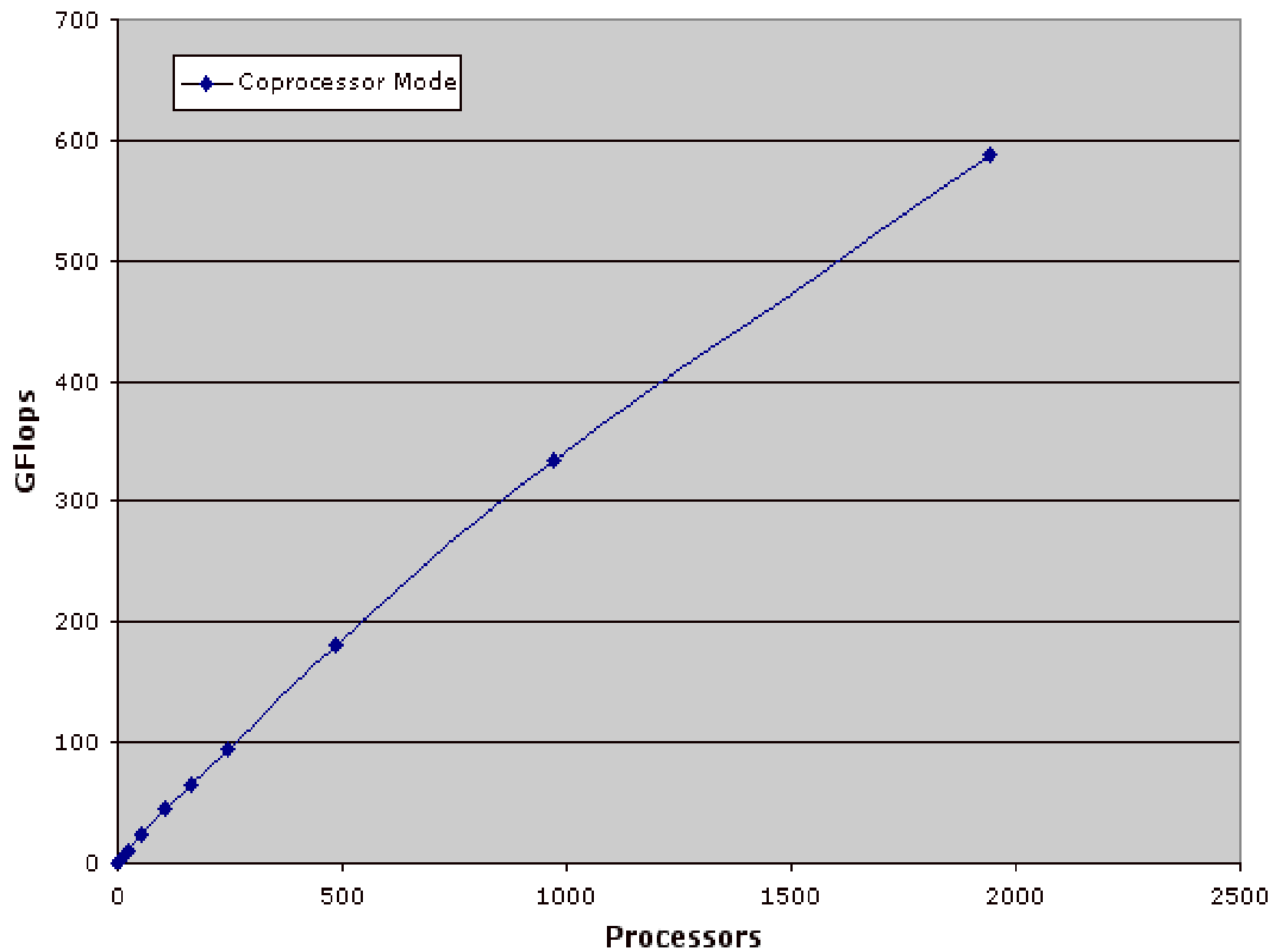


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Explicit HOMME Integration Rate: 80 km, 20 levels



Blue Gene/L Performance: ~80 km 20 level HOMME



Blue Gene/L Per-Processor Performance

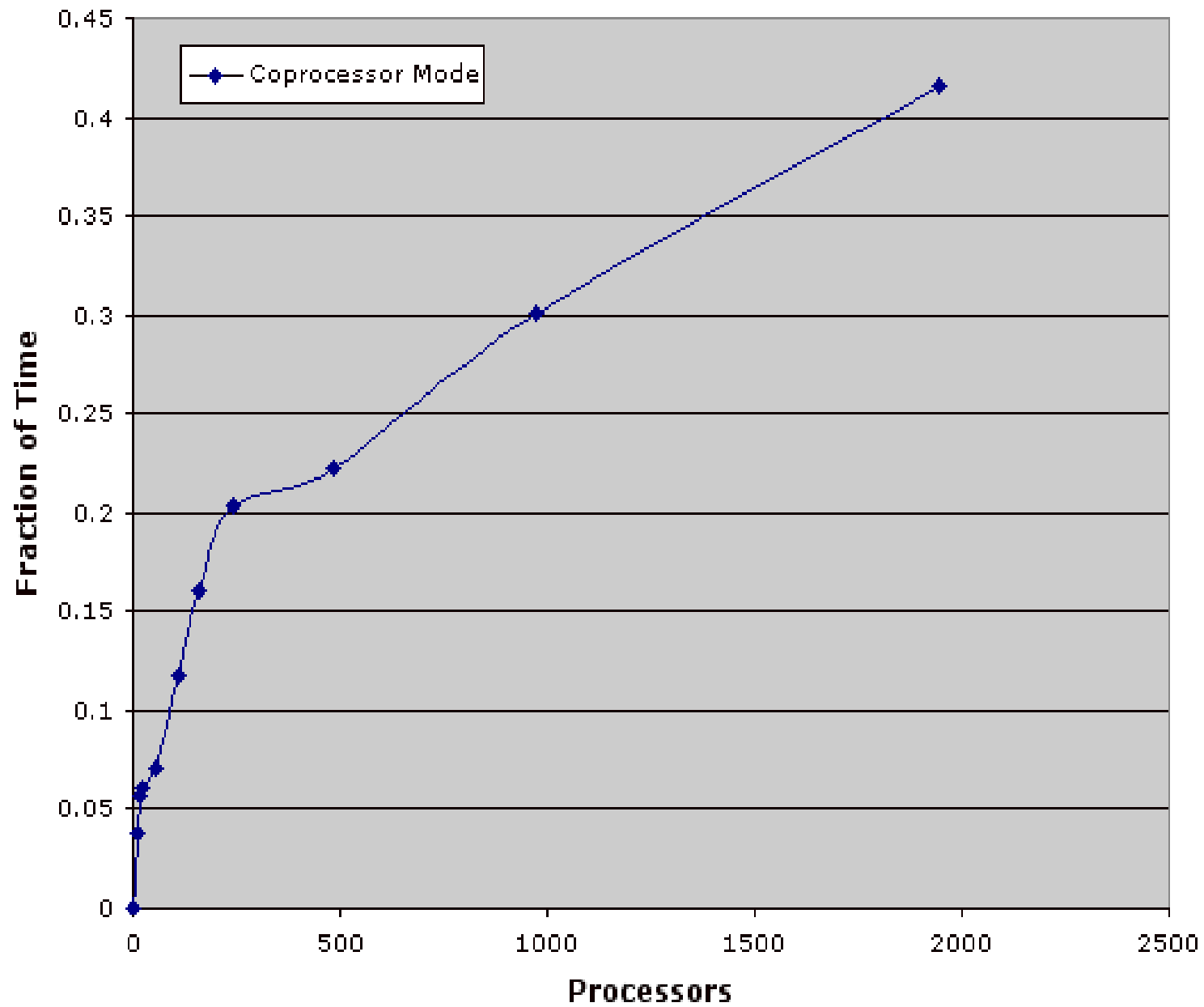
QuickTime™ and a
TIFF (Uncompressed) decompressor
are needed to see this picture.

- Ne=18, 20 level data
- Explicit primitive equations
- “Coprocessor” Mode
- 16% of 2.8 Gflops peak
- BG/L vector intrinsics
- Not clear how many double FMA’s are being generated.

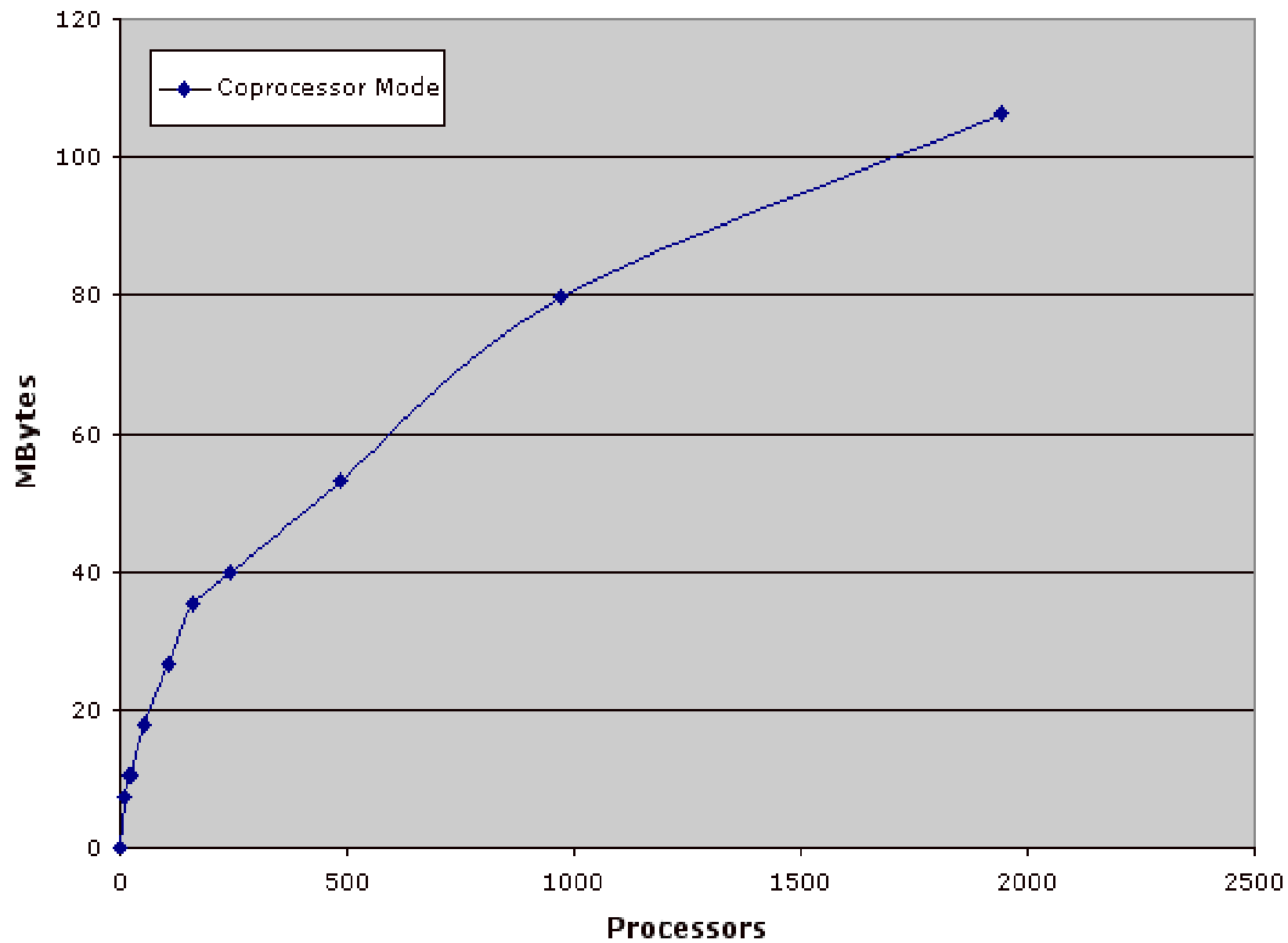


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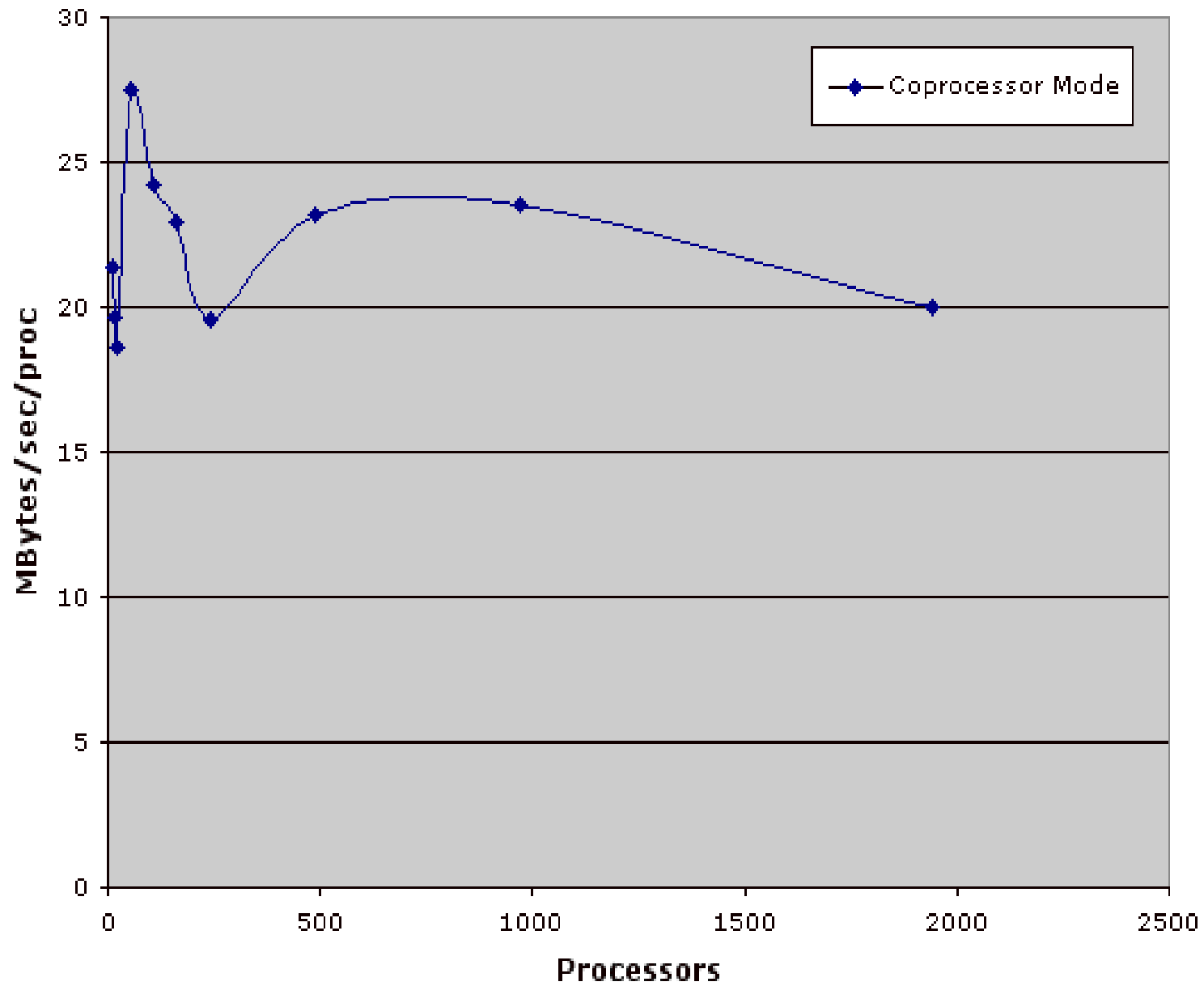
Fraction of Time Spent In Communication



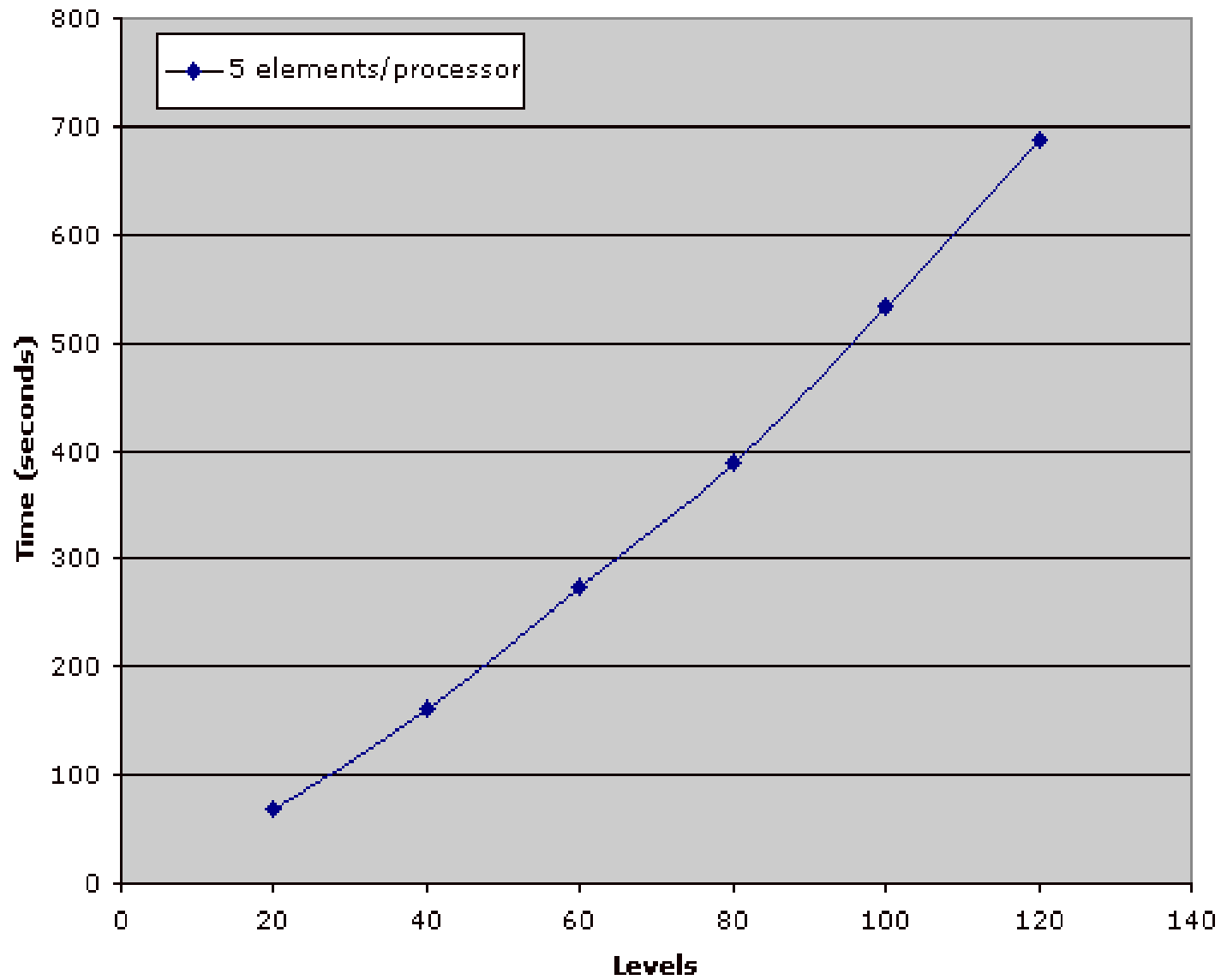
HOMME Communication Volume



Sustained Blue Gene/L Bandwidth



Scaling of Computational Cost with Number of Levels in HOMME



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BlueGene/L Collaboration

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CU Boulder



Blue Gene/L



CU Denver



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Questions?



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CRCP (super-parameterized) physics...



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CRCP or super-parameterization: What is “super” about it?

- The basic idea is to represent sub grid scales of the 3D large-scale model (with horizontal resolution ~ 100 km) by embedding a 2D cloud resolving model **in each column** of the large-scale model.
- Involves thousands of 2D cloud resolving models interacting in a way consistent with large scale dynamics.
- **Embarrassingly parallel but extremely expensive.** (150x over traditional physics)

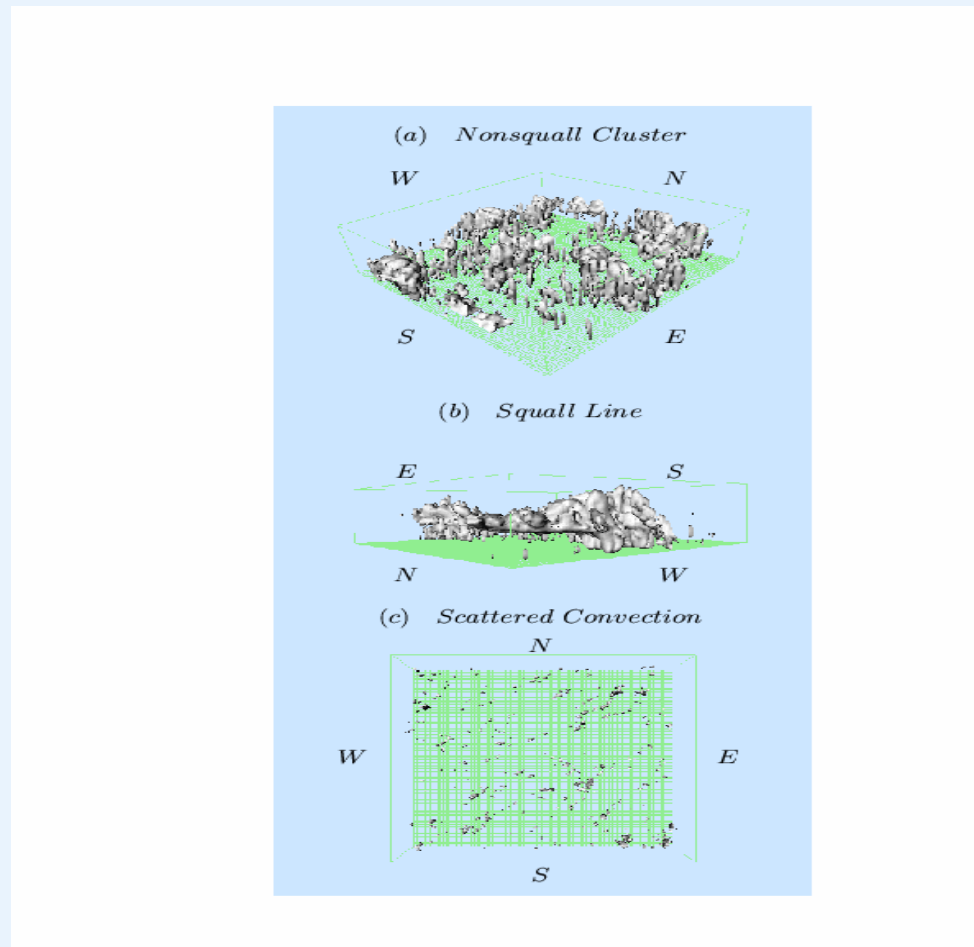


CRCP is the next step in the quest for a cloud-system-resolving AGCM

- The computational cost of the following simulations are approximately the same:
 - A **millennium**-long simulation using a traditional climate model.
 - A **few years**-long simulation using a traditional climate model with CRCP
 - A **day**-long simulation of a cloud-system-resolving AGCM O(few km)
- Cost of each separated by 3 orders of magnitude!



CRCP: 2D or not 2D: that is the question...



400 km box of clouds with
 $dx = dy = 2$ km



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Advantages of CRCP

- Allows for better representation of convection, clouds and radiative transfer and surface exchange.
- Allows for a dynamic response to changes in cloud parameters (particle sizes, aerosol characteristics, precipitation mechanisms at approximately correct length scales.
- Example: cloud top longwave cooling in response to particle size change.

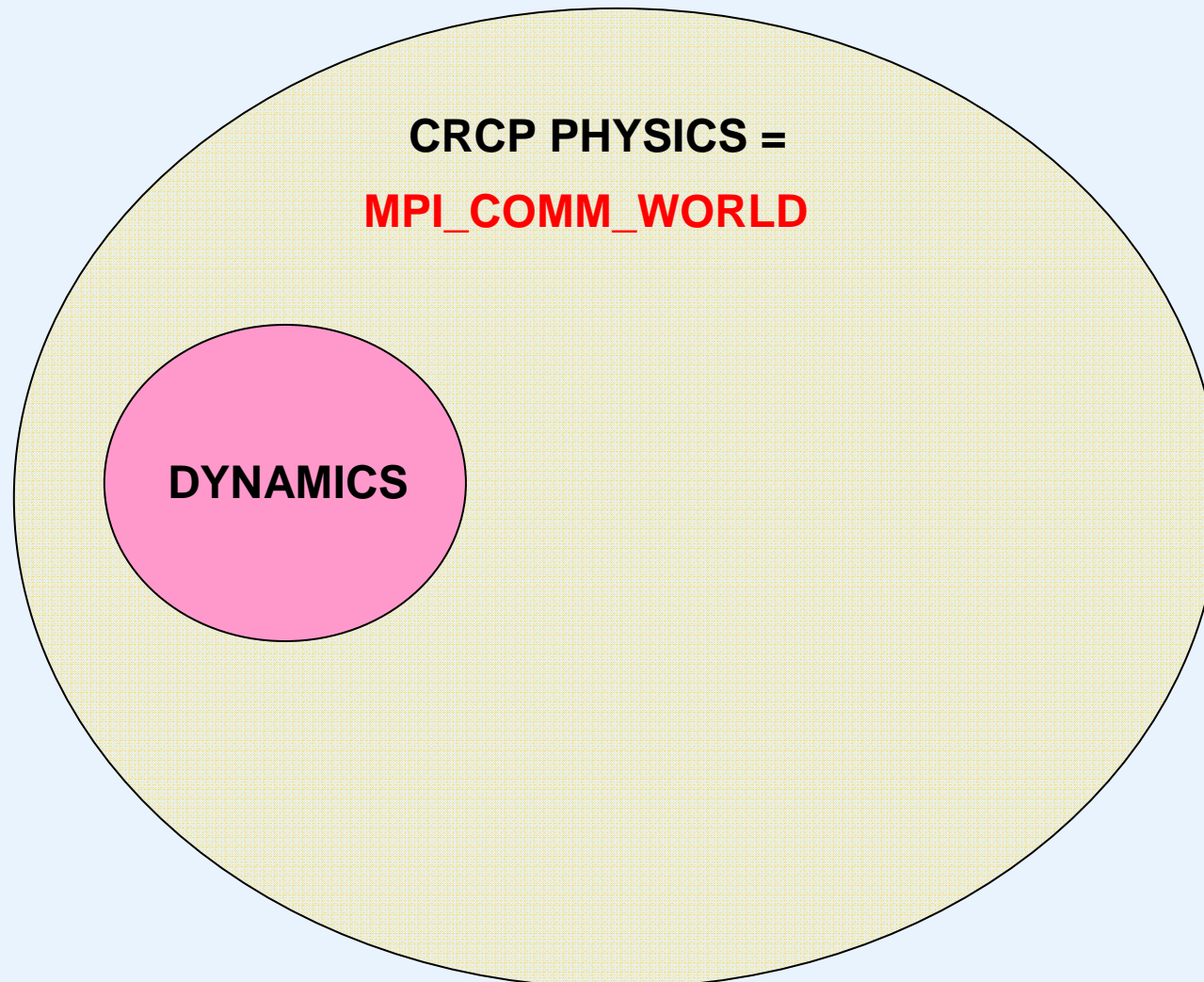


Coupling CRCP physics and dynamics together...

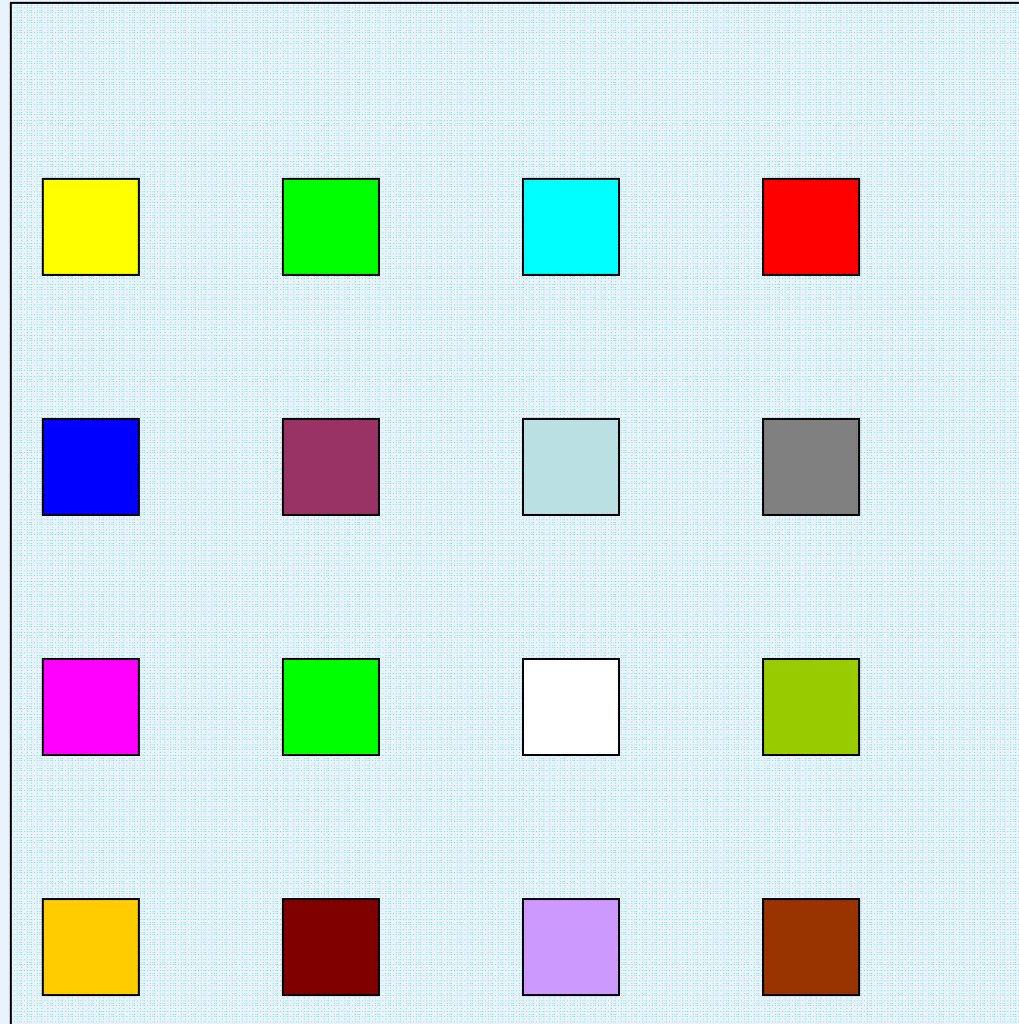


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Logical View of CRCP-HOMME Coupling on Blue Gene/L



Hardware View of CRCP-HOMME Coupling on Blue Gene/L

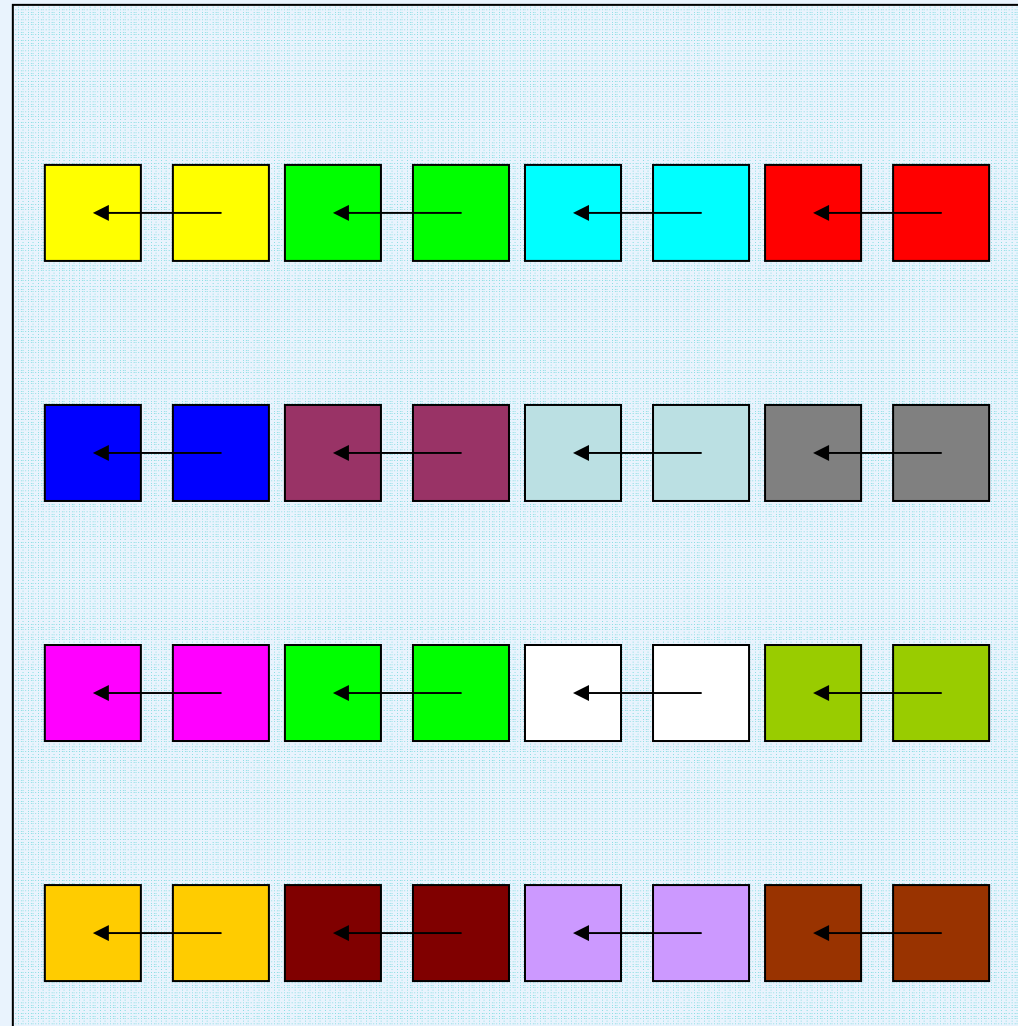


Begin with elements laid out for dynamics scheme



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Hardware View of CRCP-HOMME Coupling on Blue Gene/L

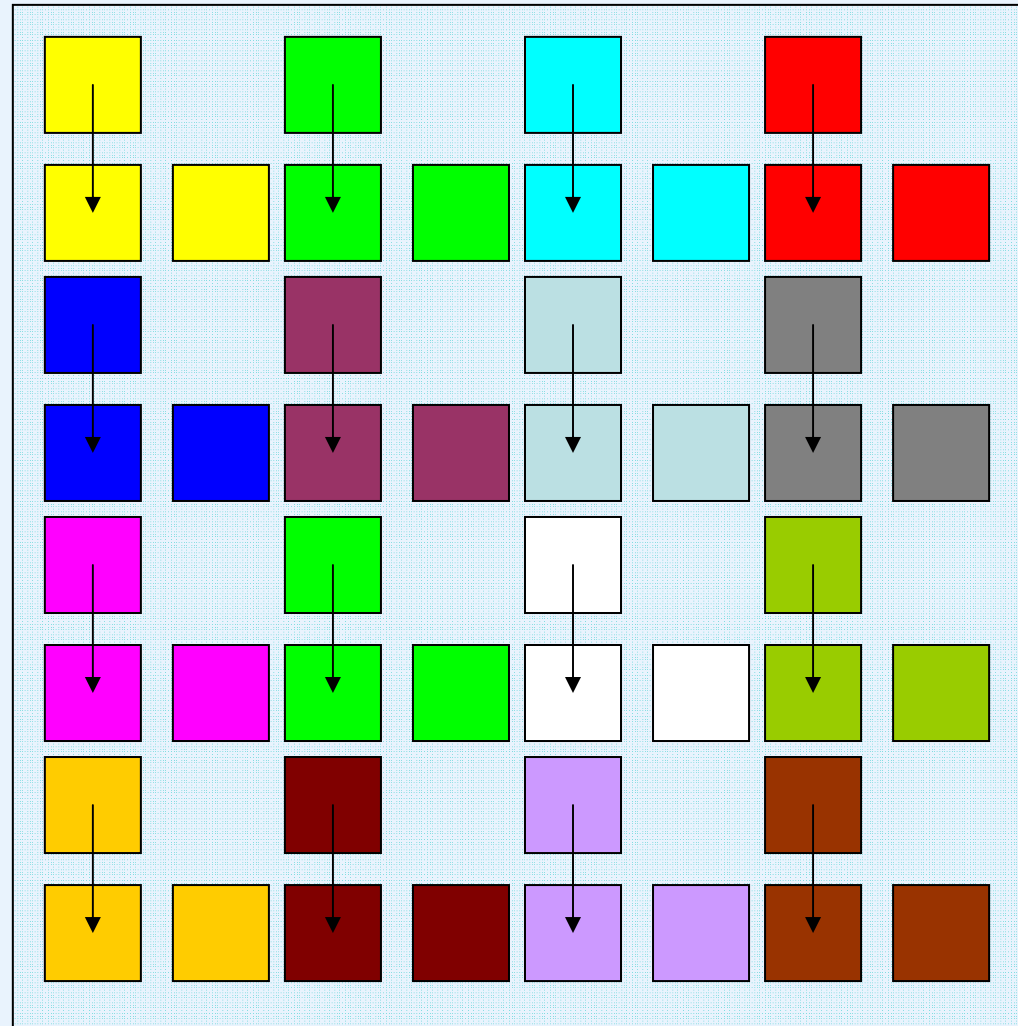


Begin scattering the columns



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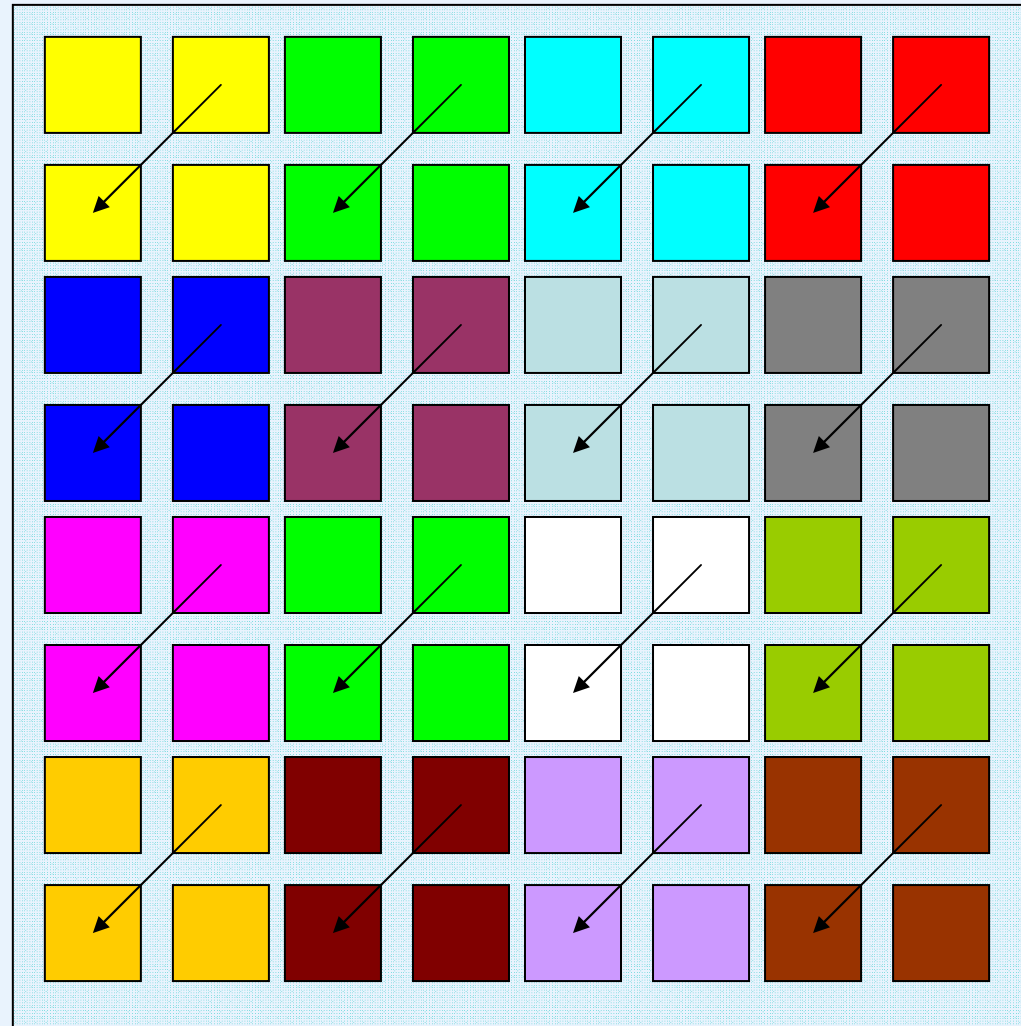


Continue scattering columns



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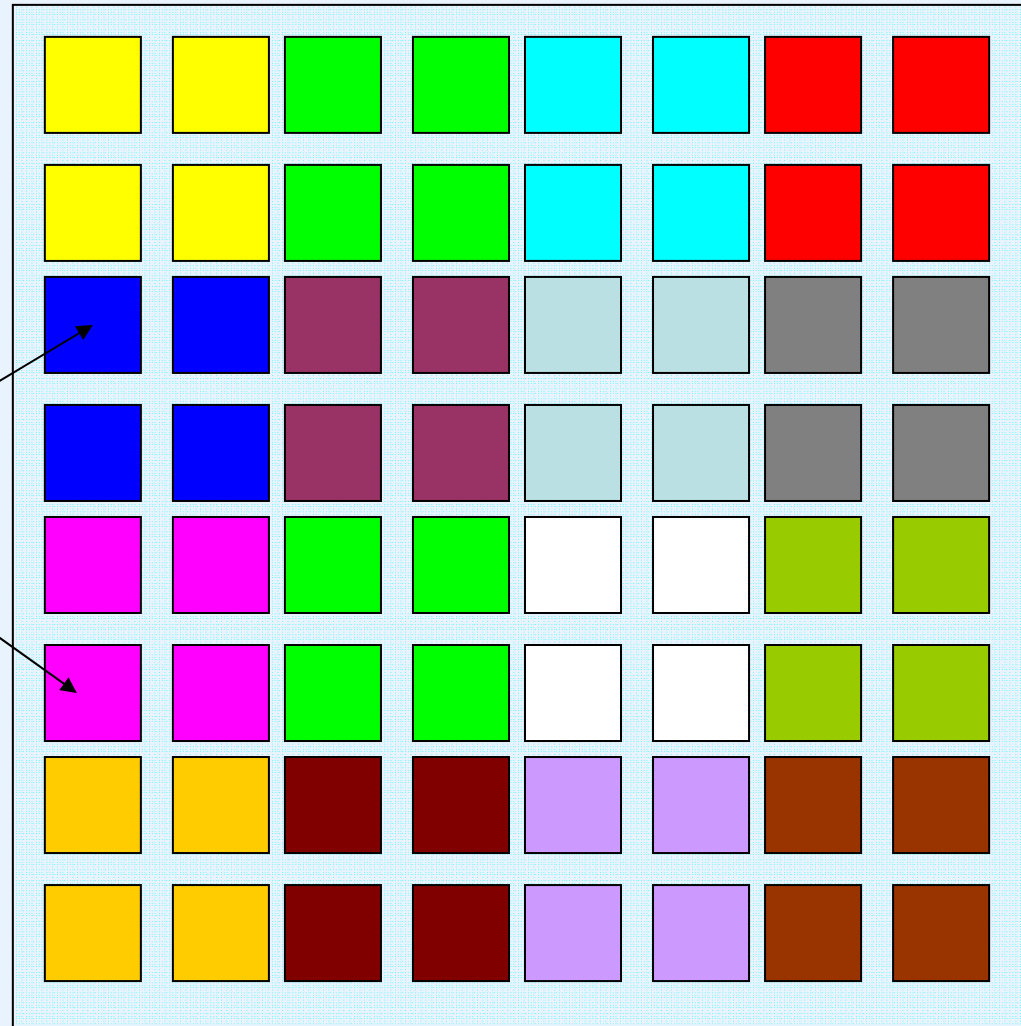
Continue scattering columns



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Hardware View of CRCP Physics Layout on Blue Gene/L

Physics Columns



Colors denote
elements

Simplified 2-d BG/L topology



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