

# Documentation for the `gluex_root_analysis` software

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## Abstract

The *GlueX* ROOT analysis software is used to analyze data in the standard PART (Physics Analysis ROOT Tree) format based on the *DSelector* software concept. In order to prepare the PART files, the raw experimental data in EVIO format are input to the *GlueX* reconstruction software that reconstructs charged particle tracks and neutral particle showers and stores this information in Reconstructed Event Storage (REST) format. Using the *ReactionFilter* plugin, the *GlueX* Analysis library builds combinations of final state particles and applies standard loose event selections to generate skims of the reconstructed data based on the event reaction supplied by the analyzer. The output of *ReactionFilter* is a ROOT file which can be read by the *DSelector* software package and provides the basis for a *GlueX* ROOT analysis, as described in this document. We will discuss basic concepts of the *DSelector*, the fields available in the PART format, and methods to perform various typical analysis activities. In the following discussion it is assumed that you have configured a working *GlueX* software environment.

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# 1 Introduction

The *GlueX* ROOT analysis software package provides an executable named `MakeDSelector` that allows the user to generate a basic framework of *DSelector* program files for the analysis of a particular reaction. This code will be different for each reaction. The location of this executable is defined by the environment variable `ROOT_ANALYSIS_HOME`, which is configured by the standard GlueX software configuration scripts. An example of how to use this is shown below:

```
MakeDSelector input-root-tree-file-name root-tree-name program-name
```

where `input-root-tree-file-name` is the name of a PART file, `root-tree-name` is the name of the TTree contained in this file and `program-name` can be any label you choose. The command above will create two files in the current directory with the name `DSelector_program-name` with the extensions `.C` and `.h` which contain the basic template to read and analyze the specified ROOT tree. The files contain the general functionality, with some comments and example code for the user to expand on.

## 2 The DSelector class structure

In the `.h` header file you can find a class defined as `DSelector_program-name` that inherits from the `DSelector` class. The `DSelector` class is defined in the *GlueX* ROOT analysis library and provides all of the methods that are necessary to read the ROOT tree. It inherits from the ROOT `TSelector` class that provides the methods `Init()`, `Process()` and `Finalize()` and the structure to loop over all of the entries in the ROOT tree; these form the base of the `.C` file generated.

The `DSelector_program-name` can be run in several ways. For testing with just one input file, it can be run using ROOT interactively, eg

```
root mytreefile.root
.x $ROOT_ANALYSIS_HOME/scripts/Load_DSelector.C
name-of-the-root-tree->Process("DSelector_program-name.C+")
```

Note that you should supply either one or two plus signs at the end of the `DSelector` file name to increase the processing speed of your analysis. If you specify `+`, the `DSelector` will only be recompiled if you modify either of the `DSelector` files. The preferred option is `++`, which will force recompilation at every execution, which can help protect against errors that occur when switching between software versions.

The `DSelector_program-name` can be run over one or more files using a root script that has the following form:

```
TChain chain("name-of-the-root-tree");
chain->Add("root-file-name");
gROOT->ProcessLine(".x $ROOT_ANALYSIS_HOME/scripts/Load_DSelector.C");
chain.Process("DSelector_program-name.C++");
```

Alternatively, the `DSelector_program-name` can be run in parallel on multiple cores with PROOF-Lite:

```
int NThreads = 16; // or as many as you want to use
TChain *chain = new TChain("name-of-the-root-tree");
chain.Add("root-file-name");
gROOT->ProcessLine(".x $ROOT_ANALYSIS_HOME/scripts/Load_DSelector.C");
DPROOFLiteManager *dproof = new DPROOFLiteManager();
dproof->Process_Chain(chain, "DSelector_program-name.C+", NThreads, "outputHistFileName",
"outputTreeFileName");
```

### 2.1 The Wrapper concept

Each piece of information is stored as a branch in the ROOT tree. Several wrapper objects are implemented, to group the branches belonging to the same reconstructed object (track/shower/combination of these) to-

gether. These wrappers are provided by the *DSelector* and are defined in `ROOT_ANALYSIS_HOME/libraries/DSelector`. There are four basic types of objects in the tree – charged particle tracks, neutral particles, combos and beam photons – which leads to the definition of four standard wrapper objects:

```
// OBJECT ARRAYS: RECONSTRUCTED
DChargedTrackHypothesis* dChargedHypoWrapper;
DNeutralParticleHypothesis* dNeutralHypoWrapper;
DBeamParticle* dBeamWrapper;
DParticleCombo* dComboWrapper;
```

The first three wrappers are used to access the reconstructed final state particles, charged and neutral, and the reconstructed initial state beam photons. The fourth wrapper represents the output of the standard analysis library used by *ReactionFilter*, which combines the initial and final state particles, with kinematic fitting by default, to generate the full event. The resulting final state particles and intermediate states (if defined) from the *ReactionFilter* are accessed by additional wrappers specific to the reaction at hand. As an example, an exclusive final state like  $\pi^+ + \pi^- + \pi^0 + p$  with the  $\pi^0$  decaying into two photons and its mass being constrained in the kinematic fit will lead to 8 additional wrappers in the generated *DSelector* that include a wrapper for the initial state beam photon and the decay of the  $\pi^0$  with its mass being constrained by the fit:

```
//Step 0
DParticleComboStep* dStep0Wrapper;
DBeamParticle* dComboBeamWrapper;
DChargedTrackHypothesis* dPiPlusWrapper;
DChargedTrackHypothesis* dPiMinusWrapper;
DChargedTrackHypothesis* dProtonWrapper;

//Step 1
DParticleComboStep* dStep1Wrapper;
DKinematicData* dDecayingPi0Wrapper;
DNeutralParticleHypothesis* dPhoton1Wrapper;
DNeutralParticleHypothesis* dPhoton2Wrapper;
```

For example, to loop over all charged particle tracks of an event that were used in forming all the available combos and perform a calculation using the value of the start counter energy-loss one might attempt something like the following:

```
// check if any charged track has a ST hit by looking at its dEdx value
int NChargedTracksHypos = (int)Get_NumChargedHypos();
int FoundStartCounterHits = 0;
for (int k=0; k<NChargedTracksHypos; k++){
    dChargedHypoWrapper->Set_ArrayIndex(k);
    float ST_dEdx = dChargedHypoWrapper->Get_dEdx_ST();
    if (ST_dEdx>0.){
        FoundStartCounterHits++; // NOTE THAT THIS CODE OVERCOUNTS THE NUMBER OF HITS!
    }
}
```

The key to this loop is that one sets the pointer of the wrapper to the right index by using the method `Set_ArrayIndex(k)`. Now that the wrapper points to the desired object one can access the desired value from that object.

Note that this code will not do what the user desires for several reasons. First, by default *ReactionFilter* will only write out the initial and final state particles used in the combos saved to the ROOT tree, unless the U1 option is specified. Second, each charged track is reconstructed under several different mass hypotheses ( $e$ ,  $\pi$ ,  $K$ ,  $p/\bar{p}$  by default), and it is these specific hypotheses which are used in the reconstructed combos that are saved to the ROOT tree. The mapping between charged particle hypotheses and reconstructed tracks is given by the `DChargedTrackHypothesis::Get_TrackID()` function, so one could in principle keep track of the number of unique track IDs, but in general this is an issue to be very careful about.

## 2.2 Init(TTree \*locTree) Method

This function is used to initialize histograms and any other class variables which will be used by the analysis code for every event. This method can and most likely will be called more than once, in particular when using a TChain with more than one root file or when using PROOF in the context of multi-threading. This method is inherited from the ROOT TSelector class and overridden in the DSelector class. The code snippet in the Init() method shown below will ensure that the code that follows after will be executed only once:

```
bool locInitializedPriorFlag = dInitializedFlag; //save whether have been initialized previously
DSelector::Init(locTree); //This must be called to initialize wrappers for each new TTree
//gDirectory now points to the output file with name dOutputFileName (if any)
if(locInitializedPriorFlag)
    return; //have already created histograms, etc. below: exit
```

Therefore any initialization of variables or histogram definitions intended for the whole analysis need to be done after this part of the code. This includes any definitions of Analysis Actions and Cut Actions as well as custom branches for an output ROOT tree.

The call to the method Get.ComboWrappers() initializes all instances of wrappers available for the analysis of the ROOT tree. This call will configure a wrapper for each particle in the reaction. For example, if the reaction is  $\gamma p \rightarrow p\pi^0\pi^+\pi^-$ ,  $\pi^0 \rightarrow \gamma\gamma$  the following pointers defined in the class header will be initialized

```
//Step 0
dComboBeamWrapper = static_cast<DBeamParticle*>(dStep0Wrapper->Get_InitialParticle());
dPiPlusWrapper = static_cast<DChargedTrackHypothesis*>(dStep0Wrapper->Get_FinalParticle(1));
dPiMinusWrapper = static_cast<DChargedTrackHypothesis*>(dStep0Wrapper->Get_FinalParticle(2));
dProtonWrapper = static_cast<DChargedTrackHypothesis*>(dStep0Wrapper->Get_FinalParticle(3));

//Step 1
dStep1Wrapper = dComboWrapper->Get_ParticleComboStep(1);
dDecayingPi0Wrapper = dStep1Wrapper->Get_InitialParticle();
dPhoton1Wrapper = static_cast<DNeutralParticleHypothesis*>(dStep1Wrapper->Get_FinalParticle(0));
dPhoton2Wrapper = static_cast<DNeutralParticleHypothesis*>(dStep1Wrapper->Get_FinalParticle(1));
```

These wrapper objects, as shown in the example above, are instances of classes and are pointers to the corresponding instances of the particles for a given combo and information about the combo itself. All instances are defined in the DSelector library and provide methods to access any data within the tree for a given combo. So for example with the line

```
TLorentzVector xv4 = dPhoton1Wrapper->Get_X4_Shower();
```

you will access the Lorentz 4-vector for the calorimeter shower associated with the first photon used in this example reaction for a given combo within the combo loop. This will be discussed in more detail below when describing the Process() method.

The full particle combination corresponding to the reconstructed reaction is described by a DParticleCombo object, which contains one or more DParticleComboStep objects. Each DParticleComboStep corresponds to an intermediate decay step in the overall reaction. For example, the reaction  $\gamma p \rightarrow p\pi^+\pi^-$  has only one step (Step0), while  $\gamma p \rightarrow p\pi^0\pi^+\pi^-$ ,  $\pi^0 \rightarrow \gamma\gamma$  has two steps, with Step0 being the primary photoproduction reaction and Step1 being the  $\pi^0 \rightarrow \gamma\gamma$  decay.

Each final state particle is either charged or neutral and is accessed through a wrapper of type DNeutralParticleHypothesis or DChargedTrackHypothesis. There are many methods associated with these two classes that provide access to all the relevant variables in the tree associated with these particles within a combo. Intermediate particles that decay into some final state particles will also be represented by a wrapper class of type DKinematicData but only if the mass of this intermediate state particle is constrained in the kinematic fit when requested by the reaction filter. If the intermediate particle mass is not constrained in the kinematic fit, then its properties (e.g. 4-momentum) can be constructed by the user based on the other reconstructed particles. All beam photons are represented by the wrapper class DBeamParticle. Note that the beam photon energies are not altered by the kinematic fitter. More details about these wrappers and their methods will be discussed further below when discussing the loop over all combos of an event.

### 2.2.1 Histogram definitions

Histograms can be defined easily as in any root script. The user will need to define histograms as needed for their analysis. Note that you should define the histogram variables in the class definition in the `.h` file before creating them in the `DSelector::Init()` function.

## 2.3 Analysis Actions

Analysis actions are a useful system to compartmentalize analysis code, which is applied to every combination in the tree. This code can either include histograms (see Section 2.3.1) or event selection cuts (see Section 2.3.2). Every analysis action is registered as an entry in the vector `dAnalysisActions`, and they are executed in the same order as specified in the `Init()` method.

```
dAnalysisActions.push_back(new DHistogramAction_NAME(...));  
OR  
dAnalysisActions.push_back(new DCutAction_NAME(...));
```

Three histograms are automatically generated and show the effect of each action on the data. Evidently, histogram actions will not alter the number of surviving events or combos.

`NumEventsSurvivedAction` Shows the number of events that passed each analysis action

`NumCombosSurvivedAction` Shows the number of combinations that passed each analysis action. This is a 2-dimensional histogram, as the number of initial combinations is different for each event.

`NumCombosSurvivedAction` Shows the total number of combinations that passed each analysis action, added for all events.

In order for the analysis actions to work as intended, the entire vector of analysis actions has to be initialized in the `Init()` method:

```
Initialize_Actions();
```

and reset for every event in the `Process()` method:

```
Reset_Actions_NewEvent();
```

Within the loop over all combinations in the tree, the actions can be executed at any time with this command:

```
if(!Execute_Actions())  
    continue;
```

This will skip the rest of your `DSelector` in case the combination does not satisfy all cuts, which may speed up the process. The `IsComboCut` flag is automatically set for these combinations.

Several analysis actions are provided in the code and are summarized below. The reader should be encouraged to extend this list.

[Explanation on how to write a completely new custom action, maybe a template generating script?](#)

### 2.3.1 Histogram Actions

Currently available histogram actions are listed in the following. Boolean variables always specify if the measured (false) or the fitted four-vectors (true) are used. The optional `string` is a unique identifier, which is only needed if the same action is used multiple times.

`AnalyzeCutActions` see 2.3.3

`ParticleComboKinematics` Creates a set of histograms for the kinematic distributions of each particle in the reaction.

```
DHistogramAction_ParticleComboKinematics(dComboWrapper, bool, "")
```

**ParticleID** Creates a set of PID histograms for each particle in the reaction

```
DHistogramAction_ParticleID(dComboWrapper, bool, "")
```

**InvariantMass** Plots the invariant mass of a particle or set of particles. 2D histograms as a function of the kinematic fit confidence level will be generated in linear and logarithmic scale. The step specifies the level in the decay chain, where 0 is the main reaction. In the example below, the mass is plotted in 100 bins between 1.0 and 2.0 GeV.

```
DHistogramAction_InvariantMass(dComboWrapper, bool, Lambda, 100, 1.0, 2.0, "")  
OR  
DHistogramAction_InvariantMass(dComboWrapper, bool, step, {Proton,Pi0}, 100, 1.0, 2.0, "")
```

**MissingMass** This action can plot the total missing mass of the reaction or the recoil mass against a particle or set of particles. In the example below, the missing mass is plotted in 100 bins between -1.0 and 1.0 GeV.

```
DHistogramAction_MissingMass(dComboWrapper, bool, 100, -1.0, 1.0, "")  
OR  
DHistogramAction_MissingMass(dComboWrapper, bool, step, Proton, 100, -1.0, 1.0, "")  
OR  
DHistogramAction_MissingMass(dComboWrapper, bool, step, {Proton,Pi0}, 100, -1.0, 1.0, "")
```

**MissingMassSquared** This plots the square of the missing mass. The syntax is the same as above.

**MissingP4** This action plots missing energy, 3 missing momentum components  $p_x, p_y, p_z$  and missing transverse momentum  $p_T$ .

```
DHistogramAction_MissingP4(dComboWrapper, bool, "")
```

**2DInvariantMass** This action can be used to plot the invariant mass of one particle combination against another one. All particles have to exist in the same reaction step.

```
DHistogramAction_2DInvariantMass(dComboWrapper, bool, step, {Pi0,Proton}, {Eta,Proton},  
100, 0, 2, 100, 0, 2, "")
```

**Dalitz** This plots the square of two invariant mass combinations against each other. The syntax is exactly the same as above.

**vanHove** This action plots the vanHove plot for three particles or three sets of particles

```
DHistogramAction_vanHove(dComboWrapper, bool, Proton, Eta, Pi0, "")  
OR  
DHistogramAction_vanHove(dComboWrapper, bool, {Proton,PiMinus}, Eta, Pi0, "")
```

**vanHoveFour** Plots the great-circle plot for 4 particles or particle systems.

**KinFitResults** Plots the  $\chi^2/NDF$  and the confidence level of the kinematic fit, the latter also in a plot with a logarithmic horizontal axis.

```
DHistogramAction_KinFitResults(dComboWrapper, "")
```

**BeamEnergy** Plots the beam energy.

```
DHistogramAction_BeamEnergy(dComboWrapper, bool, "")
```

`Energy_UnusedShowers` Plot the total energy of the unused showers in the reaction.

```
DHistogramAction_Energy_UnusedShowers(dComboWrapper, "")
```

### 2.3.2 Cut Actions

Every Cut Action will skip the rest of the code if a combination does not pass the selection and will set the flag `IsComboCut` accordingly. They are added to the `dAnalysisActions` vector exactly like Histogram Actions. Currently available cut actions are:

`ChiSqOrCL` Special cut action to compare multiple reactions, see dedicated section (Section 2.3.4)

`PIDDeltaT` Cut on the timing of one type of particle in a certain detector

```
DCutAction_PIDDeltaT(dComboWrapper, bool, 0.5, PiPlus, SYS_TOF, "")
```

`NoPIDHit` Cut specific particles that have no particle ID hit in a certain detector

```
DCutAction_NoPIDHit(dComboWrapper, PiPlus, SYS_BCAL, "")
```

`dEdx` Predefined cut on the specific energy loss  $dE/dx$ . Automatically distinguishes between heavy particles (mass  $\geq$  proton mass) and light particles and uses predefined function (might need tuning!). Only the cut on the CDC  $dE/dx$  has been implemented.

```
DCutAction_dEdx(dComboWrapper, bool, PiPlus, SYS_CDC, "")
```

`KinFitFOM` Cut all combos with smaller kinematic fit confidence level

```
DCutAction_KinFitFOM(dComboWrapper, double locMinimumConfidenceLevel, "")
```

`KinFitChiSq` Cut all combos with larger kinematic fit  $\chi^2/NDF$ .

```
DCutAction_KinFitChiSq(dComboWrapper, double locMaximumChiSq, "")
```

`MissingMass` This cuts the total missing mass of the reaction or the recoil mass against a particle or set of particles.

```
DCutAction_MissingMass(dComboWrapper, bool, 0.1, 0.2, "")
OR
DCutAction_MissingMass(dComboWrapper, bool, step, Pi0, 0.1, 0.2, "")
OR
DCutAction_MissingMass(dComboWrapper, bool, step, {Proton, Pi0}, 0.5, 1.5, "")
```

`MissingMassSquared` Cuts on the squared value of the missing mass. The syntax is the same as for the missing mass.

`MissingEnergy` Cuts on the missing energy of the reaction. Syntax is the same as for missing mass.

`InvariantMass` Cut on the invariant mass of a particle or a particle combination within the same reaction step.

```
DCutAction_InvariantMass(dComboWrapper, bool, Pi0, 0.10, 0.15, "")
OR
DCutAction_InvariantMass(dComboWrapper, bool, step, {Proton, PiMinus}, 1.0, 1.5, "")
```

`InvariantMassVeto` same as above

`BeamEnergy` Cut on beam energy interval.

```
DCutAction_BeamEnergy(dComboWrapper, bool, locMinBeamEnergy, locMaxBeamEnergy, "")
```

`TrackShowerEOverP` Cut on  $E/p$  above value for  $e^+/e^-$ , below for all other particles.

```
DCutAction_TrackShowerEOverP(dComboWrapper, bool, SYS_FCAL, Electron, 0.7, "")
```

`ShowerQuality` Requires a minimal shower quality for all photons in the detector, here cut below 0.5 for FCAL

```
DCutAction_ShowerQuality(dComboWrapper, SYS_FCAL, 0.5)
```

`Kinematics` Cut on momentum,  $\theta$  and  $\phi$  of a particle. If the lower limit is larger than the higher limit for one variable, the cut on this variable is ignored.

```
DCutAction_Kinematics(dComboWrapper, step, Proton, bool, MinP, MaxP,  
MinTheta = 1.0, MaxTheta = 0.0, MinPhi = 1.0, MaxPhi = 0.0)
```

`TrackBCALPreshowerFraction` Cut on preshower fraction, weighted by  $\sin \theta$  of the track. At the moment only implemented for  $e^+/e^-$ .

```
DCutAction_TrackBCALPreshowerFraction(dComboWrapper, bool, Electron, 0.1, "")
```

`Energy_UnusedShowers` Cut on the summed energy of unused showers.

```
DCutAction_Energy_UnusedShowers(dComboWrapper, locMaxEnergy_UnusedShowersCut, "")
```

`NumUnusedTracks` Cut on the maximum number of unused tracks.

```
DCutAction_NumUnusedTracks(dComboWrapper, locMaxUnusedTracks, "")
```

`NumUnusedShowers` Cut on the number of unused showers.

```
DCutAction_NumUnusedShowers(dComboWrapper, locMaxUnusedShowers, "")
```

`VanHoveAngle` Select a sector in the vanHove plane.

```
DCutAction_VanHoveAngle(dComboWrapper, bool, Proton, Eta, Pi0, MinAngle, MaxAngle, "")  
OR  
DCutAction_VanHoveAngle(dComboWrapper, bool, {Proton, Pi0}, Eta, Pi0, MinAngle, MaxAngle)
```

`VanHoveAngleFour` Select a part of the vanHove space for 4 particles, defined by 2 angles. Syntax similar as above.

### 2.3.3 Analyze Cut Actions

### 2.3.4 KinFit Comparison between different reactions

### 2.3.5 Analysis Utilities

`Get_IsPolarizedBeam` If the RCDB environment is set up, this function returns `true` for polarized beam runs. It also sets a flag according to PARA/PERP.

```
bool Get_IsPolarizedBeam(int locRunNumber, bool& locIsPARAFlag)
```



`Get_PolarizationAngle` If the RCDB environment is set up, this function returns the polarization angle.

```
bool Get_PolarizationAngle(int locRunNumber, int& locPolarizationAngle
```

`Get_CoherentPeak` If the RCDB environment is set up, this function returns the coherent peak energy for polarized runs, 0 for amorphous runs.

```
bool Get_CoherentPeak(int locRunNumber, double& locCoherentPeak, bool locIsPolarizedFlag)
```

`Get_BeamBunchPeriod` If the RCDB environment is set up, this returns the beam bunch period.

```
double Get_BeamBunchPeriod(int locRunNumber)
```

`Get_AccidentalScalingFactor` If the RCDB environment is set up, this returns the accidental scaling factor.

```
double Get_AccidentalScalingFactor(int locRunNumber, double locBeamEnergy, bool locIsMC)
```

`AccidentalScalingFactorError` **Caution:** Uncertainty on the accidental scaling factor is not defined at this point.

```
double Get_AccidentalScalingFactorError(int locRunNumber, double locBeamEnergy, bool locIsMC)
```

`Get_DeltaT_RF` Calculate the time difference with respect to the RF signal.

```
double Get_DeltaT_RF(int locRunNumber, locBeamX4_Measured, dComboWrapper)
```

`Get_RelativeBeamBucket` Calculate which relative beam bucket this combination is in.

```
int Get_RelativeBeamBucket(int locRunNumber, locBeamX4_Measured, dComboWrapper)
```

`Get_BeamEndpoint` Get the energy of the beam end point from CCDB.

```
double Get_BeamEndpoint(int locRunNumber)
```

`Get_ColumnTAGM` Get the tagger microscope column for this beam energy from CCDB.

```
int Get_ColumnTAGM(int locRunNumber, double locBeamEnergy)
```

`Get_CounterTAGH` Get the tagger hodoscope counter for this beam energy from CCDB.

```
int Get_CounterTAGH(int locRunNumber, double locBeamEnergy)
```

`Calc_ProdPlanePhi_Pseudoscalar` Calculates the  $\phi$  angle for beam asymmetries of pseudoscalar meson photoproduction.

```
double Calc_ProdPlanePhi_Pseudoscalar(locBeamEnergy, Proton, locMesonP4)
```

`Calc_DecayPlanePsi_Vector_2BodyDecay` Calculates decay asymmetry angle  $\psi$  for vector meson photoproduction, in the case where the vector meson decays into 2 pseudoscalar mesons (e.g.  $\phi \rightarrow 2K$ ). Returns the scattering angle  $\theta$  as well.

```
double Calc_DecayPlanePsi_Vector_2BodyDecay(locBeamEnergy, Proton, locBaryonP4, locMesonP4, locMesonProduct1P4, locDecayPlaneTheta)
```

`CalcDecayPlanePsi_Vector_3BodyDecay` Calculates decay asymmetry angle  $\psi$  for vector meson photoproduction, in the case where the vector meson decays into 3 pseudoscalar mesons (e.g.  $\omega \rightarrow 3\pi$ ). Returns the scattering angle  $\theta$  as well.

```
double Calc_DecayPlanePsi_Vector_3BodyDecay(locBeamEnergy, Proton, locBaryonP4, locMesonP4,
      locMesonProduct1P4, locMesonProduct2P4, locDecayPlaneTheta)
```

`Calc_vanHoveCoord` Calculate the 2D vanHove coordinates from 3 four-vectors.

```
std::tuple<double,double> Calc_vanHoveCoord(locXP4, locYP4, locZP4)
```

`Calc_vanHoveCoordFour` Calculate the 3D vanHove coordinates from 4 four-vectors.

```
std::tuple<double,double,double> Calc_vanHoveCoordFour(locVec1P4, locVec2P4, locVec3P4,
      locVec4P4)
```

## 2.4 Process(Long64\_t locEntry) Method

This is the main event loop which is called for each event. There are three main sections in the default code which is provided. The first is for any event-level initialization or calculations. The second comprises the loop over all available combinations in the event. The third is for additional event-level calculations and output.

The data for the event is read from the tree with the following code section:

```
//CALL THIS FIRST
DSelector::Process(locEntry); //Gets the data from the tree for the entry
```

Note that this method `Process()` is inherited from the `ROOT` class `TSelector`. This method is overridden by the `DSelector` class and calls the method `Get_Entry()`, defined in the class `DTreeInterface`, which reads the event data from file into memory. Remember that if a `TChain` is used and a new tree file is opened, the method `Init()` will be executed again first.

If you are using analysis actions, then they need to be initialized for each event:

```
/** SETUP UNIQUENESS TRACKING ****/
//ANALYSIS ACTIONS: Reset uniqueness tracking for each action
//For any actions that you are executing manually, be sure to call Reset_NewEvent() on them here
Reset_Actions_NewEvent();
dAnalyzeCutActions->Reset_NewEvent(); // manual action, must call Reset_NewEvent()
```

If you are filling histograms yourself, the combinatorics must be carefully handled to avoid double-counting, either by “uniqueness tracking” or some other method. The discussion of these techniques is the focus of another document. [\[Add reference here when it's ready\]](#)

### 2.4.1 Looping over combos in an event

At this point the loop over all combos for this event is started. The definition of a combo is combining a beam photon and the target particle with the reconstructed final state particles in the GlueX detector that can form the defined reaction. The four-momenta of all these particles are then used by the kinematic fitter in the `DReaction` plugin to evaluate the likelihood of this combination of particles to be the reaction that triggered this event. Note that the total number of reconstructed final state particles in the GlueX detector, neutrals and charged tracks for a given event can be more than is required by the reaction. Therefore there exist methods to get the number of “Unused” neutral and charged particles for the combo (`Get_NumUnusedShowers()` and `Get_NumUnusedTracks()`) that reflect the number of particles that are not used by the kinematic fitter.

Note that even if there is a unique combination of final state particles in an event, this event still will result in multiple combos if there are multiple beam photons available for this event, which is often the case for the

intense GlueX photon beam. Typically, combos are kept if the reconstructed final state particles are in time with the tagged beam photons within 3 or 4 of the beam bunches on either side of the prompt beam bunch. For example, if you are looking at the reaction  $\gamma p \rightarrow p\pi^+\pi^-$ , and find an event with only one candidate for the negatively charged pion, the positively charged pion, and the proton, and 3 tagged photons within the required timing window, then you will end up with 3 combos for that event. All necessary wrappers are configured by calling the `dComboWrapper->Set_CombIndex(UInt_t)` function, as shown below to initialize all variables within the loop with the appropriate values for the given combo.

Note that the *DSelector* can be used not only to run over the output of the *ReactionFilter* plugin, but also to write out ROOT trees which contain a set of reduced events and/or branches containing additional information. When these trees are created, if additional selections are being applied on the combos being analyzed, an event will be written out if least one combo in that event passes the selections. To keep track of which combos have been rejected and which are kept for further analysis, we use the “Combo-Cut” flag. This can be set using `dComboWrapper->Set_IsComboCut(bool)` and its value retrieved using `dComboWrapper->Get_IsComboCut()`.

An example beginning of the combo loop is shown below:

```
for(UInt_t loc_i = 0; loc_i < Get_NumCombos(); ++loc_i)
{
    //Set branch array indices for combo and all combo particles
    dComboWrapper->Set_CombIndex(loc_i);

    // Is used to indicate when combos have been cut
    if(dComboWrapper->Get_IsComboCut()) // Is false when tree originally created
        continue; // Combo has been cut previously

    // *****
    // now from here on out you can do your stuff!
    // *****
}
```

A detailed discussion of the data available through the different particle wrappers is given in Section 3. By default, code is added to configure variables to store the unique ID numbers of each particle in the combination, along with their measured and kinematically-fitted momentum 4-vectors. Equipped with the data access methods described below and the wrappers that provide the access to these methods, we can get any data from within the tree. Continuing with the earlier example of three pions and a recoil proton in the final state, we can calculate the invariant mass of the 3-pion system using either the measured momentum 4-vectors or those determined by the kinematic fitter. In the example below, we also show how to access other information, in particular the shower quality factor for neutral particles and the z-position of the associated shower in the calorimeter:

```
// These quantities are from the kinematic fitter
TLorentzVector locBeamP4 = dComboBeamWrapper->Get_P4();
TLorentzVector locPiPlusP4 = dPiPlusWrapper->Get_P4();
TLorentzVector locPiMinusP4 = dPiMinusWrapper->Get_P4();
TLorentzVector locProtonP4 = dProtonWrapper->Get_P4();
TLorentzVector locPhoton1P4 = dPhoton1Wrapper->Get_P4();
TLorentzVector locPhoton2P4 = dPhoton2Wrapper->Get_P4();

// 3-Pion Final state based on kinematic fit quantities:
TLorentzVector ThreePiFinalState = locPiPlusP4 + locPiMinusP4 + locPhoton1P4 + locPhoton2P4;

// These quantities are the measured quantities
TLorentzVector locBeamP4_Measured = dComboBeamWrapper->Get_P4_Measured();
TLorentzVector locPiPlusP4_Measured = dPiPlusWrapper->Get_P4_Measured();
TLorentzVector locPiMinusP4_Measured = dPiMinusWrapper->Get_P4_Measured();
TLorentzVector locProtonP4_Measured = dProtonWrapper->Get_P4_Measured();
TLorentzVector locPhoton1P4_Measured = dPhoton1Wrapper->Get_P4_Measured();
TLorentzVector locPhoton2P4_Measured = dPhoton2Wrapper->Get_P4_Measured();

// 3-Pion Final state based on measured quantities:
```

```

TLorentzVector ThreePiFinalState_Measured = locPiPlusP4_M + locPiMinusP4_M + locPhoton1P4_M +
    locPhoton2P4_M;

// Get the shower quality factor of the first photon and find the z-position of the shower
float qf = dPhoton1Wrapper->Get_Shower_Quality();
TLorentzVector Gamma1_Shower = dPhoton1Wrapper->Get_X4_Shower();
float Shower1Zposition = Gamma1_Shower.Z();

```

## 2.4.2 Handling Accidental Beam Photon Contributions

The reconstructed final state particle combination is associated with the timing of a particular accelerator beam bunch (which we will call the “RF time”). The measured initial state beam photon time will either be consistent, or “in-time”, with this selected beam bunch, or “out-of-time” with it. While signal events are generally always associated with in-time beam photons, there is also an in-time contribution due to combinations where the beam photon is not the real photon which created the reconstructed final state particles (for example, due to multiple in-time photons or due to the correct photon not being reconstructed). There are several methods which can be used to subtract this background contribution, the most common of which is to estimate it using out-of-time beam photon combos and subtract the background by filling histograms with appropriate weights, or propagating these weights to later analysis stages.

In this method, in-time beam photons are assigned a weight of 1, while out-of-time beam photons are assigned a weight of  $1/n$ , where  $n$  is the number of out-of-time beam bunches that are considered in the analysis. The RF time is determined from the combo using the `dComboBeamWrapper()->Get_RFTime()` method. The following code shows an example of how to determine the RF time of the beam photon used in a given combo and how to define the weight for the combo based on the RF time of the beam photon. In this particular example it is expected that 4 beam bunches on either side of the in-time beam bunch (at  $t = 0$ ) are available leading to a weight factor of  $-1/8$ . An additional run-dependent multiplicative factor should be applied, to correct for non-uniformities in the out-of-time peaks, which can be accessed through `DAnalysisUtilities::Get_AccidentalScalingFactor()`.

```

TLorentzVector locVertex = dComboBeamWrapper->Get_X4_Measured();
float locRFTime = dComboWrapper->Get_RFTime();
TLorentzVector locBeamP4 = dComboBeamWrapper->Get_P4();
.....
float DT_RF = locVertex.T() - (locRFTime + (locVertex.Z() - dTargetCenter.Z())/29.9792458);

// Now an event weight can be assigned:
//   if DT_RF = +/- 2ns within zero the beam photon is in time
//   within +-4, +-8, +-12, ... the beam photon is out of time
double weight = 1.;
if (TMath::Abs(DT_RF)>2.){ // TOFIX: should actually pull the real beam bunch spacing for this
    weight = -0.125;      // -1/8
    weight *= dAnalysisUtilities->Get_AccidentalScalingFactor(); // correct for correlation
    effects
}

// the weight then can be used to fill a histogram with the weight
// thereby automatically subtract accidental background
// for example, dEBeam->Fill(locBeamP4.E(), weight);

```

## 2.4.3 Kinematic Fit Quality

The *ReactionFilter* ROOT trees are generated with no constraint on the quality of the kinematic fit by default. This quantity can be very powerful for separating signal from background. This information can be accessed in one of two ways: by looking at the  $\chi^2$  of the kinematic fit (suggested), which should have some distribution peaking at 1 (when normalized by the number of degrees of freedom in the fit) and falling towards higher values for the signal, or the confidence level of the fit, which should have a roughly flat distribution of the signal, with the background piling up around 0. There is no preferred cut value for a general case; you

should choose a value by studying these distributions for your signal and some background contribution that you would like to reject. This information can be accessed in the following way:

```
// Kinematic fit confidence level and chi^2/d.o.f.
double chi2dof = dComboWrapper->Get_ConfidenceLevel_KinFit("") /
    dComboWrapper->Get_NDF_KinFit("");
double chi2dof = dComboWrapper->Get_ChiSq_KinFit("") / dComboWrapper->Get_NDF_KinFit("");
// reject events with a poor chi^2 for this reaction
if(chi2dof > 5.) {
    dComboWrapper->Set_IsComboCut(true); // reject this combo for future consideration
    continue;
}
```

#### 2.4.4 Long-Lived Particles

As previously discussed, particle combinations can have multiple steps in order to model the effect of additional decays of narrow resonances. These decaying particles can be short-lived ( $\omega, \eta, \dots$ ) or long-lived ( $K_S, \Lambda, \dots$ ). In the case of long-lived particles, additional event selections can be made to separate them from combinatorial background. The most accurate information on the displacement of the secondary vertex from the primary vertex is obtained from the kinematic fit, particularly when a vertex constraint is included. Two common variables to cut on are the magnitude of this displacement, or “flight distance”, and this magnitude divided by the uncertainty in the position of these vertices, or “flight significance”. Below, we give an example for the reaction  $\gamma p \rightarrow K^+ K^+ \Xi^-$  (1320), which has two detached vertices:  $\Xi^- \rightarrow \pi^- \Lambda^0$ ,  $\Lambda^0 \rightarrow p \pi^-$ .

The cases in which the decaying particle is and is not mass constrained in the fit are handled differently. If no mass constraint is applied in the fit, first obtain the production X4 (vertex spatial position and time) and the X4 for the decaying particle. The pathlength of the decaying particle is the magnitude of the difference between the two vertices. The uncertainty of the pathlength is saved directly to the tree, but must be extracted by hand. The flight significance is then the pathlength divided by the uncertainty in the pathlength.

```
TLorentzVector locProdSpacetimeVertex = dComboBeamWrapper->Get_X4();
TLorentzVector locDecayingXiX4 =
    dTreeInterface->Get_TObject<TLorentzVector>("DecayingXiMinus_X4", loc_i); // change if the
    mass is constrained in the fit
TLorentzVector locDeltaSpacetimeXi = locProdSpacetimeVertex - locDecayingXiX4;
double locPathLengthXi = locDeltaSpacetimeXi.Vect().Mag();
float locPathLengthSigmaXi = Get_Fundamental<Float_t>("DecayingXiMinus_PathLengthSigma", loc_i);
double locPathLengthSignificanceXi = locPathLengthXi/locPathLengthSigmaXi;
```

If a mass constraint on the decaying particle was applied, then a wrapper for the decaying particle is configured and should be used to obtain the vertex position instead:

```
TLorentzVector locDecayingXiX4 = dDecayingXiMinusWrapper->Get_X4();
```

For a secondary detached vertex, for example the  $\Lambda$  decay in  $\Xi^- \rightarrow \Lambda \pi^-$ , the only difference in the calculation is to compare it to the parent detached vertex:

```
TLorentzVector locDecayingLambX4 =
    dTreeInterface->Get_TObject<TLorentzVector>("DecayingLambda_X4", loc_i);
TLorentzVector locDeltaSpacetimeLamb = locDecayingXiX4 - locDecayingLambX4;
double locPathLengthLamb = locDeltaSpacetimeLamb.Vect().Mag();
float locPathLengthSigmaLamb = Get_Fundamental<Float_t>("DecayingLambda_PathLengthSigma", loc_i);
double locPathLengthSignificanceLamb = locPathLengthLamb/locPathLengthSigmaLamb;
```

## 2.4.5 Analyzing Thrown Information

When analyzing simulated data (MC), information about the particles thrown by the physics event generator is also stored for each event. The data is stored in objects of type `DMCThrown` and is also accessible through a wrapper interface. This information can be accessed in several different ways. Each `DChargedTrackHypothesis` and `DNeutralParticleHypothesis` has a member function named `Get_ThrownIndex()`, which can be passed as an argument to `dThrownWrapper->Set_ArrayIndex()`. The list of thrown particles can also be studied directly, such as in the following example:

```
//Thrown beam: just use directly
if(dThrownBeam != NULL)
    double locEnergy = dThrownBeam->Get_P4().E();

//Loop over throws
for(UInt_t loc_i = 0; loc_i < Get_NumThrown(); ++loc_i)
{
    //Set branch array indices corresponding to this particle
    dThrownWrapper->Set_ArrayIndex(loc_i);

    //Do stuff with the wrapper here ...
    //Example - fill a histogram of the momentum distribution of protons
    if(dThrownWrapper->Get_PID() == Proton)
        dProtonPmag->Fill(dThrownWrapper->Get_P4().Vect().Mag());
}
```

## 2.5 ROOT Tree Output

The *DSelector* can be used to create TTrees for further analysis, either by another *DSelector* or by another program. Two types of TTree output are support: the standard *ReactionFilter* output in PART format, which stores data mostly in various arrays of objects, and a “flat” tree with a simpler structure.

The code to handle ROOT tree output is in several locations, and generally works through the (thread-safe) `DTreeInterface` class. In the `Init()` function, besides defining the names of output files, you can optionally add branches to store additional event/combo information. The following shows an example of these features:

```
// In Init() function...

//USERS: SET OUTPUT FILE NAME //can be overridden by user in PROOF
dOutputFileName = "ggg.root"; //" for none
dOutputTreeFileName = ""; //" for none
dFlatTreeFileName = ""; //output flat tree (one combo per tree entry), "" for none
dFlatTreeName = ""; //if blank, default name will be chosen

/***** EXAMPLE USER INITIALIZATION: CUSTOM OUTPUT BRANCHES - MAIN TREE
*****/

//EXAMPLE MAIN TREE CUSTOM BRANCHES (OUTPUT ROOT FILE NAME MUST FIRST BE GIVEN!!!! (ABOVE: TOP)):
//The type for the branch must be included in the brackets
//1st function argument is the name of the branch
//2nd function argument is the name of the branch that contains the size of the array (for
    fundamentals only)
dTreeInterface->Create_Branch_Fundamental<Int_t>("my_int"); //fundamental = char, int, float,
    double, etc.
dTreeInterface->Create_Branch_FundamentalArray<Int_t>("my_int_array", "my_int");
dTreeInterface->Create_Branch_FundamentalArray<Float_t>("my_combo_array", "NumCombos");
dTreeInterface->Create_Branch_NoSplitTObject<TLorentzVector>("my_p4");
dTreeInterface->Create_Branch_ClonesArray<TLorentzVector>("my_p4_array");

/***** EXAMPLE USER INITIALIZATION: CUSTOM OUTPUT BRANCHES - FLAT TREE
```

```

*****/

//EXAMPLE FLAT TREE CUSTOM BRANCHES (OUTPUT ROOT FILE NAME MUST FIRST BE GIVEN!!!! (ABOVE: TOP)):
//The type for the branch must be included in the brackets
//1st function argument is the name of the branch
//2nd function argument is the name of the branch that contains the size of the array (for
    fundamentals only)
dFlatTreeInterface->Create_Branch_Fundamental<Int_t>("flat_my_int"); //fundamental = char, int,
    float, double, etc.
dFlatTreeInterface->Create_Branch_FundamentalArray<Int_t>("flat_my_int_array", "flat_my_int");
dFlatTreeInterface->Create_Branch_NoSplitTObject<TLorentzVector>("flat_my_p4");
dFlatTreeInterface->Create_Branch_ClonesArray<TLorentzVector>("flat_my_p4_array");

```

In the Process() function, these trees can be filled either once per event (i.e. before the combo loop)

```

/***** EXAMPLE: FILL CUSTOM OUTPUT BRANCHES
*****/
Int_t locMyInt = 7;
dTreeInterface->Fill_Fundamental<Int_t>("my_int", locMyInt);

TLorentzVector locMyP4(4.0, 3.0, 2.0, 1.0);
dTreeInterface->Fill_TObject<TLorentzVector>("my_p4", locMyP4);

for(int loc_i = 0; loc_i < locMyInt; ++loc_i)
    dTreeInterface->Fill_Fundamental<Int_t>("my_int_array", 3*loc_i, loc_i); //2nd argument =
        value, 3rd = array index

```

or once per combo (i.e. inside the combo loop)

```

/***** EXAMPLE: FILL CUSTOM OUTPUT BRANCHES
*****/
TLorentzVector locMyComboP4(8.0, 7.0, 6.0, 5.0);
//for arrays below: 2nd argument is value, 3rd is array index
//NOTE: By filling here, AFTER the cuts above, some indices won't be updated (and will be
    whatever they were from the last event)
    //So, when you draw the branch, be sure to cut on "IsComboCut" to avoid these.
dTreeInterface->Fill_Fundamental<Float_t>("my_combo_array", -2*loc_i, loc_i);
dTreeInterface->Fill_TObject<TLorentzVector>("my_p4_array", locMyComboP4, loc_i);

```

For the normal tree output, the tree should be written out at the end of Process():

```

/***** EXAMPLE: FILL CLONE OF TTREE HERE WITH CUTS APPLIED
*****/
Bool_t locIsEventCut = true;
for(UInt_t loc_i = 0; loc_i < Get_NumCombos(); ++loc_i) {
    //Set branch array indices for combo and all combo particles
    dComboWrapper->Set_ComboIndex(loc_i);
    // Is used to indicate when combos have been cut
    if(dComboWrapper->Get_IsComboCut())
        continue;
    locIsEventCut = false; // At least one combo succeeded
    break;
}
if(!locIsEventCut && dOutputTreeFileName != "")
    Fill_OutputTree();

```

For a flat tree, this format has one entry per combo, and so it should be filled and written out right at the end of the combo loop, for example:

```

/***** FILL FLAT TREE (IF DESIRED)

```



```

*****/
//FILL ANY CUSTOM BRANCHES FIRST!!
Int_t locMyInt_Flat = 7;
dFlatTreeInterface->Fill_Fundamental<Int_t>("flat_my_int", locMyInt_Flat);

TLorentzVector locMyP4_Flat(4.0, 3.0, 2.0, 1.0);
dFlatTreeInterface->Fill_TObject<TLorentzVector>("flat_my_p4", locMyP4_Flat);

for(int loc_j = 0; loc_j < locMyInt_Flat; ++loc_j)
{
    dFlatTreeInterface->Fill_Fundamental<Int_t>("flat_my_int_array", 3*loc_j, loc_j); //2nd
        argument = value, 3rd = array index
    TLorentzVector locMyComboP4_Flat(8.0, 7.0, 6.0, 5.0);
    dFlatTreeInterface->Fill_TObject<TLorentzVector>("flat_my_p4_array", locMyComboP4_Flat,
        loc_j);
}

//FILL FLAT TREE
Fill_FlatTree(); //for the active combo

```

### 3 DSelector Data Classes

All the quantities in the ROOT file are accessed through some wrapper objects defined in your `DSelector`. The variable names used in the ROOT trees and the generated `DSelector` code have distinct qualifiers to help to indicate their meaning and are explained here in more detail.

#### Variable names prefixes and suffixes

`Beam` Beam photon related quantity/ies for a given event before Kinematic Fit

`NeutralHypo` Neutral Particle related quantity/ies for a given event before Kinematic Fit

`ChargedHypo` Charged Particle related quantity/ies for a given event before Kinematic Fit

`_Measured` A measured quantity as determined by the reconstruction software

`_KinFit` A measured quantity that has been altered by the Kinematic Fitter.

`_X4` 4-vector containing position and time (`TLorentzVector`).

`_P4` 4-vector containing momentum and energy (`TLorentzVector`).

`BeamCombo` Initial state beam photon for a given combo.

`Proton` Final state proton for a given combo.

`PiPlus` Final state  $\pi^+$  for a given combo.

`PiMinus` Final state  $\pi^-$  for a given combo.

`Photon1` First final state photon for a given combo. Subsequent photons are labeled `Photon2`, `Photon3`, ...

`PiZero` Final state  $\pi^0$  for a given combo if the mass was constrained in by the kinematic fitter. Other particles, such as `Eta`, `Kshort`, and `Lambda` can also be specified.

Note, in the above example the  $\pi^0$  is not a final state particle, as it does decay into photons and only those are detected directly. Therefore, such a particle is only listed explicitly as a separate particle in the ROOT tree if its mass was constrained by the kinematic fit to a fixed value, usually the PDG value.

In order to access the variables in the tree, one must use the above-mentioned wrappers and their methods. The detailed functionality and implementations of all these wrapper methods can be found in `DBeamParticle.h`, `DChargedTrackHypothesis.h`, `DNeutralParticleHypothesis.h`, and `DParticleCombo.h` located in `glux_root_analysis` library. These methods are discussed in more detail below.



### 3.0.1 DBeamParticle.h

Most of the methods needed by the incoming photon are inherited from `DKinematicData.h`

#### PID & KINEMATIC DATA

`Get_PID()` Returns type `Particle_t` defined in `halld_recon/src/libraries/include/particleType.h`

`Get_P4()` Get the 4-momentum of this particle

`Get_X4()` Get the 4-vector position (time and position) of this particle at the production vertex from the kinematic fit

`Get_P4_Measured()` Same as above

`Get_X4_Measured()` Same as above but the measured quantities, see example code above using the wrapper `dComboBeamWrapper` to get the vertex 4-vector.

Note that since the kinematic fitter does not change the 4-momentum of the beam photon `_P4()` and `_P4_Measured()` are the same in for the incident beam particle! The vertex position still can change though.

### 3.0.2 DChargedTrackHypothesis.h

Note that this class does inherit from `DKinematicData.h` and as such has the above listed methods as well as the following ones:

#### IDENTIFIERS

`Get_TrackID()` Each physical particle has its own identifier but is fit by multiple mass hypotheses, refers back to the parent particle, returns int.

`Get_ThrownIndex()` Array index of thrown particle if match found, returns int (-1 otherwise). MC data only.

`Get_ID()` Same as `Get_TrackID()`

#### TRACKING INFO

`Get_ChiSq_Tracking()` The  $\chi^2$  from the charged particle fit (Kalman filter), returns float

`Get_NDF_Tracking()` The number of degrees of freedom in the fit, equal to the number of hits - 5, returns uint

`Get_ChiSq_DCdEdx()` The  $\chi^2$  between the measured energy loss in the drift chamber and the value expected for the given mass hypotheses, where the most appropriate value between the CDC and FDC  $dE/dx$ 's is used, returns float

`Get_NDF_DCdEdx()` The number of degrees of freedom in the fit, returns uint

`Get_dEdx_CDC()` The calculated energy lose in the CDC, returns float

`Get_dEdx_FDC()` The calculated energy loss in the FDC, returns float

#### TIMING INFO

`Get_HitTime()` The time of the hit in whatever timing detector the charged particle is matched to, returns float

`Get_RFDeltaTVar()` The variance of the  $\Delta t_{RF}$  variable using the above hit time, returns float

`Get_Beta_Timing()` The calculated  $\beta$  from the above hit time, returns float

`Get_ChiSq_Timing()` The corresponding  $\chi^2$  from the above hit time for the given mass hypothesis, returns float

`Get_NDF_Timing()` The number of degrees of freedom used in this  $\chi^2$ , returns uint

`Get_ConfidenceLevel_Timing()` The corresponding confidence level, returns float

`Get_Beta_Timing_Measured()` The same as above, but using the measured momentum, returns float

`Get_ChiSq_Timing_Measured()` The same as above, but using the measured momentum, returns float  
`idenceLevel_Timing_Measured()` The same as above, but using the measured momentum, returns float  
`Get_Detector_System_Timing()` The detector in which the matched timing hit was measured (BCAL/FCAL/SC/TOF), returns `DetectorSystem_t`  
 HIT ENERGY  
`Get_dEdx_TOF()` Measured energy loss in the TOF paddles, returns float  
`Get_dEdx_ST()` Measured energy loss in the Start Counter paddles, returns float  
`Get_Energy_BCAL()` Energy of a matched BCAL shower, returns float  
`Get_Energy_BCALPreshower()` Energy deposited in the first layer of a matched BCAL shower, returns float  
`Get_Energy_BCALLayer2()` Energy deposited in the second layer of a matched BCAL shower, returns float  
`Get_Energy_BCALLayer3()` Energy deposited in the third layer of a matched BCAL shower, returns float  
`Get_Energy_BCALLayer4()` Energy deposited in the fourth layer of a matched BCAL shower, returns float  
`Get_SigLong_BCAL()` The r.m.s. deviation (?) of the BCAL shower energy in the longitudinal direction, returns float  
`Get_SigTheta_BCAL()` The r.m.s. deviation (?) of the BCAL shower energy in the polar angle, returns float  
`Get_SigTrans_BCAL()` The r.m.s. deviation (?) of the BCAL shower energy in the transverse direction, returns float  
`Get_RMSTime_BCAL()` The r.m.s. deviation (?) of the BCAL shower time, returns float  
`Get_Energy_FCAL()` Energy of a matched FCAL shower, returns float  
`Get_E1E9_FCAL()` The  $E1/E9$  ratio for the matched FCAL shower, returns float  
`Get_E9E25_FCAL()` The  $E9/E25$  ratio for the matched FCAL shower, returns float  
`Get_SumU_FCAL()` The SumU value for the matched FCAL shower, returns float  
`Get_SumV_FCAL()` The SumV value for the matched FCAL shower, returns float  
 SHOWER MATCH  
`Get_TrackBCAL_DeltaPhi()` Difference in azimuthal positions of the shower and matched track along the face of the BCAL, returns float, 999.0 if not matched, units are radians  
`Get_TrackBCAL_DeltaZ()` Difference in z positions of the shower and matched track along the face of the BCAL, returns float, 999.0 if not matched  
`Get_TrackFCAL_DOCA()` Difference in positions of the shower and matched track along the face of the FCAL, returns float, 999.0 if not matched  
 DIRC INFO  
`Get_Track_NumPhotons_DIRC()` Number of DIRC photons associated with the track, returns float, 999.0 if no DIRC info  
`Get_Track_ThetaC_DIRC()` Measured Cherenkov angle, returns float, 999.0 if no DIRC info  
`Get_Track_Lele_DIRC()` Measured Likelihood for the electron hypothesis, returns float, 999.0 if no info  
`Get_Track_Lpi_DIRC()` Measured Likelihood for the pion hypothesis, returns float, 999.0 if no info  
`Get_Track_Lk_DIRC()` Measured Likelihood for the kaon hypothesis, returns float, 999.0 if no info  
`Get_Track_Lp_DIRC()` Measured Likelihood for the proton hypothesis, returns float, 999.0 if no info

### 3.0.3 DNeutralParticleHypothesis.h

Note that this class does inherit from DKinematicData.h and as such has those methods as well as the following ones:

#### IDENTIFIERS

`Get_NeutralID()` Each physical particle has its own number, returns int.

`Get_ThrownIndex()` Array index of thrown particle if match found, returns int (-1 otherwise). MC data only.

`Get_ID()` Same as `Get_NeutralID()`

#### TIMING

`Get_HitTime()` The reconstructed shower time, returns float

`Get_RFDeltaTVar()` The variance of the  $\Delta t_{RF}$  variable using the above shower time, returns float

`Get_Beta_Timing()` The calculated  $\beta$  from the above hit time, returns float

`Get_ChiSq_Timing()` The corresponding  $\chi^2$  from the above hit time for the given mass hypothesis, returns float

`Get_NDF_Timing()` The number of degrees of freedom used in this  $\chi^2$ , returns uint

`Get_ConfidenceLevel_Timing()` The corresponding confidence level, returns float

`Get_Beta_Timing_Measured()` The same as above, but using the measured momentum, returns float

`Get_ChiSq_Timing_Measured()` The same as above, but using the measured momentum, returns float

`Get_ConfidenceLevel_Timing_Measured()` The same as above, but using the measured momentum, returns float

`Get_Detector_System_Timing()` The detector in which the matched timing hit was measured (BCAL/FCAL/SC/TOF), returns `DetectorSystem_t`

#### SHOWER INFO

`Get_Energy_BCAL()` Energy of the BCAL shower, returns float

`Get_Energy_BCALPreshower()` Energy deposited in the first layer of the BCAL shower, returns float

`Get_Energy_BCALLayer2()` Energy deposited in the second layer of the BCAL shower, returns float

`Get_Energy_BCALLayer3()` Energy deposited in the third layer of the BCAL shower, returns float

`Get_Energy_BCALLayer4()` Energy deposited in the fourth layer of the BCAL shower, returns float

`Get_SigLong_BCAL()` The r.m.s. deviation (?) of the BCAL shower energy in the longitudinal direction, returns float

`Get_SigTheta_BCAL()` The r.m.s. deviation (?) of the BCAL shower energy in the polar angle, returns float

`Get_SigTrans_BCAL()` The r.m.s. deviation (?) of the BCAL shower energy in the transverse direction, returns float

`Get_RMSTime_BCAL()` The r.m.s. deviation (?) of the BCAL shower time, returns float

`Get_Energy_FCAL()` Energy of a FCAL shower, returns float

`Get_E1E9_FCAL()` The  $E1/E9$  ratio for the FCAL shower, returns float

`Get_E9E25_FCAL()` The  $E9/E25$  ratio for the FCAL shower, returns float

`Get_SumU_FCAL()` The SumU value for the FCAL shower, returns float

`Get_SumV_FCAL()` The SumV value for the FCAL shower, returns float

#### SHOWER MATCH

- `Get_TrackBCAL_DeltaPhi()` Difference in azimuthal positions of the shower and matched track along the face of the BCAL, returns float, 999.0 if not matched, units are radians
- `Get_TrackBCAL_DeltaZ()` Difference in z positions of the shower and matched track along the face of the BCAL, returns float, 999.0 if not matched
- `Get_TrackFCAL_DOCA()` Difference in positions of the shower and matched track along the face of the FCAL, returns float, 999.0 if not matched

As an important reminder note that in order to make the `Energy_BCALayerX` data available in the root tree the reaction filter requires to be run with the following command line parameter:

`-PANALYSIS:BCAL_VERBOSE_ROOT_OUTPUT=1`

and similarly for FCAL if the `E1E9`, `E9E25`, `SumU` and `SumV` data should be in the root tree the following command line parameter is required on the command line when running the reaction filter:

`-PANALYSIS:FCAL_VERBOSE_ROOT_OUTPUT=1`

### 3.0.4 DParticleCombo.h

#### SHOWER MATCH

`Get_RFTime_Measured()` RF bucket time for the given combination, measured at the center of the target, returns float.

`Get_RFTime()` Same as above, returns float.

#### TARGET

`Get_TargetPID()` Particle type of target, returns `Particle_t`.

`Get_TargetCenter()` Position of target center, returns `TVector3`.

#### KINFIT

`Get_ChiSq_KinFit()`  $\chi^2$  of kinematic fit of this combo to the assumed reaction, returns float.

`Get_NDF_KinFit()` Number of degrees of freedom for the above kinematic fit, returns float.

`Get_ConfidenceLevel_KinFit()` Confidence level determined from  $\chi^2$  and number of degrees of freedom, returns float.

#### UNUSED ENERGY

`Get_Energy_UnusedShowers()` Total energy of unmatched showers that are in time with the combo, but are not part of the combo, returns float.

#### UNUSED TRACKS

`Get_NumUnusedTracks()` Number of charged particle tracks in the event and not used in the combo, returns `UChar_t`.

`Get_NumUnusedShowers()` Number of unmatched showers in the event and not used in the combo, returns `UChar_t`.

#### EVENT INFO

`Get_RunNumber()` Run number of the event, returns `UInt_t`.

`Get_EventNumber()` Event number of the event, returns `ULong64_t`.

`Get_L1TriggerBits()` Trigger bit information, returns `UInt_t`.

`Get_MCWeight()` Optional event weight factor for MC events, returns float. =

### 3.0.5 DParticleComboStep.h

`Get_X4()` Vertex position and time for this step of the reaction, returns `TLorentzVector`.