

Chapel HPC Challenge Entry: 2011

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Benchmarks: John Lewis, Kristi Maschhoff, Jonathan Claridge



SC11: November 15th, 2011



What is Chapel?



- A new parallel programming language
 - Design and development led by Cray Inc.
- Overall goal: Improve programmer productivity
- Being developed as open source (BSD) at <u>SourceForge</u>
- Target Architectures:
 - multicore desktops and laptops
 - commodity clusters
 - Cray architectures (as well as those from other vendors)
- A work-in-progress





Hardware Platforms



NCCS Jaguar: Cray XT5™

- Dual AMD 6-core Istanbul Opteron processors
- Cray SeaStar[™] interconnect
- 16 GB of memory per node
- 9984 nodes

Cray Internal system Kaibab: Cray XE6™

- Dual AMD 12-core Magny-Cours Opteron processors
- Cray Gemini™ interconnect
- 32 GB of memory per node
- 84 nodes





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Life's now harder due to NUMA nodes/first-touch policies





THE SUPERCOMPUTER COMPANY

Speaking of being a work-in-progress...

- We have several performance-critical irons in the fire, many aimed at the Cray XE/XK and Cascade lines:
 - lightweight user-level tasking layers
 - a native communication layer for Cray's Gemini™ interconnect
 - a bulk copy optimizations for array slices
 - new hooks for overlapping communication and computation
 - a hierarchical locale concept for NUMA nodes and other emerging node architectures
- Look for these in our 1.5 release, April 2011





Chapel Codes for 2011



HPCC:

- 1. Stream Triad
 - EP
 - Global
- 2. Random Access (RA)
- 3. HPL
 - focusing on Schur Complement computation

Others:

- 4. SSCA #2 (an unstructured graph benchmark)
 - focusing on Kernel 4: Betweenness centrality
- 5. Adaptive Mesh Refinement (AMR) Framework

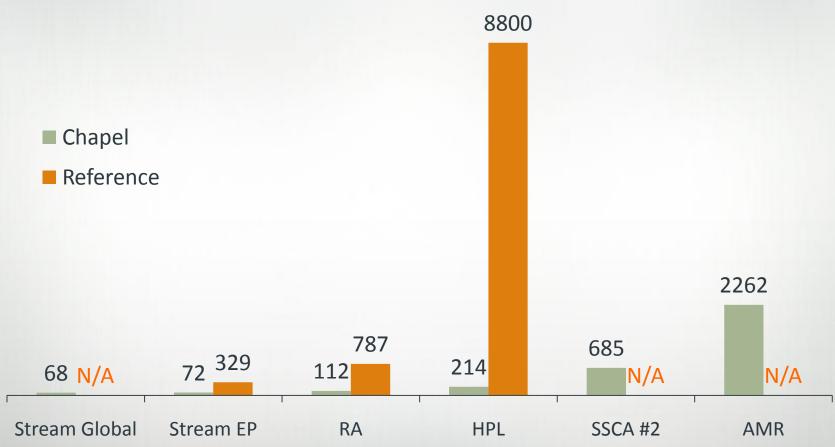




Chapel Source Code Sizes





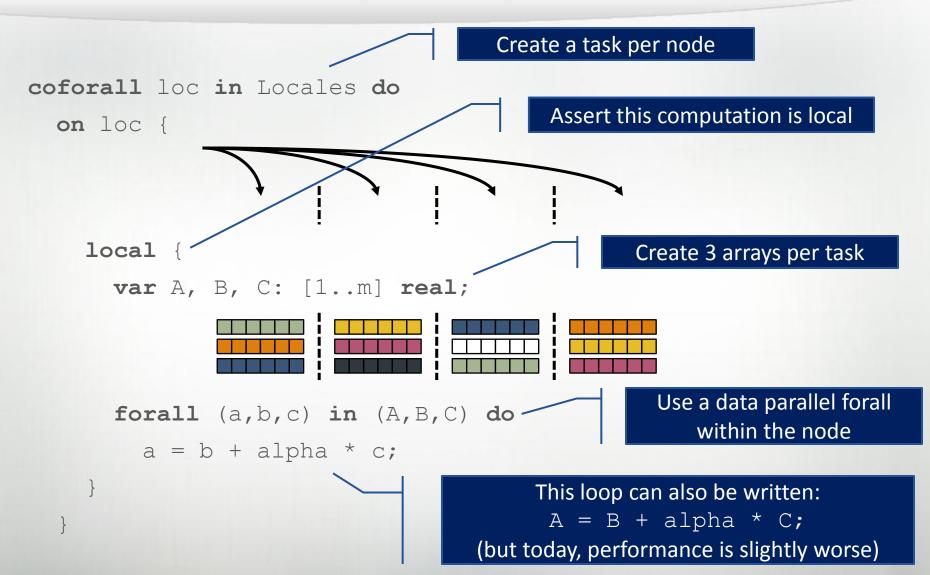








EP STREAM Triad in Chapel (Excerpts)

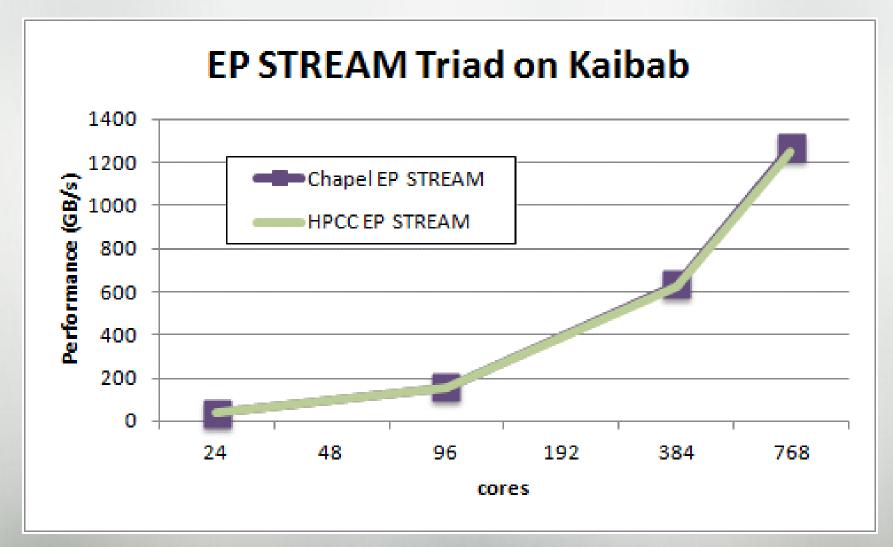








HPCC Stream Performance on Kaibab (XE6)

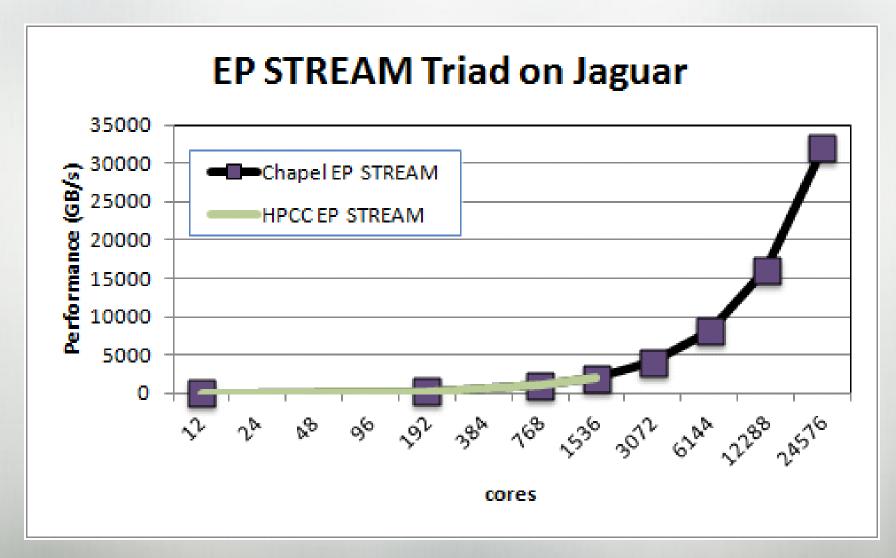








HPCC Stream Performance on Jaguar (XT5)









Global STREAM Triad in Chapel (Excerpts)

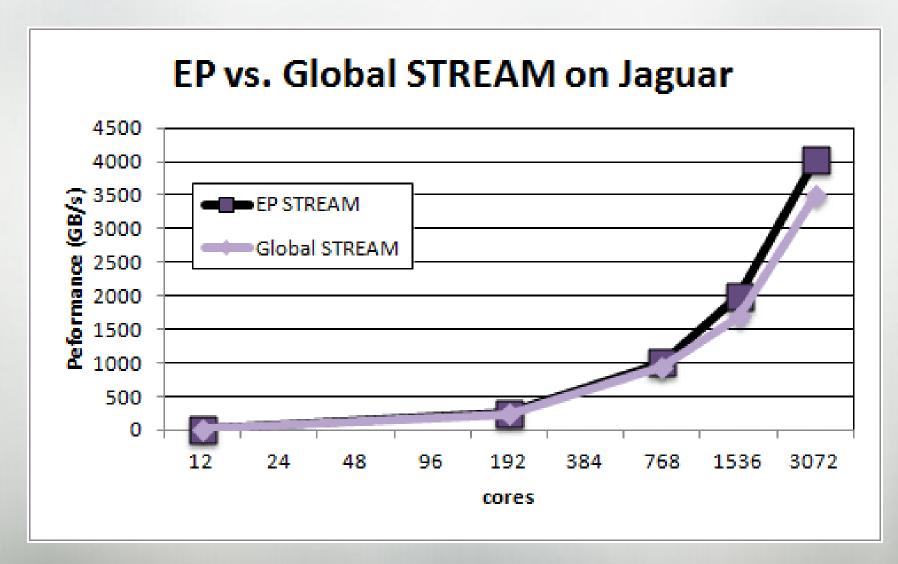
```
const ProblemSpace: domain(1, int(64))
                    dmapped Block([1..m])
                  = [1..m];
var A, B, C: [ProblemSpace] real;
forall (a,b,c) in (A,B,C) do
  a = b + alpha * c;
```





HPCC Global STREAM on Jaguar (XT5)









Declare two blocked index sets

Global Random Access in Chapel (Excerpts)

local T[myR & indexMask] ^= myR;

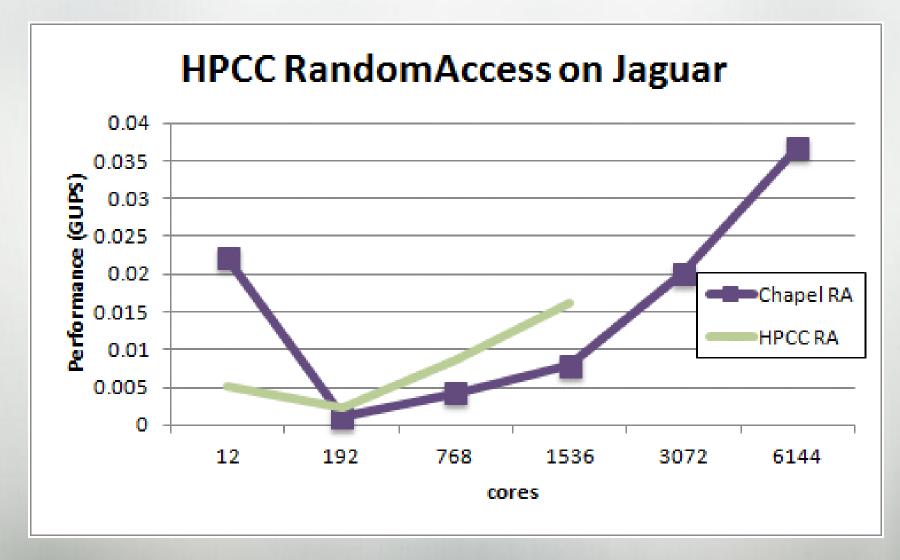
This can also be written:

```
on T[r&indexMask] do
    T[r&indexMask] ^= r;
(but again, performance is slightly worse today)
```













HPL



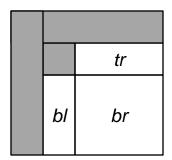
 Since 2009 entry, we have taken HPL from a shared memory implementation to distributed memory

- Have only started studying performance tuning in past month
 - Focusing initially on Schur Complement phase

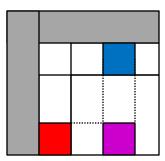




Schur Complement

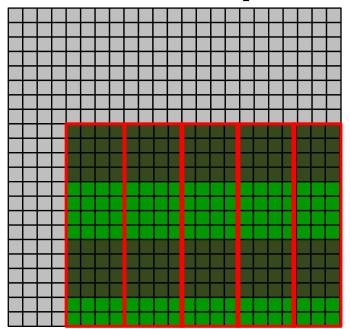


accumulate into each block in br the product of its corresponding blocks from bl and tr





Schur Complement w/ distribution



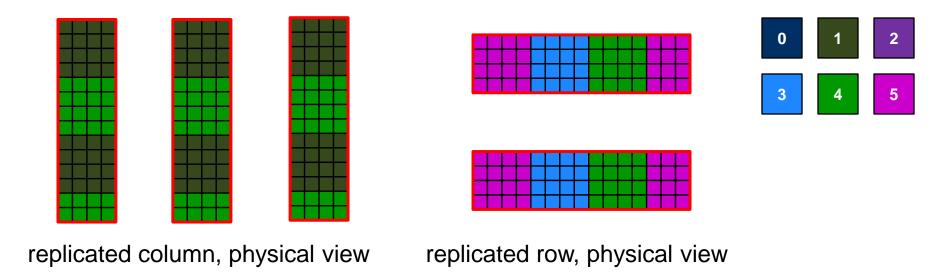
replicated col, logical view

replicated row, logical view





Schur Complement w/ distribution



- For scalability, we had to add distributions beyond Block:
 - **Block-Cyclic**
 - Replicated
 - Dimensional: meta-distribution that takes a distribution per dimension







Schur Complement (elegant version)

```
proc schurComplement(AD, BD, Rest) {
  replA[AD] = Ab[AD];
  replB[BD] = Ab[BD];
  forall (row, col) in Rest by (blkSize, blkSize) do
    local dgemm(replA[row..#blkSize, ..],
                replB[.., col..#blkSize],
                Ab[row..#blkSize, col..#blkSize]);
proc dgemm(A: [1.., 1..] elemType,
           B: [1.., 1..] elemType,
           C: [1.., 1..] elemType) {
  for i in C.domain.dim(1) do
    for j in C.domain.dim(2) do
      for k in A.domain.dim(2) do
        C[i,j] = A[i, k] * B[k, j];
```

Parallel loop over all result blocks

Dgemm of slices from replicated row/col blocks







Schur Complement (manually tuned version)

```
proc schurComplement(blk, const slct) {
  const AD dim2 = MatVectSpace.dim(2)[blk..#blkSize];
  const BD dim1 = MatVectSpace.dim(1)[blk..#blkSize];
  fillReplicants(blk, AD dim2, BD dim1);
  const RestLocal = MatVectSpace[blk+blkSize.., blk+blkSize..];
  Rest = RestLocal;
  RestByBlkSize = RestLocal by (blkSize, blkSize);
  if Rest.numIndices == 0 then
    return;
  forall (row,col) in RestByBlkSize do
    dqdriver(row, col, slct);
```

Most changes involve inserting temporaries to localize data or optimize copies until the compiler can do it automatically







Schur Complement (manually tuned version) 2

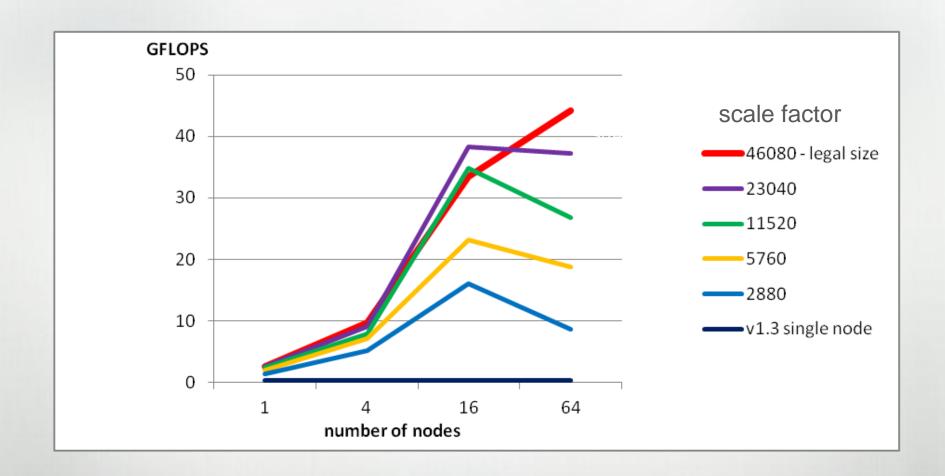
```
proc dgdriver(row, col, slct) {
  local {
    const rnq1 = Rest.dim(1) [row..#blkSize],
          rng2 = Rest.dim(2)[col..#blkSize],
          slctLoc = slct;
    dgemm(replA.localSlice(rng1, ..),
          replB.localSlice(.., rng2),
          Ab.localSlice(rng1, rng2),
          slctLoc);
proc dgemm(LreplA, LreplB, LAb, slct) {
  const r1 = LAb.domain.dim(1), r2 = LAb.domain.dim(2),
        rB = 1..blkSize;
for i in r1 do
    for k in r2 do {
      var acc: elemType = 0;
      for j in rB do
        acc -= LreplA[i,j] * LreplB[j,k];
      LAb[i,k] += acc;
```







Schur Complement Performance (XE)



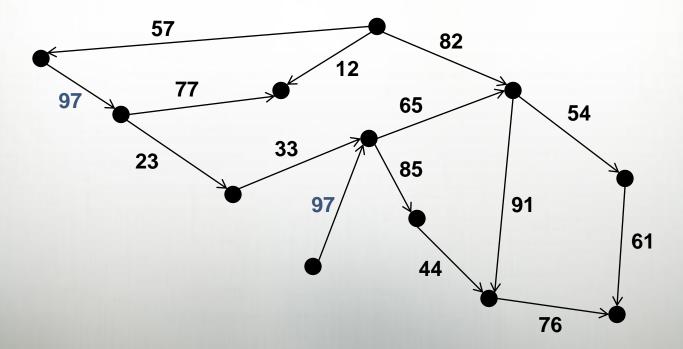






Goal: Compute Betweenness Centrality for graph

Our approach: Madduri et al.'s modification of Brandes's 2001 algorithm







SSCA#2, kernel 4 excerpts

```
Create a task per locale
do {
  coforall loc in Locales {
                                       Loop over all of the active
    on loc {
                                        vertices
       forall u in rcLocal[Active Level]. Members {
         forall v in G.Neighbors(u)
                                          On the locale owning the neighbor...
           on v
             if min distance$[v].readXX() < 0 {</pre>
                if min distance$[v] < 0 {</pre>
                  min distance $[v] = current distance c;
                  rcLocal[Next Level].Members.add(v);
                } else {
                  min distance $[v] = current distance c;
```





SSCA#2, kernel 4 excerpts

```
min distance $[v].readFF() == current distance c
           path count$[v] += path count$[u].readFF();
           children list[u].add child(v);
                                    Update our paths/children
    rcLocal[Next Level] = new Level Set(Sparse Vertex List);
    rcLocal[Next Level].previous = rcLocal[Active Level];
                                     Swap our current level set for a fresh one
    Active Remaining[here.id]
      rcLocal[Active Level].Members.numIndices:bool;
                                    See how much more local work we have
remaining = | | reduce Active Remaining;
while remaining;
                                        See if any work remains globally...
```

and continue if there is





SSCA#2 Kernel 4



 Performance is measured in Traversed Edges Per Second (TEPS)

February 2011: 22 TEPS

November 2011: 264,569 TEPS

 Improvement due to a combination of algorithmic changes and optimizations in Chapel







Adaptive Mesh Refinement Framework

Setting: Had UW Applied Math PhD Student (Jonathan Claridge) write an AMR framework in Chapel

- expertise with AMR
- no prior use of Chapel
- little-to-no experience parallel programming



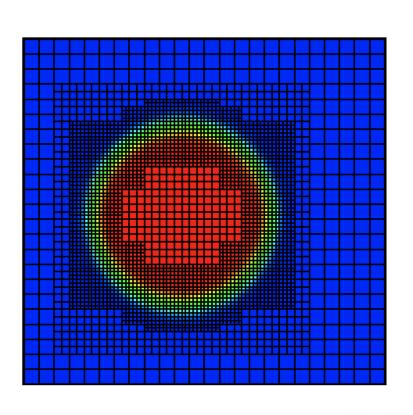


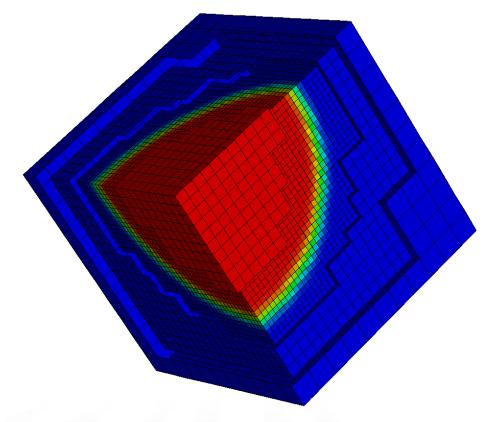




One Key Feature: Rank-independent

Uses the same code to produce results in 2D, 3D, 6D, 17D...





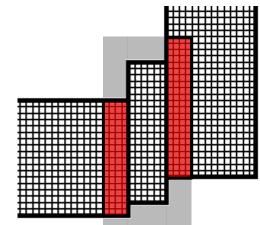






Levels: Sibling overlaps

A grid's layer of ghost cells will, in general, overlap some of its siblings. Data will be copied into these overlapped ghost cells prior to mathematical operations.



Calculating the **overlaps** between siblings:

```
var neighbors: domain(Grid);
var overlaps: [neighbors] domain(dimension, stridable=true);
for sibling in parent level.grids {
  var overlap = extended cells[sibling.cells];
  if overlap.numIndices > 0 && sibling != this {
    neighbors.add(sibling);
    overlaps[sibling] = overlap;
```









 Chapel domains made many fundamental AMR calculations very easy, even in a dimension-independent setting

- Difficult to do apples-to-apples comparisons (in terms of size or performance) with other AMR codes
 - Or even to know what would be a meaningful use of the framework to time
 - Community is generally lacking standardized AMR benchmarks





Pointers to code



Stream, RA, HPL:

- In release: \$CHPL_HOME/examples/hpcc
- Under SVN:

https://chapel.svn.sourceforge.net/svnroot/chapel/trunk/test/release/examples/hpcc/https://chapel.svn.sourceforge.net/svnroot/chapel/trunk/test/studies/hpcc/HPL

SSCA #2:

- In release: \$CHPL_HOME/examples/ssca2
- Under SVN:

https://chapel.svn.sourceforge.net/svnroot/chapel/trunk/test/release/examples/ssca2/

AMR:

• Under SVN:

https://chapel.svn.sourceforge.net/svnroot/chapel/trunk/test/studies/amr/



Summary



- Nodes are getting harder to optimize for
- Chapel compiler continues to improve
 - new optimizations
 - new distributions
 - new runtime capabilities
- Lots of work remains
 - but issues continue to seem tractable







- Mon: full-day tutorial
- Mon: 2nd annual CHUG happy hour/meet-up
- Tues: "HPC Challenge" BoF (12:15-1:15)
- Wed: "Chapel Lightning Talks" BoF (12:15-1:15)
 - I/O, education, tasking, GPUs, interoperability
- Thurs: "Punctuated Equilibrium at Exascale" Panel (5:30-7:00)
- Fri: half-day tutorial
- T-Th: Chapel posters in PGAS booth, Chapel team members staffing (T 2-4, W 10-12, W 4-6, Th 10-12)



