## **GRCHOMBO** - Using the Cluster

### How to use?

1) What are nodes and cores?

2) OpenMP vs MPI

3) What are ranks and threads?

4) How to decide#nodes/cores/ranks/threads?

5) Load balancing



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## The Cluster



### Nodes vs Cores



- Nodes have to communicate through the network
- Cores wihin he same node:
  - 1) can communicate directly (==> faster)
  - 2) have shared memory (==> faster)



### GRChombo shares boxes across CPUs



# MPI 101

MPI (Message Passing Interface) is an interface for communication between processes (ranks!) across distributed memory



### Example of 'Reduction' - global sum



Where can you find it in the code? 1) In Chombo:

Useful functions in BaseTools  $\rightarrow$  SPMD.H Other files as BoxTools  $\rightarrow$  BoxLayoutDatal.H (it's a dangerous road to go there!)

6///0

2) In GRChombo

GRChomboCore → SetupFunctions.hpp AMRInterpolator → MPIContext.hpp (and others...)

# OpenMP 101

OpenMP (Open Multi-Processing) is an interface for parallel programming (threads!) with shared memory



### Example of FOR loop for (int i = 0; i < n; ++i) {</pre> C++ // do stuff #pragma omp parallel for OpenMP for (int i = 0; i < n; ++i) {</pre> & C++ // do stuff in parallel Where can you find it in the code?

Original image by EPCC under license at https://creativecommons.org/licenses/by-nc-sa/4.0/deed.en\_US (colors inverted)

## Sample Job Script

#!/bin/bash
#SBATCH -p (partition)
#SBATCH -A (account)
#SBATCH --time=12:00:00
#SBATCH --job-name=sample

#SBATCH --nodes=10 #SBATCH --ntasks-per-node=8 #SBATCH --cpus-per-task=4

export OMP\_NUM\_THREADS=\$SLURM\_CPUS\_PER\_TASK

# other stuff? Import modules?
# (...)

mpiexec program params.txt

(this is for a slurm job script, even though similar in other managers as pbs)

#ranks/node \* #threads/rank = #cpus/node
(32 in this case)

6///0

#ranks/node \* #nodes = #ranks
(80 in this case)

# nodes# MPI ranks per node# OpenMP threads per MPI rank

(tell OpenMP about it)

To run with MPI, you might see "mpiexec / mpirun / srun / ... "

# Load Balancing

### You run a simulation. What to look for?

out.0 •			
GRAMRLevel::advance	level 0 at	time 8.75 (2.13795 M/hr).	Boxes on this rank: 1.
GRAMRLevel::advance	level 1 at	time 8.75 (2.13783 M/hr).	Boxes on this rank: 1.
GRAMRLevel::advance	level 2 at	time 8.75 (2.13772 M/hr).	Boxes on this rank: 1.
GRAMRLevel::advance	level 3 at	time 8.75 (2.13754 M/hr).	Boxes on this rank: 1.
GRAMRLevel::advance	level 4 at	time 8.75 (2.13743 M/hr).	Boxes on this rank: 1.
GRAMRLevel::advance	level 5 at	time 8.75 (2.13724 M/hr).	Boxes on this rank: 1.
GRAMRLevel::advance	level 6 at	time 8.75 (2.13705 M/hr).	Boxes on this rank: 1.
GRAMRLevel::advance	level 6 at	time 8.75391 (2.13781 M/hr	). Boxes on this rank: 1.
GRAMRLevel::advance	level 5 at	time 8.75781 (2.13856 M/hr	r). Boxes on this rank: 1.
GRAMRLevel::advance	level 6 at	time 8.75781 (2.13838 M/hr	). Boxes on this rank: 1.
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	the leve		#boxes
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#### Goals:

not too many boxes per rank
 as few ranks with no boxes as possible

**Note:** what matter most is the finest level (6 in this case), as it's the one that runs more often. The levels above can have no boxes in some ranks.

Look at pout.0  $\rightarrow$  too many boxes?

Look at last pout's  $\rightarrow$  are they empty? How many are empty?

(LB.txt might have some relevant info if you explore)



## Load Balancing

### Too many boxes? Solutions ("This is not an exact science", Miren Radia):

Option 1: increase #nodes Option 2: reduce #threads/rank (openMP) and increase #ranks/node (MPI) Option 3: decrease resolution

Option 4: increase box size

**Note 1:** OpenMP is slower than MPI in GRChombo (beyond 4 OpenMP threads, scaling is not good), so pick option 2 instead of 1 only if you want to save CPU hours, if queue waiting time is too big or simply can't ask for more nodes.

**Note 2:** sometimes you might experience error messages saying you ran out of memory. That is equivalent to having too many boxes, so proceed as above.



### Too few boxes?

Do the opposite of the options above, but typically option 1 (saves cpu hours!!) or 2 will be enough for this case.



# **Option 3: Change Resolution**

### Things you can do:

- A) Change #levels (max\_level parameter)
- B) Change N1-N3 (make the grid coarser/finer)
- C) Change N1-N3 and L accordingly (make grid smaller/bigger)
  - D) Change regridding threshold
  - E) [advanced] Change tagging criteria

### A few extra tips:

- Can you apply some symmetric boundary conditions to your problem?

- If you are doing black hole simulations, ensure ~40 points across the horizon. If you have more/less, then think of options A/B/C above.

Note: an easy way to find the horizon is to plot a contour of 'chi' (after puncture gauge settles, that is at about 0.25 for 0 spin, and closer to 0.15 for a high 0.8 spin)



## **Option 4: Change Box Sizes**

params.txt	×							
<pre>max_level = regrid_inte</pre>	= 9 erval =	00	0 64	64 64	64	64	64	0
<pre># Max and r max_grid_s: block_fact</pre>	nin box ize = 10 or = 16	size 6	es					

Box Size	% without ghosts			
4	6%			
8	19%			
16	38%			
32	60%			









### 16x16(x16)



## **Option 4: Change Box Sizes**

+ Box Size ==> - communication - #ghosts

- but + memory/rank (if you have few boxes, otherwise not true)
  - & **#boxes** ==> can't use as many MPI ranks

Recommended: 8, 16, 24 or 32 (multiples of 8, to match vectorization)

If you have very few boxes, switch to 8 If you have too many boxes, switch to 16/32



## **Key Points**

- Don't forget to set OMP\_NUM\_THREADS

Try to choose a reasonable #nodes, #ranks, #threads
||
v
Look at your pout files (or LB.txt). Too many boxes? Too few boxes?
||
v
Balance MPI / openMP / resolution / box size

