**GW** Analysis Tools

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## **Gravitational Waves Analysis Tools**

A suite of analysis tools useful for gravitational wave science. All code is written in C++, with some of the interface classes wrapped in Cython to allow for python-access.

### 1.1 Compatibility

Known to work with gcc/g++-7

Known to work with gcc/g++-9

Need nvcc - known to work with v9.1 of CUDA

### 1.2 Required Software

Required non-standard C libraries: FFTW3 ADOL-C GSL CUDA

Required non-standard Python packages: Cython

Required non-standard packages for documentation: Doxygen

### 1.3 Current Development

NOTE: currently using static parameters to share data between threads for mcmc\_gw.cpp. This could cause issues when running multiple samplers at the same time. Investigating further.

To do:

Change MCMC\_MH to use the more general threadPool class instead of a custom threadpool, incorporate job class and comparator

#### 1.4 Installation

For proper compilation, update or create the enviornment variables CPATH, LIBRARY\_PATH, and LD\_LIBRARY — \_PATH, which should point to header files and lib files, respectively. Specifically, these variables should point to the above libraries.

Also, the PYTHONPATH environment variables must point to /gw\_analysis\_tools\_py/src because I can't figure how to get this shit to work.

In the root directory of the project, run 'make' to compile source files, create the library file and create the cython modules, and create the documentation.

To just create C++/C files, run 'make c'.

Run 'make test' to build a test program that will create an executable.

## 1.5 Supported Functionality

#### 1.5.1 Waveform Generation

IMRPhenomD, IMRPhenomPv2

#### 1.5.2 Modified Gravity

ppE\_IMRPhenomD\_Inspiral ppE\_IMRPhenomD\_IMR ppE\_IMRPhenomPv2\_Inspiral ppE\_IMRPhenomPv2\_IMR

#### 1.5.3 Fisher Analysis

utilizes the above waveform templates

#### 1.5.4 MCMC Routines

Has a generic MCMC sampler, MCMC\_MH, that utilizes gaussian steps, differential evolution steps, and Fisher informed steps. Includes wrapping MCMC\_MH\_GW for GW specific sampling, currently only for one detector.

Includes log likelihood caclulation for implementation in other samplers.

### 1.6 Usage

#### 1.6.1 Environment variables

The environment variable PYTHONPATH should include the directory \$(PROJECT\_DIR)

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#### 1.6.2 Include

To include header files, use -I\$(PROJECT\_DIRECTORY)/include

#### 1.6.3 Link

To link object files, use -L\$(PROJECT\_DIRECTORY)/lib -lgwanalysistools

For Cuda code: use -lcuda -lcudart

For Cuda, may need to link to /usr/local/cuda/lib64/ (or wherever this library is on your machine)

#### 1.6.4 Python Importable Code

Two modules currently available:

1.6.4.1 gw\_analysis\_tools\_py.mcmc\_routines\_ext

Wraps the log\_likelihood functions in mcmc\_routines.cpp

1.6.4.2 gw\_analysis\_tools\_py.waveform\_generator\_ext

Wraps the fourier\_waveform function in waveform\_generator.cpp

Also contains the SNR calculation function

#### 1.6.4.3 Custom Waveforms

If adding waveforms and to have full accesibility:

Create class, using other waveforms as template – need interface to create full waveform (plus,cross polarization), and amplitude/phase

Add the option as a waveform to waveform\_generation.cpp, including the header file at the top of the waveform\_\cdot\ generation.cpp file

For autodiff Fishers – write the class as a template with double and adouble types for all variables. Then write the necessary fisher subroutines (see fisher file to determine whats necessary)

For numerical Fishers - write finite difference method, following the template of the previous waveforms

For MCMC sampling – write mcmc\_fisher\_wrapper and mcmc\_likelihood\_wrapper options and write any necessary initialization in MCMC\_MH\_GW

**Author** 

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# gw\_analysis\_tools

A suite of tools useful for doing statistical studies on gravitational wave science, including routines useful in  $MC \leftarrow MC$  studies, wave template generation, Fisher analysis, etc. Written in C++ and wrapped in Cython for access in Python.

6 gw\_analysis\_tools

# **Hierarchical Index**

## 3.1 Class Hierarchy

This inheritance list is sorted roughly, but not completely, alphabetically:

$alpha\_coeffs < T > \dots \dots$
Comparator
comparator_ac_fft
comparator_ac_serial
default_comp< jobtype >
epsilon_coeffs $<$ T $>$
fftw_outline
gen_params
GPUplan
$IMRPhenomD < T > \dots \dots$
IMRPhenomPv2< T >
ppE_IMRPhenomD_Inspiral< T >
dCS_IMRPhenomD< T >
$dCS_IMRPhenomD_log < T > \dots \dots$
$EdGB\_IMRPhenomD < T > \dots \dots$
$\label{eq:edgb_IMRPhenomD_log} EdGB\_IMRPhenomD\_log < T >  \dots \qquad \qquad$
ppE_IMRPhenomD_IMR <t>44</t>
$lambda\_parameters < T > \dots \dots$
sampler
$source\_parameters < T > \dots \dots$
sph_harm< T >
threaded_ac_jobs_fft
threaded_ac_jobs_serial
threadPool< jobtype, comparator >
ThreadPool
useful powers $\langle T \rangle$

8 Hierarchical Index

## **Class Index**

## 4.1 Class List

Here are the classes, structs, unions and interfaces with brief descriptions:

alpha_coeffs< T >
Comparator
Class to facilitate the comparing of chains for priority
comparator_ac_fft
Comparator to sort ac-jobs
comparator_ac_serial
Comparator to sort ac-jobs
$dCS_IMRPhenomD < T > 15$
$dCS\_IMRPhenomD\_log < T > \dots \dots$
default_comp< jobtype >
Default comparator for priority_queue in threadPool – no comparison
EdGB_IMRPhenomD< T >
$EdGB\_IMRPhenomD\_log < T > \dots \dots$
epsilon_coeffs< T >
fftw outline
gen_params
GPUplan
IMRPhenomD< T >
IMRPhenomPv2< T >
$lambda\_parameters < T > \dots \dots$
ppE_IMRPhenomD_IMR< T >
ppE_IMRPhenomD_Inspiral< T >
sampler
source parameters $\langle T \rangle$
sph_harm< T >
threaded_ac_jobs_fft
Class to contain spectral method jobs
threaded_ac_jobs_serial
Class to contain serial method jobs
threadPool< jobtype, comparator >
Class for creating a pool of threads to asynchronously distribute work
ThreadPool
useful_powers< T >
To speed up calculations within the for loops, we pre-calculate reoccuring powers of M*F and Pi,
since the pow() function is prohibatively slow

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# File Index

## 5.1 File List

Here is a list of all documented files with brief descriptions:

include/autocorrelation.h
include/autocorrelation_cuda.h
include/autocorrelation_cuda.hu
include/ <b>D_Z_Config.h</b>
include/detector_util.h
include/fisher.h
include/GWATConfig.h
include/IMRPhenomD.h
include/IMRPhenomP.h
include/mcmc_gw.h
include/mcmc_sampler.h
include/mcmc_sampler_internals.h
include/ppE_IMRPhenomD.h
include/threadPool.h
include/util.h
include/waveform_generator.h
include/waveform_generator_C.h
include/waveform_util.h
src/autocorrelation.cpp
src/autocorrelation_cuda.cu
src/detector_util.cpp
src/fisher.cpp
src/IMRPhenomD.cpp
src/IMRPhenomP.cpp
src/mcmc_gw.cpp
src/mcmc_sampler.cpp
src/mcmc_sampler_internals.cpp
src/ppE_IMRPhenomD.cpp
src/util.cpp
src/waveform_generator.cpp
src/wayeform_util_cpp

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## **Class Documentation**

## 6.1 alpha\_coeffs < T > Struct Template Reference

#### **Public Attributes**

- T coeff1
- T coeff2
- T coeff3
- T coeff4
- T coeff5

The documentation for this struct was generated from the following file:

• include/IMRPhenomP.h

### 6.2 Comparator Class Reference

Class to facilitate the comparing of chains for priority.

#### **Public Member Functions**

• bool operator() (int i, int j)

### 6.2.1 Detailed Description

Class to facilitate the comparing of chains for priority.

3 levels of priority: 0 (high) 1 (default) 2 (low)

The documentation for this class was generated from the following file:

• src/mcmc\_sampler.cpp

### 6.3 comparator\_ac\_fft Class Reference

comparator to sort ac-jobs

```
#include <autocorrelation.h>
```

#### **Public Member Functions**

• bool operator() (threaded\_ac\_jobs\_fft t, threaded\_ac\_jobs\_fft k)

#### 6.3.1 Detailed Description

comparator to sort ac-jobs

Starts with the longest jobs, then works down the list

The documentation for this class was generated from the following file:

• include/autocorrelation.h

### 6.4 comparator\_ac\_serial Class Reference

comparator to sort ac-jobs

```
#include <autocorrelation.h>
```

#### **Public Member Functions**

• bool operator() (threaded\_ac\_jobs\_serial t, threaded\_ac\_jobs\_serial k)

#### 6.4.1 Detailed Description

comparator to sort ac-jobs

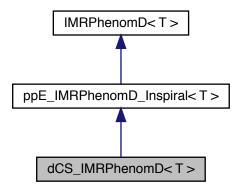
Starts with the longest jobs, then works down the list

The documentation for this class was generated from the following file:

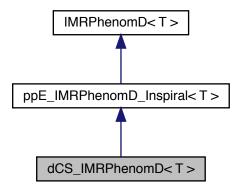
• include/autocorrelation.h

## 6.5 dCS\_IMRPhenomD < T > Class Template Reference

Inheritance diagram for dCS\_IMRPhenomD< T >:



Collaboration diagram for dCS\_IMRPhenomD< T >:



#### **Public Member Functions**

virtual int construct\_waveform (T \*frequencies, int length, std::complex< T > \*waveform, source\_parameters<</li>
 T > \*params)

Constructs the waveform as outlined by.

- virtual T dCS\_phase\_mod (source\_parameters< T > \*param)
- virtual T dCS\_phase\_factor (source\_parameters < T > \*param)
- virtual int construct\_amplitude (T \*frequencies, int length, T \*amplitude, source\_parameters< T > \*params)

  Constructs the Amplitude as outlined by IMRPhenomD.
- virtual int construct\_phase (T \*frequencies, int length, T \*phase, source\_parameters< T > \*params)

  Constructs the Phase as outlined by IMRPhenomD.

#### 6.5.1 Member Function Documentation

#### 6.5.1.1 construct\_amplitude()

Constructs the Amplitude as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output amplitude, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

#### 6.5.1.2 construct\_phase()

Constructs the Phase as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output phase, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

#### 6.5.1.3 construct\_waveform()

Constructs the waveform as outlined by.

arguments: array of frequencies, length of that array, a complex array for the output waveform, and a source\_parameters structure

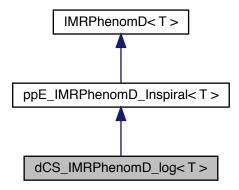
Reimplemented from IMRPhenomD< T>.

The documentation for this class was generated from the following files:

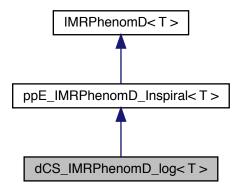
- include/ppE\_IMRPhenomD.h
- src/ppE\_IMRPhenomD.cpp

## 6.6 dCS\_IMRPhenomD\_log< T > Class Template Reference

Inheritance diagram for dCS\_IMRPhenomD\_log< T >:



Collaboration diagram for dCS\_IMRPhenomD\_log< T >:



#### **Public Member Functions**

virtual int construct\_waveform (T \*frequencies, int length, std::complex< T > \*waveform, source\_parameters<</li>
 T > \*params)

Constructs the waveform as outlined by.

- virtual T dCS\_phase\_mod (source\_parameters< T > \*param)
- virtual T dCS\_phase\_factor (source\_parameters < T > \*param)
- virtual int construct\_amplitude (T \*frequencies, int length, T \*amplitude, source\_parameters< T > \*params)

  Constructs the Amplitude as outlined by IMRPhenomD.
- virtual int construct\_phase (T \*frequencies, int length, T \*phase, source\_parameters< T > \*params)

  Constructs the Phase as outlined by IMRPhenomD.

#### 6.6.1 Member Function Documentation

#### 6.6.1.1 construct\_amplitude()

Constructs the Amplitude as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output amplitude, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

#### 6.6.1.2 construct\_phase()

Constructs the Phase as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output phase, and a source\_parameters structure

Reimplemented from IMRPhenomD< T>.

#### 6.6.1.3 construct\_waveform()

Constructs the waveform as outlined by.

arguments: array of frequencies, length of that array, a complex array for the output waveform, and a source\_parameters structure

Reimplemented from IMRPhenomD< T>.

The documentation for this class was generated from the following files:

- include/ppE\_IMRPhenomD.h
- src/ppE\_IMRPhenomD.cpp

## 6.7 default\_comp < jobtype > Class Template Reference

Default comparator for priority\_queue in threadPool – no comparison.

#include <threadPool.h>

#### **Public Member Functions**

• bool operator() (jobtype j, jobtype k)

#### 6.7.1 Detailed Description

```
template < class jobtype > class default_comp < jobtype >
```

Default comparator for priority\_queue in threadPool – no comparison.

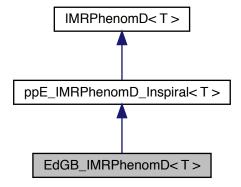
First in first out, not sorting

The documentation for this class was generated from the following file:

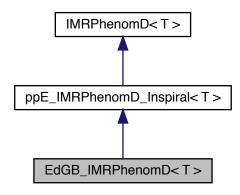
• include/threadPool.h

## 6.8 EdGB\_IMRPhenomD < T > Class Template Reference

Inheritance diagram for EdGB\_IMRPhenomD< T >:



Collaboration diagram for EdGB\_IMRPhenomD< T >:



#### **Public Member Functions**

virtual int construct\_waveform (T \*frequencies, int length, std::complex < T > \*waveform, source\_parameters < T > \*params)

Constructs the waveform as outlined by.

- virtual T EdGB\_phase\_mod (source\_parameters < T > \*param)
- virtual T EdGB phase factor (source parameters< T > \*param)
- virtual int construct\_amplitude (T \*frequencies, int length, T \*amplitude, source\_parameters< T > \*params)

  Constructs the Amplitude as outlined by IMRPhenomD.
- virtual int construct\_phase (T \*frequencies, int length, T \*phase, source\_parameters< T > \*params)

  Constructs the Phase as outlined by IMRPhenomD.

#### 6.8.1 Member Function Documentation

#### 6.8.1.1 construct\_amplitude()

Constructs the Amplitude as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output amplitude, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

#### 6.8.1.2 construct\_phase()

Constructs the Phase as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output phase, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

#### 6.8.1.3 construct\_waveform()

Constructs the waveform as outlined by.

arguments: array of frequencies, length of that array, a complex array for the output waveform, and a source\_parameters structure

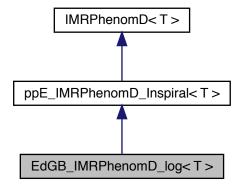
Reimplemented from IMRPhenomD< T >.

The documentation for this class was generated from the following files:

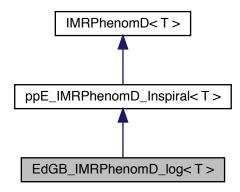
- include/ppE\_IMRPhenomD.h
- src/ppE\_IMRPhenomD.cpp

## 6.9 EdGB\_IMRPhenomD\_log < T > Class Template Reference

Inheritance diagram for EdGB\_IMRPhenomD\_log< T >:



Collaboration diagram for EdGB\_IMRPhenomD\_log< T >:



#### **Public Member Functions**

virtual int construct\_waveform (T \*frequencies, int length, std::complex < T > \*waveform, source\_parameters < T > \*params)

Constructs the waveform as outlined by.

- virtual T EdGB\_phase\_mod (source\_parameters < T > \*param)
- virtual T EdGB phase factor (source parameters< T > \*param)
- virtual int construct\_amplitude (T \*frequencies, int length, T \*amplitude, source\_parameters< T > \*params)

  Constructs the Amplitude as outlined by IMRPhenomD.
- virtual int construct\_phase (T \*frequencies, int length, T \*phase, source\_parameters< T > \*params)

  Constructs the Phase as outlined by IMRPhenomD.

#### 6.9.1 Member Function Documentation

#### 6.9.1.1 construct\_amplitude()

Constructs the Amplitude as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output amplitude, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

#### 6.9.1.2 construct\_phase()

Constructs the Phase as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output phase, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

## 6.9.1.3 construct\_waveform()

Constructs the waveform as outlined by.

arguments: array of frequencies, length of that array, a complex array for the output waveform, and a source\_parameters structure

Reimplemented from IMRPhenomD< T >.

The documentation for this class was generated from the following files:

- include/ppE\_IMRPhenomD.h
- src/ppE\_IMRPhenomD.cpp

# 6.10 epsilon\_coeffs < T > Struct Template Reference

**Public Attributes** 

- T coeff1
- T coeff2
- T coeff3
- T coeff4
- T coeff5

The documentation for this struct was generated from the following file:

• include/IMRPhenomP.h

# 6.11 fftw\_outline Struct Reference

## **Public Attributes**

- fftw\_complex \* in
- fftw\_complex \* out
- fftw\_plan p

The documentation for this struct was generated from the following file:

· include/util.h

# 6.12 gen\_params Struct Reference

## **Public Attributes**

- double mass1
- double mass2
- · double Luminosity\_Distance
- double spin1 [3]
- double spin2 [3]
- double phic =0
- double tc =0
- int \* bppe
- double \* betappe
- int Nmod
- double incl\_angle
- · double theta
- · double phi
- bool NSflag
- double f\_ref =0
- double phiRef =0
- double thetaJN = -1
- double alpha0 = 0
- double zeta\_polariz = 0
- double **phi\_aligned** = 0
- double chil = 0
- double chip = 0
- bool sky\_average
- gsl\_spline \* **Z\_DL\_spline\_ptr** =NULL
- gsl\_interp\_accel \* Z\_DL\_accel\_ptr = NULL
- std::string cosmology ="PLANCK15"

## 6.12.1 Member Data Documentation

```
6.12.1.1 betappe
double* gen_params::betappe
ppE coefficient for the phase modification - vector for multiple modifications
6.12.1.2 bppe
int* gen_params::bppe
ppE b parameter (power of the frequency) - vector for multiple modifications
6.12.1.3 f_ref
double gen_params::f_ref =0
Reference frequency for PhenomPv2
6.12.1.4 incl_angle
double gen_params::incl_angle
*angle between angular momentum and the total momentum
6.12.1.5 Luminosity_Distance
double gen_params::Luminosity_Distance
Luminosity distance to the source
6.12.1.6 mass1
double gen_params::mass1
mass of the larger body in Solar Masses
6.12.1.7 mass2
double gen_params::mass2
mass of the smaller body in Solar Masses
6.12.1.8 Nmod
int gen_params::Nmod
```

Number of phase modificatinos

```
6.12.1.9 NSflag
bool gen_params::NSflag
BOOL flag for early termination of NS binaries
6.12.1.10 phic
double gen_params::phic =0
coalescence phase of the binary
6.12.1.11 spin1
double gen_params::spin1[3]
Spin vector of the larger mass [Sx,Sy,Sz]
6.12.1.12 spin2
double gen_params::spin2[3]
Spin vector of the smaller mass [Sx,Sy,Sz]
6.12.1.13 tc
double gen_params::tc =0
coalescence time of the binary
6.12.1.14 theta
double gen_params::theta
spherical angles for the source location relative to the detector
The documentation for this struct was generated from the following file:
```

• include/util.h

# 6.13 GPUplan Struct Reference

#### **Public Attributes**

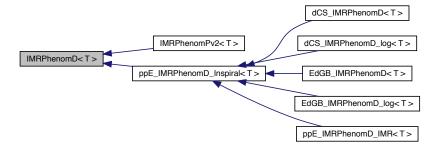
- · int device\_id
- double \* device data
- double \* host\_data
- int \* host\_lag
- int \* device\_lag
- int \* device\_lags
- int \* initial\_lag
- · cudaStream t stream

The documentation for this struct was generated from the following file:

• include/autocorrelation\_cuda.hu

# 6.14 IMRPhenomD < T > Class Template Reference

Inheritance diagram for IMRPhenomD< T >:



## **Public Member Functions**

- virtual void **fisher\_calculation** (double \*frequency, int length, gen\_params \*parameters, double \*\*amplitude\_deriv, double \*\*phase\_deriv, double \*amplitude, int \*amp\_tapes, int \*phase\_tapes)
- virtual void change\_parameter\_basis (T \*old\_param, T \*new\_param, bool sky\_average)

Convience method to change parameter basis between common Fisher parameters and the intrinsic parameters of IMRPhenomD.

• virtual void construct\_amplitude\_derivative (double \*frequencies, int length, int dimension, double \*\*amplitude derivative, source parameters< double > \*input params, int \*tapes=NULL)

Construct the derivative of the amplitude for a given source evaluated by the given frequency.

virtual void construct\_phase\_derivative (double \*frequencies, int length, int dimension, double \*\*phase\_←
derivative, source\_parameters< double > \*input\_params, int \*tapes=NULL)

Construct the derivative of the phase for a given source evaluated by the given frequency.

• virtual void amplitude tape (source parameters< double > \*input params, int \*tape)

Creates the tapes for derivatives of the amplitude.

virtual void phase\_tape (source\_parameters < double > \*input\_params, int \*tape)
 Creates the tapes for derivatives of phase.

virtual int construct\_waveform (T \*frequencies, int length, std::complex < T > \*waveform, source\_parameters < T > \*params)

Constructs the waveform as outlined by.

 $\bullet \ \ \text{virtual std::complex} < \mathsf{T} > \\ \text{construct\_waveform (T frequency, source\_parameters} < \mathsf{T} > \\ *params) \\$ 

overloaded method to evaluate the waveform for one frequency instead of an array

- virtual int construct\_amplitude (T \*frequencies, int length, T \*amplitude, source\_parameters < T > \*params)

  Constructs the Amplitude as outlined by IMRPhenomD.
- virtual int construct\_phase (T \*frequencies, int length, T \*phase, source\_parameters < T > \*params)
   Constructs the Phase as outlined by IMRPhenomD.
- virtual T build\_amp (T f, lambda\_parameters< T > \*lambda, source\_parameters< T > \*params, useful\_powers< T > \*pows, T \*amp\_coeff, T \*deltas)

constructs the IMRPhenomD amplitude for frequency f

virtual T build\_phase (T f, lambda\_parameters< T > \*lambda, source\_parameters< T > \*params, useful\_powers< T > \*pows, T \*phase\_coeff)

constructs the IMRPhenomD phase for frequency f

virtual T assign\_lambda\_param\_element (source\_parameters < T > \*source\_param, int i)

Calculate the lambda parameters from Khan et al for element i.

virtual void assign\_lambda\_param (source\_parameters< T > \*source\_param, lambda\_parameters< T > \*lambda)

Wrapper for the Lambda parameter assignment that handles the looping.

virtual void precalc\_powers\_ins (T f, T M, useful\_powers< T > \*Mf\_pows)

Pre-calculate powers of Mf, to speed up calculations for the inspiral waveform (both amplitude and phase.

virtual void precalc\_powers\_PI (useful\_powers< T > \*PI\_pows)

Pre-calculate powers of pi, to speed up calculations for the inspiral phase.

virtual void precalc powers ins phase (T f, T M, useful powers < T > \*Mf pows)

Pre-calculate powers of Mf, to speed up calculations for the inspiral phase.

virtual void precalc\_powers\_ins\_amp (T f, T M, useful\_powers < T > \*Mf\_pows)

Pre-calculate powers of Mf, to speed up calculations for the inspiral amplitude.

virtual void assign pn amplitude coeff (source parameters < T > \*source param, T \*coeff)

Calculates the static PN coeffecients for the amplitude.

 $\bullet \ \ virtual \ void \ assign\_static\_pn\_phase\_coeff \ (source\_parameters < T > *source\_param, \ T \ *coeff) \\$ 

Calculates the static PN coeffecients for the phase - coeffecients 0,1,2,3,4,7.

- virtual void assign\_nonstatic\_pn\_phase\_coeff (source\_parameters < T > \*source\_param, T \*coeff, T f)
   Calculates the dynamic PN phase coefficients 5,6.
- virtual void assign\_nonstatic\_pn\_phase\_coeff\_deriv (source\_parameters< T > \*source\_param, T \*Dcoeff, T f)

Calculates the derivative of the dynamic PN phase coefficients 5,6.

virtual void post\_merger\_variables (source\_parameters < T > \*source\_param)

Calculates the post-merger ringdown frequency and dampening frequency.

virtual T fpeak (source\_parameters < T > \*params, lambda\_parameters < T > \*lambda)

Solves for the peak frequency, where the waveform transitions from intermediate to merger-ringdown.

virtual T amp\_ins (T f, source\_parameters < T > \*param, T \*pn\_coeff, lambda\_parameters < T > \*lambda, useful\_powers < T > \*pow)

Calculates the scaled inspiral amplitude A/A0 for frequency f with precomputed powers of MF and PI.

- $\bullet \ \ virtual\ T\ Damp\_ins\ (T\ f, source\_parameters < T > *param,\ T\ *pn\_coeff,\ lambda\_parameters < T > *lambda)$ 
  - Calculates the derivative wrt frequency for the scaled inspiral amplitude A/A0 for frequency f.
- virtual T phase\_ins (T f, source\_parameters < T > \*param, T \*pn\_coeff, lambda\_parameters < T > \*lambda, useful\_powers < T > \*pow)

Calculates the inspiral phase for frequency f with precomputed powers of MF and PI for speed.

virtual T Dphase\_ins (T f, source\_parameters< T > \*param, T \*pn\_coeff, lambda\_parameters< T > \*lambda)

Calculates the derivative of the inspiral phase for frequency f.

- virtual T amp\_mr (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the scaled merger-ringdown amplitude A/A0 for frequency f.
- virtual T phase\_mr (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the merger-ringdown phase for frequency f.
- virtual T Damp\_mr (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the derivative wrt frequency for the scaled merger-ringdown amplitude A/A0 for frequency f.
- virtual T Dphase\_mr (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the derivative of the merger-ringdown phase for frequency f.
- virtual T amp\_int (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda, T \*deltas)
   Calculates the scaled intermediate range amplitude A/A0 for frequency f.
- virtual T phase\_int (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the intermediate phase for frequency f.
- virtual T Dphase\_int (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the derivative of the intermediate phase for frequency f.
- virtual void phase\_connection\_coefficients (source\_parameters < T > \*param, lambda\_parameters < T > \*lambda, T \*pn\_coeffs)

Calculates the phase connection coefficients alpha{0,1} and beta{0,1}.

- virtual T calculate\_beta1 (source\_parameters < T > \*param, lambda\_parameters < T > \*lambda, T \*pn ←
   \_coeffs)
- virtual T calculate\_beta0 (source\_parameters < T > \*param, lambda\_parameters < T > \*lambda, T \*pn ←
   \_coeffs)
- $\bullet \ \ \ virtual\ T\ \textbf{calculate\_alpha1}\ (source\_parameters < T > *param,\ lambda\_parameters < T > *lambda)$
- virtual T calculate\_alpha0 (source\_parameters< T > \*param, lambda\_parameters< T > \*lambda)
- virtual void amp\_connection\_coeffs (source\_parameters < T > \*param, lambda\_parameters < T > \*lambda, T \*pn\_coeffs, T \*coeffs)

Solves for the connection coefficients to ensure the transition from inspiral to merger ringdown is continuous and smooth.

- virtual T calculate\_delta\_parameter\_0 (T f1, T f2, T f3, T v1, T v2, T v3, T dd1, T dd3, T M)
   Calculates the delta\_0 component.
- virtual T calculate\_delta\_parameter\_1 (T f1, T f2, T f3, T v1, T v2, T v3, T dd1, T dd3, T M)
   Calculates the delta 1 component.
- virtual T calculate\_delta\_parameter\_2 (T f1, T f2, T f3, T v1, T v2, T v3, T dd1, T dd3, T M)
   Calculates the delta\_2 component.
- virtual T calculate\_delta\_parameter\_3 (T f1, T f2, T f3, T v1, T v2, T v3, T dd1, T dd3, T M)

  Calculates the delta\_3 component.
- virtual T calculate\_delta\_parameter\_4 (T f1, T f2, T f3, T v1, T v2, T v3, T dd1, T dd3, T M)
   Calculates the delta\_4 component.

## 6.14.1 Member Function Documentation

#### 6.14.1.1 amp\_ins()

Calculates the scaled inspiral amplitude A/A0 for frequency f with precomputed powers of MF and PI.

return a T

additional argument contains useful powers of MF and PI in structure userful\_powers

#### 6.14.1.2 amp\_int()

Calculates the scaled intermediate range amplitude A/A0 for frequency f.

return a T

```
6.14.1.3 amp_mr()
```

Calculates the scaled merger-ringdown amplitude A/A0 for frequency f.

return a T

## 6.14.1.4 amplitude\_tape()

Creates the tapes for derivatives of the amplitude.

For efficiency in long runs of large sets of fishers, the tapes can be precomputed and reused

#### **Parameters**

input_params	source parameters structure of the desired source
tape	tape ids

Reimplemented in ppE\_IMRPhenomD\_IMR< T>, and ppE\_IMRPhenomD\_Inspiral< T>.

## 6.14.1.5 assign\_nonstatic\_pn\_phase\_coeff()

Calculates the dynamic PN phase coefficients 5,6.

f is in Hz

## 6.14.1.6 assign\_nonstatic\_pn\_phase\_coeff\_deriv()

Calculates the derivative of the dynamic PN phase coefficients 5,6.

f is in Hz

## 6.14.1.7 build\_amp()

constructs the IMRPhenomD amplitude for frequency f

arguments: numerical parameters from Khan et al lambda\_parameters structure, source\_parameters structure, useful\_powers<T> structure, PN parameters for the inspiral portions of the waveform, and the delta parameters for the intermediate region, numerically solved for using the amp\_connection\_coeffs function

## 6.14.1.8 build\_phase()

constructs the IMRPhenomD phase for frequency f

arguments: numerical parameters from Khan et al lambda\_parameters structure, source\_parameters structure, useful\_powers structure, PN parameters for the inspiral portions of the waveform

## 6.14.1.9 calculate\_delta\_parameter\_0()

Calculates the delta\_0 component.

Solved in Mathematica and imported to C

## 6.14.1.10 calculate\_delta\_parameter\_1()

Calculates the delta\_1 component.

Solved in Mathematica and imported to C

#### 6.14.1.11 calculate\_delta\_parameter\_2()

Calculates the delta\_2 component.

Solved in Mathematica and imported to C

## 6.14.1.12 calculate\_delta\_parameter\_3()

Calculates the delta\_3 component.

Solved in Mathematica and imported to C

## 6.14.1.13 calculate\_delta\_parameter\_4()

Calculates the delta\_4 component.

Solved in Mathematica and imported to C

#### 6.14.1.14 change\_parameter\_basis()

Convience method to change parameter basis between common Fisher parameters and the intrinsic parameters of IMRPhenomD.

Takes input array of old parameters and ouputs array of transformed parameters

#### **Parameters**

old_param	array of old params, order {A0, tc, phic, chirpmass, eta, spin1, spin2}
new_param	output new array: order {m1,m2,DL, spin1,spin2,phic,tc}

#### 6.14.1.15 construct\_amplitude()

Constructs the Amplitude as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output amplitude, and a source\_parameters structure

## Parameters

frequencies	T array of frequencies the waveform is to be evaulated at
length	integer length of the input array of frequencies and the output array
amplitude	output T array for the amplitude
params	Structure of source parameters to be initilized before computation

 $\label{eq:continuous} Reimplemented in \ EdGB\_IMRPhenomD < T>, \ EdGB\_IMRPhenomD\_log < T>, \ dCS\_IMRPhenomD < T>, \ and \ dCS\_IMRPhenomD\_log < T>.$ 

## 6.14.1.16 construct\_amplitude\_derivative()

```
int length,
int dimension,
double ** amplitude_derivative,
source_parameters< double > * input_params,
int * tapes = NULL ) [virtual]
```

Construct the derivative of the amplitude for a given source evaluated by the given frequency.

Order of output: dh/d \theta : \theta \el {A0,tc, phic, chirp mass, eta, symmetric spin, antisymmetric spin}

#### **Parameters**

frequencies	input array of frequency
length	length of the frequency array
amplitude_derivative	< dimension of the fisher output array for all the derivatives double[dimension][length]
input_params	Source parameters structure for the source
tapes	int array of tape ids, if NULL, these will be calculated

Reimplemented in ppE\_IMRPhenomD\_IMR< T >, and ppE\_IMRPhenomD\_Inspiral< T >.

## 6.14.1.17 construct\_phase()

Constructs the Phase as outlined by IMRPhenomD.

arguments: array of frequencies, length of that array, T array for the output phase, and a source\_parameters structure

#### **Parameters**

frequencies	T array of frequencies the waveform is to be evaluated at
length	integer length of the input and output arrays
phase	output T array for the phasee
params	structure of source parameters to be calculated before computation

 $\label{eq:continuous} Reimplemented in \ EdGB\_IMRPhenomD < T>, \ EdGB\_IMRPhenomD\_log < T>, \ dCS\_IMRPhenomD < T>, \ and \ dCS\_IMRPhenomD\_log < T>.$ 

## 6.14.1.18 construct\_phase\_derivative()

```
template<class T >
void IMRPhenomD< T >::construct_phase_derivative (
```

```
double * frequencies,
int length,
int dimension,
double ** phase_derivative,
source_parameters< double > * input_params,
int * tapes = NULL ) [virtual]
```

Construct the derivative of the phase for a given source evaluated by the given frequency.

Order of output: dh/d \theta: \theta \el {A0,tc, phic, chirp mass, eta, symmetric spin, antisymmetric spin}

#### **Parameters**

frequencies	input array of frequency
length	length of the frequency array
phase_derivative	< dimension of the fisher output array for all the derivatives double[dimension][length]
input_params	Source parameters structure for the source
tapes	int array of tape ids, if NULL, these will be calculated

Reimplemented in ppE\_IMRPhenomD\_IMR< T >, and ppE\_IMRPhenomD\_Inspiral< T >.

```
6.14.1.19 construct_waveform() [1/2]
```

Constructs the waveform as outlined by.

arguments: array of frequencies, length of that array, a complex array for the output waveform, and a source\_parameters structure

#### **Parameters**

frequencies	T array of frequencies the waveform is to be evaluated at
length	integer length of the array of frequencies and the waveform
waveform	complex T array for the waveform to be output

 $\label{eq:continuous} Reimplemented in \ EdGB\_IMRPhenomD < T>, \ EdGB\_IMRPhenomD\_log < T>, \ dCS\_IMRPhenomD < T>, \ and \ dCS\_IMRPhenomD\_log < T>.$ 

```
6.14.1.20 construct_waveform() [2/2]
```

```
template<class T >
std::complex< T > IMRPhenomD< T >::construct_waveform (
```

```
T frequency,
source_parameters< T > * params ) [virtual]
```

overloaded method to evaluate the waveform for one frequency instead of an array

#### **Parameters**

```
frequency T array of frequencies the waveform is to be evaluated at
```

#### 6.14.1.21 Damp\_ins()

Calculates the derivative wrt frequency for the scaled inspiral amplitude A/A0 for frequency f.

This is an analytic derivative for the smoothness condition on the amplitude connection

return a T

## 6.14.1.22 Damp\_mr()

Calculates the derivative wrt frequency for the scaled merger-ringdown amplitude A/A0 for frequency f.

This is an analytic derivative for the smoothness condition on the amplitude connection

The analytic expression was obtained from Mathematica - See the mathematica folder for code

return a T

## 6.14.1.23 Dphase\_ins()

Calculates the derivative of the inspiral phase for frequency f.

For phase continuity and smoothness return a T

Reimplemented in ppE\_IMRPhenomD\_Inspiral< T >.

#### 6.14.1.24 Dphase\_int()

Calculates the derivative of the intermediate phase for frequency f.

For phase continuity and smoothness return a T

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

#### 6.14.1.25 Dphase\_mr()

Calculates the derivative of the merger-ringdown phase for frequency f.

For phase continuity and smoothness return a T

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

#### 6.14.1.26 fpeak()

Solves for the peak frequency, where the waveform transitions from intermediate to merger-ringdown.

returns Hz

## 6.14.1.27 phase\_connection\_coefficients()

Calculates the phase connection coefficients alpha{0,1} and beta{0,1}.

Note: these coefficients are stored in the lambda parameter structure, not a separate array

#### 6.14.1.28 phase\_ins()

Calculates the inspiral phase for frequency f with precomputed powers of MF and PI for speed.

#### return a T

extra argument of precomputed powers of MF and pi, contained in the structure useful\_powers < T>

Reimplemented in ppE IMRPhenomD Inspiral< T >.

#### 6.14.1.29 phase\_int()

Calculates the intermediate phase for frequency f.

#### return a T

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

### 6.14.1.30 phase\_mr()

Calculates the merger-ringdown phase for frequency f.

#### return a T

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

## 6.14.1.31 phase\_tape()

Creates the tapes for derivatives of phase.

For efficiency in long runs of large sets of fishers, the tapes can be precomputed and reused

#### **Parameters**

input_params	source parameters structure of the desired source
tape	tape ids

Reimplemented in ppE IMRPhenomD IMR< T >, and ppE IMRPhenomD Inspiral< T >.

#### 6.14.1.32 post\_merger\_variables()

Calculates the post-merger ringdown frequency and dampening frequency.

Returns in Hz - assigns fRD to var[0] and fdamp to var[1]

#### 6.14.1.33 precalc\_powers\_ins()

Pre-calculate powers of Mf, to speed up calculations for the inspiral waveform (both amplitude and phase.

It seems the pow() function is very slow, so to speed things up, powers of Mf will be precomputed and passed to the functions within the frequency loops

## 6.14.1.34 precalc\_powers\_ins\_amp()

Pre-calculate powers of Mf, to speed up calculations for the inspiral amplitude.

It seems the pow() function is very slow, so to speed things up, powers of Mf will be precomputed and passed to the functions within the frequency loops

#### 6.14.1.35 precalc\_powers\_ins\_phase()

Pre-calculate powers of Mf, to speed up calculations for the inspiral phase.

It seems the pow() function is very slow, so to speed things up, powers of Mf will be precomputed and passed to the functions within the frequency loops

#### 6.14.1.36 precalc\_powers\_PI()

Pre-calculate powers of pi, to speed up calculations for the inspiral phase.

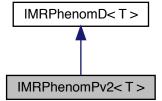
It seems the pow() function is very slow, so to speed things up, powers of PI will be precomputed and passed to the functions within the frequency loops

The documentation for this class was generated from the following files:

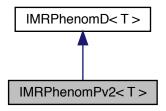
- include/IMRPhenomD.h
- src/IMRPhenomD.cpp

# 6.15 IMRPhenomPv2< T > Class Template Reference

Inheritance diagram for IMRPhenomPv2< T >:



Collaboration diagram for IMRPhenomPv2< T >:



#### **Public Member Functions**

- virtual T alpha (T omega, T q, T chi2l, T chi2)
- virtual T epsilon (T omega, T q, T chi2l, T chi2)
- virtual void calculate\_euler\_coeffs (alpha\_coeffs< T > \*acoeffs, epsilon\_coeffs< T > \*ecoeffs, source\_parameters< T > \*params)

Pre calculate euler angle coefficients.

- virtual T d (int I, int mp, int m, T s)
- virtual int construct\_waveform (T \*frequencies, int length, std::complex< T > \*waveform\_plus, std
   ::complex< T > \*waveform\_cross, source\_parameters< T > \*params)

Constructs the waveform for IMRPhenomPv2 - uses IMRPhenomD, then twists up.

- virtual void WignerD (T d2[5], T dm2[5], useful\_powers< T > \*pows, source\_parameters< T > \*params)
- virtual void calculate\_twistup (T alpha, std::complex< T > \*hp\_factor, std::complex< T > \*hc\_factor, T d2[5], T dm2[5], sph\_harm< T > \*sph\_harm)
- virtual void calculate\_euler\_angles (T \*alpha, T \*epsilon, useful\_powers< T > \*pows, alpha\_coeffs< T > \*acoeffs, epsilon\_coeffs< T > \*ecoeffs)
- virtual void PhenomPv2\_Param\_Transform (source\_parameters< T > \*params)
- virtual void PhenomPv2\_Param\_Transform\_J (source\_parameters< T > \*params)
- virtual T L2PN (T eta, useful\_powers< T > \*pow)

#### 6.15.1 Member Function Documentation

## 6.15.1.1 calculate\_euler\_coeffs()

Pre calculate euler angle coefficients.

Straight up stolen from LALsuite

#### 6.15.1.2 construct\_waveform()

Constructs the waveform for IMRPhenomPv2 - uses IMRPhenomD, then twists up.

arguments: array of frequencies, length of that array, a complex array for the output waveform, and a source\_parameters structure

#### **Parameters**

frequencies	T array of frequencies the waveform is to be evaluated at
length	integer length of the array of frequencies and the waveform
waveform_plus	complex T array for the plus polariaztion waveform to be output
waveform_cross	complex T array for the cross polarization waveform to be output

#### 6.15.1.3 PhenomPv2\_Param\_Transform()

```
\label{template} $$\operatorname{IMRPhenomPv2} T > :: PhenomPv2\_Param\_Transform ($$\operatorname{source\_parameters} T > * params ) [virtual]
```

/Brief Parameter transformtion to precalculate needed parameters for PhenomP from source parameters

Pretty much stolen verbatim from lalsuite

#### 6.15.1.4 PhenomPv2\_Param\_Transform\_J()

/Brief Parameter transformtion to precalculate needed parameters for PhenomP from source parameters – assumed inclination of total angular momentum J is given, not orbital angular momentum (in source frame (Lhat == zhat)

Pretty much stolen verbatim from lalsuite

The documentation for this class was generated from the following files:

- include/IMRPhenomP.h
- src/IMRPhenomP.cpp

# 6.16 lambda\_parameters < T > Struct Template Reference

## **Public Attributes**

- T rho [4]
- T v2
- T gamma [4]
- T sigma [5]
- T beta [5]
- T alpha [7]

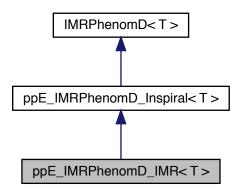
The documentation for this struct was generated from the following file:

• include/IMRPhenomD.h

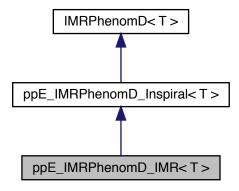
# 6.17 $ppE_IMRPhenomD_IMR < T > Class Template Reference$

#include <ppE\_IMRPhenomD.h>

Inheritance diagram for ppE\_IMRPhenomD\_IMR< T >:



Collaboration diagram for ppE\_IMRPhenomD\_IMR< T >:



#### **Public Member Functions**

- virtual T Dphase\_mr (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the derivative of the merger-ringdown phase for frequency f.
- virtual T phase\_mr (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the merger-ringdown phase for frequency f.
- virtual T phase\_int (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the intermediate phase for frequency f.
- virtual T Dphase\_int (T f, source\_parameters < T > \*param, lambda\_parameters < T > \*lambda)
   Calculates the derivative of the intermediate phase for frequency f.
- virtual void **fisher\_calculation** (double \*frequency, int length, gen\_params \*parameters, double \*\*amplitude\_deriv, double \*\*phase\_deriv, double \*amplitude, int \*amp\_tapes, int \*phase\_tapes)
- virtual void amplitude\_tape (source\_parameters < double > \*input\_params, int \*tape)
   Creates the tapes for derivatives of the amplitude.
- virtual void phase\_tape (source\_parameters< double > \*input\_params, int \*tape)
   Creates the tapes for derivatives of phase.
- virtual void construct\_amplitude\_derivative (double \*frequencies, int length, int dimension, double \*\*amplitude\_derivative, source\_parameters< double > \*input\_params, int \*tapes=NULL)

Construct the derivative of the amplitude for a given source evaluated by the given frequency.

virtual void construct\_phase\_derivative (double \*frequencies, int length, int dimension, double \*\*phase\_←
derivative, source\_parameters< double > \*input\_params, int \*tapes=NULL)

Construct the derivative of the phase for a given source evaluated by the given frequency.

## 6.17.1 Detailed Description

```
template < class T > class ppE_IMRPhenomD_IMR < T >
```

Class that extends the IMRPhenomD waveform to include non-GR terms in the full phase. This is an appropriate waveform choice for propagation effects

## 6.17.2 Member Function Documentation

## 6.17.2.1 amplitude\_tape()

Creates the tapes for derivatives of the amplitude.

For efficiency in long runs of large sets of fishers, the tapes can be precomputed and reused

#### **Parameters**

input_params	source parameters structure of the desired source
tape	tape ids

Reimplemented from ppE\_IMRPhenomD\_Inspiral< T >.

## 6.17.2.2 construct\_amplitude\_derivative()

Construct the derivative of the amplitude for a given source evaluated by the given frequency.

Order of output: dh/d \theta : \theta \el {A0,tc, phic, chirp mass, eta, symmetric spin, antisymmetric spin}

## **Parameters**

frequencies	input array of frequency
length	length of the frequency array
amplitude_derivative	< dimension of the fisher output array for all the derivatives double[dimension][length]
input_params	Source parameters structure for the source
tapes	int array of tape ids, if NULL, these will be calculated

Reimplemented from ppE\_IMRPhenomD\_Inspiral< T >.

#### 6.17.2.3 construct\_phase\_derivative()

Construct the derivative of the phase for a given source evaluated by the given frequency.

Order of output: dh/d \theta: \theta \el {A0,tc, phic, chirp mass, eta, symmetric spin, antisymmetric spin}

#### **Parameters**

frequencies	input array of frequency
length	length of the frequency array
phase_derivative	< dimension of the fisher output array for all the derivatives double[dimension][length]
input_params	Source parameters structure for the source
tapes	int array of tape ids, if NULL, these will be calculated

Reimplemented from ppE\_IMRPhenomD\_Inspiral< T >.

## 6.17.2.4 Dphase\_int()

Calculates the derivative of the intermediate phase for frequency f.

For phase continuity and smoothness return a T

Reimplemented from IMRPhenomD< T >.

## 6.17.2.5 Dphase\_mr()

Calculates the derivative of the merger-ringdown phase for frequency f.

For phase continuity and smoothness return a T

Reimplemented from IMRPhenomD < T >.

#### 6.17.2.6 phase\_int()

Calculates the intermediate phase for frequency f.

return a T

Reimplemented from IMRPhenomD< T >.

#### 6.17.2.7 phase\_mr()

Calculates the merger-ringdown phase for frequency f.

return a T

Reimplemented from IMRPhenomD< T >.

#### 6.17.2.8 phase\_tape()

Creates the tapes for derivatives of phase.

For efficiency in long runs of large sets of fishers, the tapes can be precomputed and reused

## **Parameters**

input_params	source parameters structure of the desired source	
tape	tape ids	

Reimplemented from ppE\_IMRPhenomD\_Inspiral< T >.

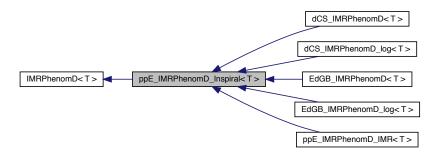
The documentation for this class was generated from the following files:

- include/ppE\_IMRPhenomD.h
- src/ppE\_IMRPhenomD.cpp

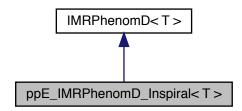
# 6.18 ppE\_IMRPhenomD\_Inspiral < T > Class Template Reference

#include <ppE\_IMRPhenomD.h>

Inheritance diagram for ppE\_IMRPhenomD\_Inspiral< T >:



Collaboration diagram for ppE\_IMRPhenomD\_Inspiral< T >:



#### **Public Member Functions**

virtual T phase\_ins (T f, source\_parameters < T > \*param, T \*pn\_coeff, lambda\_parameters < T > \*lambda, useful\_powers < T > \*pow)

Overloaded method for the inspiral portion of the phase.

virtual T Dphase\_ins (T f, source\_parameters< T > \*param, T \*pn\_coeff, lambda\_parameters< T > \*lambda)

Calculates the derivative of the inspiral phase for frequency f.

- virtual void **fisher\_calculation** (double \*frequency, int length, gen\_params \*parameters, double \*\*amplitude\_deriv, double \*\*phase\_deriv, double \*amplitude, int \*amp\_tapes, int \*phase\_tapes)
- virtual void amplitude tape (source parameters< double > \*input params, int \*tape)

Creates the tapes for derivatives of the amplitude.

virtual void phase\_tape (source\_parameters < double > \*input\_params, int \*tape)

Creates the tapes for derivatives of phase.

 virtual void construct\_amplitude\_derivative (double \*frequencies, int length, int dimension, double \*\*amplitude\_derivative, source\_parameters< double > \*input\_params, int \*tapes=NULL)

Construct the derivative of the amplitude for a given source evaluated by the given frequency.

virtual void construct\_phase\_derivative (double \*frequencies, int length, int dimension, double \*\*phase\_←
derivative, source\_parameters< double > \*input\_params, int \*tapes=NULL)

Construct the derivative of the phase for a given source evaluated by the given frequency.

## 6.18.1 Detailed Description

```
\label{template} \begin{split} \text{template} \! < \! \text{class T} \! > \\ \text{class ppE\_IMRPhenomD\_Inspiral} \! < \! \text{T} \! > \end{split}
```

Class that extends the IMRPhenomD waveform to include non-GR terms in the inspiral portion of the phase. This is an appropriate waveform choice for generation effects, but not necessarily for propagation effects

#### 6.18.2 Member Function Documentation

## 6.18.2.1 amplitude\_tape()

Creates the tapes for derivatives of the amplitude.

For efficiency in long runs of large sets of fishers, the tapes can be precomputed and reused

#### **Parameters**

input_params	source parameters structure of the desired source
tape	tape ids

Reimplemented from IMRPhenomD< T >.

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

## 6.18.2.2 construct\_amplitude\_derivative()

Construct the derivative of the amplitude for a given source evaluated by the given frequency.

Order of output: dh/d \theta : \theta \el {A0,tc, phic, chirp mass, eta, symmetric spin, antisymmetric spin}

#### **Parameters**

frequencies	input array of frequency	
length	length of the frequency array	
amplitude_derivative	< dimension of the fisher output array for all the derivatives double[dimension][length]	
input_params	Source parameters structure for the source	
tapes	int array of tape ids, if NULL, these will be calculated	

Reimplemented from IMRPhenomD< T >.

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

#### 6.18.2.3 construct\_phase\_derivative()

Construct the derivative of the phase for a given source evaluated by the given frequency.

Order of output: dh/d \theta: \theta \el {A0,tc, phic, chirp mass, eta, symmetric spin, antisymmetric spin}

#### **Parameters**

frequencies	input array of frequency	
length	length of the frequency array	
phase_derivative	< dimension of the fisher output array for all the derivatives double[dimension][length]	
input_params	input_params Source parameters structure for the source	
tapes	int array of tape ids, if NULL, these will be calculated	

Reimplemented from IMRPhenomD< T >.

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

## 6.18.2.4 Dphase\_ins()

Calculates the derivative of the inspiral phase for frequency f.

For phase continuity and smoothness return a T

Reimplemented from IMRPhenomD< T >.

#### 6.18.2.5 phase\_tape()

Creates the tapes for derivatives of phase.

For efficiency in long runs of large sets of fishers, the tapes can be precomputed and reused

#### **Parameters**

input_params	source parameters structure of the desired source
tape	tape ids

Reimplemented from IMRPhenomD< T >.

Reimplemented in ppE\_IMRPhenomD\_IMR< T >.

The documentation for this class was generated from the following files:

- include/ppE\_IMRPhenomD.h
- src/ppE\_IMRPhenomD.cpp

# 6.19 sampler Struct Reference

## **Public Attributes**

- int types\_of\_steps = 4
- double \*\* step\_prob
- double \*\* prob\_boundaries
- double \* chain\_temps
- bool \* waiting
- int \* chain\_pos
- · double swp\_freq
- · int chain N
- int numThreads
- int N\_steps
- int dimension
- bool fisher\_exist
- bool \* de\_primed
- int \* priority

- double \*\*\* output
- bool pool
- int progress =0
- bool show\_progress
- · int num threads
- · int history\_length
- · int history\_update
- int \* current\_hist\_pos
- double \*\*\* history
- double \* current\_likelihoods
- int \* check\_stepsize\_freq
- double \* max\_target\_accept\_ratio
- double \* min\_target\_accept\_ratio
- int \* gauss last accept ct
- int \* gauss\_last\_reject\_ct
- int \* de\_last\_accept\_ct
- int \* de\_last\_reject\_ct
- int \* fish\_last\_accept\_ct
- int \* fish\_last\_reject\_ct
- double \*\* randgauss\_width
- double \*\*\* fisher\_vecs
- double \*\* fisher vals
- int \* fisher\_update\_ct
- int fisher\_update\_number
- std::function< double(double \*, int, int)> lp
- std::function< double(double \*, int, int)> II
- std::function< void(double \*, int, double \*\*, int)> fish
- gsl\_rng \*\* rvec
- int \* nan counter
- int \* num gauss
- int \* num\_fish
- int \* num de
- int \* num\_mmala
- double time\_elapsed\_cpu
- double time\_elapsed\_wall
- double time\_elapsed\_cpu\_ac
- double time\_elapsed\_wall\_ac
- int \* fish\_accept\_ct
- int \* fish\_reject\_ct
- int \* de accept ct
- int \* de reject ct
- int \* gauss\_accept\_ct
- int \* gauss\_reject\_ct
- int \* mmala\_accept\_ct
- int \* mmala\_reject\_ct
- int \* swap\_accept\_ct
- int \* swap\_reject\_ct
- int \* step\_accept\_ct
- int \* step\_reject\_ct
- double \*\*\* II\_Ip\_output

The documentation for this struct was generated from the following file:

• include/mcmc\_sampler\_internals.h

# 6.20 source\_parameters < T > Struct Template Reference

## **Static Public Member Functions**

• static source\_parameters < T > populate\_source\_parameters (gen\_params \*param\_in)

Builds the structure that shuttles source parameters between functions -updated version to incorporate structure argument.

static source\_parameters < T > populate\_source\_parameters\_old (T mass1, T mass2, T Luminosity\_
 —
 Distance, T \*spin1, T \*spin2, T phi\_c, T t\_c, bool sky\_average)

Builds the structure that shuttles source parameters between functions- outdated in favor of structure argument.

## **Public Attributes**

- T mass1
- T mass2
- T M
- T q
- T spin1z
- T spin2z
- T spin1x
- T spin2x
- T spin1y
- T spin2y
- · T chirpmass
- Teta
- T chi s
- T chi a
- T chi\_eff
- T chi\_pn
- T DL
- T delta\_mass
- TfRD
- T fdamp
- T f1
- T f3
- T f1\_phase
- Tf2\_phase
- T phic
- Ttc
- T A0
- Ts
- T chil
- T chip
- Tf\_ref
- T phi\_aligned
- Tincl\_angle
- T phiRef
- T alpha0
- T thetaJN
- T zeta polariz
- T \* betappe
- int \* bppe
- int Nmod

- T phi
- T theta
- T SP
- · TSL
- bool sky\_average
- gsl\_spline \* **Z\_DL\_spline\_ptr** = NULL
- gsl\_interp\_accel \* **Z\_DL\_accel\_ptr** = NULL
- std::string cosmology

#### 6.20.1 Member Function Documentation

#### 6.20.1.1 populate\_source\_parameters()

Builds the structure that shuttles source parameters between functions -updated version to incorporate structure argument.

Populates the structure that is passed to all generation methods - contains all relavent source parameters

#### 6.20.1.2 populate\_source\_parameters\_old()

Builds the structure that shuttles source parameters between functions- outdated in favor of structure argument.

Populates the structure that is passed to all generation methods - contains all relavent source parameters

#### **Parameters**

mass1	mass of the larger body - in Solar Masses
mass2	mass of the smaller body - in Solar Masses
Luminosity_Distance	Luminosity Distance in Mpc
spin2	spin vector of the larger body {sx,sy,sz}
phi_c	spin vector of the smaller body {sx,sy,sz}
t_c	coalescence phase
sky_average	coalescence time

## 6.20.2 Member Data Documentation

T source\_parameters< T >::delta\_mass

Delta mass comibination

```
6.20.2.1 chi_a
template<class T>
T source_parameters< T >::chi_a
Antisymmetric spin combination
6.20.2.2 chi_eff
template<class T>
T source_parameters< T >::chi_eff
Effective spin
6.20.2.3 chi_pn
template<class T>
T source_parameters< T >::chi_pn
PN spin
6.20.2.4 chi_s
template<class T>
T source_parameters< T >::chi_s
Symmetric spin combination
6.20.2.5 chirpmass
{\tt template}{<}{\tt class} \ {\tt T}{>}
T source_parameters< T >::chirpmass
Chirp mass of the binary
6.20.2.6 delta_mass
{\tt template}{<}{\tt class} \ {\tt T}{>}
```

```
6.20.2.7 DL
```

```
template<class T>
T source_parameters< T >::DL
```

Luminoisity Distance

```
6.20.2.8 eta
```

```
template<class T>
T source_parameters< T >::eta
```

Symmetric mass ratio

#### 6.20.2.9 f1

```
template<class T>
T source_parameters< T >::f1
```

Transition Frequency 1 for the amplitude

```
6.20.2.10 f1_phase
```

```
template<class T>
T source_parameters< T >::fl_phase
```

Transition frequency 1 for the phase

#### 6.20.2.11 f2\_phase

```
template<class T>
T source_parameters< T >::f2_phase
```

Transition frequency 2 for the phase

## 6.20.2.12 f3

```
template<class T>
T source_parameters< T >::f3
```

Transition Frequency 2 for the amplitude

## 6.20.2.13 fdamp

```
template<class T>
T source_parameters< T >::fdamp
```

Dampening frequency after merger

```
6.20.2.14 fRD
template<class T>
T source_parameters< T >::fRD
Ringdown frequency after merger
6.20.2.15 M
template<class T>
T source_parameters< T >::M
Total mass
6.20.2.16 mass1
template<class T>
T source_parameters< T >::mass1
mass of the larger component
6.20.2.17 mass2
template<class T>
T source_parameters< T >::mass2
mass of the smaller component
6.20.2.18 Nmod
template<class T>
int source_parameters< T >::Nmod
Number of modifications to phase
6.20.2.19 phic
template<class T>
T source_parameters< T >::phic
Coalescence phase
6.20.2.20 spin1x
```

 ${\tt template}{<}{\tt class} \ {\tt T}{>}$ 

T source\_parameters< T >::spin1x

x-Spin component of the larger body

```
6.20.2.21 spin1y
template < class T >
T source_parameters< T >::spin1y
y-Spin component of the larger body
6.20.2.22 spin1z
{\tt template}{<}{\tt class} \ {\tt T}{>}
T source_parameters< T >::spin1z
z-Spin component of the larger body
6.20.2.23 spin2x
template < class T >
T source_parameters< T >::spin2x
x-Spin component of the smaller body
6.20.2.24 spin2y
template<class T>
T source_parameters< T >::spin2y
y-Spin component of the smaller body
6.20.2.25 spin2z
template<class T>
T source_parameters< T >::spin2z
z-Spin component of the smaller body
6.20.2.26 tc
template<class T>
T source_parameters< T >::tc
```

# Coalescence time

The documentation for this struct was generated from the following files:

- · include/util.h
- src/util.cpp

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# 6.21 sph\_harm < T > Struct Template Reference

# **Public Attributes**

- std::complex< T > Y22
- std::complex< T > Y21
- std::complex < T > Y20
- std::complex< T > Y2m1
- std::complex < T > Y2m2

The documentation for this struct was generated from the following file:

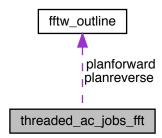
· include/util.h

# 6.22 threaded\_ac\_jobs\_fft Class Reference

Class to contain spectral method jobs.

```
#include <autocorrelation.h>
```

Collaboration diagram for threaded\_ac\_jobs\_fft:



# **Public Attributes**

- double \*\* data
- int \* length
- int \* start
- int \* end
- int dimension
- fftw\_outline \* planforward
- fftw\_outline \* planreverse
- int \* lag
- double \* target

# 6.22.1 Detailed Description

Class to contain spectral method jobs.

#### 6.22.2 Member Data Documentation

```
6.22.2.1 dimension
```

int  $threaded_ac_jobs_fft::dimension$ 

Read only - end index

6.22.2.2 end

int\* threaded\_ac\_jobs\_fft::end

Read only - start index

6.22.2.3 lag

int\* threaded\_ac\_jobs\_fft::lag

fftw plan to use for spectral method

6.22.2.4 length

 $\verb|int*| threaded_ac_jobs_fft:: length|$ 

Read only - Data to use - full chain

6.22.2.5 planforward

fftw\_outline\* threaded\_ac\_jobs\_fft::planforward

Read only - dimension being analyzed

6.22.2.6 planreverse

fftw\_outline\* threaded\_ac\_jobs\_fft::planreverse

fftw plan to use for spectral method

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# 6.22.2.7 start

```
int* threaded_ac_jobs_fft::start
```

Read only - length of total data

# 6.22.2.8 target

```
double* threaded_ac_jobs_fft::target
```

# READ AND WRITE - final lag

The documentation for this class was generated from the following file:

• include/autocorrelation.h

# 6.23 threaded\_ac\_jobs\_serial Class Reference

Class to contain serial method jobs.

```
#include <autocorrelation.h>
```

# **Public Attributes**

- double \*\* data
- int \* length
- int \* start
- int \* end
- int dimension
- int \* lag
- double \* target

# 6.23.1 Detailed Description

Class to contain serial method jobs.

#### 6.23.2 Member Data Documentation

# 6.23.2.1 dimension

int threaded\_ac\_jobs\_serial::dimension

# Read only – end index

```
6.23.2.2 end
int* threaded_ac_jobs_serial::end
Read only - start index
6.23.2.3 lag
int* threaded_ac_jobs_serial::lag
Read only - dimension being analyzed
6.23.2.4 length
int* threaded_ac_jobs_serial::length
```

6.23.2.5 start

int\* threaded\_ac\_jobs\_serial::start

Read only - length of total data

Read only - Data to use - full chain

6.23.2.6 target

double\* threaded\_ac\_jobs\_serial::target

READ AND WRITE - final lag

The documentation for this class was generated from the following file:

• include/autocorrelation.h

# 6.24 ThreadPool Class Reference

**Public Member Functions** 

- ThreadPool (std::size\_t numThreads)
- void enqueue (int i)
- void enqueue\_swap (int i)
- void public\_stop ()

The documentation for this class was generated from the following file:

• src/mcmc\_sampler.cpp

64 Class Documentation

# 6.25 ThreadPool Class Reference

#### **Public Member Functions**

- ThreadPool (std::size\_t numThreads)
- void enqueue (int i)
- void enqueue\_swap (int i)
- void public\_stop ()

The documentation for this class was generated from the following file:

• src/mcmc\_sampler.cpp

# 6.26 useful\_powers < T > Struct Template Reference

To speed up calculations within the for loops, we pre-calculate reoccuring powers of M\*F and Pi, since the pow() function is prohibatively slow.

```
#include <util.h>
```

# **Public Attributes**

- T MFthird
- T MFsixth
- ▼ MF7sixth
- T MF2third
- T MF4third
- T MF5third
- T MFsquare
- T MF7third
- T MF8third
- T MFcube
- TMFminus\_5third
- · double Plsquare
- · double Plcube
- · double PIthird
- · double Pl2third
- double Pl4third
- double PI5third
- double PI7third
- · double Plminus\_5third

# 6.26.1 Detailed Description

```
template < class T > struct useful_powers < T >
```

To speed up calculations within the for loops, we pre-calculate reoccuring powers of M\*F and Pi, since the pow() function is prohibatively slow.

Powers of PI are initialized once, and powers of MF need to be calculated once per for loop (if in the inspiral portion). use the functions precalc\_powers\_ins\_amp, precalc\_powers\_ins\_phase, precalc\_powers\_pi to initialize

The documentation for this struct was generated from the following file:

· include/util.h

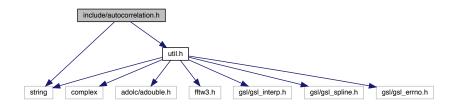
# **Chapter 7**

# **File Documentation**

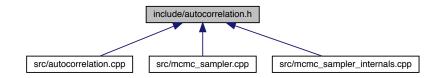
# 7.1 include/autocorrelation.h File Reference

```
#include <string>
#include "util.h"
```

Include dependency graph for autocorrelation.h:



This graph shows which files directly or indirectly include this file:



# Classes

• class threaded\_ac\_jobs\_fft

Class to contain spectral method jobs.

· class threaded\_ac\_jobs\_serial

Class to contain serial method jobs.

• class comparator\_ac\_fft

comparator to sort ac-jobs

· class comparator\_ac\_serial

comparator to sort ac-jobs

#### **Functions**

• void write\_auto\_corr\_file\_from\_data\_file (std::string autocorr\_filename, std::string datafile, int length, int dimension, int num\_segments, double target\_corr, int num\_threads)

- void write\_auto\_corr\_file\_from\_data (std::string autocorr\_filename, double \*\*data, int length, int dimension, int num\_segments, double target\_corr, int num\_threads)
- void auto\_corr\_from\_data (double \*\*data, int length, int dimension, int \*\*output, int num\_segments, double target corr, int num\_threads)

Calculates the autocorrelation length for a set of data for a number of segments for each dimension – completely host code, utilitizes FFTW3 for longer chuncks of the chains.

- void threaded\_ac\_spectral (int thread, threaded\_ac\_jobs\_fft job)
- · void threaded ac serial (int thread, threaded ac jobs serial job)
- double auto\_correlation\_serial (double \*arr, int length, int start, double target)
- void auto\_correlation\_spectral (double \*chain, int length, double \*autocorr, fftw\_outline \*plan\_forw, fftw\_outline \*plan\_rev)
- void auto\_correlation\_spectral (double \*chain, int length, int start, double \*autocorr, fftw\_outline \*plan\_forw, fftw outline \*plan rev)

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT – accepts FFTW plan as argument for plan reuse and multi-threaded applications.

void auto\_correlation\_spectral (double \*chain, int length, double \*autocorr)

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT.

- double auto correlation (double \*arr, int length, double tolerance)
- double auto\_correlation\_serial\_old (double \*arr, int length)
- double auto\_correlation\_grid\_search (double \*arr, int length, int box\_num=10, int final\_length=50, double target\_length=.01)

Grid search method of computing the autocorrelation.

• double auto correlation internal (double \*arr, int length, int lag, double ave)

Internal function to compute the auto correlation for a given lag.

void auto\_corr\_intervals\_outdated (double \*data, int length, double \*output, int num\_segments, double accuracy)

Function that computes the autocorrelation length on an array of data at set intervals to help determine convergence.

- void write\_auto\_corr\_file\_from\_data (std::string autocorr\_filename, double \*\*output, int intervals, int dimension, int N\_steps)
- void **write\_auto\_corr\_file\_from\_data\_file** (std::string autocorr\_filename, std::string output\_file, int intervals, int dimension, int N\_steps)

# 7.1.1 Detailed Description

Autocorrelation header file

#### 7.1.2 Function Documentation

### 7.1.2.1 auto\_corr\_from\_data()

Calculates the autocorrelation length for a set of data for a number of segments for each dimension – completely host code, utilitizes FFTW3 for longer chuncks of the chains.

Takes in the data from a sampler, shape data[N\_steps][dimension]

Outputs lags that correspond to the target\_corr – shape output[dimension][num\_segments]

#### **Parameters**

	data	Input data
	length	length of input data
	dimension	dimension of data
out	output	array that stores the auto-corr lengths – array[num_segments]
	num_segments	number of segements to compute the auto-corr length
	target_corr	Autocorrelation for which the autocorrelation length is defined (lag of autocorrelation for which it equals the target_corr)
	num_threads	Total number of threads to use

# 7.1.2.2 auto\_corr\_intervals\_outdated()

Function that computes the autocorrelation length on an array of data at set intervals to help determine convergence.

outdated version - new version uses FFTs

# **Parameters**

	data	Input data
	length	length of input data
out	output	array that stores the auto-corr lengths – array[num_segments]
	num_segments number of segements to compute the auto-corr length	
	accuracy	longer chains are computed numerically, this specifies the tolerance

#### 7.1.2.3 auto\_correlation\_grid\_search()

Grid search method of computing the autocorrelation.

Hopefully more reliable than the box-search method, which can sometimes get caught in a recursive loop when the stepsize isn't tuned, but also faster than the basic linear, serial search

#### **Parameters**

arr	Input array to use for autocorrelation	
length	Length of input array	
box_num	number of boxes to use for each iteration, default is 10	
final_length	number of elements per box at which the grid search ends and the serial calculation begins	
target_length	target correlation that corresponds to the returned lag	

#### 7.1.2.4 auto\_correlation\_internal()

Internal function to compute the auto correlation for a given lag.

# 7.1.2.5 auto\_correlation\_spectral() [1/2]

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT – accepts FFTW plan as argument for plan reuse and multi-threaded applications.

Based on the Wiener-Khinchin Theorem.

Algorithm used from https://lingpipe-blog.com/2012/06/08/autocorrelation-fft-kiss-eigen/

NOTE the length used in initializing the fftw plans should be L = pow(2, std::ceil( std::log2(length) ) ) – the plans are padded so the total length is a power of two

Option to provide starting index for multi-dimension arrays in collapsed to one dimension

length is the length of the segment to be analyzed, not necessarily the dimension of the chain

#### 7.1.2.6 auto\_correlation\_spectral() [2/2]

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT.

Based on the Wiener-Khinchin Theorem.

Algorithm used from https://lingpipe-blog.com/2012/06/08/autocorrelation-fft-kiss-eigen/

# 7.1.2.7 write\_auto\_corr\_file\_from\_data()

```
void write_auto_corr_file_from_data (
    std::string autocorr_filename,
    double ** data,
    int length,
    int dimension,
    int num_segments,
    double target_corr,
    int num_threads)
```

# **Parameters**

length	length of input data
dimension	dimension of data
num_segments	number of segements to compute the auto-corr length
target_corr	Autocorrelation for which the autocorrelation length is defined (lag of autocorrelation for which it equals the target_corr)
num_threads	Total number of threads to use

### 7.1.2.8 write\_auto\_corr\_file\_from\_data\_file()

```
std::string datafile,
int length,
int dimension,
int num_segments,
double target_corr,
int num_threads)
```

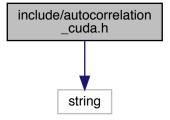
#### **Parameters**

length	length of input data
dimension	dimension of data
num_segments	number of segements to compute the auto-corr length
target_corr	Autocorrelation for which the autocorrelation length is defined (lag of autocorrelation for which it equals the target_corr)
num_threads	Total number of threads to use

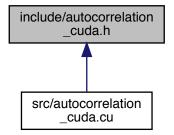
# 7.2 include/autocorrelation\_cuda.h File Reference

#include <string>

Include dependency graph for autocorrelation\_cuda.h:



This graph shows which files directly or indirectly include this file:



# **Macros**

#define THREADS\_PER\_BLOCK 512

#### **Functions**

void write\_file\_auto\_corr\_from\_data\_file\_accel (std::string acfile, std::string chains\_file, int dimension, int N
 \_ steps, int num\_segments, double target\_corr)

Write data file for autocorrelation lengths of the data given a data file name, as written by the mcmc\_sampler.

 void write\_file\_auto\_corr\_from\_data\_accel (std::string acfile, double \*\*output, int dimension, int N\_steps, int num\_segments, double target\_corr)

Write data file given output chains, as formatted by the mcmc sampler.

• void <a href="mailto:auto\_corr\_from\_data\_accel">accel</a> (double \*\*output, int dimension, int N\_steps, int num\_segments, double target corr, double \*\*autocorr)

Find autocorrelation of data at different points in the chain length and output to autocorr.

void launch\_ac\_gpu (int device, int element, double \*\*data, int length, int dimension, double target\_corr, int num\_segments)

Launch the GPU kernel, formatted for the thread pool.

void ac\_gpu\_wrapper (int thread, int job\_id)

Wrapper function for the thread pool.

• void auto correlation spectral accel (double \*chains, int length, double \*autocorr)

#### 7.2.1 Detailed Description

Header file for CUDA accelerated algorithms

Currently, no algorithms are used in any other parts of the project, so if CUDA or CUDA-enabled devices are not available, this file can be skipped in compilation by commenting out the OBJECTSCUDA line in the makefile

# 7.2.2 Function Documentation

#### 7.2.2.1 ac\_gpu\_wrapper()

Wrapper function for the thread pool.

#### **Parameters**

thread	Host thread
job⊷	Job ID
_id	

#### 7.2.2.2 auto\_corr\_from\_data\_accel()

Find autocorrelation of data at different points in the chain length and output to autocorr.

#### **Parameters**

	output	Chain data input
	dimension	Dimension of the data
	N_steps	Number of steps in the data
	num_segments	number of segments to calculate the autocorrelation length
	target_corr	Target correlation ratio
out	autocorr	Autocorrelation lengths for the different segments

# 7.2.2.3 write\_file\_auto\_corr\_from\_data\_accel()

```
void write_file_auto_corr_from_data_accel (
    std::string acfile,
    double ** output,
    int dimension,
    int N_steps,
    int num_segments,
    double target_corr )
```

Write data file given output chains, as formatted by the mcmc\_sampler.

#### **Parameters**

acfile	Output autocorrelation filename
output	Chain data from MCMC_sampler
dimension	Dimension of the data
N_steps	Number of steps in the chain
num_segments	Number of segments to check the autocorrelation length for each dimension
target_corr	Target correlation ratio to use for the correlation length calculation

# 7.2.2.4 write\_file\_auto\_corr\_from\_data\_file\_accel()

```
std::string chains_file,
int dimension,
int N_steps,
int num_segments,
double target_corr )
```

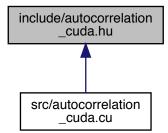
Write data file for autocorrelation lengths of the data given a data file name, as written by the mcmc\_sampler.

#### **Parameters**

acfile	Filename of the autocorrelation data
chains_file	Filename of the data file for the chains
dimension	Dimension of the data
N_steps	Number of steps in the chain
num_segments	Number of segments to check the autocorrelation length for each dimension
target_corr	Target correlation ratio to use for the correlation length calculation

# 7.3 include/autocorrelation\_cuda.hu File Reference

This graph shows which files directly or indirectly include this file:



#### Classes

• struct GPUplan

# **Functions**

- \_\_device\_\_ \_\_host\_\_ void auto\_corr\_internal (double \*arr, int length, int lag, double average, double \*corr, int start\_id)
  - Internal function to calculate the autocorrelation for a given lag Customized for the thread pool architecture, with extra arguments because of the way the memory is allocated.
- \_\_global\_\_ void auto\_corr\_internal\_kernal (double \*arr, int length, double average, int \*rho\_index, double target\_corr, double var, int start\_id)

Internal function to launch the CUDA kernel for a range of autocorrelations.

- void allocate\_gpu\_plan (GPUplan \*plan, int data\_length, int dimension, int num\_segments)

  Allocates memory for autocorrelation—GPU structure.
- void deallocate\_gpu\_plan (GPUplan \*plan, int data\_length, int dimension, int num\_segments)

  Deallocates memory for the autocorrelation—GPU structure.
- void copy\_data\_to\_device (GPUplan \*plan, double \*\*input\_data, int data\_length, int dimension, int num\_ segments)

Copy data to device before starting kernels.

# 7.3.1 Function Documentation

#### 7.3.1.1 allocate\_gpu\_plan()

Allocates memory for autocorrelation-GPU structure.

#### **Parameters**

plan	Structure for GPU plan
data_length	Length of data
dimension	Dimension of the data
num_segments	Number of segments to calculate the autocorrelation length

# 7.3.1.2 auto\_corr\_internal()

Internal function to calculate the autocorrelation for a given lag Customized for the thread pool architecture, with extra arguments because of the way the memory is allocated.

### **Parameters**

	arr	Input array of data	
	length	Length of input array	
	lag	Lag to be used to calculate the correlation	
	average	Average of the array arr	
out	corr	output correlation	
	start_id	ID of location to start calculation - input arrary arr is assumed to be contiguented that the contiguented in the contiguented	
		dimensions	

#### 7.3.1.3 auto\_corr\_internal\_kernal()

Internal function to launch the CUDA kernel for a range of autocorrelations.

Correlation function used:

```
 rho(lag) = 1 \ / \ (length - lag) \ (arr[i+lag]-average) \ (arr[i]-average) \\ target\_corr = rho(rho\_index)/rho(0) = rho(rho\_index)/var
```

# **Parameters**

	arr	Input array of data
	length	Length of data array
	average	Average of input data
out	rho_index	Index of the lag that results ina correlation ratio target_corr
	target_corr	Target correlation ratio rho(lag)/rho(0) = target_corr
	var	Variance rho(0)
	start_id	Starting index to use for the data array arr

# 7.3.1.4 copy\_data\_to\_device()

Copy data to device before starting kernels.

#### **Parameters**

plan	GPU plan
input_data	Input chain data
data_length	Length of data
dimension	Dimension of the data
num_segments	Number of segments to calculate the autocorrelation length

# 7.3.1.5 deallocate\_gpu\_plan()

Deallocates memory for the autocorrelation–GPU structure.

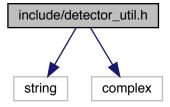
# **Parameters**

plan	Structure for the GPU plan
data_length	Length of data
dimension	Dimension of the data
num_segments	Number of segments to calculate the autocorrelation length

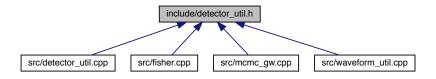
# 7.4 include/detector\_util.h File Reference

```
#include <string>
#include <complex>
last data dependency graph for detector
```

Include dependency graph for detector\_util.h:



This graph shows which files directly or indirectly include this file:



#### **Functions**

• void populate\_noise (double \*frequencies, std::string detector, double \*noise\_root, int length=0)

Function to populate the squareroot of the noise curve for various detectors.

• double aLIGO analytic (double f)

Analytic function approximating the PSD for aLIGO.

std::complex< double > Q (double theta, double phi, double iota)

Utility for the overall amplitude and phase shift for spin-aligned systems.

• double right interferometer cross (double theta, double phi)

Response function of a 90 deg interferometer for cross polarization.

double right\_interferometer\_plus (double theta, double phi)

Response function of a 90 deg interferometer for plus polarization.

• double Hanford\_O1\_fitted (double f)

Numerically fit PSD to the Hanford Detector's O1.

void celestial\_horizon\_transform (double RA, double DEC, double gps\_time, std::string detector, double \*phi, double \*theta)

Transform from celestial coordinates to local horizontal coords.

• void derivative\_celestial\_horizon\_transform (double RA, double DEC, double gps\_time, std::string detector, double \*dphi\_dRA, double \*dtheta\_dRA, double \*dphi\_dDEC, double \*dtheta\_dDEC)

Numerical derivative of the transformation.

double DTOA (double theta1, double theta2, std::string detector1, std::string detector2)

calculate difference in time of arrival (DTOA) for a given source location and 2 different detectors

• double radius\_at\_lat (double latitude, double elevation)

#### **Variables**

- const double H\_LAT = 0.81079526383
- const double H LONG =-2.08405676917
- const double H azimuth offset = 2.199
- const double H radius = 6367299.93401105
- const double H elevation = 142.554
- const double L\_LAT =0.53342313506
- const double L\_LONG =-1.58430937078
- const double L azimuth offset = 3.4557
- const double L\_radius = 6372795.50144497
- const double L elevation = -6.574
- const double V\_LAT = 0.76151183984
- const double **V\_LONG** =0.18333805213
- const double V\_azimuth\_offset = 1.239
- const double **V\_radius** = 6374824.24470673
- const double V\_elevation = 51.884
- const double RE polar =6357e3
- const double RE equatorial = 6378e3

#### 7.4.1 Detailed Description

Header file for all detector-specific utilities

#### 7.4.2 Function Documentation

### 7.4.2.1 aLIGO\_analytic()

```
double aLIGO_analytic ( \label{eq:double_f} \mbox{double } f \mbox{ )}
```

Analytic function approximating the PSD for aLIGO.

CITE (Will?)

# 7.4.2.2 celestial\_horizon\_transform()

Transform from celestial coordinates to local horizontal coords.

```
(RA,DEC) -> (altitude, azimuth)
```

Need gps\_time of transformation, as the horizontal coords change in time

detector is used to specify the lat and long of the local frame

# Parameters

RA	in RAD
DEC	in RAD
phi	in RAD
theta	in RAD

# 7.4.2.3 derivative\_celestial\_horizon\_transform()

Numerical derivative of the transformation.

Planned for use in Fisher calculations, but not currently implemented anywhere

#### **Parameters**

RA	in RAD
DEC	in RAD

# 7.4.2.4 DTOA()

calculate difference in time of arrival (DTOA) for a given source location and 2 different detectors

#### **Parameters**

theta1	spherical polar angle for detector 1 in RAD
theta2	spherical polar angle for detector 2 in RAD
detector1	name of detector one
detector2	name of detector two

# 7.4.2.5 Hanford\_O1\_fitted()

```
double Hanford_O1_fitted ( double f )
```

Numerically fit PSD to the Hanford Detector's O1.

CITE (Yunes?)

# 7.4.2.6 populate\_noise()

Function to populate the squareroot of the noise curve for various detectors.

If frequencies are left as NULL, standard frequency spacing is applied and the frequencies are returned, in which case the frequencies argument becomes an output array

Detector names must be spelled exactly

Detectors include: aLIGO\_analytic, Hanford\_O1\_fitted

#### **Parameters**

frequencies	double array of frquencies (NULL)
detector	String to designate the detector noise curve to be used
noise_root	ouptput double array for the square root of the PSD of the noise of the specified detector
length	integer length of the output and input arrays

# 7.4.2.7 Q()

Utility for the overall amplitude and phase shift for spin-aligned systems.

For spin aligned, all the extrinsic parameters have the effect of an overall amplitude modulation and phase shift

# 7.4.2.8 radius\_at\_lat()

/brief Analytic approximation of the radius from the center of earth to a given location

Just the raidus as a function of angles, modelling an oblate spheroid

#### **Parameters**

latitude	latitude in degrees
elevation	elevation in meters

### 7.4.2.9 right\_interferometer\_cross()

Response function of a 90 deg interferometer for cross polarization.

Theta and phi are local, horizontal coordinates relative to the detector

# 7.4.2.10 right\_interferometer\_plus()

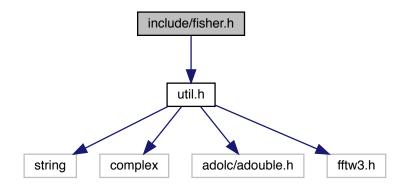
```
double right_interferometer_plus ( \label{eq:constraint} \mbox{double } theta, \\ \mbox{double } phi \mbox{ )}
```

Response function of a 90 deg interferometer for plus polarization.

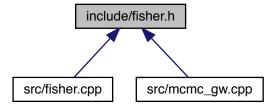
Theta and phi are local, horizontal coordinates relative to the detector

# 7.5 include/fisher.h File Reference

```
#include "util.h"
Include dependency graph for fisher.h:
```



This graph shows which files directly or indirectly include this file:



#### **Functions**

void fisher (double \*frequency, int length, string generation\_method, string detector, double \*\*output, int dimension, gen\_params \*parameters, int \*amp\_tapes=NULL, int \*phase\_tapes=NULL, double \*noise=N←ULL)

Calculates the fisher matrix for the given arguments.

• void calculate\_derivatives (double \*\*amplitude\_deriv, double \*\*phase\_deriv, double \*amplitude, double \*frequencies, int length, string detector, string gen\_method, gen\_params \*parameters)

Abstraction layer for handling the case separation for the different waveforms.

 void fisher\_autodiff (double \*frequency, int length, string generation\_method, string detector, double \*\*output, int dimension, gen\_params \*parameters, int \*amp\_tapes=NULL, int \*phase\_tapes=NULL, double \*noise=NULL)

Calculates the fisher matrix for the given arguments to within numerical error using automatic differention - slower than the numerical version.

#### 7.5.1 Function Documentation

# 7.5.1.1 calculate\_derivatives()

Abstraction layer for handling the case separation for the different waveforms.

### 7.5.1.2 fisher()

Calculates the fisher matrix for the given arguments.

# **Parameters**

length	if 0, standard frequency range for the detector is used
output	double [dimension][dimension]
amp_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method
phase_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method

# 7.5.1.3 fisher\_autodiff()

Calculates the fisher matrix for the given arguments to within numerical error using automatic differention - slower than the numerical version.

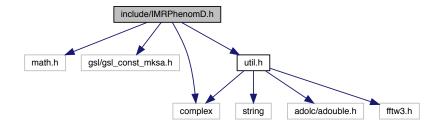
# **Parameters**

length	if 0, standard frequency range for the detector is used
output	double [dimension][dimension]
amp_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method
phase_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method

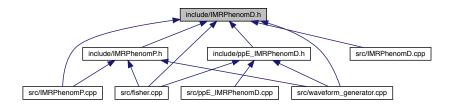
# 7.6 include/IMRPhenomD.h File Reference

```
#include <math.h>
#include <gsl/gsl_const_mksa.h>
#include <complex>
#include "util.h"
```

Include dependency graph for IMRPhenomD.h:



This graph shows which files directly or indirectly include this file:



# Classes

- struct lambda\_parameters < T >
- class IMRPhenomD< T >

# **Variables**

• const double lambda\_num\_params [19][11]

# 7.6.1 Detailed Description

Header file for utilities

# 7.6.2 Variable Documentation

# 7.6.2.1 lambda\_num\_params

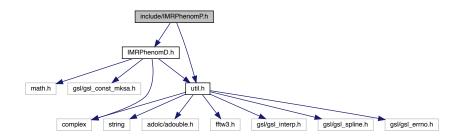
const double lambda\_num\_params[19][11]

Numerically calibrated parameters from arXiv:1508.07253 see the table in the data directory for labeled version

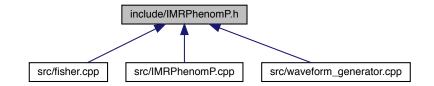
# 7.7 include/IMRPhenomP.h File Reference

```
#include "IMRPhenomD.h"
#include "util.h"
```

Include dependency graph for IMRPhenomP.h:



This graph shows which files directly or indirectly include this file:



### Classes

- struct alpha\_coeffs< T >
- struct epsilon\_coeffs< T >
- class IMRPhenomPv2< T >

# 7.7.1 Detailed Description

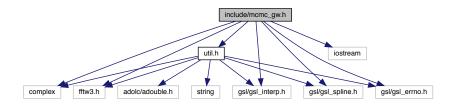
Header file for IMRPhenomP functions

Currently, only Pv2 is supported.

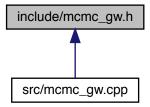
Wrapped around IMRPhenomD

# 7.8 include/mcmc\_gw.h File Reference

```
#include <complex>
#include <fftw3.h>
#include "util.h"
#include <iostream>
#include <gsl/gsl_interp.h>
#include <gsl/gsl_spline.h>
#include <gsl/gsl_errno.h>
Include dependency graph for mcmc gw.h:
```



This graph shows which files directly or indirectly include this file:



### **Functions**

• double maximized\_coal\_log\_likelihood\_IMRPhenomD (double \*frequencies, int length, std::complex< double > \*data, double \*noise, double SNR, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, bool NSflag, fftw\_outline \*plan)

Function to calculate the log Likelihood as defined by -1/2 (d-h|d-h) maximized over the extrinsic parameters phic and tc.

- double maximized\_coal\_log\_likelihood\_IMRPhenomD (double \*frequencies, size\_t length, double \*real\_
   data, double \*imag\_data, double \*noise, double SNR, double chirpmass, double symmetric\_mass\_ratio,
   double spin1, double spin2, bool NSflag)
- double maximized\_coal\_log\_likelihood\_IMRPhenomD (double \*frequencies, size\_t length, double \*real\_
   data, double \*imag\_data, double \*noise, double SNR, double chirpmass, double symmetric\_mass\_ratio,
   double spin1, double spin2, bool NSflag, fftw\_outline \*plan)
- double maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param (double \*frequencies, int length, std
   ::complex< double > \*data, double \*noise, double chirpmass, double symmetric\_mass\_ratio, double spin1,
   double spin2, double Luminosity\_Distance, double theta, double phi, double iota, bool NSflag, fftw\_outline
   \*plan)

- double maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param (double \*frequencies, size\_t length, double \*real\_data, double \*imag\_data, double \*noise, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, double Luminosity\_Distance, double theta, double phi, double iota, bool NSflag)
- double maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param (double \*frequencies, size\_t length, double \*real\_data, double \*imag\_data, double \*noise, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, double Luminosity\_Distance, double theta, double phi, double iota, bool NSflag, fftw outline \*plan)
- double maximized\_Log\_Likelihood\_aligned\_spin\_internal (std::complex< double > \*data, double \*psd, double \*frequencies, std::complex< double > \*detector\_response, size\_t length, fftw\_outline \*plan)

Maximized match over coalescence variables - returns log likelihood NOT NORMALIZED for aligned spins.

• double Log\_Likelihood (std::complex< double > \*data, double \*psd, double \*frequencies, size\_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw\_outline \*plan)

Unmarginalized log of the likelihood.

• double maximized\_Log\_Likelihood\_unaligned\_spin\_internal (std::complex< double > \*data, double \*psd, double \*frequencies, std::complex< double > \*hplus, std::complex< double > \*hcross, size\_t length, fftw outline \*plan)

log likelihood function that maximizes over extrinsic parameters tc, phic, D, and phiRef, the reference frequency - for unaligned spins

double maximized\_Log\_Likelihood (std::complex < double > \*data, double \*psd, double \*frequencies, size ←
 \_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw\_outline \*plan)

routine to maximize over all extrinsic quantities and return the log likelihood

- double maximized\_Log\_Likelihood (double \*data\_real, double \*data\_imag, double \*psd, double \*frequencies, size\_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw outline \*plan)
- double maximized\_coal\_Log\_Likelihood (std::complex < double > \*data, double \*psd, double \*frequencies, size\_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw\_outline \*plan, double \*tc, double \*phic)

Function to maximize only over coalescence variables to and phic, returns the maximum values used.

- double maximized\_coal\_Log\_Likelihood\_internal (std::complex< double > \*data, double \*psd, double \*frequencies, std::complex< double > \*detector\_response, size\_t length, fftw\_outline \*plan, double \*tc, double \*phic)
- double Log\_Likelihood\_internal (std::complex < double > \*data, double \*psd, double \*frequencies, std
   ::complex < double > \*detector\_response, int length, fftw\_outline \*plan)

Internal function for the unmarginalized log of the likelihood.

void MCMC\_MH\_GW (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension, int chain\_id), int numThreads, bool pool, bool show\_prog, int num\_detectors, std::complex< double > \*\*data, double \*\*noise\_psd, double \*\*frequencies, int \*data\_length, double gps\_time, std::string \*detector, int Nmod, int \*bppe, std::string generation\_method, std::string statistics\_filename, std::string chain\_filename, std::string auto corr filename, std::string checkpoint filename)

Wrapper for the MCMC\_MH function, specifically for GW analysis.

void continue\_MCMC\_MH\_GW (std::string start\_checkpoint\_file, double \*\*\*output, int dimension, int N\_← steps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension, int chain\_id), int numThreads, bool pool, bool show\_prog, int num\_detectors, std::complex< double > \*\*data, double \*\*noise\_psd, double \*\*frequencies, int \*data\_length, double gps\_time, std::string \*detector, int Nmod, int \*bppe, std::string generation\_method, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string final\_checkpoint\_filename)

Takes in an MCMC checkpoint file and continues the chain.

• void MCMC\_method\_specific\_prep (std::string generation\_method, int dimension, double \*seeding\_var, bool local seeding)

Unpacks MCMC parameters for method specific initiation.

void MCMC\_fisher\_wrapper (double \*param, int dimension, double \*\*output, int chain\_id)

Fisher function for MCMC for GW.

double MCMC\_likelihood\_wrapper (double \*param, int dimension, int chain\_id)

log likelihood function for MCMC for GW

# 7.8.1 Detailed Description

Header file for the Graviational Wave specific MCMC routines

#### 7.8.2 Function Documentation

#### 7.8.2.1 continue\_MCMC\_MH\_GW()

```
void continue_MCMC_MH_GW (
            std::string start_checkpoint_file,
             double *** output,
             int dimension,
             int N_steps,
             int swp_freq,
             double(*)(double *param, int dimension, int chain_id) log_prior,
             int numThreads,
             bool pool,
             bool show_prog,
             int num_detectors,
             std::complex< double > ** data,
             double ** noise_psd,
             double ** frequencies,
             int * data_length,
             double gps_time,
             std::string * detectors,
             int Nmod,
             int * bppe,
             std::string generation_method,
             std::string statistics_filename,
             std::string chain_filename,
             std::string auto_corr_filename,
             std::string final_checkpoint_filename )
```

Takes in an MCMC checkpoint file and continues the chain.

Obviously, the user must be sure to correctly match the dimension, number of chains, the generation\_method, the prior function, the data, psds, freqs, and the detectors (number and name), and the gps\_time to the previous run, otherwise the behavior of the sampler is undefined.

numThreads and pool do not necessarily have to be the same

### 7.8.2.2 Log\_Likelihood()

Unmarginalized log of the likelihood.

### 7.8.2.3 Log\_Likelihood\_internal()

```
double Log_Likelihood_internal (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    std::complex< double > * detector_response,
    int length,
    fftw_outline * plan )
```

Internal function for the unmarginalized log of the likelihood.

```
.5 * ((h | h) - 2(D | h))
```

#### 7.8.2.4 maximized\_coal\_Log\_Likelihood()

```
double maximized_coal_Log_Likelihood (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    size_t length,
    gen_params * params,
    std::string detector,
    std::string generation_method,
    fftw_outline * plan,
    double * tc,
    double * phic )
```

Function to maximize only over coalescence variables to and phic, returns the maximum values used.

# 7.8.2.5 maximized\_coal\_log\_likelihood\_IMRPhenomD() [1/3]

Function to calculate the log Likelihood as defined by -1/2 (d-h|d-h) maximized over the extrinsic parameters phic and tc.

frequency array must be uniform spacing - this shouldn't be a problem when working with real data as DFT return uniform spacing

#### **Parameters**

```
chirpmass in solar masses
```

# 7.8.2.6 maximized\_coal\_log\_likelihood\_IMRPhenomD() [2/3]

#### **Parameters**

chirpmass in solar masses

# $\textbf{7.8.2.7} \quad maximized\_coal\_log\_likelihood\_IMRPhenomD() \ \ \, \texttt{[3/3]}$

#### **Parameters**

chirpmass in solar masses

#### 7.8.2.8 maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param() [1/3]

 ${\tt double\ maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param\ (}$ 

```
double * frequencies,
int length,
std::complex< double > * data,
double * noise,
double chirpmass,
double symmetric_mass_ratio,
double spin1,
double spin2,
double Luminosity_Distance,
double theta,
double phi,
double iota,
bool NSflag,
fftw_outline * plan )
```

#### **Parameters**

chirpmass in solar masses

# 7.8.2.9 maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param() [2/3]

# Parameters

chirpmass in solar masses

# 7.8.2.10 maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param() [3/3]

```
double chirpmass,
double symmetric_mass_ratio,
double spin1,
double spin2,
double Luminosity_Distance,
double theta,
double phi,
double iota,
bool NSflag,
fftw_outline * plan )
```

#### **Parameters**

```
chirpmass in solar masses
```

# 7.8.2.11 maximized\_Log\_Likelihood()

```
double maximized_Log_Likelihood (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    size_t length,
    gen_params * params,
    std::string detector,
    std::string generation_method,
    fftw_outline * plan )
```

routine to maximize over all extrinsic quantities and return the log likelihood

 $\label{local-phic} \begin{array}{l} \textbf{IMRPhenomD} - \textbf{maximizes over DL}, \textbf{phic}, \textbf{tc}, \\ \textbf{hota}, \\ \textbf{hota$ 

### 7.8.2.12 maximized\_Log\_Likelihood\_aligned\_spin\_internal()

```
double maximized_Log_Likelihood_aligned_spin_internal (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    std::complex< double > * detector_response,
    size_t length,
    fftw_outline * plan )
```

Maximized match over coalescence variables - returns log likelihood NOT NORMALIZED for aligned spins.

Note: this function is not properly normalized for an absolute comparison. This is made for MCMC sampling, so to minimize time, constant terms like (Data|Data), which would cancel in the Metropolis-Hasting ratio, are left out for efficiency

#### 7.8.2.13 maximized\_Log\_Likelihood\_unaligned\_spin\_internal()

```
double maximized_Log_Likelihood_unaligned_spin_internal (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    std::complex< double > * hplus,
    std::complex< double > * hcross,
    size_t length,
    fftw_outline * plan )
```

log likelihood function that maximizes over extrinsic parameters tc, phic, D, and phiRef, the reference frequency - for unaligned spins

Ref: arXiv 1603.02444v2

# 7.8.2.14 MCMC\_fisher\_wrapper()

Fisher function for MCMC for GW.

Wraps the fisher calculation in src/fisher.cpp and unpacks parameters correctly for common GW analysis

Supports all the method/parameter combinations found in MCMC\_MH\_GW

# 7.8.2.15 MCMC\_likelihood\_wrapper()

log likelihood function for MCMC for GW

Wraps the above likelihood functions and unpacks parameters correctly for common GW analysis

Supports all the method/parameter combinations found in MCMC\_MH\_GW

### 7.8.2.16 MCMC\_method\_specific\_prep()

```
void MCMC_method_specific_prep (
    std::string generation_method,
    int dimension,
    double * seeding_var,
    bool local_seeding )
```

Unpacks MCMC parameters for method specific initiation.

Populates seeding vector if non supplied, populates mcmc\_Nmod, populates mcmc\_log\_beta, populates mcmc\_cintrinsic

#### 7.8.2.17 MCMC\_MH\_GW()

```
void MCMC_MH_GW (
             double *** output,
             int dimension,
             int N_steps,
             int chain N.
             double * initial_pos,
             double * seeding_var,
             double * chain_temps,
             int swp_freq,
             double(*)(double *param, int dimension, int chain_id) log_prior,
             int numThreads.
             bool pool,
             bool show_prog,
             int num_detectors,
             std::complex< double > ** data,
             double ** noise_psd,
             double ** frequencies,
             int * data_length,
             double gps_time,
             std::string * detectors,
             int Nmod,
             int * bppe,
             std::string generation_method,
             std::string statistics filename.
             std::string chain_filename,
             std::string auto_corr_filename,
             std::string checkpoint_file )
```

Wrapper for the MCMC\_MH function, specifically for GW analysis.

Handles the details of setting up the MCMC sampler and wraps the fisher and log likelihood to conform to the format of the sampler

*NOTE* – This sampler is NOT thread safe. There is global memory declared for each call to MCMC\_MH\_GW, so separate samplers should not be run in the same process space

Supported parameter combinations:

```
IMRPhenomD - 4 dimensions - In chirpmass, eta, chi1, chi2
```

```
IMRPhenomD - 7 dimensions - In D_L, tc, phic, In chirpmass, eta, chi1, chi2
```

IMRPhenomD - 8 dimensions - cos inclination, RA, DEC, In D\_L, In chirpmass, eta, chi1, chi2

 $\frac{dCS\_IMRPhenomD\_log}{dCS\_IMRPhenomD\_log} - 8 \ dimensions - cos \ inclination, \ RA, \ DEC, \ In \ D\_L, \ In \ chirpmass, \ eta, \ chi1, \ chi2, \ In \ \ lapha^2 \ (the \ coupling \ parameter)$ 

dCS\_IMRPhenomD- 8 dimensions – cos inclination, RA, DEC, In D\_L, In chirpmass, eta, chi1, chi2,  $\alpha^2$  (the coupling parameter)

dCS\_IMRPhenomD\_root\_alpha- 8 dimensions – cos inclination, RA, DEC, In D\_L, In chirpmass, eta, chi1, chi2, \sqrt \alpha (in km) (the coupling parameter)

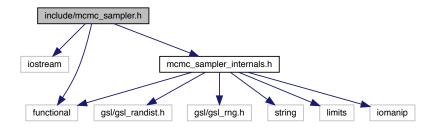
IMRPhenomPv2 - 9 dimensions - cos J\_N, In chirpmass, eta, |chi1|, |chi1|, theta\_1, theta\_2, phi\_1, phi\_2

#### **Parameters**

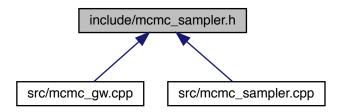
statistics_filename	Filename to output sampling statistics, if empty string, not output	
chain_filename	Filename to output data (chain 0 only), if empty string, not output	
auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output	
checkpoint_file	Filename to output data for checkpoint, if empty string, not saved	

# 7.9 include/mcmc\_sampler.h File Reference

```
#include <iostream>
#include <functional>
#include "mcmc_sampler_internals.h"
Include dependency graph for mcmc_sampler.h:
```



This graph shows which files directly or indirectly include this file:



### **Functions**

- void mcmc step threaded (int j)
- void mcmc\_swap\_threaded (int i, int j)

void continue\_MCMC\_MH (std::string start\_checkpoint\_file, double \*\*\*output, int N\_steps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension), double(\*log\_likelihood)(double \*param, int dimension), void(\*fisher)(double \*param, int dimension, double \*\*fisher), int numThreads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string end\_checkpoint\_file)

void MCMC MH loop (sampler \*sampler)

Internal function that runs the actual loop for the sampler.

- void MCMC\_MH (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension), double(\*log\_likelihood)(double \*param, int dimension), void(\*fisher)(double \*param, int dimension, double \*\*fisher), int numThreads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_\(\cup \) filename, std::string auto\_corr\_filename, std::string checkpoint\_filename)
- void MCMC\_MH (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension, int chain\_id), double(\*log\_likelihood)(double \*param, int dimension, int chain\_id), void(\*fisher)(double \*param, int dimension, double \*\*fisher, int chain\_id), int numThreads, bool pool, bool show\_prog, std
  ::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string checkpoint
  \_filename)
- void MCMC\_MH\_internal (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, std::function< double(double \*, int, int)> log\_ comprior, std::function< double(double \*, int, int)> log\_likelihood, std::function< void(double \*, int, double \*\*, int)>fisher, int numThreads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_comprise filename, std::string auto corr filename, std::string checkpoint filename)

Generic sampler, where the likelihood, prior are parameters supplied by the user.

void continue\_MCMC\_MH\_internal (std::string start\_checkpoint\_file, double \*\*\*output, int N\_steps, int swp
 \_freq, std::function< double(double \*, int, int)> log\_prior, std::function< double(double \*, int, int)> log\_
 likelihood, std::function< void(double \*, int, double \*\*, int)>fisher, int numThreads, bool pool, bool show
 \_prog, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string end\_checkpoint\_file)

Routine to take a checkpoint file and begin a new chain at said checkpoint.

### 7.9.1 Detailed Description

Header file for mcmc sampler

### 7.9.2 Function Documentation

#### 7.9.2.1 continue\_MCMC\_MH() [1/2]

std::string statistics\_filename,
std::string chain\_filename,
std::string auto\_corr\_filename,
std::string end\_checkpoint\_file )

### **Parameters**

	start_checkpoint_file	File for starting checkpoint	
out	output	output array, dimensions: output[chain_N][N_steps][dimension]	
	N_steps	Number of new steps to take	
	swp_freq	frequency of swap attempts between temperatures	
	log_prior	Funcion pointer for the log_prior	
	log_likelihood	Function pointer for the log_likelihood	
	fisher	Function pointer for the fisher - if NULL, fisher steps are not used	
	numThreads	Number of threads to use	
	pool	Boolean for whether to use deterministic'' vsstochastic" sampling	
	show_prog	Boolean for whether to show progress or not (turn off for cluster runs	
	statistics_filename	Filename to output sampling statistics, if empty string, not output	
	chain_filename	Filename to output data (chain 0 only), if empty string, not output	
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output	
	end_checkpoint_file	Filename to output data for checkpoint at the end of the continued run, if empty string, not saved	

### **7.9.2.2 continue\_MCMC\_MH()** [2/2]

	start_checkpoint_file	File for starting checkpoint
out	output	output array, dimensions: output[chain_N][N_steps][dimension]
	N_steps	Number of new steps to take
	swp_freq	frequency of swap attempts between temperatures
	log_prior	Funcion pointer for the log_prior
	log_likelihood	Function pointer for the log_likelihood
	fisher	Function pointer for the fisher - if NULL, fisher steps are not used
	numThreads	Number of threads to use
	pool Boolean for whether to use deterministic'' vsstochastic" sa	
	show_prog Boolean for whether to show progress or not (turn off for cluster runs	
	statistics_filename	Filename to output sampling statistics, if empty string, not output

### **Parameters**

chain_filename	Filename to output data (chain 0 only), if empty string, not output
auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
end_checkpoint_file	Filename to output data for checkpoint at the end of the continued run, if empty string, not saved

### 7.9.2.3 continue\_MCMC\_MH\_internal()

```
void continue_MCMC_MH_internal (
    std::string start_checkpoint_file,
    double *** output,
    int N_steps,
    int swp_freq,
    std::function< double(double *, int, int) > log_prior,
    std::function< double(double *, int, int) > log_likelihood,
    std::function< void(double *, int, double **, int) > fisher,
    int numThreads,
    bool pool,
    bool show_prog,
    std::string statistics_filename,
    std::string auto_corr_filename,
    std::string auto_corr_filename,
    std::string end_checkpoint_file)
```

Routine to take a checkpoint file and begin a new chain at said checkpoint.

See MCMC\_MH\_internal for more details of parameters (pretty much all the same)

	start_checkpoint_file	File for starting checkpoint
out	output	output array, dimensions: output[chain_N][N_steps][dimension]
	N_steps	Number of new steps to take
	swp_freq	frequency of swap attempts between temperatures
	log_prior	std::function for the log_prior function – takes double *position, int dimension, int chain_id
	log_likelihood std::function for the log_likelihood function – takes double *position, in dimension, int chain id	
	fisher	std::function for the fisher function – takes double *position, int dimension, double **output_fisher, int chain_id
	numThreads	Number of threads to use
	pool	Boolean for whether to use deterministic'' vsstochastic" sampling
	show_prog	Boolean for whether to show progress or not (turn off for cluster runs
	statistics_filename	Filename to output sampling statistics, if empty string, not output
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	end_checkpoint_file	Filename to output data for checkpoint at the end of the continued run, if empty string, not saved

# **7.9.2.4** MCMC\_MH() [1/2]

```
void MCMC_MH (
             double *** output,
             int dimension,
             int N_steps,
             int chain_N,
             double * initial_pos,
             double * seeding_var,
             double * chain_temps,
             int swp_freq,
             double(*)(double *param, int dimension) log_prior,
             double(*)(double *param, int dimension) log_likelihood,
             void(*)(double *param, int dimension, double **fisher) fisher,
             int numThreads,
             bool pool,
             bool show_prog,
             std::string statistics_filename,
             std::string chain_filename,
             std::string auto_corr_filename,
             std::string checkpoint_filename )
```

out	output	Output chains, shape is double[chain_N, N_steps,dimension]
	dimension	dimension of the parameter space being explored
	N_steps	Number of total steps to be taken, per chain
	chain_N	Number of chains
	initial_pos	Initial position in parameter space - shape double[dimension]
	seeding_var	Variance of the normal distribution used to seed each chain higher than 0 - shape double[dimension]
	chain_temps	Double array of temperatures for the chains
	swp_freq	the frequency with which chains are swapped
	log_prior	Funcion pointer for the log_prior
	log_likelihood	Function pointer for the log_likelihood
	fisher	Function pointer for the fisher - if NULL, fisher steps are not used
	numThreads	Number of threads to use (=1 is single threaded)
	pool	boolean to use stochastic chain swapping (MUST have >2 threads)
	show_prog	boolean whether to print out progress (for example, should be set to `'false'' if submitting to a cluster)
	statistics_filename Filename to output sampling statistics, if empty string, not output	
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	checkpoint_filename	Filename to output data for checkpoint, if empty string, not saved

```
7.9.2.5 MCMC_MH() [2/2]
```

```
void MCMC\_MH (
```

```
double *** output,
int dimension,
int N_steps,
int chain_N,
double * initial_pos,
double * seeding_var,
double * chain_temps,
int swp_freq,
double(*)(double *param, int dimension, int chain_id) log_prior,
double(*)(double *param, int dimension, int chain_id) log_likelihood,
void(*)(double *param, int dimension, double **fisher, int chain_id) fisher,
int numThreads,
bool pool,
bool show_prog,
std::string statistics_filename,
std::string chain_filename,
std::string auto_corr_filename,
std::string checkpoint_filename )
```

#### **Parameters**

out	output	Output chains, shape is double[chain_N, N_steps,dimension]
	dimension	dimension of the parameter space being explored
	N_steps	Number of total steps to be taken, per chain
	chain_N	Number of chains
	initial_pos	Initial position in parameter space - shape double[dimension]
	seeding_var	Variance of the normal distribution used to seed each chain higher than 0 - shape double[dimension]
	chain_temps	Double array of temperatures for the chains
	swp_freq	the frequency with which chains are swapped
	log_prior	Funcion pointer for the log_prior
	log_likelihood	Function pointer for the log_likelihood
	fisher	Function pointer for the fisher - if NULL, fisher steps are not used
	numThreads	Number of threads to use (=1 is single threaded)
	pool	boolean to use stochastic chain swapping (MUST have >2 threads)
	show_prog	boolean whether to print out progress (for example, should be set to `'false" if submitting to a cluster)
	statistics_filename	Filename to output sampling statistics, if empty string, not output
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	checkpoint_filename	Filename to output data for checkpoint, if empty string, not saved

# 7.9.2.6 MCMC\_MH\_internal()

```
double * seeding_var,
double * chain_temps,
int swp_freq,
std::function< double(double *, int, int)> log_prior,
std::function< double(double *, int, int)> log_likelihood,
std::function< void(double *, int, double **, int)> fisher,
int numThreads,
bool pool,
bool show_prog,
std::string statistics_filename,
std::string auto_corr_filename,
std::string checkpoint_file)
```

Generic sampler, where the likelihood, prior are parameters supplied by the user.

Base of the sampler, generic, with user supplied quantities for most of the samplers properties

Uses the Metropolis-Hastings method, with the option for Fisher/MALA steps if the Fisher routine is supplied.

3 modes to use -

single threaded (numThreads = 1) runs single threaded

multi-threaded `'deterministic'' (numThreads>1; pool = false) progresses each chain in parallel for swp\_freq steps, then waits for all threads to complete before swapping temperatures in sequenctial order (j, j+1) then (j+1, j+2) etc (sequenctially)

multi-threaded `'stochastic'' (numThreads>2; pool = true) progresses each chain in parallel by queueing each temperature and evaluating them in the order they were submitted. Once finished, the threads are queued to swap, where they swapped in the order they are submitted. This means the chains are swapped randomly, and the chains do NOT finish at the same time. The sampler runs until the the 0th chain reaches the step number

Note on limits: In the prior function, if a set of parameters should be disallowed, return -std::numeric\_ $\leftarrow$  limits<double>::infinity() - (this is in the limits> file in std)

Format for the auto\_corr file (compatable with csv, dat, txt extensions): each row is a dimension of the cold chain, with the first row being the lengths used for the auto-corr calculation:

lengths: length1, length2...

dim1: length1, length2 ...

Format for the chain file (compatable with csv, dat, txt extensions): each row is a step, each column a dimension:

Step1: dim1, dim2, ...

Step2: dim1, dim2, ...

Statistics\_filename : should be txt extension

checkpoint\_file: This file saves the final position of all the chains, as well as other metadata, and can be loaded by the function <FUNCTION> to continue the chain from the point it left off. Not meant to be read by humans, the data order is custom to this software library. An empty string ("") means no checkpoint will be saved. For developers, the contents are:

dimension, # of chains

temps of chains

Stepping widths of all chains

Final position of all chains

### **Parameters**

out	output	Output chains, shape is double[chain_N, N_steps,dimension]
	dimension	dimension of the parameter space being explored
	N_steps	Number of total steps to be taken, per chain
	chain_N	Number of chains
	initial_pos	Initial position in parameter space - shape double[dimension]
	seeding_var	Variance of the normal distribution used to seed each chain higher than 0 - shape double[dimension]
	chain_temps	Double array of temperatures for the chains
	swp_freq	the frequency with which chains are swapped
	log_prior	std::function for the log_prior function – takes double *position, int dimension, int chain_id
	log_likelihood	std::function for the log_likelihood function – takes double *position, int dimension, int chain_id
	fisher	std::function for the fisher function – takes double *position, int dimension, double **output_fisher, int chain_id
	numThreads	Number of threads to use (=1 is single threaded)
	pool	boolean to use stochastic chain swapping (MUST have >2 threads)
	show_prog	boolean whether to print out progress (for example, should be set to `'false'' if submitting to a cluster)
	statistics_filename	Filename to output sampling statistics, if empty string, not output
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	checkpoint_file	Filename to output data for checkpoint, if empty string, not saved

### 7.9.2.7 MCMC\_MH\_loop()

```
void MCMC_MH_loop (
          sampler * sampler )
```

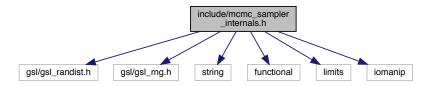
Internal function that runs the actual loop for the sampler.

# 7.10 include/mcmc\_sampler\_internals.h File Reference

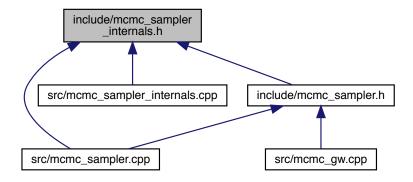
```
#include <gsl/gsl_randist.h>
#include <gsl/gsl_rng.h>
#include <string>
#include <functional>
#include <limits>
```

#include <iomanip>

Include dependency graph for mcmc\_sampler\_internals.h:



This graph shows which files directly or indirectly include this file:



#### Classes

· struct sampler

### **Functions**

- int mcmc\_step (sampler \*sampler, double \*current\_param, double \*next\_param, int chain\_number) interface function between the sampler and the internal step functions
- void gaussian\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param, int chain\_id) Straight gaussian step.
- void fisher\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param, int chain\_index) Fisher informed gaussian step.
- void **update fisher** (sampler \*sampler, double \*current param, int chain index)
- void mmala\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param)

  MMALA informed step Currently not supported.
- void diff\_ev\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param, int chain\_id) differential evolution informed step
- void chain\_swap (sampler \*sampler, double \*\*\*output, int step\_num, int \*swp\_accepted, int \*swp\_rejected) subroutine to perform chain comparison for parallel tempering

- int single\_chain\_swap (sampler \*sampler, double \*chain1, double \*chain2, int T1\_index, int T2\_index) subroutine to actually swap two chains
- void assign\_probabilities (sampler \*sampler, int chain\_index)

update and initiate probabilities for each variety of step

- void allocate\_sampler\_mem (sampler \*sampler)
- void deallocate\_sampler\_mem (sampler \*sampler)
- void update history (sampler \*sampler, double \*new params, int chain index)
- void write\_stat\_file (sampler \*sampler, std::string filename)
- void write\_checkpoint\_file (sampler \*sampler, std::string filename)

Routine that writes metadata and final positions of a sampler to a checkpoint file.

void load\_checkpoint\_file (std::string check\_file, sampler \*sampler)

load checkpoint file into sampler struct

- void assign\_ct\_p (sampler \*sampler, int step, int chain\_index)
- void assign\_ct\_m (sampler \*sampler, int step, int chain\_index)

#### **Variables**

const double limit\_inf = -std::numeric\_limits<double>::infinity()

### 7.10.1 Detailed Description

Internal functions of the generic MCMC sampler (nothing specific to GW)

### 7.10.2 Function Documentation

### 7.10.2.1 assign\_probabilities()

update and initiate probabilities for each variety of step

Type 0: Gaussian step

Type 1: Differential Evolution step

Type 2: MMALA step (currently not supported)

Type 3: Fisher step

### 7.10.2.2 chain\_swap()

subroutine to perform chain comparison for parallel tempering

The total output file is passed, and the chains are swapped sequentially

This is the routine for 'Deterministic' sampling (parallel or sequential, but not pooled)

### **Parameters**

sampler	sampler struct	
output	output vector containing chains	
step_num	current step number	

### 7.10.2.3 diff\_ev\_step()

differential evolution informed step

Differential evolution uses the past history of the chain to inform the proposed step:

Take the difference of two random, accepted previous steps and step along that with some step size, determined by a gaussian

### **Parameters**

	sampler	Sampler struct
	current_param	current position in parameter space
out	proposed_param	Proposed position in parameter space

# 7.10.2.4 fisher\_step()

Fisher informed gaussian step.

	sampler	Sampler struct
	current_param	current position in parameter space
out	proposed_param	Proposed position in parameter space

### 7.10.2.5 gaussian\_step()

Straight gaussian step.

#### **Parameters**

	sampler	Sampler struct
	current_param	current position in parameter space
out	proposed_param	Proposed position in parameter space

# 7.10.2.6 load\_checkpoint\_file()

load checkpoint file into sampler struct

NOTE – allocate\_sampler called in function – MUST deallocate manually

NOTE - sampler->chain\_temps allocated internally - MUST free manually

### 7.10.2.7 mmala\_step()

MMALA informed step - Currently not supported.

# **Parameters**

sampler		Sampler struct
	current_param	current position in parameter space
out	proposed_param	Proposed position in parameter space

# 7.10.2.8 single\_chain\_swap()

```
int single_chain_swap (
          sampler * sampler,
```

```
double * chain1,
double * chain2,
int T1_index,
int T2_index )
```

subroutine to actually swap two chains

This is the more general subroutine, which just swaps the two chains passed to the function

#### **Parameters**

sampler	sampler structure
chain1	parameter position of chain that could be changed
chain2	chain that is not swapped, but provides parameters to be swapped by the other chain
T1_index	number of chain swappe in chain_temps
T2_index	number of chain swapper in chain_temps

### 7.10.2.9 write\_checkpoint\_file()

Routine that writes metadata and final positions of a sampler to a checkpoint file.

# 7.10.3 Variable Documentation

```
7.10.3.1 limit_inf
```

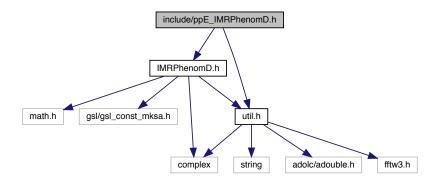
```
const double limit_inf = -std::numeric_limits<double>::infinity()
```

Structure storing everything that defines an instance of the sampler

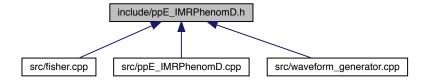
# 7.11 include/ppE\_IMRPhenomD.h File Reference

```
#include "IMRPhenomD.h"
#include "util.h"
```

Include dependency graph for ppE\_IMRPhenomD.h:



This graph shows which files directly or indirectly include this file:



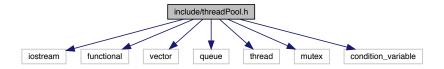
### Classes

- class ppE\_IMRPhenomD\_Inspiral < T >
- class ppE IMRPhenomD IMR< T >
- class dCS\_IMRPhenomD\_log< T >
- class dCS\_IMRPhenomD< T >
- class EdGB\_IMRPhenomD\_log< T >
- class EdGB\_IMRPhenomD< T >

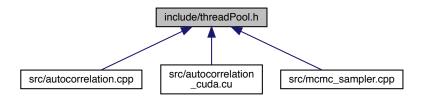
### 7.12 include/threadPool.h File Reference

```
#include <iostream>
#include <functional>
#include <vector>
#include <queue>
#include <thread>
#include <mutex>
```

#include <condition\_variable>
Include dependency graph for threadPool.h:



This graph shows which files directly or indirectly include this file:



### **Classes**

- class default\_comp< jobtype >
   Default comparator for priority\_queue in threadPool no comparison.
- class threadPool< jobtype, comparator >

Class for creating a pool of threads to asynchronously distribute work.

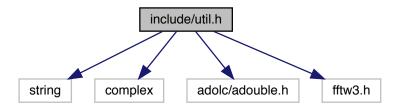
### 7.12.1 Detailed Description

Header file (declarations and definitions because of template functions) for the implementation of a generic thread pool

# 7.13 include/util.h File Reference

```
#include <string>
#include <complex>
#include "adolc/adouble.h"
#include <fftw3.h>
#include <gsl/gsl_interp.h>
#include <gsl/gsl_spline.h>
```

#include <gsl/gsl\_errno.h>
Include dependency graph for util.h:



This graph shows which files directly or indirectly include this file:



### Classes

- · struct fftw\_outline
- struct sph\_harm< T >
- struct gen\_params
- struct useful\_powers< T >

To speed up calculations within the for loops, we pre-calculate reoccuring powers of M\*F and Pi, since the pow() function is prohibatively slow.

struct source\_parameters< T >

### **Macros**

- #define PBSTR "||||||||||||
- #define PBWIDTH 60

### **Functions**

- void initiate\_LumD\_Z\_interp (gsl\_interp\_accel \*\*Z\_DL\_accel\_ptr, gsl\_spline \*\*Z\_DL\_spline\_ptr)
   Function that uses the GSL libraries to interpolate pre-calculated Z-D\_L data.
- void free\_LumD\_Z\_interp (gsl\_interp\_accel \*\*Z\_DL\_accel\_ptr, gsl\_spline \*\*Z\_DL\_spline\_ptr)

  Frees the allocated interpolation function.
- adouble Z\_from\_DL\_interp (adouble DL, gsl\_interp\_accel \*Z\_DL\_accel\_ptr, gsl\_spline \*Z\_DL\_spline\_ptr)
- $\bullet \ \ double \ \textbf{Z\_from\_DL\_interp} \ \ (double \ \textbf{DL}, \ gsl\_interp\_accel \ *\textbf{Z\_DL\_accel\_ptr}, \ gsl\_spline \ *\textbf{Z\_DL\_spline\_ptr})$
- double Z\_from\_DL (double DL, std::string cosmology)

Calculates the redshift given the luminosity distance.

double DL\_from\_Z (double Z, std::string cosmology)

Calculates the luminosity distance given the redshift.

double cosmology interpolation function (double x, double \*coeffs, int interp degree)

Custom interpolation function used in the cosmology calculations.

double cosmology\_lookup (std::string cosmology)

Helper function for mapping cosmology name to an internal index.

adouble Z from DL (adouble DL, std::string cosmology)

Calculates the redshift given the luminosity distance adouble version for ADOL-C implementation.

adouble DL\_from\_Z (adouble Z, std::string cosmology)

Calculates the luminosity distance given the redshift adouble version for ADOL-C implementation.

• adouble cosmology interpolation function (adouble x, double \*coeffs, int interp degree)

Custom interpolation function used in the cosmology calculations adouble version for ADOL-C.

void printProgress (double percentage)

routine to print the progress of a process to the terminal as a progress bar

· void initiate likelihood function (fftw outline \*plan, int length)

Allocate memory for FFTW3 methods used in a lot of inner products input is a locally defined structure that houses all the pertinent data.

void allocate FFTW3 mem inverse (fftw outline \*plan, int length)

Allocate memory for FFTW3 methods used in a lot of inner products –INVERSE input is a locally defined structure that houses all the pertinent data.

void deactivate\_likelihood\_function (fftw\_outline \*plan)

deallocates the memory used for FFTW routines

double \*\* allocate 2D array (int dim1, int dim2)

Utility to malloc 2D array.

void deallocate\_2D\_array (double \*\*array, int dim1, int dim2)

Utility to free malloc'd 2D array.

double \*\*\* allocate 3D array (int dim1, int dim2, int dim3)

Utility to malloc 3D array.

void deallocate\_3D\_array (double \*\*\*array, int dim1, int dim2, int dim3)

Utility to free malloc'd 2D array.

• void read\_file (std::string filename, double \*\*output, int rows, int cols)

Utility to read in data.

• void read\_file (std::string filename, double \*output)

Utility to read in data (single dimension vector)

void read\_LOSC\_data\_file (std::string filename, double \*output, double \*data\_start\_time, double \*duration, double \*fs)

Read data file from LIGO Open Science Center.

• void read\_LOSC\_PSD\_file (std::string filename, double \*\*output, int rows, int cols)

Read PSD file from LIGO Open Science Center.

• void allocate\_LOSC\_data (std::string \*data\_files, std::string psd\_file, int num\_detectors, int psd\_length, int data\_file\_length, double trigger\_time, std::complex< double > \*\*data, double \*\*psds, double \*\*freqs)

Prepare data for MCMC directly from LIGO Open Science Center.

- void free\_LOSC\_data (std::complex < double > \*\*data, double \*\*psds, double \*\*freqs, int num\_detectors, int length)
- void tukey\_window (double \*window, int length, double alpha)

Tukey window function for FFTs.

void write\_file (std::string filename, double \*\*input, int rows, int cols)

Utility to write 2D array to file.

void write\_file (std::string filename, double \*input, int length)

Utility to write 1D array to file.

• double calculate\_eta (double mass1, double mass2)

Calculates the symmetric mass ration from the two component masses.

- adouble calculate\_eta (adouble mass1, adouble mass2)
- double calculate\_chirpmass (double mass1, double mass2)

Calculates the chirp mass from the two component masses.

- adouble calculate\_chirpmass (adouble mass1, adouble mass2)
- double calculate mass1 (double chirpmass, double eta)

Calculates the larger mass given a chirp mass and symmetric mass ratio.

- adouble calculate\_mass1 (adouble chirpmass, adouble eta)
- double calculate mass2 (double chirpmass, double eta)

Calculates the smaller mass given a chirp mass and symmetric mass ratio.

- adouble calculate\_mass2 (adouble chirpmass, adouble eta)
- void celestial\_horizon\_transform (double RA, double DEC, double gps\_time, double LONG, double LAT, double \*phi, double \*theta)

Utility to transform from celestial coord RA and DEC to local horizon coord for detector response functions.

double gps\_to\_GMST (double gps\_time)

Utility to transform from gps time to GMST https://aa.usno.navy.mil/faq/docs/GAST.php.

double gps\_to\_JD (double gps\_time)

Utility to transform from gps to JD.

void transform\_cart\_sph (double \*cartvec, double \*sphvec)

utility to transform a vector from cartesian to spherical (radian)

void transform\_sph\_cart (double \*sphvec, double \*cartvec)

utility to transform a vector from spherical (radian) to cartesian

template<class T >

T trapezoidal\_sum\_uniform (double delta\_x, int length, T \*integrand)

Trapezoidal sum rule to approximate discrete integral - Uniform spacing.

template < class T >

T trapezoidal sum (double \*delta x, int length, T \*integrand)

Trapezoidal sum rule to approximate discrete integral - Non-Uniform spacing.

template < class T >

T simpsons\_sum (double delta\_x, int length, T \*integrand)

Simpsons sum rule to approximate discrete integral - Uniform spacing.

• long factorial (long num)

Local function to calculate a factorial.

double pow\_int (double base, int power)

Local power function, specifically for integer powers.

- adouble pow\_int (adouble base, int power)
- template<class T >

std::complex < T > cpolar (T mag, T phase)

template<class T >

std::complex< T > XLALSpinWeightedSphericalHarmonic (T theta, T phi, int s, int I, int m)

double cbrt\_internal (double base)

Fucntion that just returns the cuberoot.

adouble cbrt\_internal (adouble base)

Fucntion that just returns the cuberoot ADOL-C doesn't have the cbrt function (which is faster), so have to use the power function.

### **Variables**

- const double gamma E = 0.5772156649015328606065120900824024310421
- const double c = 299792458.
- const double G =6.674e-11\*(1.98855e30)
- const double MSOL SEC =4.925491025543575903411922162094833998e-6
- const double MPC\_SEC = 3.085677581491367278913937957796471611e22/c

# 7.13.1 Detailed Description

General utilities (functions and structures) independent of modelling method

#### 7.13.2 Function Documentation

### 7.13.2.2 allocate\_3D\_array()

Utility to malloc 3D array.

### 7.13.2.3 allocate\_LOSC\_data()

```
void allocate_LOSC_data (
    std::string * data_files,
    std::string psd_file,
    int num_detectors,
    int psd_length,
    int data_file_length,
    double trigger_time,
    std::complex< double > ** data,
    double ** psds,
    double ** freqs )
```

Prepare data for MCMC directly from LIGO Open Science Center.

Trims data for Tobs (determined by PSD file) 3/4\*Tobs in front of trigger, and 1/4\*Tobs behind

Currently, default to sampling frequency and observation time set by PSD - cannot be customized

Output is in order of PSD columns - string vector of detectos MUST match order of PSD cols

Output shapes—psds = [num\_detectors][psd\_length] data = [num\_detectors][psd\_length]

freqs = [num\_detectors][psd\_length]

Total observation time = 1/( freq[i] - freq[i-1]) (from PSD file)

Sampling frequency fs = max frequency from PSD file

ALLOCATES MEMORY - must be freed to prevent memory leak

#### **Parameters**

	data_files	Vector of strings for each detector file from LOSC
	psd_file	String of psd file from LOSC
	num_detectors	Number of detectors to use
	psd_length	Length of the PSD file (number of rows of DATA)
	data_file_length	Length of the data file (number of rows of DATA)
	trigger_time	Time for the signal trigger (GPS)
out	data	Output array of data for each detector
out	psds	Output array of psds for each detector
out	freqs	Output array of freqs for each detector

### 7.13.2.4 calculate\_chirpmass()

Calculates the chirp mass from the two component masses.

The output units are whatever units the input masses are

#### 7.13.2.5 calculate\_mass1()

Calculates the larger mass given a chirp mass and symmetric mass ratio.

Units of the output match the units of the input chirp mass

### 7.13.2.6 calculate\_mass2()

Calculates the smaller mass given a chirp mass and symmetric mass ratio.

Units of the output match the units of the input chirp mass

### 7.13.2.7 celestial\_horizon\_transform()

Utility to transform from celestial coord RA and DEC to local horizon coord for detector response functions.

Outputs are the spherical polar angles defined by North as 0 degrees azimuth and the normal to the earth as 0 degree polar

### **Parameters**

	RA	Right acsension (rad)
	DEC	Declination (rad)
	gps_time	GPS time
	LONG	Longitude (rad)
	LAT	Latitude (rad)
out	phi	horizon azimuthal angle (rad)
out	theta	horizon polar angle (rad)

### 7.13.2.8 cosmology\_interpolation\_function()

Custom interpolation function used in the cosmology calculations.

Power series in half power increments of x, up to 11/2. powers of x

# 7.13.2.9 deallocate\_2D\_array()

Utility to free malloc'd 2D array.

# 7.13.2.10 deallocate\_3D\_array()

Utility to free malloc'd 2D array.

```
7.13.2.11 DL_from_Z()
```

```
double DL_from_Z ( \label{eq:cosmology} \mbox{double $Z$,} \\ \mbox{std::string $cosmology$ )}
```

Calculates the luminosity distance given the redshift.

Based on Astropy.cosmology calculations – see python script in the ./data folder of the project – numerically calculated given astropy.cosmology's definitions ( http://docs.astropy.org/en/stable/cosmology/) and used scipy.optimize to fit to a power series, stepping in half powers of Z. These coefficients are then output to a header file (D\_Z\_config.h) which are used here to calculate distance. Custom cosmologies etc can easily be acheived by editing the python script D\_Z\_config.py, the c++ functions do not need modification. They use whatever data is available in the header file. If the functional form of the fitting function changes, these functions DO need to change.

5 cosmological models are available (this argument must be spelled exactly):

PLANCK15, PLANCK13, WMAP9, WMAP7, WMAP5

```
7.13.2.12 free_LOSC_data()
```

/brief Free data allocated by prep LOSC data function

#### 7.13.2.13 initiate\_LumD\_Z\_interp()

Function that uses the GSL libraries to interpolate pre-calculated Z-D\_L data.

Initiates the required functions - GSL interpolation requires allocating memory before hand

### 7.13.2.14 pow\_int()

Local power function, specifically for integer powers.

Much faster than the std version, because this is only for integer powers

### 7.13.2.15 printProgress()

routine to print the progress of a process to the terminal as a progress bar

Call everytime you want the progress printed

Utility to read in data.

Takes filename, and assigns to output[rows][cols]

File must be comma separated doubles

### **Parameters**

	filename	input filename, relative to execution directory
out	output	array to store output, dimensions rowsXcols
	rows	first dimension
	cols	second dimension

Utility to read in data (single dimension vector)

Takes filename, and assigns to output[i\*rows + cols]

Output vector must be long enough, no check is done for the length

File must be comma separated doubles

	filename	input filename, relative to execution directory
out	output	output array, assumed to have the proper length of total items

### 7.13.2.18 read\_LOSC\_data\_file()

```
void read_LOSC_data_file (
    std::string filename,
    double * output,
    double * data_start_time,
    double * duration,
    double * fs )
```

Read data file from LIGO Open Science Center.

Convenience function for cutting off the first few lines of text

### **Parameters**

	filename	input filename
out	output	Output data
out	data_start_time	GPS start time of the data in file
out	duration	Duration of the signal
out	fs	Sampling frequency of the data

### 7.13.2.19 read\_LOSC\_PSD\_file()

```
void read_LOSC_PSD_file (
          std::string filename,
          double ** output,
          int rows,
          int cols )
```

Read PSD file from LIGO Open Science Center.

Convenience function for cutting off the first few lines of text

# 7.13.2.20 simpsons\_sum()

Simpsons sum rule to approximate discrete integral - Uniform spacing.

More accurate than the trapezoidal rule, but must be uniform

### 7.13.2.21 transform\_cart\_sph()

utility to transform a vector from cartesian to spherical (radian)

order:

```
cart: x, y, z
```

spherical: r, polar, azimuthal

### 7.13.2.22 transform\_sph\_cart()

utility to transform a vector from spherical (radian) to cartesian

order:

```
cart: x, y, z
```

spherical: r, polar, azimuthal

### 7.13.2.23 trapezoidal\_sum()

Trapezoidal sum rule to approximate discrete integral - Non-Uniform spacing.

This version is slower than the uniform version, but will handle non-uniform spacing

#### 7.13.2.24 trapezoidal\_sum\_uniform()

Trapezoidal sum rule to approximate discrete integral - Uniform spacing.

This version is faster than the general version, as it has half the function calls

Something may be wrong with this function - had an overall offset for real data that was fixed by using the simpsons rule - not sure if this was because of a boost in accuracy or because something is off with the trapezoidal sum

### 7.13.2.25 tukey\_window()

Tukey window function for FFTs.

As defined by https://en.wikipedia.org/wiki/Window\_function

Utility to write 2D array to file.

Grid of data, comma separated

Grid has rows rows and cols columns

### **Parameters**

filename	Filename of output file, relative to execution directory
input	Input 2D array pointer array[rows][cols]
rows	First dimension of array
cols	second dimension of array

```
7.13.2.27 write_file() [2/2]
```

Utility to write 1D array to file.

Single column of data

filename	Filename of output file, relative to execution directory	
input	input 1D array pointer array[length]	
length	length of array	

### 7.13.2.28 XLALSpinWeightedSphericalHarmonic()

### Shamelessly stolen from LALsuite

#### **Parameters**

theta	polar angle (rad)
phi	azimuthal angle (rad)
s	spin weight
1	mode number l
m	mode number m

### 7.13.2.29 Z\_from\_DL()

Calculates the redshift given the luminosity distance.

Based on Astropy.cosmology calculations – see python script in the ./data folder of the project – numerically calculated given astropy.cosmology's definitions ( http://docs.astropy.org/en/stable/cosmology/) and used scipy.optimize to fit to a power series, stepping in half powers of DL. These coefficients are then output to a header file (D\_Z\_config.h) which are used here to calculate redshift. Custom cosmologies etc can easily be acheived by editing the python script D\_Z\_config.py, the c++ functions do not need modification. They use whatever data is available in the header file.

5 cosmological models are available (this argument must be spelled exactly, although case insensitive):

PLANCK15, PLANCK13, WMAP9, WMAP7, WMAP5

Function that returns Z from a given luminosity Distance – only Planck15

adouble version for ADOL-C calculations

```
7.13.2.31 Z_from_DL_interp() [2/2]
double Z_from_DL_interp (
              double DL,
              {\tt gsl\_interp\_accel} \ * \ {\tt Z\_DL\_accel\_ptr},
              {\tt gsl\_spline} \ * \ {\it Z\_DL\_spline\_ptr} \ )
Function that returns Z from a given luminosity Distance – only Planck15
7.13.3 Variable Documentation
7.13.3.1 c
const double c = 299792458.
Speed of light m/s
7.13.3.2 G
const double G = 6.674e - 11*(1.98855e30)
Gravitational constant in m**3/(s**2 SolMass)
7.13.3.3 gamma_E
const double gamma_E = 0.5772156649015328606065120900824024310421
Euler number
7.13.3.4 MPC_SEC
const double MPC_SEC = 3.085677581491367278913937957796471611e22/c
consts.kpc.to('m')*1000/c Mpc in sec
7.13.3.5 MSOL_SEC
```

#### Generated by Doxygen

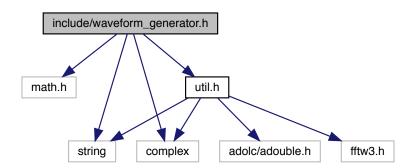
G/c\*\*3 seconds per solar mass

const double MSOL\_SEC =4.925491025543575903411922162094833998e-6

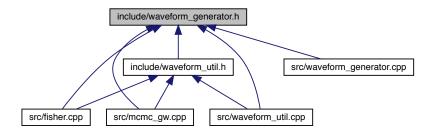
# 7.14 include/waveform\_generator.h File Reference

```
#include <math.h>
#include "util.h"
#include <complex>
#include <string>
```

Include dependency graph for waveform\_generator.h:



This graph shows which files directly or indirectly include this file:



#### **Functions**

- int fourier\_waveform (double \*frequencies, int length, std::complex< double > \*waveform\_plus, std
   ::complex< double > \*waveform\_cross, std::string generation\_method, gen\_params \*parameters)
- int fourier\_waveform (double \*frequencies, int length, double \*waveform\_plus\_real, double \*waveform
   \_plus\_imag, double \*waveform\_cross\_real, double \*waveform\_cross\_imag, std::string generation\_method,
   gen\_params \*parameters)
- int **fourier\_waveform** (double \*frequencies, int length, std::complex< double > \*waveform, std::string generation\_method, gen\_params \*parameters)
- int **fourier\_waveform** (double \*frequencies, int length, double \*waveform\_real, double \*waveform\_imag, std::string generation\_method, gen\_params \*parameters)
- int **fourier\_amplitude** (double \*frequencies, int length, double \*amplitude, std::string generation\_method, gen\_params \*parameters)
- int **fourier\_phase** (double \*frequencies, int length, double \*phase, std::string generation\_method, gen\_params \*parameters)

# 7.15 include/waveform\_generator\_C.h File Reference

#### **Functions**

- int fourier\_waveformC (double \*frequencies, int length, double \*waveform\_plus\_real, double \*waveform\_cross\_real, double \*waveform\_cross\_imag, char \*generation\_method, double mass1, double mass2, double DL, double spin1x, double spin1y, double spin1z, double spin2x, double spin2x, double spin2y, double spin2z, double tc, double f\_ref, double phiRef, double \*ppE\_beta, int \*ppE\_b, int Nmod, double incl\_angle, double theta, double phi)
- int **fourier\_amplitudeC** (double \*frequencies, int length, double \*amplitude, char \*generation\_method, double mass1, double mass2, double DL, double spin1x, double spin1y, double spin1z, double spin2x, double spin2x, double incl\_angle, double theta, double phi)
- int **fourier\_phaseC** (double \*frequencies, int length, double \*phase, char \*generation\_method, double mass1, double mass2, double DL, double spin1x, double spin1y, double spin1z, double spin2x, double spin2x, double spin2z, double phic, double tc, double f\_ref, double phiRef, double \*ppE\_beta, int \*ppE\_b, int Nmod, double incl\_angle, double theta, double phi)
- void initiate LumD Z interp C ()
- void free LumD Z interp C ()

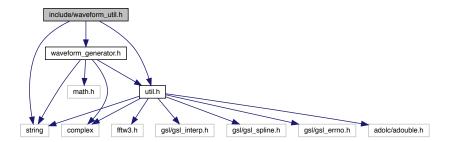
### 7.15.1 Detailed Description

Header file for the C wrapping of the waveform\_generation.cpp

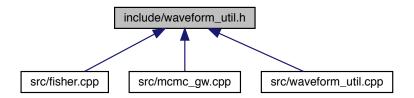
# 7.16 include/waveform\_util.h File Reference

```
#include "waveform_generator.h"
#include "util.h"
#include <string>
```

Include dependency graph for waveform\_util.h:



This graph shows which files directly or indirectly include this file:



### **Functions**

double data\_snr\_maximized\_extrinsic (double \*frequencies, int length, std::complex< double > \*data, double \*psd, std::string detector, std::string generation\_method, gen\_params \*param)

Utility to calculate the snr of a fourier transformed data stream while maximizing over the coalescence parameters phic and tc.

- double data\_snr\_maximized\_extrinsic (double \*frequencies, int length, double \*data\_real, double \*data\_
  imag, double \*psd, std::string detector, std::string generation\_method, gen\_params \*param)
  - Light wrapper for the data\_snr\_maximized\_extrinsic method.
- double calculate\_snr (std::string detector, std::complex< double > \*waveform, double \*frequencies, int length)

Caclulates the snr given a detector and waveform (complex) and frequencies.

- int fourier\_detector\_response (double \*frequencies, int length, std::complex< double > \*hplus, std
   ::complex< double > \*hcross, std::complex< double > \*detector\_response, double theta, double phi, std
   ::string detector)
- int fourier\_detector\_response (double \*frequencies, int length, std::complex< double > \*response, std
  ::string detector, std::string generation method, gen\_params \*parameters)

Function to produce the detector response caused by impinging gravitational waves from a quasi-circular binary.

• int fourier\_detector\_amplitude\_phase (double \*frequencies, int length, double \*amplitude, double \*phase, std::string detector, std::string generation\_method, gen\_params \*parameters)

Calculates the amplitude (magnitude) and phase (argument) of the response of a given detector.

# 7.16.1 Detailed Description

Header file for waveform specific utilites

### 7.16.2 Function Documentation

### 7.16.2.1 calculate\_snr()

Caclulates the snr given a detector and waveform (complex) and frequencies.

This function computes the un-normalized snr: \sqrt( ( H | H ) )

#### **Parameters**

detector	detector name - must match the string of populate_noise precisely
waveform	complex waveform
frequencies	double array of frequencies that the waveform is evaluated at
length	length of the above two arrays

### 7.16.2.2 data\_snr\_maximized\_extrinsic() [1/2]

Utility to calculate the snr of a fourier transformed data stream while maximizing over the coalescence parameters phic and tc.

The gen\_params structure holds the parameters for the template to be used (the maximimum likelihood parameters)

#### **Parameters**

frequencies	Frequencies used by data
length	length of the data
data	input data in the fourier domain
psd	PSD for the detector that created the data
detector	Name of the detector –See noise_util for options
generation_method	Generation method for the template – See waveform_generation.cpp for options
param	gen_params structure for the template

# 7.16.2.3 data\_snr\_maximized\_extrinsic() [2/2]

Light wrapper for the data\_snr\_maximized\_extrinsic method.

Splits the data into real and imaginary, so all the arguments are C-safe

#### **Parameters**

frequencies	Frequencies used by data	
length	length of the data	
data_real	input data in the fourier domain – real part	
data_imag	input data in the fourier domain – imaginary part	
psd	PSD for the detector that created the data	
detector	Name of the detector –See noise_util for options	
generation_method	Generation method for the template – See waveform_generation.cpp for options	
param	gen_params structure for the template	

### 7.16.2.4 fourier\_detector\_amplitude\_phase()

Calculates the amplitude (magnitude) and phase (argument) of the response of a given detector.

This is for general waveforms, and will work for precessing waveforms

Not as fast as non-precessing, but that can't be helped. MUST include plus/cross polarizations

### 7.16.2.5 fourier\_detector\_response() [1/2]

	frequencies	array of frequencies corresponding to waveform
	length	length of frequency/waveform arrays
	hcross	precomputed cross polarization of the waveform
out	detector_response	detector response
	theta	polar angle (rad) theta in detector frame
	phi	azimuthal angle (rad) phi in detector frame
	detector	detector - list of supported detectors in noise_util

### 7.16.2.6 fourier\_detector\_response() [2/2]

Function to produce the detector response caused by impinging gravitational waves from a quasi-circular binary.

By using the structure parameter, the function is allowed to be more flexible in using different method of waveform generation - not all methods use the same parameters

This puts the responsibility on the user to pass the necessary parameters

Detector options include classic interferometers like LIGO/VIRGO (coming soon: ET and LISA)

This is a wrapper that combines generation with response functions: if producing mulitple responses for one waveform (ie stacking Hanford, Livingston, and VIRGO), it will be considerably more efficient to calculate the waveform once, then combine each response manually

#### **Parameters**

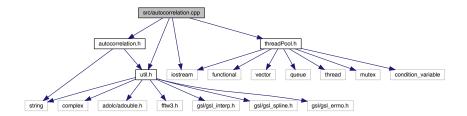
	frequencies	double array of frequencies for the waveform to be evaluated at
	length	integer length of all the arrays
out	response	complex array for the output plus polarization waveform
	generation_method	String that corresponds to the generation method - MUST BE SPELLED EXACTLY
	parameters	structure containing all the source parameters

# 7.17 README.dox File Reference

# 7.18 src/autocorrelation.cpp File Reference

```
#include "autocorrelation.h"
#include "util.h"
#include "threadPool.h"
#include <iostream>
```

Include dependency graph for autocorrelation.cpp:



### **Macros**

#define MAX SERIAL 200000

#### **Functions**

- void write\_auto\_corr\_file\_from\_data\_file (std::string autocorr\_filename, std::string datafile, int length, int dimension, int num\_segments, double target\_corr, int num\_threads)
- void write\_auto\_corr\_file\_from\_data (std::string autocorr\_filename, double \*\*data, int length, int dimension, int num\_segments, double target\_corr, int num\_threads)
- void auto\_corr\_from\_data (double \*\*data, int length, int dimension, int \*\*output, int num\_segments, double target\_corr, int num\_threads)

Calculates the autocorrelation length for a set of data for a number of segments for each dimension – completely host code, utilitizes FFTW3 for longer chuncks of the chains.

- void threaded\_ac\_spectral (int thread, threaded\_ac\_jobs\_fft job)
- void threaded\_ac\_serial (int thread, threaded\_ac\_jobs\_serial job)
- double auto\_correlation\_serial (double \*arr, int length, int start, double target)
- void auto\_correlation\_spectral (double \*chain, int length, double \*autocorr, fftw\_outline \*plan\_forw, fftw\_outline \*plan\_rev)
- void auto\_correlation\_spectral (double \*chain, int length, int start, double \*autocorr, fftw\_outline \*plan\_forw, fftw\_outline \*plan\_rev)

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT – accepts FFTW plan as argument for plan reuse and multi-threaded applications.

void auto\_correlation\_spectral (double \*chain, int length, double \*autocorr)

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT.

- double auto\_correlation (double \*arr, int length, double tolerance)
- double auto\_correlation\_serial\_old (double \*arr, int length)

Grid search method of computing the autocorrelation.

• double auto correlation internal (double \*arr, int length, int lag, double ave)

Internal function to compute the auto correlation for a given lag.

void auto\_corr\_intervals\_outdated (double \*data, int length, double \*output, int num\_segments, double accuracy)

Function that computes the autocorrelation length on an array of data at set intervals to help determine convergence.

- void write\_auto\_corr\_file\_from\_data (std::string auto\_corr\_filename, double \*\*output, int intervals, int dimension, int N steps)
- void write\_auto\_corr\_file\_from\_data\_file (std::string auto\_corr\_filename, std::string output\_file, int intervals, int dimension, int N\_steps)

## 7.18.1 Detailed Description

Turns out calculating the autocorrelation is more complicated if you want to do it fast, so it gets its own file now

## 7.18.2 Macro Definition Documentation

## 7.18.2.1 MAX\_SERIAL

```
#define MAX_SERIAL 200000
```

Max length of array to use serial calculation

## 7.18.3 Function Documentation

#### 7.18.3.1 auto\_corr\_from\_data()

Calculates the autocorrelation length for a set of data for a number of segments for each dimension – completely host code, utilitizes FFTW3 for longer chuncks of the chains.

Takes in the data from a sampler, shape data[N\_steps][dimension]

Outputs lags that correspond to the target\_corr – shape output[dimension][num\_segments]

#### **Parameters**

	data	Input data	
	length	length of input data	
	dimension	dimension of data	
out	output	array that stores the auto-corr lengths – array[num_segments]	
	num_segments	segments number of segements to compute the auto-corr length	
	target_corr	Autocorrelation for which the autocorrelation length is defined (lag of autocorrelation for which it equals the target_corr)	
	num_threads	Total number of threads to use	

#### 7.18.3.2 auto\_corr\_intervals\_outdated()

Function that computes the autocorrelation length on an array of data at set intervals to help determine convergence.

outdated version - new version uses FFTs

#### **Parameters**

	data	Input data
	length	length of input data
out	output	array that stores the auto-corr lengths – array[num_segments]
	num_segments	number of segements to compute the auto-corr length
	accuracy	longer chains are computed numerically, this specifies the tolerance

## 7.18.3.3 auto\_correlation\_grid\_search()

Grid search method of computing the autocorrelation.

Hopefully more reliable than the box-search method, which can sometimes get caught in a recursive loop when the stepsize isn't tuned, but also faster than the basic linear, serial search

### **Parameters**

arr	Input array to use for autocorrelation
length	Length of input array
box_num	number of boxes to use for each iteration, default is 10
final_length	number of elements per box at which the grid search ends and the serial calculation begins
target_length	target correlation that corresponds to the returned lag

## 7.18.3.4 auto\_correlation\_internal()

```
int length,
int lag,
double ave )
```

Internal function to compute the auto correlation for a given lag.

#### 7.18.3.5 auto\_correlation\_spectral() [1/2]

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT – accepts FFTW plan as argument for plan reuse and multi-threaded applications.

Based on the Wiener-Khinchin Theorem.

```
Algorithm used from https://lingpipe-blog.com/2012/06/08/autocorrelation-fft-kiss-eigen/
```

NOTE the length used in initializing the fftw plans should be L = pow(2, std::ceil( std::log2(length) ) ) – the plans are padded so the total length is a power of two

Option to provide starting index for multi-dimension arrays in collapsed to one dimension

length is the length of the segment to be analyzed, not necessarily the dimension of the chain

#### 7.18.3.6 auto\_correlation\_spectral() [2/2]

Faster approximation of the autocorrelation of a chain. Implements FFT/IFFT.

Based on the Wiener-Khinchin Theorem.

Algorithm used from https://lingpipe-blog.com/2012/06/08/autocorrelation-fft-kiss-eigen/

## 7.18.3.7 write\_auto\_corr\_file\_from\_data()

```
void write_auto_corr_file_from_data (
    std::string autocorr_filename,
    double ** data,
    int length,
    int dimension,
    int num_segments,
    double target_corr,
    int num_threads)
```

#### **Parameters**

length	length of input data
dimension	dimension of data
num_segments	number of segements to compute the auto-corr length
target_corr	Autocorrelation for which the autocorrelation length is defined (lag of autocorrelation for which it equals the target_corr)
num_threads	Total number of threads to use

#### 7.18.3.8 write\_auto\_corr\_file\_from\_data\_file()

```
void write_auto_corr_file_from_data_file (
    std::string autocorr_filename,
    std::string datafile,
    int length,
    int dimension,
    int num_segments,
    double target_corr,
    int num_threads )
```

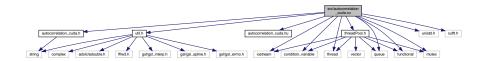
#### **Parameters**

length	length of input data
dimension	dimension of data
num_segments	number of segements to compute the auto-corr length
target_corr	Autocorrelation for which the autocorrelation length is defined (lag of autocorrelation for which it equals the target_corr)
num_threads	Total number of threads to use

# 7.19 src/autocorrelation\_cuda.cu File Reference

```
#include "autocorrelation_cuda.h"
#include "autocorrelation_cuda.hu"
#include "util.h"
#include <iostream>
#include <condition_variable>
#include <thread>
#include <queue>
#include <functional>
#include <mutex>
#include <unistd.h>
#include <threadPool.h>
#include <cufft.h>
```

Include dependency graph for autocorrelation\_cuda.cu:



#### **Functions**

\_\_device\_\_ \_\_host\_\_ void auto\_corr\_internal (double \*arr, int length, int lag, double average, double \*corr, int start\_id)

Internal function to calculate the autocorrelation for a given lag Customized for the thread pool architecture, with extra arguments because of the way the memory is allocated.

• \_\_global\_\_ void auto\_corr\_internal\_kernal (double \*arr, int length, double average, int \*rho\_index, double target\_corr, double var, int start\_id)

Internal function to launch the CUDA kernel for a range of autocorrelations.

void write\_file\_auto\_corr\_from\_data\_file\_accel (std::string acfile, std::string chains\_file, int dimension, int N
 \_ steps, int num\_segments, double target\_corr)

Write data file for autocorrelation lengths of the data given a data file name, as written by the mcmc\_sampler.

• void write\_file\_auto\_corr\_from\_data\_accel (std::string acfile, double \*\*chains, int dimension, int N\_steps, int num segments, double target corr)

Write data file given output chains, as formatted by the mcmc\_sampler.

 void auto\_corr\_from\_data\_accel (double \*\*output, int dimension, int N\_steps, int num\_segments, double target\_corr, double \*\*autocorr)

Find autocorrelation of data at different points in the chain length and output to autocorr.

void ac\_gpu\_wrapper (int thread, int job\_id)

Wrapper function for the thread pool.

• void <a href="mailto:launch\_ac\_gpu">launch\_ac\_gpu</a> (int device, int element, double \*\*data, int length, int dimension, double target\_corr, int num segments)

Launch the GPU kernel, formatted for the thread pool.

• void allocate gpu plan (GPUplan \*plan, int data length, int dimension, int num segments)

Allocates memory for autocorrelation-GPU structure.

• void deallocate\_gpu\_plan (GPUplan \*plan, int data\_length, int dimension, int num\_segments)

Deallocates memory for the autocorrelation-GPU structure.

Copy data to device before starting kernels.

## **Variables**

• GPUplan \* plans\_global

#### 7.19.1 Function Documentation

## 7.19.1.1 ac\_gpu\_wrapper()

Wrapper function for the thread pool.

## **Parameters**

thread	Host thread
job⊷	Job ID
_id	

## 7.19.1.2 allocate\_gpu\_plan()

Allocates memory for autocorrelation-GPU structure.

#### **Parameters**

plan	Structure for GPU plan
data_length	Length of data
dimension	Dimension of the data
num_segments	Number of segments to calculate the autocorrelation length

## 7.19.1.3 auto\_corr\_from\_data\_accel()

Find autocorrelation of data at different points in the chain length and output to autocorr.

#### **Parameters**

	output	Chain data input
	dimension	Dimension of the data
	N_steps	Number of steps in the data
	num_segments	number of segments to calculate the autocorrelation length
	target_corr	Target correlation ratio
out	autocorr	Autocorrelation lengths for the different segments

#### 7.19.1.4 auto\_corr\_internal()

Internal function to calculate the autocorrelation for a given lag Customized for the thread pool architecture, with extra arguments because of the way the memory is allocated.

#### **Parameters**

	arr	Input array of data
	length	Length of input array
	lag	Lag to be used to calculate the correlation
	average	Average of the array arr
out	corr	output correlation
	start_id	ID of location to start calculation – input arrary arr is assumed to be contiguous for multiple
		dimensions

## 7.19.1.5 auto\_corr\_internal\_kernal()

Internal function to launch the CUDA kernel for a range of autocorrelations.

Correlation function used:

```
\label{eq:rho(lag) = 1 / (length - lag) \sum (arr[i+lag]-average) (arr[i]-average)} \\
```

```
target\_corr = rho(rho\_index)/rho(0) = rho(rho\_index)/var
```

#### **Parameters**

	arr	Input array of data
	length	Length of data array
	average	Average of input data
out	rho_index	Index of the lag that results ina correlation ratio target_corr
	target_corr Target correlation ratio rho(lag)/rho(0) = target_corr	
	var	Variance rho(0)
	start_id	Starting index to use for the data array arr

## 7.19.1.6 copy\_data\_to\_device()

Copy data to device before starting kernels.

## **Parameters**

plan	GPU plan
input_data	Input chain data
data_length	Length of data
dimension	Dimension of the data
num_segments	Number of segments to calculate the autocorrelation length

## 7.19.1.7 deallocate\_gpu\_plan()

Deallocates memory for the autocorrelation-GPU structure.

### **Parameters**

plan	Structure for the GPU plan
data_length	Length of data
dimension	Dimension of the data
num_segments	Number of segments to calculate the autocorrelation length

## 7.19.1.8 write\_file\_auto\_corr\_from\_data\_accel()

```
void write_file_auto_corr_from_data_accel (
    std::string acfile,
    double ** chains,
    int dimension,
    int N_steps,
```

```
int num_segments,
double target_corr )
```

Write data file given output chains, as formatted by the mcmc\_sampler.

#### **Parameters**

acfile	Output autocorrelation filename	
chains	Chain data from MCMC_sampler	
dimension	Dimension of the data	
N_steps	Number of steps in the chain	
num_segments	Number of segments to check the autocorrelation length for each dimension	
target_corr	Target correlation ratio to use for the correlation length calculation	

## 7.19.1.9 write\_file\_auto\_corr\_from\_data\_file\_accel()

```
void write_file_auto_corr_from_data_file_accel (
    std::string acfile,
    std::string chains_file,
    int dimension,
    int N_steps,
    int num_segments,
    double target_corr )
```

Write data file for autocorrelation lengths of the data given a data file name, as written by the mcmc\_sampler.

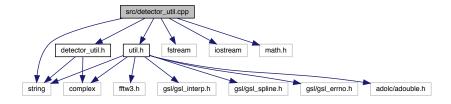
## **Parameters**

acfile	Filename of the autocorrelation data
chains_file	Filename of the data file for the chains
dimension	Dimension of the data
N_steps	Number of steps in the chain
num_segments	Number of segments to check the autocorrelation length for each dimension
target_corr	Target correlation ratio to use for the correlation length calculation

# 7.20 src/detector\_util.cpp File Reference

```
#include "detector_util.h"
#include "util.h"
#include <fstream>
#include <iostream>
#include <string>
#include <math.h>
```

Include dependency graph for detector\_util.cpp:



#### **Functions**

• void populate\_noise (double \*frequencies, std::string detector, double \*noise\_root, int length)

Function to populate the squareroot of the noise curve for various detectors.

• double aLIGO\_analytic (double f)

Analytic function approximating the PSD for aLIGO.

• double Hanford\_O1\_fitted (double f)

Numerically fit PSD to the Hanford Detector's O1.

std::complex< double > Q (double theta, double phi, double iota)

Utility for the overall amplitude and phase shift for spin-aligned systems.

• double right\_interferometer\_plus (double theta, double phi)

Response function of a 90 deg interferometer for plus polarization.

double right\_interferometer\_cross (double theta, double phi)

Response function of a 90 deg interferometer for cross polarization.

void celestial\_horizon\_transform (double RA, double DEC, double gps\_time, std::string detector, double \*phi, double \*theta)

Transform from celestial coordinates to local horizontal coords.

• void derivative\_celestial\_horizon\_transform (double RA, double DEC, double gps\_time, std::string detector, double \*dphi\_dRA, double \*dtheta\_dRA, double \*dphi\_dDEC, double \*dtheta\_dDEC)

Numerical derivative of the transformation.

double DTOA (double theta1, double theta2, std::string detector1, std::string detector2)
 calculate difference in time of arrival (DTOA) for a given source location and 2 different detectors

• double radius\_at\_lat (double latitude, double elevation)

## 7.20.1 Detailed Description

Routines to construct noise curves for various detectors and for detector specific utilities for response functions and coordinate transformations

#### 7.20.2 Function Documentation

#### 7.20.2.1 aLIGO\_analytic()

```
double aLIGO_analytic ( \label{eq:double_f} \mbox{double } f \mbox{ )}
```

Analytic function approximating the PSD for aLIGO.

CITE (Will?)

## 7.20.2.2 celestial\_horizon\_transform()

Transform from celestial coordinates to local horizontal coords.

```
(RA,DEC) -> (altitude, azimuth)
```

Need gps\_time of transformation, as the horizontal coords change in time

detector is used to specify the lat and long of the local frame

## Parameters

RA	in RAD
DEC	in RAD
phi	in RAD
theta	in RAD

## 7.20.2.3 derivative\_celestial\_horizon\_transform()

Numerical derivative of the transformation.

Planned for use in Fisher calculations, but not currently implemented anywhere

#### **Parameters**

RA	in RAD
DEC	in RAD

## 7.20.2.4 DTOA()

calculate difference in time of arrival (DTOA) for a given source location and 2 different detectors

#### **Parameters**

theta1	spherical polar angle for detector 1 in RAD
theta2	spherical polar angle for detector 2 in RAD
detector1	name of detector one
detector2	name of detector two

## 7.20.2.5 Hanford\_O1\_fitted()

```
double Hanford_O1_fitted ( double f )
```

Numerically fit PSD to the Hanford Detector's O1.

CITE (Yunes?)

## 7.20.2.6 populate\_noise()

Function to populate the squareroot of the noise curve for various detectors.

If frequencies are left as NULL, standard frequency spacing is applied and the frequencies are returned, in which case the frequencies argument becomes an output array

Detector names must be spelled exactly

Detectors include: aLIGO\_analytic, Hanford\_O1\_fitted

#### **Parameters**

frequencies	double array of frquencies (NULL)	
detector	String to designate the detector noise curve to be used	
noise_root	noise_root ouptput double array for the square root of the PSD of the noise of the specified detector	
length	integer length of the output and input arrays	

#### 7.20.2.7 Q()

Utility for the overall amplitude and phase shift for spin-aligned systems.

For spin aligned, all the extrinsic parameters have the effect of an overall amplitude modulation and phase shift

## 7.20.2.8 radius\_at\_lat()

/brief Analytic approximation of the radius from the center of earth to a given location

Just the raidus as a function of angles, modelling an oblate spheroid

#### **Parameters**

latitude	latitude in degrees
elevation	elevation in meters

### 7.20.2.9 right\_interferometer\_cross()

Response function of a 90 deg interferometer for cross polarization.

Theta and phi are local, horizontal coordinates relative to the detector

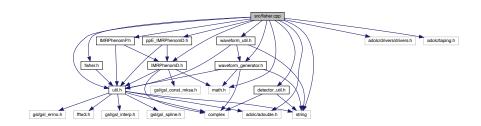
#### 7.20.2.10 right\_interferometer\_plus()

Response function of a 90 deg interferometer for plus polarization.

Theta and phi are local, horizontal coordinates relative to the detector

## 7.21 src/fisher.cpp File Reference

```
#include <fisher.h>
#include <adolc/adouble.h>
#include <adolc/drivers/drivers.h>
#include <adolc/taping.h>
#include <math.h>
#include <string>
#include "util.h"
#include "detector_util.h"
#include "IMRPhenomD.h"
#include "IMRPhenomP.h"
#include "yppE_IMRPhenomD.h"
#include "waveform_generator.h"
#include "waveform_util.h"
Include dependency graph for fisher.cpp:
```



## **Functions**

• void fisher (double \*frequency, int length, string generation\_method, string detector, double \*\*output, int dimension, gen\_params \*parameters, int \*amp\_tapes, int \*phase\_tapes, double \*noise)

Calculates the fisher matrix for the given arguments.

• void calculate\_derivatives (double \*\*amplitude\_deriv, double \*\*phase\_deriv, double \*amplitude, double \*frequencies, int length, string detector, string gen\_method, gen\_params \*parameters)

Abstraction layer for handling the case separation for the different waveforms.

• void fisher\_autodiff (double \*frequency, int length, string generation\_method, string detector, double \*\*output, int dimension, gen\_params \*parameters, int \*amp\_tapes, int \*phase\_tapes, double \*noise)

Calculates the fisher matrix for the given arguments to within numerical error using automatic differention - slower than the numerical version.

## 7.21.1 Detailed Description

All subroutines associated with waveform differentiation and Fisher analysis

## 7.21.2 Function Documentation

## 7.21.2.1 calculate\_derivatives()

Abstraction layer for handling the case separation for the different waveforms.

#### 7.21.2.2 fisher()

Calculates the fisher matrix for the given arguments.

## **Parameters**

length	if 0, standard frequency range for the detector is used
output	double [dimension][dimension]
amp_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method
phase_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method

## 7.21.2.3 fisher\_autodiff()

Calculates the fisher matrix for the given arguments to within numerical error using automatic differention - slower than the numerical version.

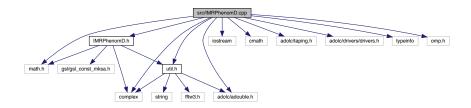
#### **Parameters**

length	if 0, standard frequency range for the detector is used
output	double [dimension][dimension]
amp_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method
phase_tapes	if speed is required, precomputed tapes can be used - assumed the user knows what they're doing, no checks done here to make sure that the number of tapes matches the requirement by the generation_method

# 7.22 src/IMRPhenomD.cpp File Reference

```
#include "IMRPhenomD.h"
#include "util.h"
#include <math.h>
#include <iostream>
#include <complex>
#include <cmath>
#include <adolc/adouble.h>
#include <adolc/taping.h>
#include <adolc/drivers/drivers.h>
#include <typeinfo>
#include <omp.h>
```

Include dependency graph for IMRPhenomD.cpp:



#### **Macros**

• #define omp ignore

## **Variables**

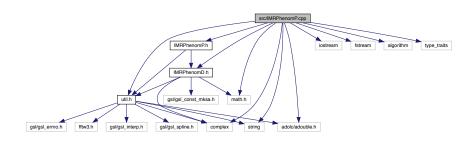
• double log\_64 = 4.15888308336

## 7.22.1 Detailed Description

File that includes all the low level functions that go into constructing the waveform

## 7.23 src/IMRPhenomP.cpp File Reference

```
#include "IMRPhenomP.h"
#include <iostream>
#include <fstream>
#include <string>
#include <complex>
#include "IMRPhenomD.h"
#include "util.h"
#include <adolc/adouble.h>
#include <math.h>
#include <algorithm>
#include <type_traits>
Include dependency graph for IMRPhenomP.cpp:
```



## **Macros**

- #define ROTATEZ(angle, vx, vy, vz)
- #define **ROTATEY**(angle, vx, vy, vz)

#### **Variables**

• const double **sqrt\_6** = 2.44948974278317788

## 7.23.1 Detailed Description

Source code for IMRPhenomP

## 7.23.2 Macro Definition Documentation

#### 7.23.2.1 ROTATEY

#### Value:

```
tmp1 = vx*cos(angle) + vz*sin(angle);\
tmp2 = - vx*sin(angle) + vz*cos(angle);\
vx = tmp1;\
vz = tmp2
```

## 7.23.2.2 ROTATEZ

## Value:

```
tmp1 = vx*cos(angle) - vy*sin(angle);\
tmp2 = vx*sin(angle) + vy*cos(angle);\
vx = tmp1;\
vy = tmp2
```

## 7.24 src/mcmc\_gw.cpp File Reference

```
#include "mcmc_gw.h"
#include "waveform_generator.h"
#include "util.h"
#include "detector_util.h"
#include "waveform_util.h"
#include "fisher.h"
#include "mcmc_sampler.h"
#include <iostream>
#include <vector>
#include <complex>
#include <fftw3.h>
#include <algorithm>
#include <gsl/gsl_interp.h>
#include <gsl/gsl_spline.h>
#include <gsl/gsl_errno.h>
Include dependency graph for mcmc_gw.cpp:
```



#### **Functions**

double maximized\_coal\_log\_likelihood\_IMRPhenomD (double \*frequencies, int length, std::complex< double > \*data, double \*noise, double SNR, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, bool NSflag, fftw\_outline \*plan)

Function to calculate the log Likelihood as defined by -1/2 (d-h|d-h) maximized over the extrinsic parameters phic and tc.

- double maximized\_coal\_log\_likelihood\_IMRPhenomD (double \*frequencies, size\_t length, double \*real\_
   data, double \*imag\_data, double \*noise, double SNR, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, bool NSflag)
- double maximized\_coal\_log\_likelihood\_IMRPhenomD (double \*frequencies, size\_t length, double \*real\_
   data, double \*imag\_data, double \*noise, double SNR, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, bool NSflag, fftw\_outline \*plan)
- double maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param (double \*frequencies, int length, std
   ::complex < double > \*data, double \*noise, double chirpmass, double symmetric\_mass\_ratio, double spin1,
   double spin2, double Luminosity\_Distance, double theta, double phi, double iota, bool NSflag, fftw\_outline
   \*plan)
- double maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param (double \*frequencies, size\_t length, double \*real\_data, double \*imag\_data, double \*noise, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, double Luminosity\_Distance, double theta, double phi, double iota, bool NSflag)
- double maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param (double \*frequencies, size\_t length, double \*real\_data, double \*imag\_data, double \*noise, double chirpmass, double symmetric\_mass\_ratio, double spin1, double spin2, double Luminosity\_Distance, double theta, double phi, double iota, bool NSflag, fftw\_outline \*plan)
- double maximized\_Log\_Likelihood (std::complex < double > \*data, double \*psd, double \*frequencies, size ←
   \_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw\_outline \*plan)

routine to maximize over all extrinsic quantities and return the log likelihood

double maximized\_Log\_Likelihood (double \*data\_real, double \*data\_imag, double \*psd, double \*frequencies, size\_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw outline \*plan)

double maximized\_coal\_Log\_Likelihood (std::complex < double > \*data, double \*psd, double \*frequencies, size\_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw\_outline \*plan, double \*tc, double \*phic)

Function to maximize only over coalescence variables to and phic, returns the maximum values used.

- double maximized\_coal\_Log\_Likelihood\_internal (std::complex< double > \*data, double \*psd, double \*frequencies, std::complex< double > \*detector\_response, size\_t length, fftw\_outline \*plan, double \*tc, double \*phic)
- double Log\_Likelihood (std::complex< double > \*data, double \*psd, double \*frequencies, size\_t length, gen\_params \*params, std::string detector, std::string generation\_method, fftw\_outline \*plan)

Unmarginalized log of the likelihood.

double maximized\_Log\_Likelihood\_aligned\_spin\_internal (std::complex < double > \*data, double \*psd, double \*frequencies, std::complex < double > \*detector\_response, size\_t length, fftw\_outline \*plan)

Maximized match over coalescence variables - returns log likelihood NOT NORMALIZED for aligned spins.

double maximized\_Log\_Likelihood\_unaligned\_spin\_internal (std::complex< double > \*data, double \*psd, double \*frequencies, std::complex< double > \*hplus, std::complex< double > \*hcross, size\_t length, fftw\_outline \*plan)

log likelihood function that maximizes over extrinsic parameters tc, phic, D, and phiRef, the reference frequency - for unaligned spins

double Log\_Likelihood\_internal (std::complex< double > \*data, double \*psd, double \*frequencies, std
 ::complex< double > \*detector\_response, int length, fftw\_outline \*plan)

Internal function for the unmarginalized log of the likelihood.

void MCMC\_MH\_GW (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension, int chain\_id), int numThreads, bool pool, bool show\_prog, int num\_detectors, std::complex< double > \*\*data, double \*\*noise\_psd, double \*\*frequencies, int \*data\_length, double gps\_time, std::string \*detectors, int Nmod, int \*bppe, std::string generation\_method, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string checkpoint\_file)

Wrapper for the MCMC\_MH function, specifically for GW analysis.

void continue\_MCMC\_MH\_GW (std::string start\_checkpoint\_file, double \*\*\*output, int dimension, int N\_← steps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension, int chain\_id), int numThreads, bool pool, bool show\_prog, int num\_detectors, std::complex< double > \*\*data, double \*\*noise\_psd, double \*\*frequencies, int \*data\_length, double gps\_time, std::string \*detectors, int Nmod, int \*bppe, std::string generation\_method, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string final\_checkpoint\_filename)

Takes in an MCMC checkpoint file and continues the chain.

void MCMC\_method\_specific\_prep (std::string generation\_method, int dimension, double \*seeding\_var, bool local\_seeding)

Unpacks MCMC parameters for method specific initiation.

void MCMC\_fisher\_wrapper (double \*param, int dimension, double \*\*output, int chain\_id)

Fisher function for MCMC for GW.

• double MCMC likelihood wrapper (double \*param, int dimension, int chain id)

log likelihood function for MCMC for GW

#### 7.24.1 Detailed Description

Routines for implementation in MCMC algorithms specific to GW CBC analysis

#### 7.24.2 Function Documentation

#### 7.24.2.1 continue\_MCMC\_MH\_GW()

```
void continue_MCMC_MH_GW (
             std::string start_checkpoint_file,
             double *** output,
              int dimension,
              int N_steps,
              int swp_freq,
              double(*)(double *param, int dimension, int chain_id) log_prior,
              int numThreads.
              bool pool,
              bool show_prog,
              int num_detectors,
              std::complex< double > ** data,
              double ** noise_psd,
              double ** frequencies,
              int * data_length,
              double gps_time,
              std::string * detectors,
              int Nmod,
              int * bppe,
              \verb|std::string| \textit{generation}_{\texttt{method}},
              std::string statistics_filename,
              std::string chain_filename,
              std::string auto_corr_filename,
              std::string final_checkpoint_filename )
```

Takes in an MCMC checkpoint file and continues the chain.

Obviously, the user must be sure to correctly match the dimension, number of chains, the generation\_method, the prior function, the data, psds, freqs, and the detectors (number and name), and the gps\_time to the previous run, otherwise the behavior of the sampler is undefined.

numThreads and pool do not necessarily have to be the same

#### 7.24.2.2 Log\_Likelihood()

Unmarginalized log of the likelihood.

#### 7.24.2.3 Log\_Likelihood\_internal()

```
double Log_Likelihood_internal (
          std::complex< double > * data,
          double * psd,
          double * frequencies,
          std::complex< double > * detector_response,
          int length,
          fftw_outline * plan )
```

Internal function for the unmarginalized log of the likelihood.

```
.5 * ((h | h) - 2(D | h))
```

#### 7.24.2.4 maximized\_coal\_Log\_Likelihood()

```
double maximized_coal_Log_Likelihood (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    size_t length,
    gen_params * params,
    std::string detector,
    std::string generation_method,
    fftw_outline * plan,
    double * tc,
    double * phic )
```

Function to maximize only over coalescence variables to and phic, returns the maximum values used.

## 7.24.2.5 maximized\_coal\_log\_likelihood\_IMRPhenomD() [1/3]

Function to calculate the log Likelihood as defined by -1/2 (d-h|d-h) maximized over the extrinsic parameters phic and tc.

frequency array must be uniform spacing - this shouldn't be a problem when working with real data as DFT return uniform spacing

## **Parameters**

```
chirpmass in solar masses
```

## 7.24.2.6 maximized\_coal\_log\_likelihood\_IMRPhenomD() [2/3]

#### **Parameters**

chirpmass	in solar masses

## **7.24.2.7** maximized\_coal\_log\_likelihood\_IMRPhenomD() [3/3]

#### **Parameters**

```
chirpmass in solar masses
```

#### 7.24.2.8 maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param() [1/3]

```
{\tt double\ maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param\ (}
```

```
double * frequencies,
int length,
std::complex< double > * data,
double * noise,
double chirpmass,
double symmetric_mass_ratio,
double spin1,
double spin2,
double Luminosity_Distance,
double theta,
double phi,
double iota,
bool NSflag,
fftw_outline * plan )
```

#### **Parameters**

chirpmass in solar masses

## 7.24.2.9 maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param() [2/3]

## Parameters

chirpmass in solar masses

## 7.24.2.10 maximized\_coal\_log\_likelihood\_IMRPhenomD\_Full\_Param() [3/3]

```
double chirpmass,
double symmetric_mass_ratio,
double spin1,
double spin2,
double Luminosity_Distance,
double theta,
double phi,
double iota,
bool NSflag,
fftw_outline * plan )
```

#### **Parameters**

chirpmass	in solar masses

## 7.24.2.11 maximized\_Log\_Likelihood()

```
double maximized_Log_Likelihood (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    size_t length,
    gen_params * params,
    std::string detector,
    std::string generation_method,
    fftw_outline * plan )
```

routine to maximize over all extrinsic quantities and return the log likelihood

IMRPhenomD – maximizes over DL, phic, tc, \iota, \phi, \theta IMRPhenomP – maximizes over DL, phic,tc, \psi, \phi , \theta

### 7.24.2.12 maximized\_Log\_Likelihood\_aligned\_spin\_internal()

```
double maximized_Log_Likelihood_aligned_spin_internal (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    std::complex< double > * detector_response,
    size_t length,
    fftw_outline * plan )
```

Maximized match over coalescence variables - returns log likelihood NOT NORMALIZED for aligned spins.

Note: this function is not properly normalized for an absolute comparison. This is made for MCMC sampling, so to minimize time, constant terms like (Data|Data), which would cancel in the Metropolis-Hasting ratio, are left out for efficiency

#### 7.24.2.13 maximized\_Log\_Likelihood\_unaligned\_spin\_internal()

```
double maximized_Log_Likelihood_unaligned_spin_internal (
    std::complex< double > * data,
    double * psd,
    double * frequencies,
    std::complex< double > * hplus,
    std::complex< double > * hcross,
    size_t length,
    fftw_outline * plan )
```

log likelihood function that maximizes over extrinsic parameters tc, phic, D, and phiRef, the reference frequency - for unaligned spins

Ref: arXiv 1603.02444v2

## 7.24.2.14 MCMC\_fisher\_wrapper()

Fisher function for MCMC for GW.

Wraps the fisher calculation in src/fisher.cpp and unpacks parameters correctly for common GW analysis

Supports all the method/parameter combinations found in MCMC\_MH\_GW

## 7.24.2.15 MCMC\_likelihood\_wrapper()

log likelihood function for MCMC for GW

Wraps the above likelihood functions and unpacks parameters correctly for common GW analysis

Supports all the method/parameter combinations found in MCMC\_MH\_GW

### 7.24.2.16 MCMC\_method\_specific\_prep()

```
void MCMC_method_specific_prep (
    std::string generation_method,
    int dimension,
    double * seeding_var,
    bool local_seeding )
```

Unpacks MCMC parameters for method specific initiation.

Populates seeding vector if non supplied, populates mcmc\_Nmod, populates mcmc\_log\_beta, populates mcmc\_← intrinsic

#### 7.24.2.17 MCMC\_MH\_GW()

```
void MCMC_MH_GW (
             double *** output,
             int dimension,
             int N_steps,
             int chain N.
             double * initial_pos,
             double * seeding_var,
             double * chain_temps,
             int swp_freq,
             double(*)(double *param, int dimension, int chain_id) log_prior,
             int numThreads.
             bool pool,
             bool show_prog,
             int num_detectors,
             std::complex< double > ** data,
             double ** noise_psd,
             double ** frequencies,
             int * data_length,
             double gps_time,
             std::string * detectors,
             int Nmod,
             int * bppe,
             std::string generation_method,
             std::string statistics filename.
             std::string chain_filename,
             std::string auto_corr_filename,
             std::string checkpoint_file )
```

Wrapper for the MCMC\_MH function, specifically for GW analysis.

Handles the details of setting up the MCMC sampler and wraps the fisher and log likelihood to conform to the format of the sampler

*NOTE* – This sampler is NOT thread safe. There is global memory declared for each call to MCMC\_MH\_GW, so separate samplers should not be run in the same process space

Supported parameter combinations:

```
IMRPhenomD - 4 dimensions - In chirpmass, eta, chi1, chi2
```

```
IMRPhenomD - 7 dimensions - In D_L, tc, phic, In chirpmass, eta, chi1, chi2
```

IMRPhenomD - 8 dimensions - cos inclination, RA, DEC, In D\_L, In chirpmass, eta, chi1, chi2

dCS\_IMRPhenomD\_log - 8 dimensions – cos inclination, RA, DEC, In D\_L, In chirpmass, eta, chi1, chi2, In  $\alpha^2$  (the coupling parameter)

dCS\_IMRPhenomD- 8 dimensions – cos inclination, RA, DEC, In D\_L, In chirpmass, eta, chi1, chi2,  $\alpha^2$  (the coupling parameter)

dCS\_IMRPhenomD\_root\_alpha- 8 dimensions – cos inclination, RA, DEC, In D\_L, In chirpmass, eta, chi1, chi2, \sqrt \alpha (in km) (the coupling parameter)

```
IMRPhenomPv2 - 9 dimensions - cos J_N, In chirpmass, eta, |chi1|, |chi1|, theta_1, theta_2, phi_1, phi_2
```

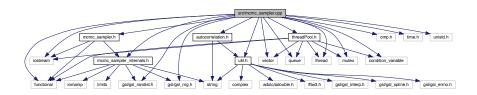
#### **Parameters**

statistics_filename	Filename to output sampling statistics, if empty string, not output
chain_filename	Filename to output data (chain 0 only), if empty string, not output
auto_corr_filename Filename to output auto correlation in some interval, if empty string, not output	
checkpoint_file	Filename to output data for checkpoint, if empty string, not saved

# 7.25 src/mcmc\_sampler.cpp File Reference

```
#include "mcmc_sampler.h"
#include "autocorrelation.h"
#include "util.h"
#include "mcmc_sampler_internals.h"
#include "threadPool.h"
#include <iostream>
#include <gsl/gsl_rng.h>
#include <gsl/gsl_randist.h>
#include <omp.h>
#include <time.h>
#include <condition_variable>
#include <mutex>
#include <thread>
#include <vector>
#include <queue>
#include <functional>
#include <unistd.h>
```

Include dependency graph for mcmc\_sampler.cpp:



## Classes

class Comparator

Class to facilitate the comparing of chains for priority.

class ThreadPool

## **Macros**

• #define omp ignore

#### **Functions**

• void MCMC\_MH\_internal (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, std::function< double(double \*, int, int)> log\_c prior, std::function< double(double \*, int, int)> log\_likelihood, std::function< void(double \*, int, double \*\*, int)>fisher, int numThreads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_c filename, std::string auto\_corr\_filename, std::string checkpoint\_file)

Generic sampler, where the likelihood, prior are parameters supplied by the user.

void continue\_MCMC\_MH\_internal (std::string start\_checkpoint\_file, double \*\*\*output, int N\_steps, int swp
 \_freq, std::function< double(double \*, int, int)> log\_prior, std::function< double(double \*, int, int)> log\_
 likelihood, std::function< void(double \*, int, double \*\*, int)>fisher, int numThreads, bool pool, bool show
 \_prog, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string end\_checkpoint\_file)

Routine to take a checkpoint file and begin a new chain at said checkpoint.

void MCMC\_MH\_loop (sampler \*sampler)

Internal function that runs the actual loop for the sampler.

- · void mcmc step threaded (int j)
- void mcmc\_swap\_threaded (int i, int j)
- void MCMC\_MH (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension), double(\*log\_likelihood)(double \*param, int dimension), void(\*fisher)(double \*param, int dimension, double \*\*fisher), int numThreads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_
  filename, std::string auto\_corr\_filename, std::string checkpoint\_file)
- void MCMC\_MH (double \*\*\*output, int dimension, int N\_steps, int chain\_N, double \*initial\_pos, double \*seeding\_var, double \*chain\_temps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension, int chain\_id), double(\*log\_likelihood)(double \*param, int dimension, int chain\_id), void(\*fisher)(double \*param, int dimension, double \*\*fisher, int chain\_id), int numThreads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string checkpoint\_file)
- void continue\_MCMC\_MH (std::string start\_checkpoint\_file, double \*\*\*output, int N\_steps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension, int chain\_id), double(\*log\_likelihood)(double \*param, int dimension, int chain\_id), void(\*fisher)(double \*param, int dimension, double \*\*fisher, int chain\_id), int num to Threads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string end\_checkpoint\_file)
- void continue\_MCMC\_MH (std::string start\_checkpoint\_file, double \*\*\*output, int N\_steps, int swp\_freq, double(\*log\_prior)(double \*param, int dimension), double(\*log\_likelihood)(double \*param, int dimension), void(\*fisher)(double \*param, int dimension, double \*\*fisher), int numThreads, bool pool, bool show\_prog, std::string statistics\_filename, std::string chain\_filename, std::string auto\_corr\_filename, std::string end\_checkpoint\_file)

#### **Variables**

- const gsl\_rng\_type \* T
- gsl\_rng \* **r**
- sampler \* samplerptr
- ThreadPool \* poolptr

## 7.25.1 Detailed Description

Source file for the sampler foundation

Source file for generic MCMC sampler. Sub routines that are application agnostic are housed in mcmc\_sampler ← \_internals

## 7.25.2 Function Documentation

## 7.25.2.1 continue\_MCMC\_MH() [1/2]

#### **Parameters**

	start_checkpoint_file	File for starting checkpoint
out	output	output array, dimensions: output[chain_N][N_steps][dimension]
	N_steps	Number of new steps to take
	swp_freq	frequency of swap attempts between temperatures
	log_prior	Funcion pointer for the log_prior
	log_likelihood	Function pointer for the log_likelihood
	fisher	Function pointer for the fisher - if NULL, fisher steps are not used
	numThreads	Number of threads to use
	pool	Boolean for whether to use deterministic'' vsstochastic" sampling
	show_prog	Boolean for whether to show progress or not (turn off for cluster runs
	statistics_filename	Filename to output sampling statistics, if empty string, not output
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	end_checkpoint_file	Filename to output data for checkpoint at the end of the continued run, if empty string, not saved

## 7.25.2.2 continue\_MCMC\_MH() [2/2]

```
double(*)(double *param, int dimension) log_likelihood,
void(*)(double *param, int dimension, double **fisher) fisher,
int numThreads,
bool pool,
bool show_prog,
std::string statistics_filename,
std::string chain_filename,
std::string auto_corr_filename,
std::string end_checkpoint_file)
```

#### **Parameters**

	start_checkpoint_file	File for starting checkpoint
out	output	output array, dimensions: output[chain_N][N_steps][dimension]
	N_steps	Number of new steps to take
	swp_freq	frequency of swap attempts between temperatures
	log_prior	Funcion pointer for the log_prior
	log_likelihood	Function pointer for the log_likelihood
	fisher	Function pointer for the fisher - if NULL, fisher steps are not used
	numThreads	Number of threads to use
	pool	Boolean for whether to use deterministic'' vsstochastic" sampling
	show_prog	Boolean for whether to show progress or not (turn off for cluster runs
	statistics_filename	Filename to output sampling statistics, if empty string, not output
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	end_checkpoint_file	Filename to output data for checkpoint at the end of the continued run, if empty string, not saved

## 7.25.2.3 continue\_MCMC\_MH\_internal()

```
void continue_MCMC_MH_internal (
    std::string start_checkpoint_file,
    double *** output,
    int N_steps,
    int swp_freq,
    std::function< double(double *, int, int)> log_prior,
    std::function< double(double *, int, int)> log_likelihood,
    std::function< void(double *, int, double **, int)> fisher,
    int numThreads,
    bool pool,
    bool show_prog,
    std::string statistics_filename,
    std::string auto_corr_filename,
    std::string end_checkpoint_file)
```

Routine to take a checkpoint file and begin a new chain at said checkpoint.

See MCMC\_MH\_internal for more details of parameters (pretty much all the same)

#### **Parameters**

	start_checkpoint_file	File for starting checkpoint
out	output	output array, dimensions: output[chain_N][N_steps][dimension]
	N_steps	Number of new steps to take
	swp_freq	frequency of swap attempts between temperatures
	log_prior	std::function for the log_prior function – takes double *position, int dimension, int chain_id
	log_likelihood	std::function for the log_likelihood function – takes double *position, int dimension, int chain_id
	fisher	std::function for the fisher function – takes double *position, int dimension, double **output_fisher, int chain_id
	numThreads	Number of threads to use
	pool	Boolean for whether to use deterministic'' vsstochastic" sampling
	show_prog	Boolean for whether to show progress or not (turn off for cluster runs
	statistics_filename	Filename to output sampling statistics, if empty string, not output
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	end_checkpoint_file	Filename to output data for checkpoint at the end of the continued run, if empty string, not saved

## **7.25.2.4** MCMC\_MH() [1/2]

```
void MCMC_MH (
            double *** output,
             int dimension,
             int N_steps,
             int chain_N,
             double * initial_pos,
             double * seeding_var,
             double * chain_temps,
             int swp_freq,
             double(*)(double *param, int dimension) log_prior,
             double(*)(double *param, int dimension) log_likelihood,
             void(*)(double *param, int dimension, double **fisher) fisher,
             int numThreads,
             bool pool,
             bool show_prog,
             std::string statistics_filename,
             std::string chain_filename,
             std::string auto_corr_filename,
             std::string checkpoint_file )
```

## **Parameters**

out	output	Output chains, shape is double[chain_N, N_steps,dimension]	
	dimension	dimension of the parameter space being explored	
	N_steps	Number of total steps to be taken, per chain	
	chain_N	Number of chains	
	initial_pos	Initial position in parameter space - shape double[dimension]	

#### **Parameters**

seeding_var	Variance of the normal distribution used to seed each chain higher than 0 - shape double[dimension]
chain_temps	Double array of temperatures for the chains
swp_freq	the frequency with which chains are swapped
log_prior	Funcion pointer for the log_prior
log_likelihood	Function pointer for the log_likelihood
fisher	Function pointer for the fisher - if NULL, fisher steps are not used
numThreads	Number of threads to use (=1 is single threaded)
pool	boolean to use stochastic chain swapping (MUST have >2 threads)
show_prog	boolean whether to print out progress (for example, should be set to `'false'' if submitting to a cluster)
statistics_filename	Filename to output sampling statistics, if empty string, not output
chain_filename	Filename to output data (chain 0 only), if empty string, not output
auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
checkpoint_file	Filename to output data for checkpoint, if empty string, not saved

## **7.25.2.5** MCMC\_MH() [2/2]

```
void MCMC_MH (
             double *** output,
            int dimension,
            int N_steps,
             int chain_N,
             double * initial_pos,
             double * seeding_var,
             double * chain_temps,
             int swp_freq,
             double(*)(double *param, int dimension, int chain_id) log_prior,
             double(*)(double *param, int dimension, int chain_id) log_likelihood,
             void(*)(double *param, int dimension, double **fisher, int chain_id) fisher,
             int numThreads,
             bool pool,
             bool show_prog,
             std::string statistics_filename,
             std::string chain_filename,
             std::string auto_corr_filename,
             std::string checkpoint_file )
```

## **Parameters**

out	output	Output chains, shape is double[chain_N, N_steps,dimension]
	dimension	dimension of the parameter space being explored
	N_steps	Number of total steps to be taken, per chain
	chain_N	Number of chains
	initial_pos	Initial position in parameter space - shape double[dimension]
	seeding_var	Variance of the normal distribution used to seed each chain higher than 0 - shape double[dimension]
	chain_temps	Double array of temperatures for the chains

#### **Parameters**

swp_freq	the frequency with which chains are swapped
log_prior	Funcion pointer for the log_prior
log_likelihood	Function pointer for the log_likelihood
fisher	Function pointer for the fisher - if NULL, fisher steps are not used
numThreads	Number of threads to use (=1 is single threaded)
pool	boolean to use stochastic chain swapping (MUST have >2 threads)
show_prog	boolean whether to print out progress (for example, should be set to `'false'' if submitting to a cluster)
statistics_filename	Filename to output sampling statistics, if empty string, not output
chain_filename	Filename to output data (chain 0 only), if empty string, not output
auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
checkpoint_file	Filename to output data for checkpoint, if empty string, not saved

### 7.25.2.6 MCMC\_MH\_internal()

```
void MCMC_MH_internal (
             double *** output,
             int dimension,
             int N_steps,
             int chain_N,
             double * initial_pos,
             double * seeding_var,
             double * chain_temps,
             int swp_freq,
             std::function< double(double *, int, int) > log_prior,
             std::function< double(double *, int, int) > log_likelihood,
             std::function< void(double *, int, double **, int) > fisher,
             int numThreads,
             bool pool,
             bool show_prog,
             std::string statistics_filename,
             std::string chain_filename,
             std::string auto_corr_filename,
             std::string checkpoint_file )
```

Generic sampler, where the likelihood, prior are parameters supplied by the user.

Base of the sampler, generic, with user supplied quantities for most of the samplers properties

Uses the Metropolis-Hastings method, with the option for Fisher/MALA steps if the Fisher routine is supplied.

3 modes to use -

single threaded (numThreads = 1) runs single threaded

multi-threaded `'deterministic'' (numThreads>1 ; pool = false) progresses each chain in parallel for swp\_freq steps, then waits for all threads to complete before swapping temperatures in sequenctial order (j, j+1) then (j+1, j+2) etc (sequenctially)

multi-threaded `'stochastic' (numThreads>2; pool = true) progresses each chain in parallel by queueing each temperature and evaluating them in the order they were submitted. Once finished, the threads are queued to swap, where they swapped in the order they are submitted. This means the chains are swapped randomly, and the chains do NOT finish at the same time. The sampler runs until the the 0th chain reaches the step number

Note on limits: In the prior function, if a set of parameters should be disallowed, return -std::numeric $\_\leftarrow$  limits<double>::infinity() - (this is in the limits> file in std)

Format for the auto\_corr file (compatable with csv, dat, txt extensions): each row is a dimension of the cold chain, with the first row being the lengths used for the auto-corr calculation:

lengths: length1, length2...

dim1: length1, length2...

Format for the chain file (compatable with csv, dat, txt extensions): each row is a step, each column a dimension:

Step1: dim1, dim2, ...

Step2: dim1, dim2, ...

Statistics\_filename : should be txt extension

checkpoint\_file: This file saves the final position of all the chains, as well as other metadata, and can be loaded by the function <FUNCTION> to continue the chain from the point it left off. Not meant to be read by humans, the data order is custom to this software library. An empty string ("") means no checkpoint will be saved. For developers, the contents are:

dimension, # of chains

temps of chains

Stepping widths of all chains

Final position of all chains

## Parameters

arameter	5	
out	output	Output chains, shape is double[chain_N, N_steps,dimension]
	dimension	dimension of the parameter space being explored
	N_steps	Number of total steps to be taken, per chain
	chain_N	Number of chains
	initial_pos	Initial position in parameter space - shape double[dimension]
	seeding_var	Variance of the normal distribution used to seed each chain higher than 0 - shape double[dimension]
	chain_temps	Double array of temperatures for the chains
	swp_freq	the frequency with which chains are swapped
	log_prior	std::function for the log_prior function – takes double *position, int dimension, int chain_id
	log_likelihood	std::function for the log_likelihood function – takes double *position, int dimension, int chain_id
	fisher	std::function for the fisher function – takes double *position, int dimension, double **output_fisher, int chain_id
	numThreads	Number of threads to use (=1 is single threaded)
	pool	boolean to use stochastic chain swapping (MUST have >2 threads)
	show_prog	boolean whether to print out progress (for example, should be set to `'false'' if submitting to a cluster)
enerated by	y <b>sytatisti</b> cs_filename	Filename to output sampling statistics, if empty string, not output
	chain_filename	Filename to output data (chain 0 only), if empty string, not output
	auto_corr_filename	Filename to output auto correlation in some interval, if empty string, not output
	checkpoint file	Filename to output data for checkpoint, if empty string, not saved

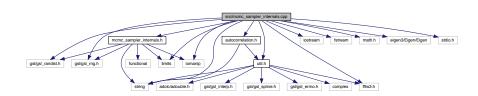
#### 7.25.2.7 MCMC\_MH\_loop()

Internal function that runs the actual loop for the sampler.

## 7.26 src/mcmc\_sampler\_internals.cpp File Reference

```
#include "mcmc_sampler_internals.h"
#include "autocorrelation.h"
#include "util.h"
#include <iostream>
#include <fstream>
#include <string>
#include <math.h>
#include <gsl/gsl_randist.h>
#include <gsl/gsl_rng.h>
#include <limits>
#include <iomanip>
#include <fftw3.h>
#include <stdio.h>
```

Include dependency graph for mcmc\_sampler\_internals.cpp:



## **Functions**

- int mcmc\_step (sampler \*sampler, double \*current\_param, double \*next\_param, int chain\_number) interface function between the sampler and the internal step functions
- void gaussian\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param, int chain\_id) Straight gaussian step.
- void fisher\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param, int chain\_index) Fisher informed gaussian step.
- void update\_fisher (sampler \*sampler, double \*current\_param, int chain\_index)
- void mmala\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param)

  MMALA informed step Currently not supported.
- void diff\_ev\_step (sampler \*sampler, double \*current\_param, double \*proposed\_param, int chain\_id) differential evolution informed step
- void chain\_swap (sampler \*sampler, double \*\*\*output, int step\_num, int \*swp\_accepted, int \*swp\_rejected) subroutine to perform chain comparison for parallel tempering

- int single\_chain\_swap (sampler \*sampler, double \*chain1, double \*chain2, int T1\_index, int T2\_index) subroutine to actually swap two chains
- void assign\_probabilities (sampler \*sampler, int chain\_index)

update and initiate probabilities for each variety of step

- void allocate\_sampler\_mem (sampler \*sampler)
- void deallocate\_sampler\_mem (sampler \*sampler)
- void update\_history (sampler \*sampler, double \*new\_params, int chain\_index)
- void write\_stat\_file (sampler \*sampler, std::string filename)
- void write checkpoint file (sampler \*sampler, std::string filename)

Routine that writes metadata and final positions of a sampler to a checkpoint file.

- void load\_checkpoint\_file (std::string check\_file, sampler \*sampler)
   load checkpoint file into sampler struct
- void assign\_ct\_p (sampler \*sampler, int step, int chain\_index)
- void assign\_ct\_m (sampler \*sampler, int step, int chain\_index)

# 7.26.1 Detailed Description

File containing definitions for all the internal, generic mcmc subroutines

#### 7.26.2 Function Documentation

#### 7.26.2.1 assign\_probabilities()

update and initiate probabilities for each variety of step

Type 0: Gaussian step

Type 1: Differential Evolution step

Type 2: MMALA step (currently not supported)

Type 3: Fisher step

#### 7.26.2.2 chain\_swap()

subroutine to perform chain comparison for parallel tempering

The total output file is passed, and the chains are swapped sequentially

This is the routine for 'Deterministic' sampling (parallel or sequential, but not pooled)

#### **Parameters**

sampler	sampler struct
output	output vector containing chains
step_num	current step number

# 7.26.2.3 diff\_ev\_step()

differential evolution informed step

Differential evolution uses the past history of the chain to inform the proposed step:

Take the difference of two random, accepted previous steps and step along that with some step size, determined by a gaussian

#### **Parameters**

sampler		sampler	Sampler struct
		current_param	current position in parameter space
	out <i>proposed_param</i>		Proposed position in parameter space

# 7.26.2.4 fisher\_step()

Fisher informed gaussian step.

	sampler	Sampler struct
	current_param	current position in parameter space
out	proposed_param	Proposed position in parameter space

## 7.26.2.5 gaussian\_step()

Straight gaussian step.

#### **Parameters**

sampler		sampler	Sampler struct
		current_param	current position in parameter space
out <i>proposed_param</i>		proposed_param	Proposed position in parameter space

# 7.26.2.6 load\_checkpoint\_file()

load checkpoint file into sampler struct

NOTE – allocate\_sampler called in function – MUST deallocate manually

NOTE - sampler->chain\_temps allocated internally - MUST free manually

# 7.26.2.7 mmala\_step()

MMALA informed step - Currently not supported.

# Parameters

	sampler		Sampler struct
		current_param	current position in parameter space
0	ut	proposed_param	Proposed position in parameter space

# 7.26.2.8 single\_chain\_swap()

```
int single_chain_swap (
          sampler * sampler,
```

```
double * chain1,
double * chain2,
int T1_index,
int T2_index )
```

subroutine to actually swap two chains

This is the more general subroutine, which just swaps the two chains passed to the function

#### **Parameters**

sampler	sampler structure
chain1	parameter position of chain that could be changed
chain2	chain that is not swapped, but provides parameters to be swapped by the other chain
T1_index	number of chain swappe in chain_temps
T2_index	number of chain swapper in chain_temps

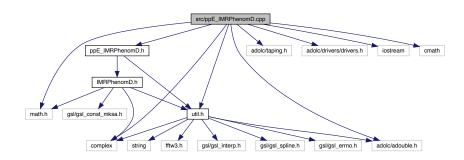
# 7.26.2.9 write\_checkpoint\_file()

Routine that writes metadata and final positions of a sampler to a checkpoint file.

# 7.27 src/ppE\_IMRPhenomD.cpp File Reference

```
#include "ppE_IMRPhenomD.h"
#include <math.h>
#include <adolc/adouble.h>
#include <adolc/taping.h>
#include <adolc/drivers/drivers.h>
#include <iostream>
#include <cmath>
#include <complex>
#include "util.h"
```

Include dependency graph for ppE\_IMRPhenomD.cpp:



# 7.27.1 Detailed Description

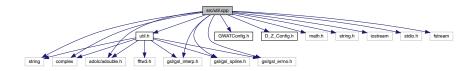
File for the implementation of the ppE formalism for testing GR

Extends the IMRPhenomD template to include non-GR phase terms

Supported waveforms: ppE Inspiral, ppE IMR, dCS, EdGB

# 7.28 src/util.cpp File Reference

```
#include "util.h"
#include "GWATConfig.h"
#include "D_Z_Config.h"
#include <math.h>
#include <string>
#include <complex>
#include <complex>
#include <iostream>
#include <fstream>
#include <adolc/adouble.h>
#include <gsl/gsl_interp.h>
#include <gsl/gsl_spline.h>
#include <gsl/gsl_errno.h>
Include dependency graph for util.cpp:
```



# **Functions**

- void initiate\_LumD\_Z\_interp (gsl\_interp\_accel \*\*Z\_DL\_accel\_ptr, gsl\_spline \*\*Z\_DL\_spline\_ptr)

  Function that uses the GSL libraries to interpolate pre-calculated Z-D\_L data.
- void free\_LumD\_Z\_interp (gsl\_interp\_accel \*\*Z\_DL\_accel\_ptr, gsl\_spline \*\*Z\_DL\_spline\_ptr)

  Frees the allocated interpolation function.
- adouble Z\_from\_DL\_interp (adouble DL, gsl\_interp\_accel \*Z\_DL\_accel\_ptr, gsl\_spline \*Z\_DL\_spline\_ptr)
- double Z\_from\_DL\_interp (double DL, gsl\_interp\_accel \*Z\_DL\_accel\_ptr, gsl\_spline \*Z\_DL\_spline\_ptr)
- double Z\_from\_DL (double DL, std::string cosmology)

Calculates the redshift given the luminosity distance.

adouble Z from DL (adouble DL, std::string cosmology)

Calculates the redshift given the luminosity distance adouble version for ADOL-C implementation.

- double DL\_from\_Z (double Z, std::string cosmology)
  - Calculates the luminosity distance given the redshift.
- adouble DL\_from\_Z (adouble Z, std::string cosmology)

Calculates the luminosity distance given the redshift adouble version for ADOL-C implementation.

• double cosmology\_interpolation\_function (double x, double \*coeffs, int interp\_degree)

Custom interpolation function used in the cosmology calculations.

adouble cosmology\_interpolation\_function (adouble x, double \*coeffs, int interp\_degree)

Custom interpolation function used in the cosmology calculations adouble version for ADOL-C.

double cosmology lookup (std::string cosmology)

Helper function for mapping cosmology name to an internal index.

void printProgress (double percentage)

routine to print the progress of a process to the terminal as a progress bar

void initiate likelihood function (fftw outline \*plan, int length)

Allocate memory for FFTW3 methods used in a lot of inner products input is a locally defined structure that houses all the pertinent data.

void allocate\_FFTW3\_mem\_inverse (fftw\_outline \*plan, int length)

Allocate memory for FFTW3 methods used in a lot of inner products –INVERSE input is a locally defined structure that houses all the pertinent data.

void deactivate likelihood function (fftw outline \*plan)

deallocates the memory used for FFTW routines

double calculate\_chirpmass (double mass1, double mass2)

Calculates the chirp mass from the two component masses.

- adouble calculate chirpmass (adouble mass1, adouble mass2)
- double calculate\_eta (double mass1, double mass2)

Calculates the symmetric mass ration from the two component masses.

- adouble calculate eta (adouble mass1, adouble mass2)
- double calculate mass1 (double chirpmass, double eta)

Calculates the larger mass given a chirp mass and symmetric mass ratio.

- adouble calculate mass1 (adouble chirpmass, adouble eta)
- double calculate\_mass2 (double chirpmass, double eta)

Calculates the smaller mass given a chirp mass and symmetric mass ratio.

- adouble calculate\_mass2 (adouble chirpmass, adouble eta)
- long factorial (long num)

Local function to calculate a factorial.

double pow int (double base, int power)

Local power function, specifically for integer powers.

- adouble pow\_int (adouble base, int power)
- · double cbrt internal (double base)

Fucntion that just returns the cuberoot.

adouble cbrt\_internal (adouble base)

Fucntion that just returns the cuberoot ADOL-C doesn't have the cbrt function (which is faster), so have to use the power function.

double \*\* allocate\_2D\_array (int dim1, int dim2)

Utility to malloc 2D array.

void deallocate\_2D\_array (double \*\*array, int dim1, int dim2)

Utility to free malloc'd 2D array.

double \*\*\* allocate\_3D\_array (int dim1, int dim2, int dim3)

Utility to malloc 3D array.

• void deallocate\_3D\_array (double \*\*\*array, int dim1, int dim2, int dim3)

Utility to free malloc'd 2D array.

void read file (std::string filename, double \*\*output, int rows, int cols)

Utility to read in data.

• void read file (std::string filename, double \*output)

Utility to read in data (single dimension vector)

• void read\_LOSC\_data\_file (std::string filename, double \*output, double \*data\_start\_time, double \*duration, double \*fs)

Read data file from LIGO Open Science Center.

- void read\_LOSC\_PSD\_file (std::string filename, double \*\*output, int rows, int cols)
  - Read PSD file from LIGO Open Science Center.
- void allocate\_LOSC\_data (std::string \*data\_files, std::string psd\_file, int num\_detectors, int psd\_length, int data\_file\_length, double trigger\_time, std::complex< double > \*\*data, double \*\*psds, double \*\*freqs)

Prepare data for MCMC directly from LIGO Open Science Center.

- void free\_LOSC\_data (std::complex< double > \*\*data, double \*\*psds, double \*\*freqs, int num\_detectors, int length)
- void tukey window (double \*window, int length, double alpha)

Tukev window function for FFTs.

void write\_file (std::string filename, double \*\*input, int rows, int cols)

Utility to write 2D array to file.

void write\_file (std::string filename, double \*input, int length)

Utility to write 1D array to file.

void celestial\_horizon\_transform (double RA, double DEC, double gps\_time, double LONG, double LAT, double \*phi, double \*theta)

Utility to transform from celestial coord RA and DEC to local horizon coord for detector response functions.

double gps\_to\_GMST (double gps\_time)

Utility to transform from gps time to GMST https://aa.usno.navy.mil/faq/docs/GAST.php.

double gps\_to\_JD (double gps\_time)

Utility to transform from gps to JD.

void transform\_cart\_sph (double \*cartvec, double \*sphvec)

utility to transform a vector from cartesian to spherical (radian)

void transform\_sph\_cart (double \*sphvec, double \*cartvec)

utility to transform a vector from spherical (radian) to cartesian

• template<class T >

```
std::complex< T > cpolar (T mag, T phase)
```

template<class T >

 $std::complex < T > XLALSpinWeightedSphericalHarmonic \ (T \ theta, \ T \ phi, \ int \ s, \ int \ I, \ int \ m)$ 

- template std::complex< double > XLALSpinWeightedSphericalHarmonic< double > (double, double, int, int, int)
- template std::complex< adouble > XLALSpinWeightedSphericalHarmonic< adouble > (adouble, adouble, int, int, int)
- template std::complex< double > cpolar< double > (double, double)
- template std::complex< adouble > cpolar< adouble > (adouble, adouble)

#### 7.28.1 Detailed Description

General utilities that are not necessarily specific to any part of the project at large

#### 7.28.2 Function Documentation

#### 7.28.2.1 allocate\_2D\_array()

Utility to malloc 2D array.

#### 7.28.2.2 allocate\_3D\_array()

Utility to malloc 3D array.

# 7.28.2.3 allocate\_LOSC\_data()

Prepare data for MCMC directly from LIGO Open Science Center.

Trims data for Tobs (determined by PSD file) 3/4\*Tobs in front of trigger, and 1/4\*Tobs behind

Currently, default to sampling frequency and observation time set by PSD – cannot be customized

Output is in order of PSD columns – string vector of detectos MUST match order of PSD cols

Output shapes—psds = [num\_detectors][psd\_length] data = [num\_detectors][psd\_length]

freqs = [num\_detectors][psd\_length]

Total observation time = 1/( freq[i] - freq[i-1]) (from PSD file)

Sampling frequency fs = max frequency from PSD file

ALLOCATES MEMORY - must be freed to prevent memory leak

data_files		Vector of strings for each detector file from LOSC
	psd_file	String of psd file from LOSC
	num_detectors	Number of detectors to use
	psd_length	Length of the PSD file (number of rows of DATA)
	data_file_length	Length of the data file (number of rows of DATA)
	trigger_time	Time for the signal trigger (GPS)
out	data	Output array of data for each detector
out	psds	Output array of psds for each detector
out	freqs	Output array of freqs for each detector

#### 7.28.2.4 calculate\_chirpmass()

Calculates the chirp mass from the two component masses.

The output units are whatever units the input masses are

#### 7.28.2.5 calculate\_mass1()

Calculates the larger mass given a chirp mass and symmetric mass ratio.

Units of the output match the units of the input chirp mass

# 7.28.2.6 calculate\_mass2()

Calculates the smaller mass given a chirp mass and symmetric mass ratio.

Units of the output match the units of the input chirp mass

# 7.28.2.7 celestial\_horizon\_transform()

Utility to transform from celestial coord RA and DEC to local horizon coord for detector response functions.

Outputs are the spherical polar angles defined by North as 0 degrees azimuth and the normal to the earth as 0 degree polar

	RA	Right acsension (rad)
	DEC	Declination (rad)
Generated b	y Doxygenme	GPS time
	LONG	Longitude (rad)
	LAT	Latitude (rad)
011#	nhi	horizon azimuthal angle (rad)

#### 7.28.2.8 cosmology\_interpolation\_function()

```
double cosmology_interpolation_function ( \label{eq:cosmology} \mbox{double } x, \\ \mbox{double } * coeffs, \\ \mbox{int } interp\_degree \ )
```

Custom interpolation function used in the cosmology calculations.

Power series in half power increments of x, up to 11/2. powers of x

#### 7.28.2.9 deallocate\_2D\_array()

Utility to free malloc'd 2D array.

#### 7.28.2.10 deallocate 3D array()

Utility to free malloc'd 2D array.

# 7.28.2.11 DL\_from\_Z()

Calculates the luminosity distance given the redshift.

Based on Astropy.cosmology calculations – see python script in the ./data folder of the project – numerically calculated given astropy.cosmology's definitions ( http://docs.astropy.org/en/stable/cosmology/) and used scipy.optimize to fit to a power series, stepping in half powers of Z. These coefficients are then output to a header file (D\_Z\_config.h) which are used here to calculate distance. Custom cosmologies etc can easily be acheived by editing the python script D\_Z\_config.py, the c++ functions do not need modification. They use whatever data is available in the header file. If the functional form of the fitting function changes, these functions DO need to change.

5 cosmological models are available (this argument must be spelled exactly):

PLANCK15, PLANCK13, WMAP9, WMAP7, WMAP5

#### 7.28.2.12 free\_LOSC\_data()

/brief Free data allocated by prep\_LOSC\_data function

#### 7.28.2.13 initiate\_LumD\_Z\_interp()

Function that uses the GSL libraries to interpolate pre-calculated Z-D\_L data.

Initiates the requried functions – GSL interpolation requires allocating memory before hand

#### 7.28.2.14 pow\_int()

Local power function, specifically for integer powers.

Much faster than the std version, because this is only for integer powers

# 7.28.2.15 printProgress()

routine to print the progress of a process to the terminal as a progress bar

Call everytime you want the progress printed

```
7.28.2.16 read_file() [1/2]
```

```
void read_file (
          std::string filename,
          double ** output,
          int rows,
          int cols )
```

Utility to read in data.

Takes filename, and assigns to output[rows][cols]

File must be comma separated doubles

#### **Parameters**

	filename	input filename, relative to execution directory
out output array to store output, dimensions rowsXco		array to store output, dimensions rowsXcols
	rows	first dimension
	cols	second dimension

Utility to read in data (single dimension vector)

Takes filename, and assigns to output[i\*rows + cols]

Output vector must be long enough, no check is done for the length

File must be comma separated doubles

#### **Parameters**

	filename	input filename, relative to execution directory
out	output	output array, assumed to have the proper length of total items

# 7.28.2.18 read\_LOSC\_data\_file()

Read data file from LIGO Open Science Center.

Convenience function for cutting off the first few lines of text

	filename	input filename
out	output	Output data
out	data_start_time	GPS start time of the data in file
out	duration	Duration of the signal
out	fs	Sampling frequency of the data

```
7.28.2.19 read_LOSC_PSD_file()
void read_LOSC_PSD_file (
              std::string filename,
              double ** output,
              int rows,
              int cols )
Read PSD file from LIGO Open Science Center.
Convenience function for cutting off the first few lines of text
7.28.2.20 transform_cart_sph()
void transform_cart_sph (
              double * cartvec,
              double * sphvec )
utility to transform a vector from cartesian to spherical (radian)
order:
cart: x, y, z
spherical: r, polar, azimuthal
7.28.2.21 transform_sph_cart()
void transform_sph_cart (
              double * sphvec,
              double * cartvec )
utility to transform a vector from spherical (radian) to cartesian
order:
cart: x, y, z
spherical: r, polar, azimuthal
7.28.2.22 tukey_window()
void tukey_window (
              double * window,
              int length,
              double alpha )
Tukey window function for FFTs.
As defined by https://en.wikipedia.org/wiki/Window_function
```

```
7.28.2.23 write_file() [1/2]
```

```
void write_file (
          std::string filename,
          double ** input,
          int rows,
          int cols )
```

Utility to write 2D array to file.

Grid of data, comma separated

Grid has rows rows and cols columns

#### **Parameters**

filename	Filename of output file, relative to execution directory
input	Input 2D array pointer array[rows][cols]
rows	First dimension of array
cols	second dimension of array

```
7.28.2.24 write_file() [2/2]
```

Utility to write 1D array to file.

Single column of data

# **Parameters**

filename	Filename of output file, relative to execution directory	
input	input 1D array pointer array[length]	
length length of array		

# 7.28.2.25 XLALSpinWeightedSphericalHarmonic()

Shamelessly stolen from LALsuite

#### **Parameters**

theta	polar angle (rad)	
phi	azimuthal angle (rad)	
s	spin weight	
1	mode number l	
m	mode number m	

#### 7.28.2.26 Z\_from\_DL()

Calculates the redshift given the luminosity distance.

Based on Astropy.cosmology calculations – see python script in the ./data folder of the project – numerically calculated given astropy.cosmology's definitions ( http://docs.astropy.org/en/stable/cosmology/) and used scipy.optimize to fit to a power series, stepping in half powers of DL. These coefficients are then output to a header file (D\_Z\_config.h) which are used here to calculate redshift. Custom cosmologies etc can easily be acheived by editing the python script D\_Z\_config.py, the c++ functions do not need modification. They use whatever data is available in the header file.

5 cosmological models are available (this argument must be spelled exactly, although case insensitive):

PLANCK15, PLANCK13, WMAP9, WMAP7, WMAP5

Function that returns Z from a given luminosity Distance – only Planck15

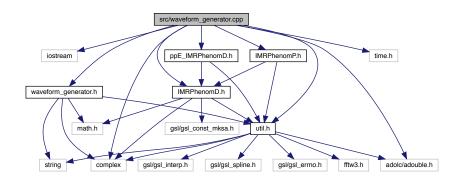
adouble version for ADOL-C calculations

Function that returns Z from a given luminosity Distance – only Planck15

# 7.29 src/waveform\_generator.cpp File Reference

```
#include <iostream>
#include "waveform_generator.h"
#include "IMRPhenomD.h"
#include "IMRPhenomP.h"
#include "ppE_IMRPhenomD.h"
#include "util.h"
#include <complex>
#include <time.h>
#include <adolc/adouble.h>
```

Include dependency graph for waveform\_generator.cpp:



# **Functions**

- int fourier\_waveform (double \*frequencies, int length, std::complex< double > \*waveform\_plus, std

  ::complex< double > \*waveform\_cross, string generation\_method, gen\_params \*parameters)
  - Function to produce the plus/cross polarizations of an quasi-circular binary.
- int fourier\_waveform (double \*frequencies, int length, double \*waveform\_plus\_real, double \*waveform—
   \_plus\_imag, double \*waveform\_cross\_real, double \*waveform\_cross\_imag, string generation\_method,
   gen\_params \*parameters)
- int fourier\_waveform (double \*frequencies, int length, std::complex< double > \*waveform, string generation\_method, gen\_params \*parameters)
  - Function to produce the (2,2) mode of an quasi-circular binary.
- int fourier\_waveform (double \*frequencies, int length, double \*waveform\_real, double \*waveform\_imag, string generation\_method, gen\_params \*parameters)
- int fourier\_amplitude (double \*frequencies, int length, double \*amplitude, string generation\_method, gen\_params \*parameters)
  - Function to produce the amplitude of the (2,2) mode of an quasi-circular binary.
- int fourier\_phase (double \*frequencies, int length, double \*phase, string generation\_method, gen\_params \*parameters)

Function to produce the phase of the (2,2) mode of an quasi-circular binary.

#### 7.29.1 Detailed Description

File that handles the construction of the (2,2) waveform as described by IMRPhenomD by Khan et. al.

Builds a waveform for given DETECTOR FRAME parameters

# 7.29.2 Function Documentation

# 7.29.2.1 fourier\_amplitude()

Function to produce the amplitude of the (2,2) mode of an quasi-circular binary.

By using the structure parameter, the function is allowed to be more flexible in using different method of waveform generation - not all methods use the same parameters

#### **Parameters**

frequencies	double array of frequencies for the waveform to be evaluated at	
length	integer length of all the arrays	
amplitude	output array for the amplitude	
generation_method	String that corresponds to the generation method - MUST BE SPELLED EXACTLY	

#### 7.29.2.2 fourier\_phase()

Function to produce the phase of the (2,2) mode of an quasi-circular binary.

By using the structure parameter, the function is allowed to be more flexible in using different method of waveform generation - not all methods use the same parameters

frequencies	double array of frequencies for the waveform to be evaluated at	
length	integer length of all the arrays	
phase	output array for the phase	
generation_method  String that corresponds to the generation method - MUST BE SPELLED EXACT		

#### 7.29.2.3 fourier\_waveform() [1/4]

Function to produce the plus/cross polarizations of an quasi-circular binary.

By using the structure parameter, the function is allowed to be more flexible in using different method of waveform generation - not all methods use the same parameters

This puts the responsibility on the user to pass the necessary parameters

NEED TO OUTLINE OPTIONS FOR EACH METHOD IN DEPTH

**NEW PHASE OPTIONS for** 

#### PHENOMD ONLY:

If phic is assigned, the reference frequency and reference phase are IGNORED.

If Phic is unassigned, a reference phase AND a reference frequency are looked for. If no options are found, both are set to 0.

If to is assigned, it is used.

If tc is unassigned, the waveform is shifted so the merger happens at 0.

PhenomPv2:

PhiRef and f\_ref are required, phic is not an option.

tc, if specified, is used with the use of interpolation. If not, tc is set such that coalescence happens at t=0

#### **Parameters**

	frequencies	double array of frequencies for the waveform to be evaluated at
	length	integer length of all the arrays
out	waveform_plus	complex array for the output plus polarization waveform
out	waveform_cross	complex array for the output cross polarization waveform
	generation_method	String that corresponds to the generation method - MUST BE SPELLED EXACTLY
	parameters	structure containing all the source parameters

#### 7.29.2.4 fourier\_waveform() [2/4]

```
int length,
double * waveform_plus_real,
double * waveform_cross_real,
double * waveform_cross_imag,
string generation_method,
gen_params * parameters )
```

#### **Parameters**

frequencies	double array of frequencies for the waveform to be evaluated at	
length	integer length of all the arrays	
waveform_plus_real	complex array for the output waveform	
waveform_plus_imag	complex array for the output waveform	
waveform_cross_real	complex array for the output waveform	
waveform_cross_imag	complex array for the output waveform	
generation_method	String that corresponds to the generation method - MUST BE SPELLED EXACTLY	
parameters	structure containing all the source parameters	

# **7.29.2.5** fourier\_waveform() [3/4]

Function to produce the (2,2) mode of an quasi-circular binary.

By using the structure parameter, the function is allowed to be more flexible in using different method of waveform generation - not all methods use the same parameters

# **Parameters**

frequencies	double array of frequencies for the waveform to be evaluated at	
length	integer length of all the arrays	
waveform	complex array for the output waveform	
generation_method	String that corresponds to the generation method - MUST BE SPELLED EXACTLY	
parameters	structure containing all the source parameters	

# 7.29.2.6 fourier\_waveform() [4/4]

```
double * waveform_real,
double * waveform_imag,
string generation_method,
gen_params * parameters )
```

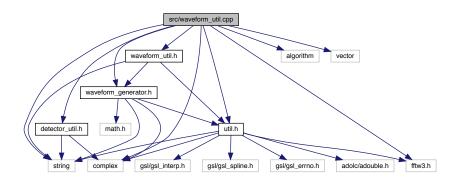
#### **Parameters**

frequencies	double array of frequencies for the waveform to be evaluated at	
length	integer length of all the arrays	
waveform_real	complex array for the output waveform	
waveform_imag	complex array for the output waveform	
generation_method	String that corresponds to the generation method - MUST BE SPELLED EXACTLY	
parameters	structure containing all the source parameters	

# 7.30 src/waveform\_util.cpp File Reference

```
#include "waveform_util.h"
#include "util.h"
#include "waveform_generator.h"
#include "detector_util.h"
#include <fftw3.h>
#include <algorithm>
#include <complex>
#include <vector>
#include <string>
```

Include dependency graph for waveform\_util.cpp:



#### **Functions**

- double data\_snr\_maximized\_extrinsic (double \*frequencies, int length, std::complex< double > \*data, double \*psd, std::string detector, std::string generation\_method, gen\_params \*param)
  - Utility to calculate the snr of a fourier transformed data stream while maximizing over the coalescence parameters phic and tc.
- double data\_snr\_maximized\_extrinsic (double \*frequencies, int length, double \*data\_real, double \*data\_← imag, double \*psd, std::string detector, std::string generation\_method, gen\_params \*param)
  - Light wrapper for the data\_snr\_maximized\_extrinsic method.

• double calculate\_snr (std::string detector, std::complex< double > \*waveform, double \*frequencies, int length)

Caclulates the snr given a detector and waveform (complex) and frequencies.

- int fourier\_detector\_response (double \*frequencies, int length, std::complex< double > \*hplus, std
   ::complex< double > \*hcross, std::complex< double > \*detector\_response, double theta, double phi, std
   ::string detector)

Function to produce the detector response caused by impinging gravitational waves from a quasi-circular binary.

• int fourier\_detector\_amplitude\_phase (double \*frequencies, int length, double \*amplitude, double \*phase, std::string detector, std::string generation\_method, gen\_params \*parameters)

Calculates the amplitude (magnitude) and phase (argument) of the response of a given detector.

#### 7.30.1 Detailed Description

Utilities for waveforms - SNR calculation and detector response

includes snr and detector response

#### 7.30.2 Function Documentation

#### 7.30.2.1 calculate\_snr()

Caclulates the snr given a detector and waveform (complex) and frequencies.

This function computes the un-normalized snr: \sqrt( ( H | H ) )

detector	detector name - must match the string of populate_noise precisely	
waveform	complex waveform	
frequencies	ies double array of frequencies that the waveform is evaluated at	
length	length of the above two arrays	

```
7.30.2.2 data_snr_maximized_extrinsic() [1/2]
```

```
int length,
std::complex< double > * data,
double * psd,
std::string detector,
std::string generation_method,
gen_params * param )
```

Utility to calculate the snr of a fourier transformed data stream while maximizing over the coalescence parameters phic and tc.

The gen\_params structure holds the parameters for the template to be used (the maximimum likelihood parameters)

#### **Parameters**

frequencies	Frequencies used by data	
length	length of the data	
data	input data in the fourier domain	
psd	PSD for the detector that created the data	
detector	Name of the detector –See noise_util for options	
generation_method	Generation method for the template – See waveform_generation.cpp for options	
param	gen_params structure for the template	

#### 7.30.2.3 data\_snr\_maximized\_extrinsic() [2/2]

 $\label{light-wrapper-for-the-data\_snr\_maximized\_extrinsic method.}$ 

Splits the data into real and imaginary, so all the arguments are C-safe

frequencies	Frequencies used by data	
length	length of the data	
data_real	input data in the fourier domain – real part	
data_imag	input data in the fourier domain – imaginary part	
psd	PSD for the detector that created the data	
detector	Name of the detector –See noise_util for options	
generation_method	Generation method for the template – See waveform_generation.cpp for options	
param	gen_params structure for the template	

#### 7.30.2.4 fourier\_detector\_amplitude\_phase()

Calculates the amplitude (magnitude) and phase (argument) of the response of a given detector.

This is for general waveforms, and will work for precessing waveforms

Not as fast as non-precessing, but that can't be helped. MUST include plus/cross polarizations

#### 7.30.2.5 fourier\_detector\_response() [1/2]

#### **Parameters**

	frequencies	array of frequencies corresponding to waveform
	length	length of frequency/waveform arrays
	hcross	precomputed cross polarization of the waveform
out	detector_response	detector response
	theta	polar angle (rad) theta in detector frame
	phi	azimuthal angle (rad) phi in detector frame
	detector	detector - list of supported detectors in noise_util

# 7.30.2.6 fourier\_detector\_response() [2/2]

Function to produce the detector response caused by impinging gravitational waves from a quasi-circular binary.

By using the structure parameter, the function is allowed to be more flexible in using different method of waveform generation - not all methods use the same parameters

This puts the responsibility on the user to pass the necessary parameters

Detector options include classic interferometers like LIGO/VIRGO (coming soon: ET and LISA)

This is a wrapper that combines generation with response functions: if producing mulitple responses for one waveform (ie stacking Hanford, Livingston, and VIRGO), it will be considerably more efficient to calculate the waveform once, then combine each response manually

	frequencies	double array of frequencies for the waveform to be evaluated at
	length	integer length of all the arrays
out	response	complex array for the output plus polarization waveform
	generation_method	String that corresponds to the generation method - MUST BE SPELLED EXACTLY
	parameters	structure containing all the source parameters

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