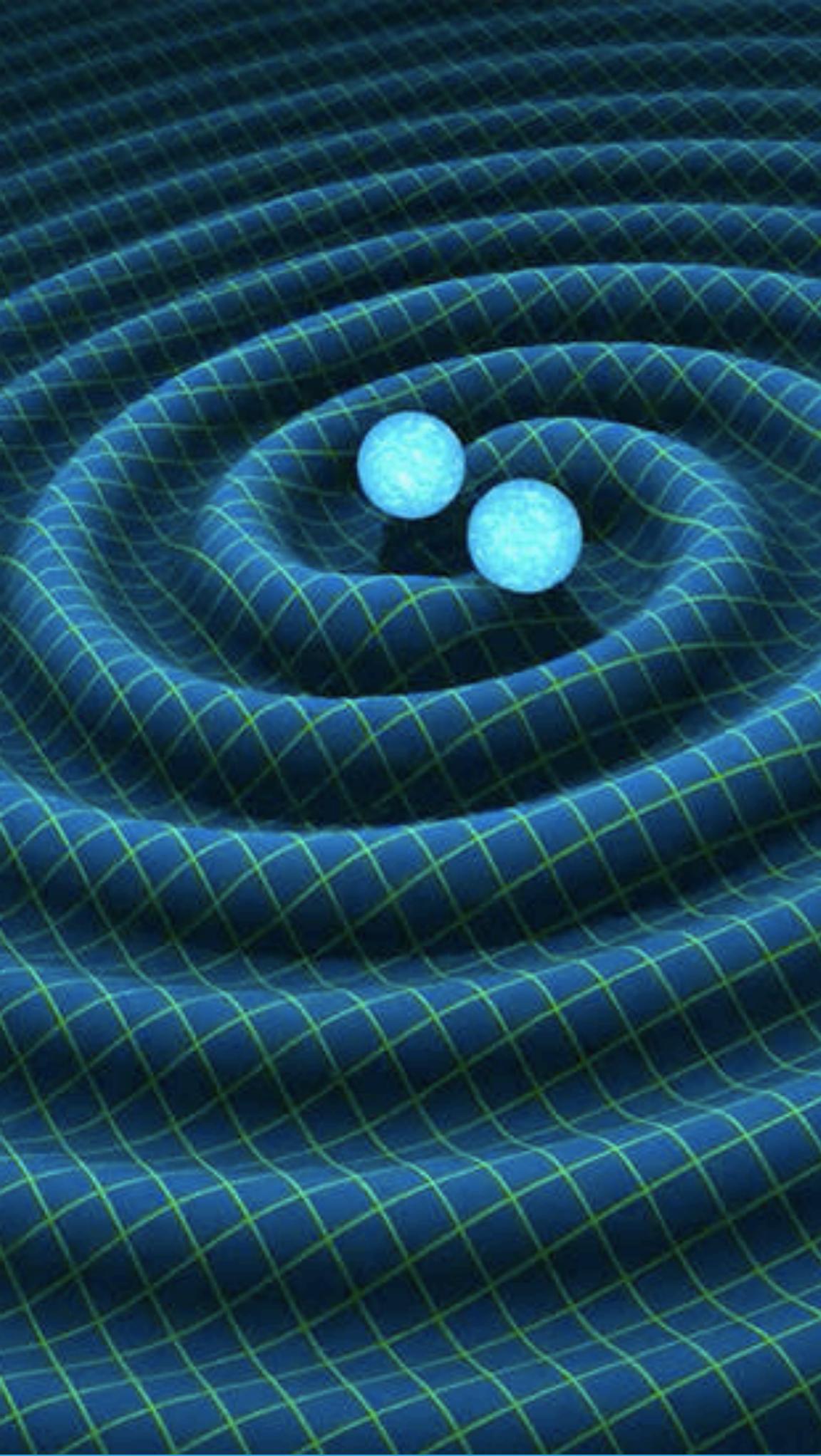




KATY CLOUGH

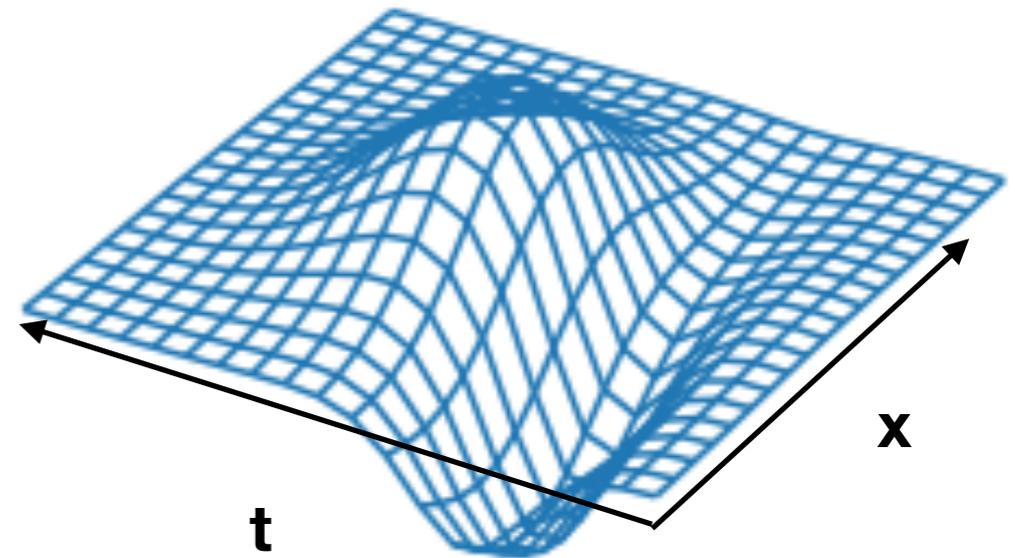
INTRO TO GRCHOMBO: THE BIG PICTURE



NUMERICAL RELATIVITY: BIG PICTURE

GR IN 2 MINUTES

$$ds^2 = f(x, t) dt^2 + g(x, t) dx^2 + 2 h(x, t) dt dx$$



$$ds^2 = \begin{pmatrix} dt & dx \end{pmatrix} \underbrace{\begin{pmatrix} f(x, t) & h(x, t) \\ h(x, t) & g(x, t) \end{pmatrix}}_{\text{“The spacetime metric” } g_{ab}(t, \vec{x})} \begin{pmatrix} dt \\ dx \end{pmatrix}$$

GR IN 2 MINUTES

“Matter tells spacetime how to curve...”

$$\mathbf{R}_{ab} - \mathbf{R}/2 \mathbf{g}_{ab} = 8\pi \mathbf{T}_{ab}$$

$\mathbf{f}(\partial^2 \mathbf{g}_{ab}, \partial \mathbf{g}_{ab}, \mathbf{g}_{ab})$

“Curvature”

“Energy-Momentum”

Can rearrange into form (using ADM decomposition):

$$\partial_t(\partial_t \mathbf{g}_{ab}) = \mathbf{f}(\partial_{xx} \mathbf{g}_{ab}, \partial_x \mathbf{g}_{ab}, \partial_t \mathbf{g}_{ab}, \mathbf{g}_{ab}, \mathbf{T}_{ab})$$

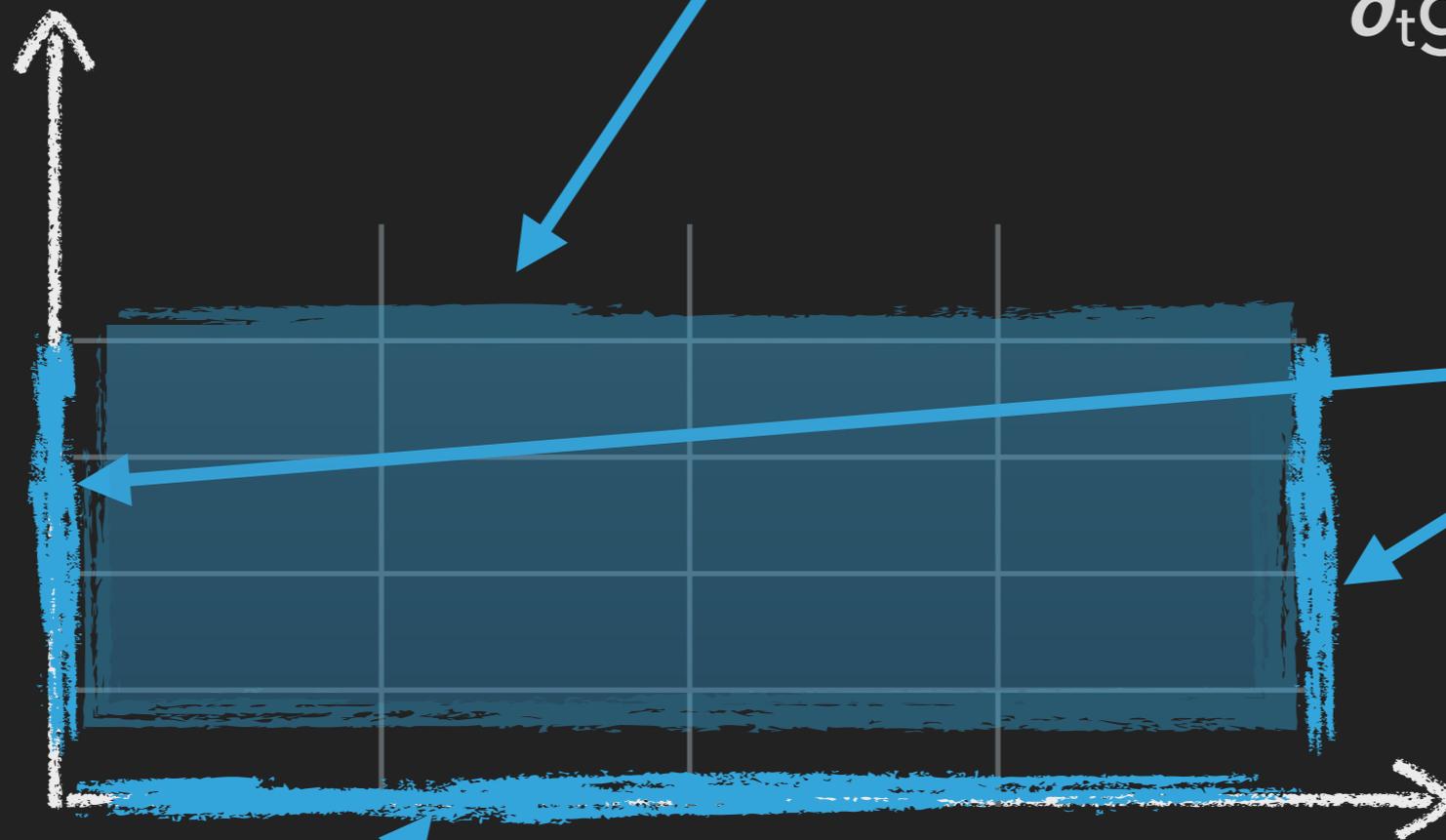
where $\partial_t \mathbf{g}_{ab} \sim \mathbf{K}_{ab}$

NR IN 2 MINUTES

"local time"

Fill using Einstein equation

$$\partial_{tt}g_{ab} = f(\partial_{xx}g_{ab}, \partial_xg_{ab}, \partial_tg_{ab}, g_{ab}, T_{ab})$$



boundary conditions
(∂_xg_{ab}, g_{ab})

"space"

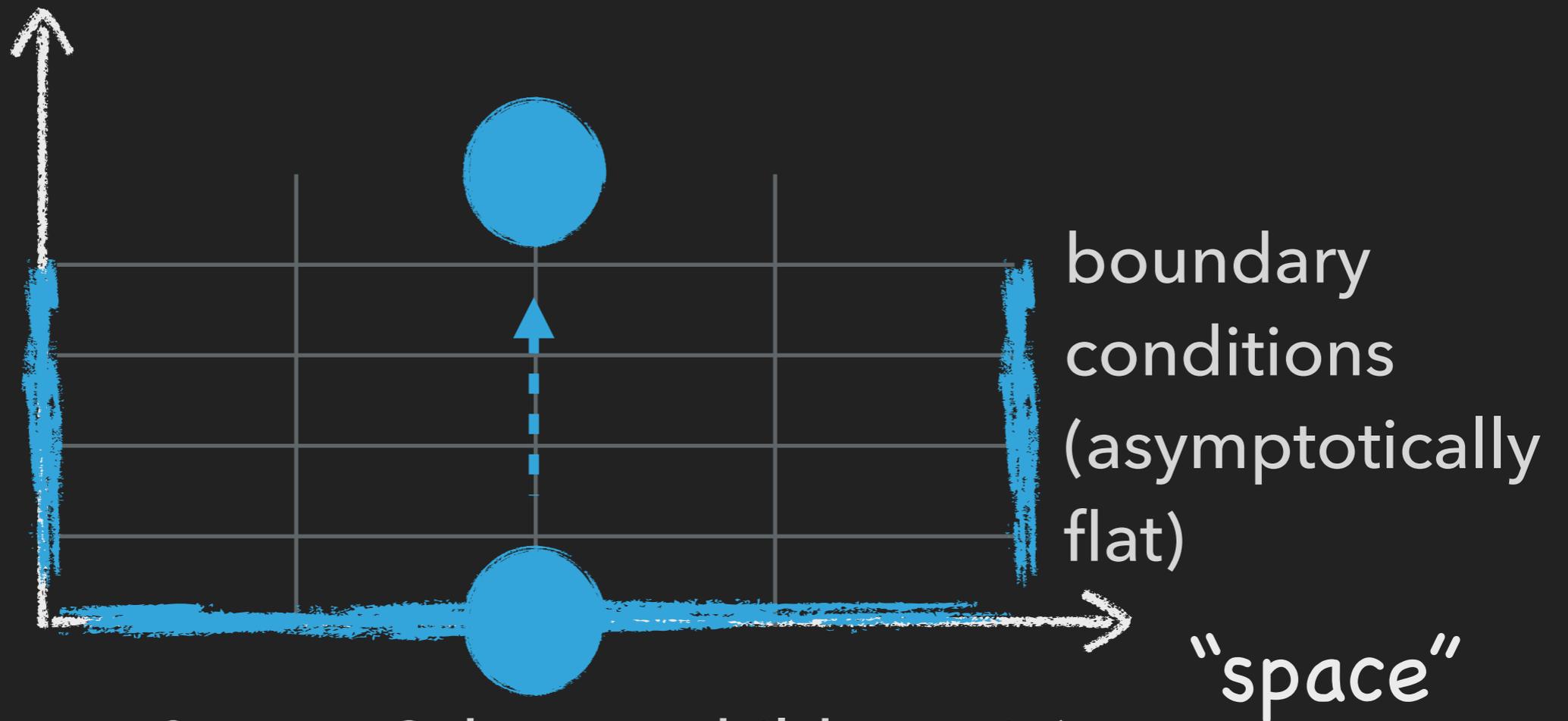
initial data (∂_tg_{ab}, g_{ab})

NR IN 2 MINUTES

"local time"

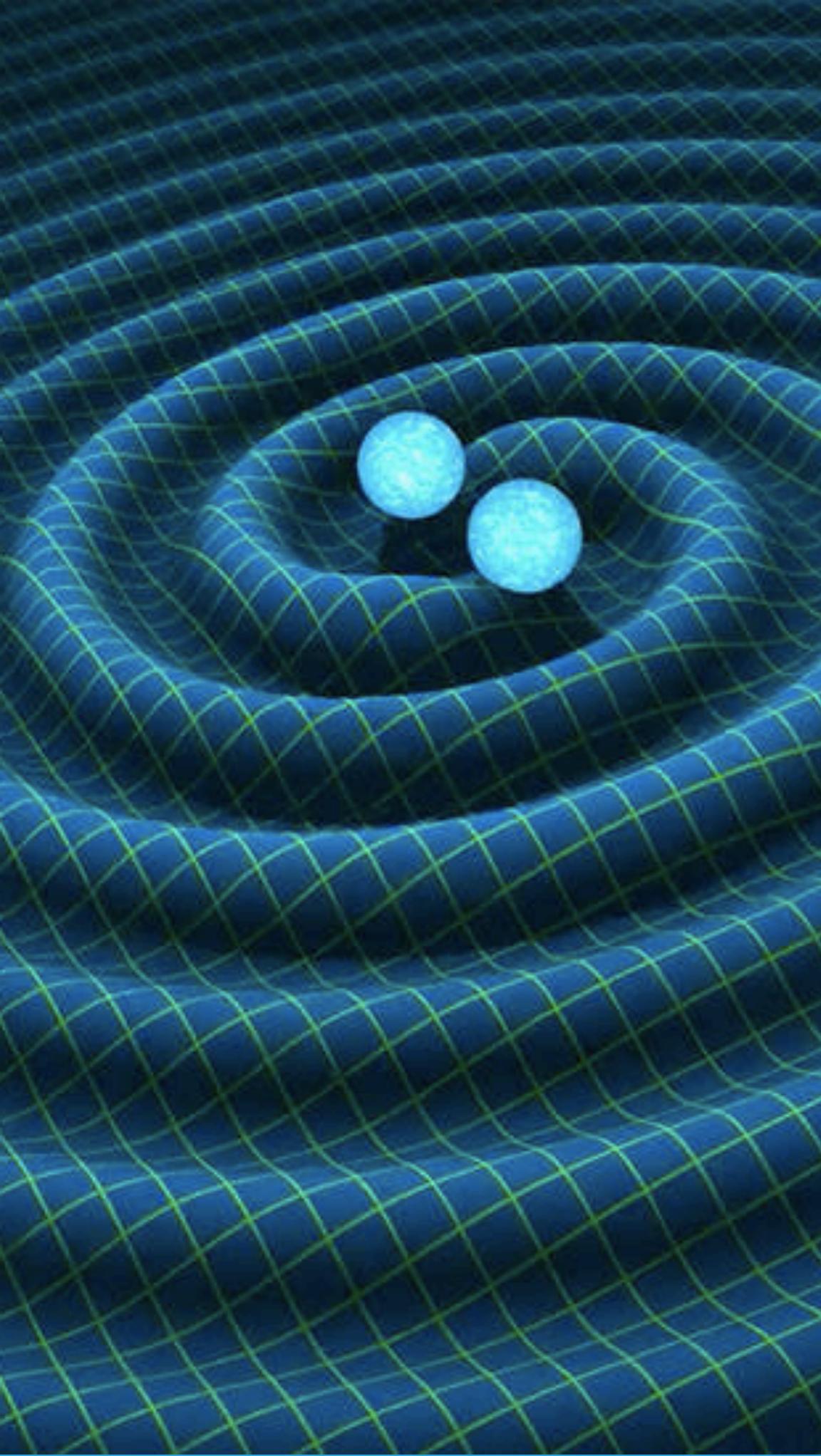
Fill using Einstein equation

$$\partial_{tt}g_{ab} = 0 \text{ (boring!) } *$$



initial data ($\partial_t g_{ab} = 0, g_{ab} = \text{Schwarzschild metric}$)

* in practise since the NR coordinates are not typically the Schwarzschild ones, we see some gauge evolution



GRCHOMBO: BIG PICTURE (FOCUS ON PROGRAM FLOW)

GRCHOMBO IS A BIG CODE

The screenshot shows the GitHub repository page for GRChombo/GRChombo. The repository is public and has 40 stars, 17 watchers, and 31 forks. The main branch is 'main', and there are 46 branches and 1 tag. The repository contains a list of files and folders, including .github/workflows, Doxygen, Examples, InstallNotes, Source, Tests, .clang-format, .gitignore, .pre-commit-config.yaml, GNUmakefile, LICENSE, README.md, and run_clang_format. The repository is described as an AMR based open-source code for numerical relativity simulations. It has 1 release, the latest being 'GRChombo: An adaptable nu...' on 10 Dec 2021. There are no packages published, and 14 contributors are listed.

GRChombo / GRChombo Public Unpin Watch 17 Fork 31 Star 40

[Code](#) [Issues 33](#) [Pull requests 7](#) [Discussions](#) [Actions](#) [Projects](#) [Wiki](#) [Security](#) [Insights](#) [Settings](#)

[main](#) [46 branches](#) [1 tag](#) [Go to file](#) [Add file](#) [Code](#) [About](#)

KAClough Merge pull request #211 from GRChombo/bugfix/210 231753c 6 days ago 355 commits

.github/workflows	Add github action to build [GR]Chombo with clang	4 months ago
Doxygen	Add GitHub action to update Doxygen documentation (#195)	6 months ago
Examples	Add fancier FOR macro replacing FOR1, FOR2, ...	10 months ago
InstallNotes	Add github action to build [GR]Chombo with clang	4 months ago
Source	Fix absolute restart_file paths bug	7 days ago
Tests	Alternative fix for segfault in SphericalExtractionTest	4 months ago
.clang-format	Added clang-format configuration file.	4 years ago
.gitignore	Ignoring files from more than one simultaneous vim edits	2 years ago
.pre-commit-config.yaml	Add pre-commit configuration file (#198)	5 months ago
GNUmakefile	Decrease make verbosity	2 years ago
LICENSE	Initial commit	4 years ago
README.md	Update README.md	3 months ago
run_clang_format	Added clang-format configuration file.	4 years ago

About

An AMR based open-source code for numerical relativity simulations.

- [Readme](#)
- [BSD-3-Clause License](#)
- [40 stars](#)
- [17 watching](#)
- [31 forks](#)

Releases 1

GRChombo: An adaptable nu... Latest
on 10 Dec 2021

Packages

No packages published
[Publish your first package](#)

Contributors 14

THREE LEVELS : CHOMBO / GRCHOMBO / BINARYBH

- ▶ Chombo - overall program flow relevant to any initial value problem - AMR, AMRLevel, ChomboParameters

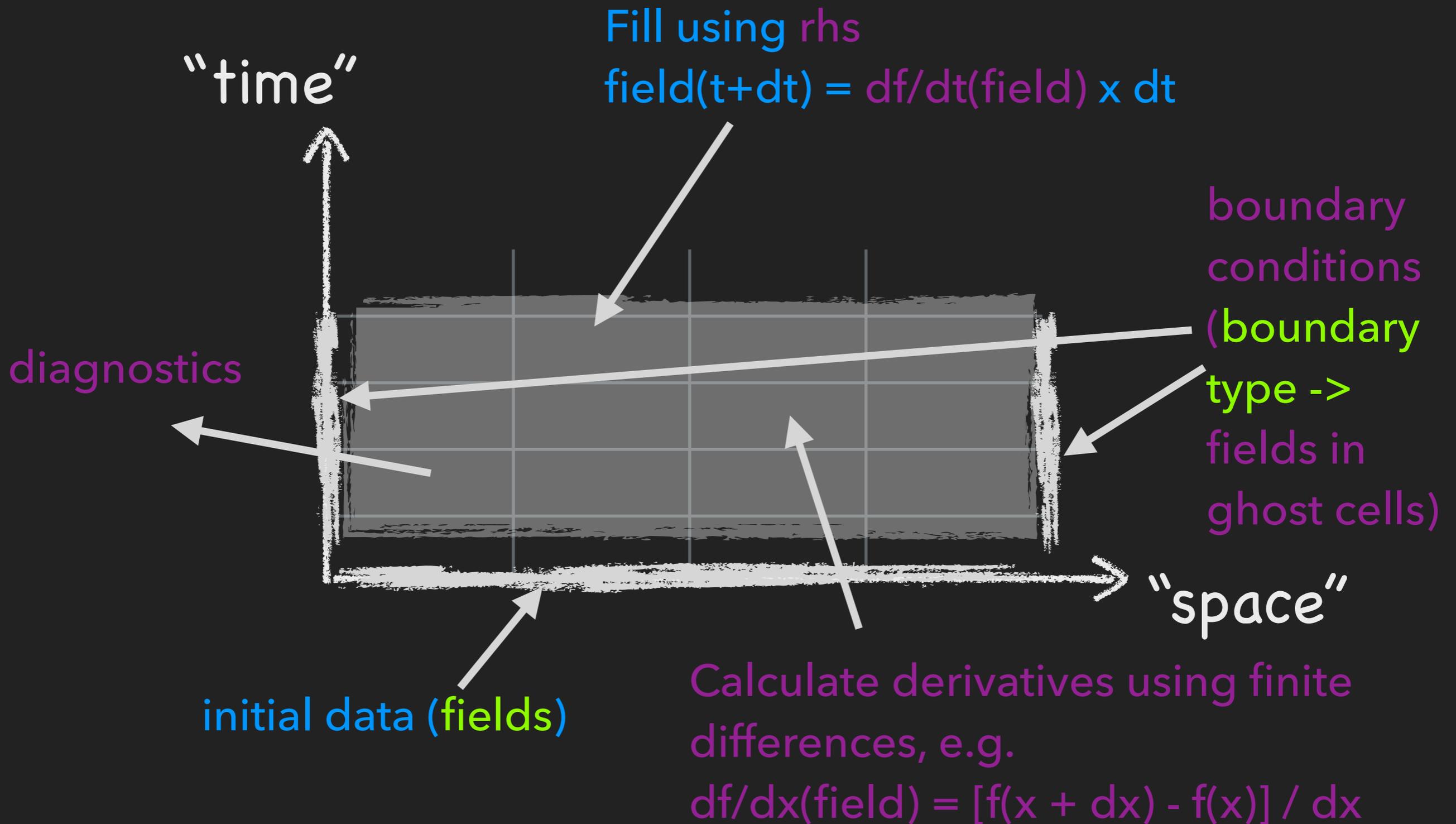


- ▶ GRChombo - specific physics actions common to most GR problems - GRAMR, GRAMRLevel, SimulationParametersBase

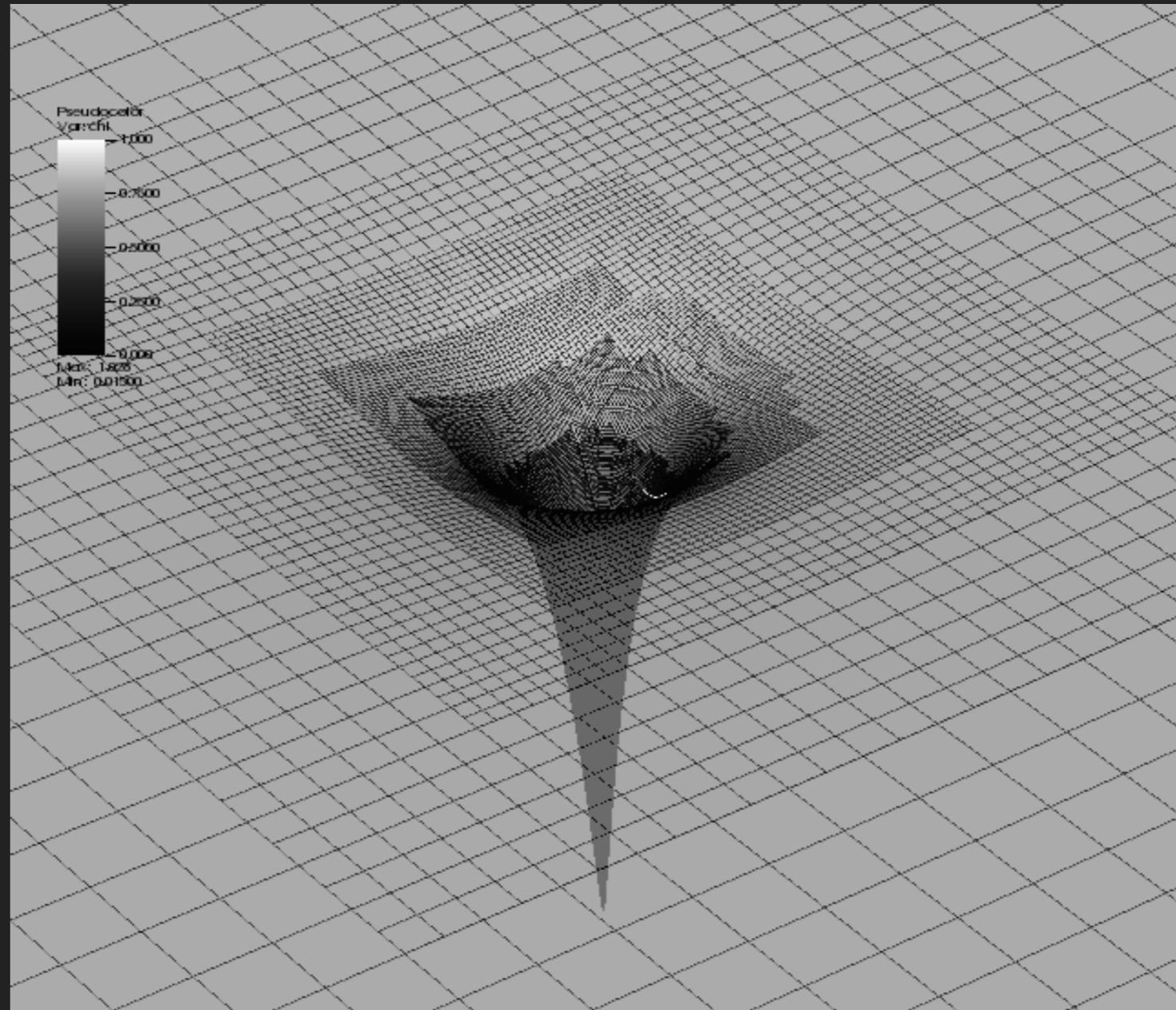


- ▶ BinaryBH - specific actions relevant to the Binary BH example - BHAMR, BinaryBHLevel, SimulationParameters

CHOMBO / GRCHOMBO / BINARYBH

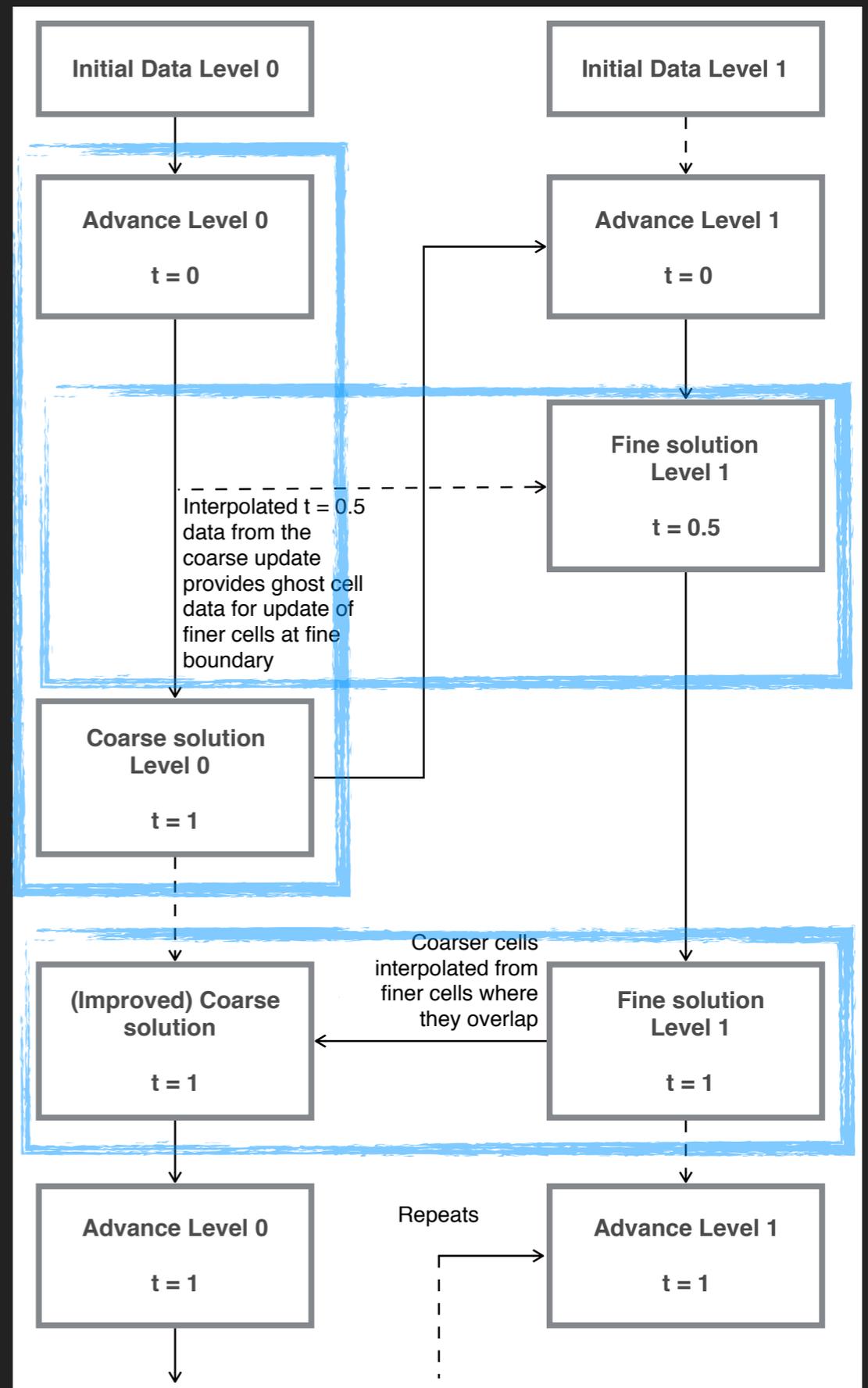
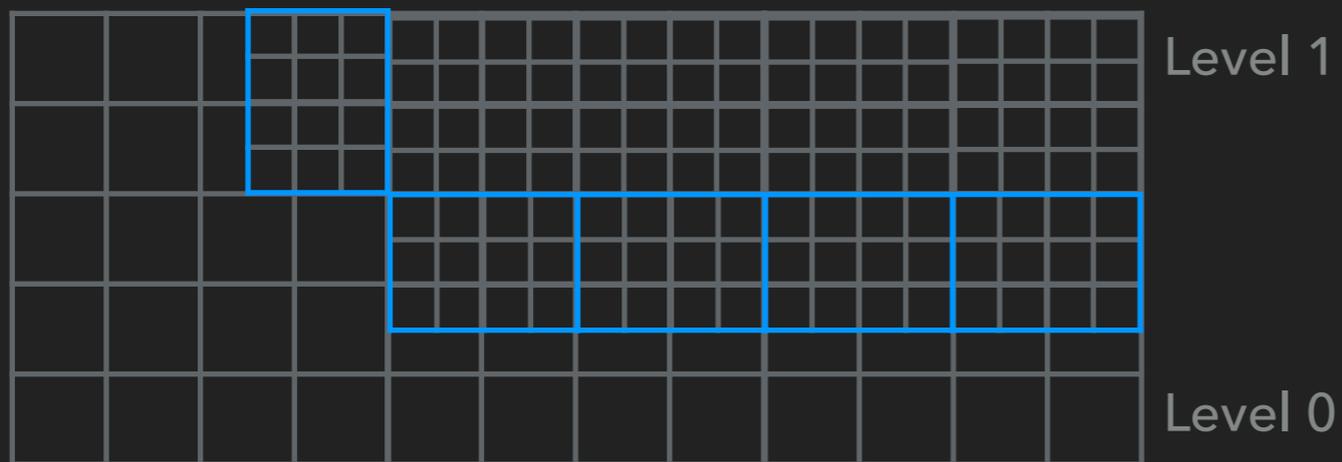


CHOMBO DEALS WITH THE ADAPTIVE MESH REFINEMENT (AMR)



AMR TIME STEPPING

- ▶ Each step is not really a single step but a series of Runge Kutta (RK4) substeps
- ▶ Data from coarser level is interpolated in both space and time to fill finer level **ghost cells** at level boundaries
- ▶ Level 0 is not finalised until Level 1 is => coarser levels have to wait for finer ones to end, so each level is processed in serial



WHERE ARE THE KEY CHOMBO FILES?

GRChombo / Chombo

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Code Issues 0 Pull requests 0 Actions Projects 0 Security Insights Settings

Branch: master **Chombo / lib / src / AMRTimeDependent /** Create new file Upload files Find file History

mirenradia Change checkpoint writing default on stop Latest commit 916f1d5 4 days ago

..

multidim	Initial commit of Chombo-3.2	2 years ago
AMR.H	Add hook in AMR::run to allow evolution stop	last month
AMR.cpp	Change checkpoint writing default on stop	4 days ago
AMRLevel.H	Change checkpoint writing default on stop	4 days ago
AMRLevel.cpp	Updated lib/src/AMRTimeDependent, lib/src/AMRTools, lib/src/BaseTools...	11 months ago
AMRLevelFactory.H	Initial commit of Chombo-3.2	2 years ago
ARK4.H	Initial commit of Chombo-3.2	2 years ago
ARK4DenseOutput.H	Initial commit of Chombo-3.2	2 years ago

WHERE ARE THE KEY GRCHOMBO FILES?

File	Commit Message	Time
..		
BoundaryConditions.cpp	Add fancier FOR macro replacing FOR1,FOR2,...	10 months ago
BoundaryConditions.hpp	Allow building GRChombo with NAMESPACE=TRUE (#151)	15 months ago
ChomboParameters.hpp	Fix absolute restart_file paths bug	7 days ago
DefaultLevelFactory.hpp	Allow building GRChombo with NAMESPACE=TRUE (#151)	15 months ago
EmptyDiagnosticVariables.hpp	Allow building GRChombo with NAMESPACE=TRUE (#151)	15 months ago
GRAMR.cpp	Move AMRInterpolator::fill_ghosts to GRAMR::fill_multilevel_ghosts	14 months ago
GRAMR.hpp	Move AMRInterpolator::fill_ghosts to GRAMR::fill_multilevel_ghosts	14 months ago
GRAMRLevel.cpp	Tidy up parameters files	13 months ago
GRAMRLevel.hpp	Tidy up parameters files	13 months ago
GRLevelData.cpp	Rewrite GRLevelData::plus with OpenMP and SIMD (#148)	14 months ago
GRLevelData.hpp	Rewrite GRLevelData::plus with OpenMP and SIMD (#148)	14 months ago
SetupFunctions.hpp	Fix absolute restart_file paths bug	7 days ago
SimulationParametersBase.hpp	Add fancier FOR macro replacing FOR1,FOR2,...	10 months ago
UserVariables.inc.hpp	Allow building GRChombo with NAMESPACE=TRUE (#151)	15 months ago
VariableType.hpp	Enable AMRInterpolator to interp diagnostic vars	2 years ago

WHERE ARE THE KEY BINARYBH FILES?

GRChombo / GRChombo

Watch 14 Star 20 Fork 22

Code Issues 2 Pull requests 4 Actions Projects 0 Wiki Security Insights Settings

Branch: master GRChombo / Examples / BinaryBH / Create new file Upload files Find file History

mirenradia Make all examples use formulation parameter Latest commit 9be2d07 12 days ago

File	Commit Message	Time Ago
BinaryBHLevel.cpp	Make all examples use formulation parameter	12 days ago
BinaryBHLevel.hpp	Improve the params for the binary merger and add puncture tracking	14 days ago
GNUmakefile	Run dos2unix and update .gitignore	3 months ago
Main_BinaryBH.cpp	Improve the params for the binary merger and add puncture tracking	14 days ago
SimulationParameters.hpp	Add params for puncture level	14 days ago
UserVariables.hpp	Amend example files	2 years ago
params.txt	Add params for puncture level	14 days ago
params_expensive.txt	Added parameters for choosing extraction modes and whether to write t...	9 months ago
params_very_cheap.txt	Allow plot files to be turned off	12 months ago

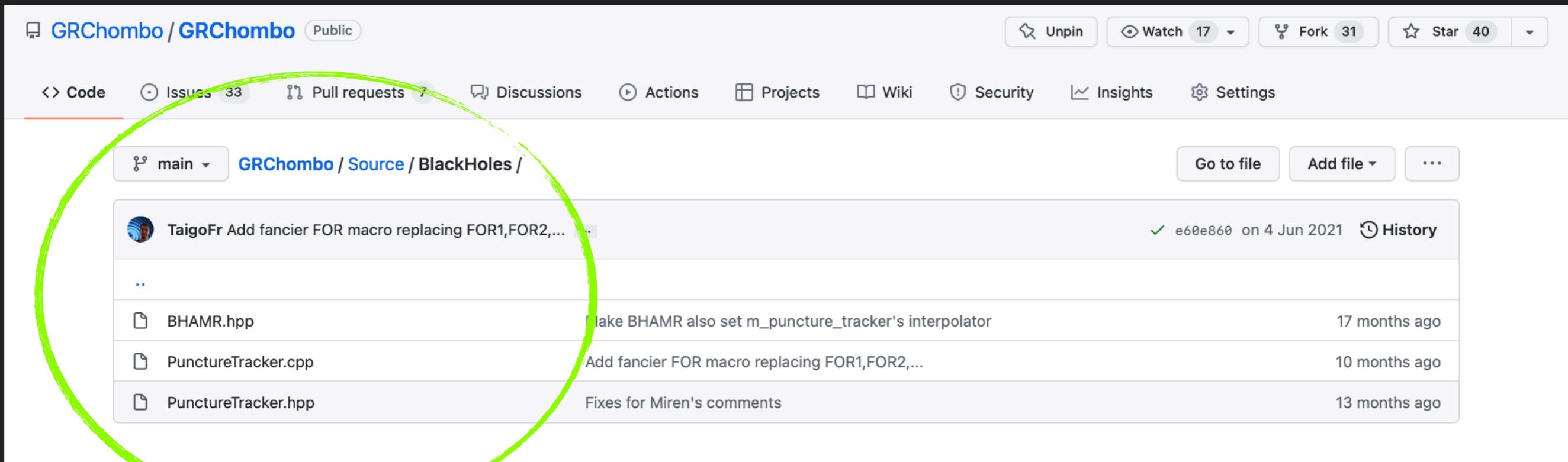
WHERE ARE THE KEY BINARYBH FILES?

The screenshot shows the GitHub repository for GRChombo, specifically the 'Source / InitialConditions / BlackHoles' directory. The commit history is displayed, showing a commit by KAClough on 12 Feb 2018. The files listed in the commit are:

File	Description	Time
BinaryBH.hpp	Adding the binary black hole initial conditions.	2 years ago
BinaryBH.impl.hpp	Separating out vars and tidy up comments	2 years ago
BoostedBH.hpp	Adding the binary black hole initial conditions.	2 years ago
BoostedBH.impl.hpp	Adding the binary black hole initial conditions.	2 years ago
KerrBH.hpp	Separating out vars and tidy up comments	2 years ago
KerrBH.impl.hpp	Separating out vars and tidy up comments	2 years ago

NB: These files are in "Source" as they are likely to be used for many examples **without modification**. If you are using something very problem specific, you may want to put it in the Example folder.

WHERE ARE THE KEY BINARYBH FILES?



The screenshot shows the GitHub repository page for GRChombo / GRChombo. The breadcrumb navigation indicates the current path is `main` / `GRChombo` / `Source` / `BlackHoles`. A green circle highlights the commit history for this directory. The commit history shows the following files and their commit messages:

File	Commit Message	Time Ago
..	..	
BHAMR.hpp	Make BHAMR also set m_puncture_tracker's interpolator	17 months ago
PunctureTracker.cpp	Add fancier FOR macro replacing FOR1, FOR2, ...	10 months ago
PunctureTracker.hpp	Fixes for Miren's comments	13 months ago

NB: These files are in "Source" as they are likely to be used for many examples **without modification**. If you are using something very problem specific, you may want to put it in the Example folder.

STRUCTURE OF AMR

- ▶ Does setup (for restart or using initial data) and runs evolution
- ▶ Knows about all of the levels, each function generally cycles through each level from coarse to fine
- ▶ Contains hooks for physics class actions (occurring in **GRAMRLevel** / **BinaryBHLevel**)



E.G. AMR::RUN() DOES THE EVOLUTION

```
769 //-----  
770 // go baby go  
771 void AMR::run(Real a_max_time, int a_max_step)  
772 {  
773     CH_TIME("AMR::run");  
774  
775     CH_assert(isDefined());  
776     CH_assert(isSetUp());  
777  
778     if (m_verbosity >= 3)  
779     {  
780         pout() << "AMR::coarseTimeStep:" << endl;  
781         pout() << "max_time = " << a_max_time << endl;  
782         pout() << "max_step = " << a_max_step << endl;  
783     }  
784
```

This function runs the evolution, after the amr object has been defined and set up (which happens in the Main_BinaryBH.cpp file)

```
1810 // write physics class header data  
1811 m_amrlevels[0]->writePlotHeader(handle);  
1812  
1813 // write physics class per-level data  
1814 for (int level = 0; level <= m_finest_level; ++level)  
1815 {  
1816     m_amrlevels[level]->prePlotLevel();  
1817     m_amrlevels[level]->writePlotLevel(handle);  
1818 }  
1819
```

Call the same function on each AMRLevel in turn

This is a hook we added to manipulate data pre plots

STRUCTURE OF GRAMR

- ▶ Inherits all functionality from AMR
- ▶ Adds in our GR specific tools, e.g. AMRInterpolator*
- ▶ Only contains things that happen globally across the grid, so actually not that much. Most actions are local to a level.



(*OK, so this is not GR specific, but it did not exist in Chombo so we built it, and now it lives in GRChombo because we don't want to hack the Chombo code too much.)

STRUCTURE OF BHAMR

- ▶ Inherits all functionality from GRAMR
- ▶ Adds in BBH specific tools, e.g. Puncture Tracking
- ▶ Again not that long!



STRUCTURE OF BHAMR

```
5
6 #ifndef BHAMR_HPP_
7 #define BHAMR_HPP_
8
9 #include "GRAMR.hpp"
10 #include "PunctureTracker.hpp"
11
12 /// A child of Chombo's AMR class to interface with tools which require
13 /// access to the whole AMR hierarchy, and those of GRAMR
14 /**
15  * This object inherits from GRAMR and adds tools required for BH spacetimes
16  */
17 class BHAMR : public GRAMR
18 {
19     public:
20         PunctureTracker m_puncture_tracker;
21
22         BHAMR() {}
23
24         void set_interpolator(AMRInterpolator<Lagrange<4>> *a_interpolator) override
25         {
26             GRAMR::set_interpolator(a_interpolator);
27             m_puncture_tracker.set_interpolator(a_interpolator);
28         }
29 };
30
31 #endif /* BHAMR_HPP_ */
```

Inheritance of GRAMR functions (and so also AMR)

BH puncture tracking control added

Setting interpolator now needs to also set it for puncture tracking, (note this **overrides** the one in GRAMR)

ALL THIS COMES TOGETHER IN MAIN_BINARYBH.CPP

```
int runGRChombo(int argc, char *argv[])
{
    // Load the parameter file and construct the SimulationParameter class
    // To add more parameters edit the SimulationParameters file.
    char *in_file = argv[1];
    GRParmParse pp(argc - 2, argv + 2, NULL, in_file);
    SimulationParameters sim_params(pp);

    if (sim_params.just_check_params)
        return 0;

#ifdef USE_TWOPUNCTURES
    TPAMR bh_amr;
    bh_amr.set_two_punctures_parameters(sim_params.tp_params);
    // Run TwoPunctures solver
    bh_amr.m_two_punctures.Run();
#else
    BHAMR bh_amr;
#endif

    // must be before 'setupAMRObject' to define punctures for tagging criteria
    if (sim_params.track_punctures)
    {
        // the tagging criterion used in this example means that the punctures
        // should be on the max level but let's fill ghosts on the level below
        // too just in case
        int puncture_tracker_min_level = sim_params.max_level - 1;
        bh_amr.m_puncture_tracker.initial_setup(
            {sim_params.bh1_params.center, sim_params.bh2_params.center},
            "punctures", sim_params.data_path, puncture_tracker_min_level);
    }
}
```

Make a BHAMR object

Setup puncture tracking which lives in BHAMR

ALL THIS COMES TOGETHER IN MAIN_BINARYBH.CPP

```
// The line below selects the problem that is simulated
// (To simulate a different problem, define a new child of AMRLevel
// and an associated LevelFactory)
DefaultLevelFactory<BinaryBHLevel> binary_bh_level_fact(bh_amr, sim_params);
setupAMRObject(bh_amr, binary_bh_level_fact);

// call this after amr object setup so grids known
// and need it to stay in scope throughout run
AMRInterpolator<Lagrange<4>> interpolator(
    bh_amr, sim_params.origin, sim_params.dx, sim_params.boundary_params,
    sim_params.verbosity);
bh_amr.set_interpolator(
    &interpolator); // also sets puncture_tracker interpolator

// must be after interpolator is set
if (sim_params.track_punctures)
    bh_amr.m_puncture_tracker.restart_punctures();

bh_amr.run(sim_params.stop_time, sim_params.max_steps);

auto now = Clock::now();
auto duration = std::chrono::duration_cast<Minutes>(now - start_time);
pout() << "Total simulation time (mins): " << duration.count() << ".\n";

bh_amr.conclude();

CH_TIMER_REPORT(); // Report results when running with Chombo timers.

return 0;
```

Setup using AMR functions

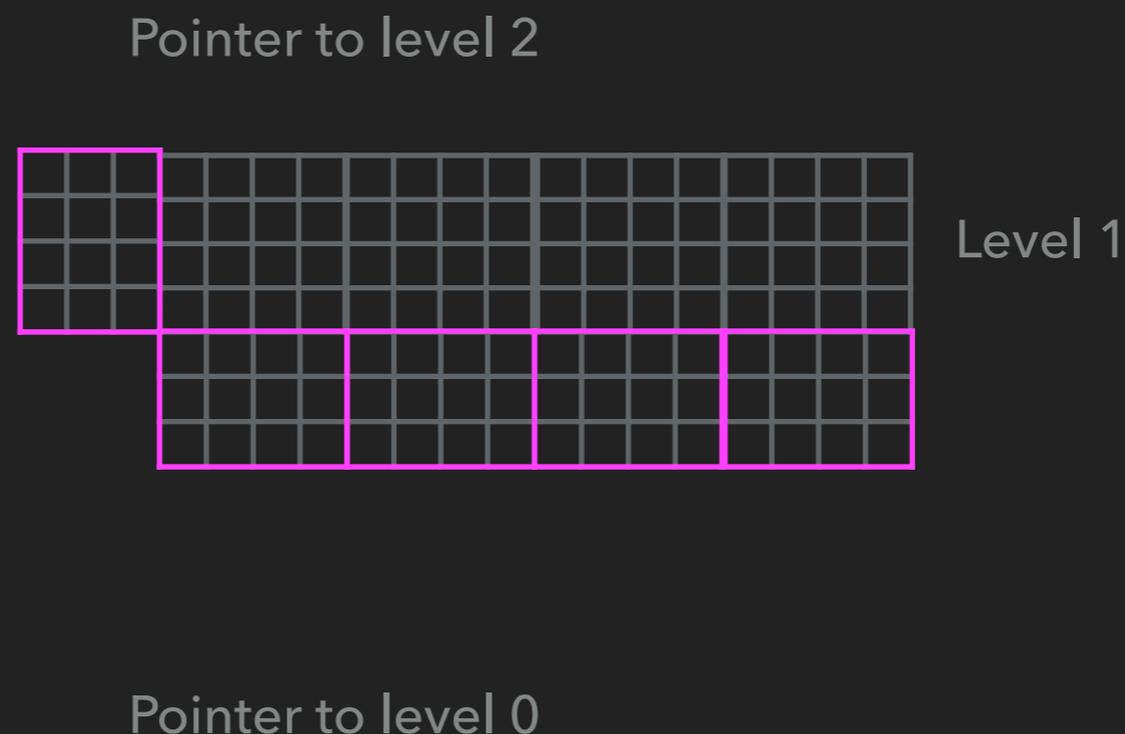
Setup interpolator using function in BHAMR

AMR run function

AMR conclude function

STRUCTURE OF AMRLEVEL

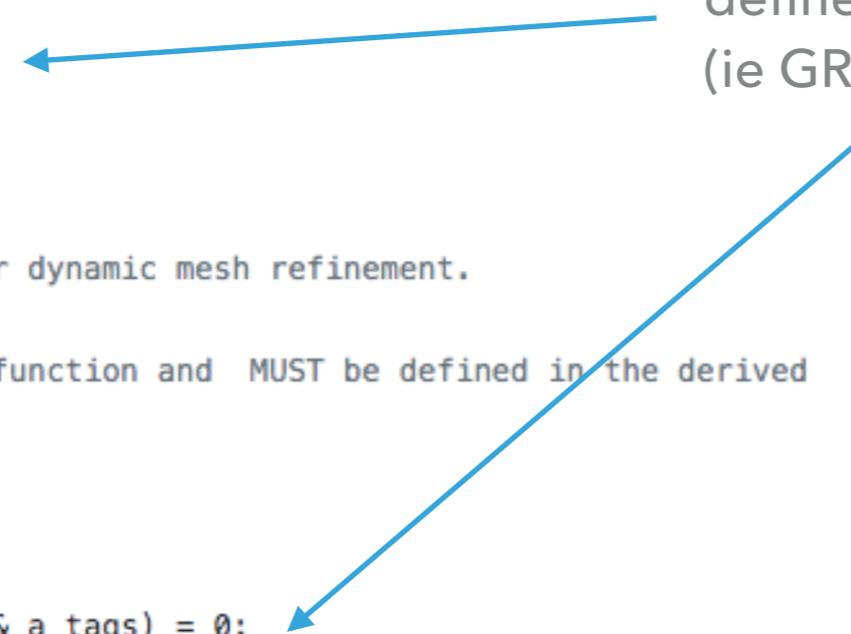
- ▶ Knows about its own level data, and has a pointer to the coarser and finer levels above and below it
- ▶ Abstract base class to be overwritten by a “physics class”
i.e. **GRAMRLevel** / **BinaryBHLevel**



STRUCTURE OF AMRLEVEL

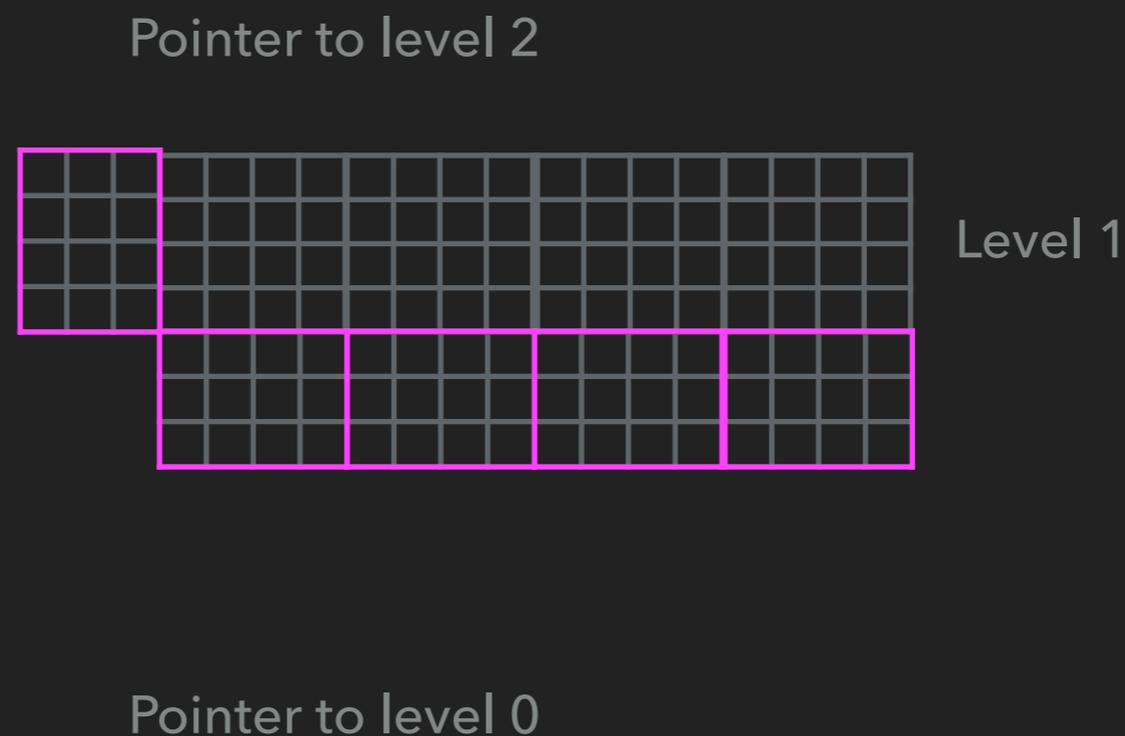
```
140  /**
141     Things to do after advancing this level by one time step.
142
143     This is a pure virtual function and MUST be defined in the derived
144     class.
145
146  */
147  virtual
148     void postTimeStep() = 0;
149
150  ///
151  /**
152     Creates tagged cells for dynamic mesh refinement.
153
154     This is a pure virtual function and MUST be defined in the derived
155     class.
156
157  */
158  virtual
159     void tagCells(IntVectSet& a_tags) = 0;
160
161  ///
162  /**
163     Creates tagged cells for mesh refinement at initialization.
164
165     This is a pure virtual function and MUST be defined in the derived
166     class.
```

Virtual functions which must be defined in the physics class (ie GRAMRLevel / BinaryBHLevel)



STRUCTURE OF GRAMRLEVEL

- ▶ Inherits from `AMRLevel` and overwrites virtual functions where these are common to most GR simulations
- ▶ Contains hooks for example specific actions (occurring in `BinaryBHLevel`, prefixed by "specific")



STRUCTURE OF GRAMRLEVEL

```
163 // things to do after a timestep
164 void GRAMRLevel::postTimeStep()
165 {
166     if (m_verbosity)
167         pout() << "GRAMRLevel::postTimeStep " << m_level << endl;
168
169     if (m_finer_level_ptr != nullptr)
170     {
171         GRAMRLevel *finer_gr_amr_level_ptr = gr_cast(m_finer_level_ptr);
172         finer_gr_amr_level_ptr->m_coarse_average.averageToCoarse(
173             m_state_new, finer_gr_amr_level_ptr->m_state_new);
174         // Synchronise times to avoid floating point errors for finer levels
175         finer_gr_amr_level_ptr->time(m_time);
176     }
177
178     specificPostTimeStep();
179
180     // enforce symmetric BCs - this is required after the averaging
181     // and potentially after specificPostTimeStep actions
182     fillBdyGhosts(m_state_new);
183
184     if (m_verbosity)
185         pout() << "GRAMRLevel::postTimeStep " << m_level << " finished" << endl;
186 }
187
188 // create tags
189 void GRAMRLevel::tagCells(IntVectSet &a_tags)
190 {
```

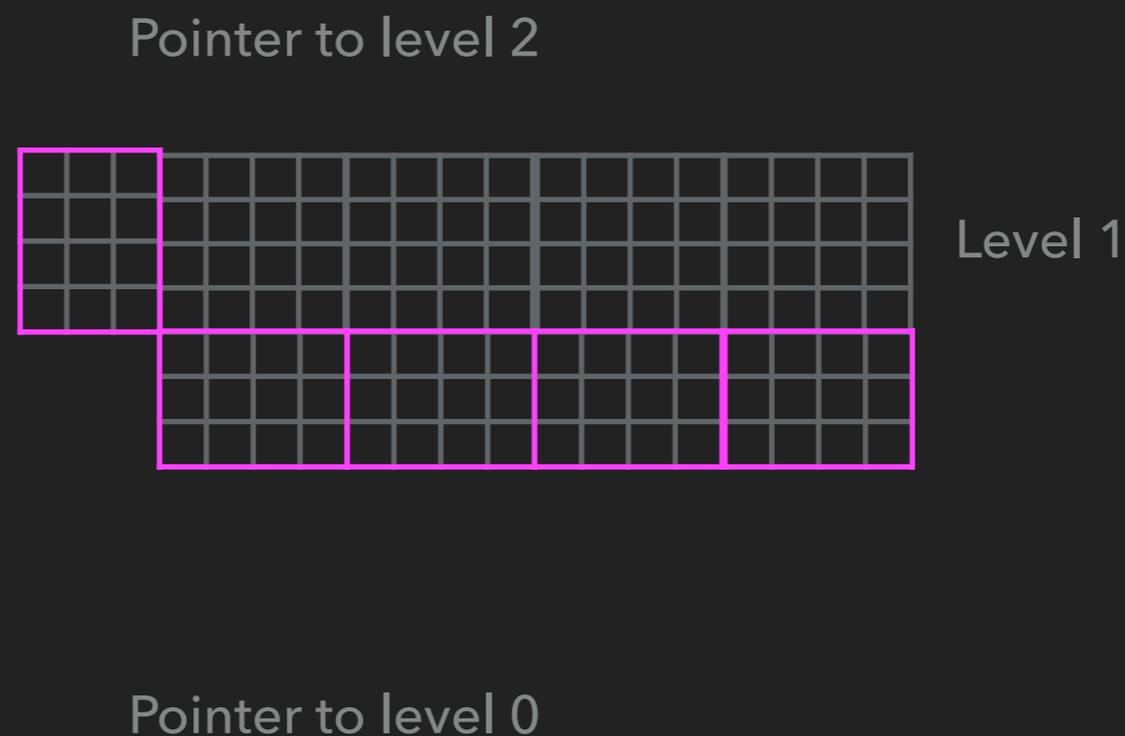
Overrides the virtual function in
AMRLevel

Communication with finer/coarser
level via pointers, e.g. here for the
overwriting of underlying coarser
cells

Hook for example specific actions
e.g. in BinaryBHLevel

STRUCTURE OF BINARYBHLEVEL

- ▶ Inherits all functionality from GRAMRLevel, overwrites virtual functions where these are specific to BinaryBH example
- ▶ Adds in required BBH specific functions via the hooks like `specificPostTimeStep()`



STRUCTURE OF BINARYBHLEVEL

```
138 void BinaryBHLevel::specificPostTimeStep()
139 {
140     CH_TIME("BinaryBHLevel::specificPostTimeStep");
141     if (m_p.activate_extraction == 1)
142     {
143         // Populate the Weyl Scalar values on the grid
144         fillAllGhosts();
145         BoxLoops::loop(Weyl4(m_p.extraction_params.extraction_center, m_dx),
146                       m_state_new, m_state_new, EXCLUDE_GHOST_CELLS);
147
148         // Do the extraction on the min extraction level
149         if (m_level == m_p.extraction_params.min_extraction_level)
150         {
151             CH_TIME("WeylExtraction");
152             // Now refresh the interpolator and do the interpolation
153             m_gr_amr.m_interpolator->refresh();
154             WeylExtraction my_extraction(m_p.extraction_params, m_dt, m_time,
155                                         m_restart_time);
156             my_extraction.execute_query(m_gr_amr.m_interpolator);
157         }
158     }
159
160     // do puncture tracking on requested level
161     if (m_p.track_punctures == 1 && m_level == m_p.puncture_tracking_level)
162     {
163         CH_TIME("PunctureTracking");
164         // only do the write out for every coarsest level timestep
```

Here is the hook we saw in GRAMRLevel!

After each timestep calculate the Weyl scalar (happens on all levels)

m_level tells us which level we are so conditional on this restricts action to that level

KEY FUNCTIONS THAT WE SPECIFY IN BHBINARYLEVEL

function	required / optional	Comment
initialdata()	required	define vars on initial grid
computeTaggingCriterion()	required	criterion for refinement
specificEvalRHS()	required	define evolution dvar/dt
specificPostTimestep()	optional	after level completes dt update
specificAdvance()	optional	happens in RK4 substeps
specificUpdateODE()	optional	happens in RK4 substeps
postRestart()	optional	done after checkpoint restart
preCheckpointLevel()	optional	before output checkpoint
prePlotLevel()	optional	before output plot file

FIND THIS AND MORE DETAILS IN THE WIKI!

The screenshot shows a GitHub repository page for 'GRChombo / GRChombo'. The repository is public and has 17 watchers, 31 forks, and 40 stars. The navigation bar includes links for Code, Issues (33), Pull requests (7), Discussions, Actions, Projects, Wiki (highlighted), Security, and Insights. The main heading is 'Structure of the code (aka where is everything?)', edited by K Clough on 15 Nov 2021. The page content explains that GRChombo is a 'big code' and provides hints on navigating the code structure. It mentions a PDF guide and useful resources. A 'Hierarchy of GRChombo' section follows, stating the code has three main levels. A 'Contents' sidebar on the right lists links for Home, Capabilities, License, Citation, Doxygen documentation, Getting started, Prerequisites, Compiling Chombo, Compiling GRChombo, Running examples, and Running the BinaryBH example.

GRChombo / GRChombo Public

Unpin Watch 17 Fork 31 Star 40

<> Code Issues 33 Pull requests 7 Discussions Actions Projects Wiki Security Insights

Structure of the code (aka where is everything?)

Edit New Page

K Clough edited this page on 15 Nov 2021 · 23 revisions

Yes, we know, GRChombo is a *big* code. At first the number of files will seem overwhelming, but with time you will start to learn where to find things and the structure will make (some) sense.

On this page we provide some hints on how to find your way around the code, but in the end you just have to dive in and learn as you go.

A useful pdf guide on this topic (with nicer pictures) from the latest GRChombo training day can be found in [Useful resources](#). One should also look at the guides on C++ classes, inheritance and templating, which are used extensively in the code - some basic knowledge of these concepts is assumed below.

Hierarchy of GRChombo

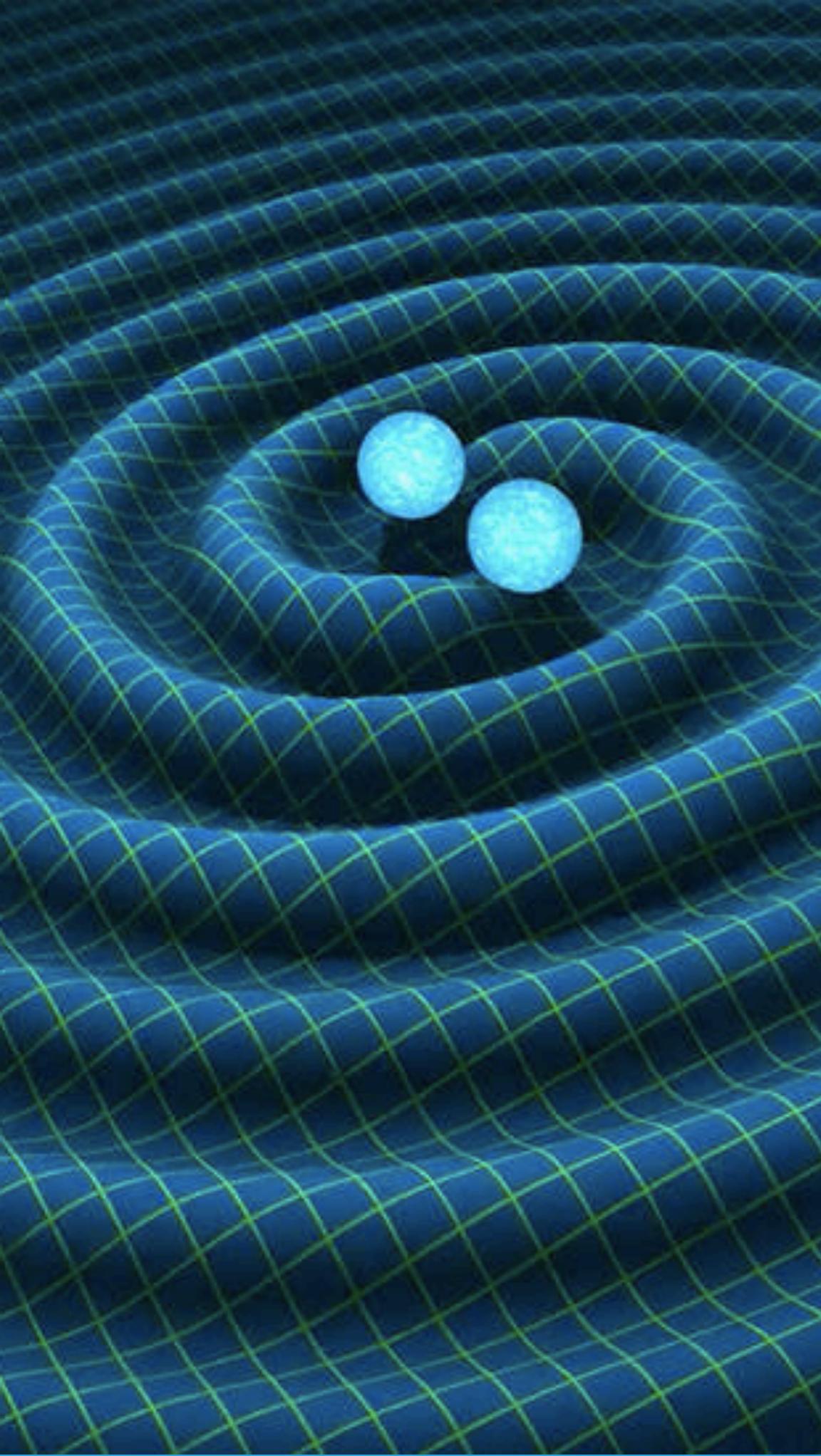
The code is designed to have 3 main levels in its hierarchy, as follows:

1. Specific Example related files, e.g. for BinaryBH - specific actions relevant to the BinaryBH example - key classes include BHAMR, BinaryBHLevel, SimulationParameters. Also important are the namespaces UserVariables and DiagnosticVariables in the BinaryBH examples folder and BinaryBH (the initial data).

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QUESTIONS?