

# News on Rho

PANDA Collaboration Meeting , Bochum

11. 9. 2013

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# Major Changes (by Ralf Kliemt)

- Rename all [Rho classes](#) to [Rho\\*](#)
- [RhoCandidate](#) handled fully with pointers
- [Unclutter Rho classes](#) → delete obsolete/unnecessary stuff
- Remove unnecessary virtual or semivirtual layers
- [Structure fitter interfaces](#)
- Put data objects from pid to pnddata & rename
- Reorganize PndAnalysis
- [New Tutorial + Rho class Docu in PANDA Wiki](#)

# New Tutorial – August 2013

- Available in PANDA-Wiki
- Supposed to be all-time-running tutorial

[Edit](#) [Attach](#) [Printable](#)

[Computing.PandaRootRhoTutorial](#) r1.15 - (

## Simulation and Analysis in PandaRoot with RHO (Updated: Sep. 4, 2013; Tested with rev 21585)

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### Preface/Requirements

This tutorial aims to demonstrate, how simulation and analysis can be performed with the [PandaRoot](#) framework. The example channel is

*p pbar -> Psi(2S)-> J/Psi (-> mu+mu-) pi+ pi-*

The analysis part will cover

- Data
  - Partic
  - Comb
- <http://panda-wiki.gsi.de/cgi-bin/view/Computing/PandaRootRhoTutorial>

# ... and additional Documentation!

Edit Attach Printable

[Computing.PandaRootAnalysisJuly13](#)

## Documentation of Rho Classes

### Index

Here you can find information about:

- [PndAnalysis](#) - [Interface](#) - [Lists and Keys](#) - [PID Algo Names](#) - [MC Truth Match](#)
- [RhoCandidate](#) - [Interface](#)
- [RhoCandList](#) - [Interface](#)
- [Particle Selectors](#) - [Kinematic Selectors](#) - [PID Selectors](#)
- [Fitters](#) - [Vertex Fitting](#) - [4C Fitting](#) - [Kinematic Fitting](#)

A tutorial for Rho with an example simulation and analysis can be found [on this Wiki page](#).

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# PndAnalysis: Data Access/PID

- Data access via PndAnalysis

```
FairRunAna *fRun = new FairRunAna();
fRun->SetInputFile("pid_complete.root");

PndAnalysis *ana = new PndAnalysis();
RhoCandList eplus, eminus, muplus, muminus, mct;
TString myPidAlgosElectron = "PidAlgoEmcBayes;PidAlgoDrc"
TString myPidAlgosMuon     = "PidAlgoMdtHardCuts"

...
while ( ana->GetEvent() )
{
    ana->FillList( eplus,    "ElectronTightPlus",    myPidAlgosElectron );
    ana->FillList( eminus,   "ElectronTightMinus",   myPidAlgosElectron );
    ana->FillList( muplus,   "MuonTightPlus",        myPidAlgosMuon );
    ana->FillList( muminus,  "MuonTightMinus",       myPidAlgosMuon );
    ana->FillList( mct,      "McTruth" );
    ...
}
```

*With standard pid macros  
only pid file needed as input,  
also including MC info now!*

- Features:

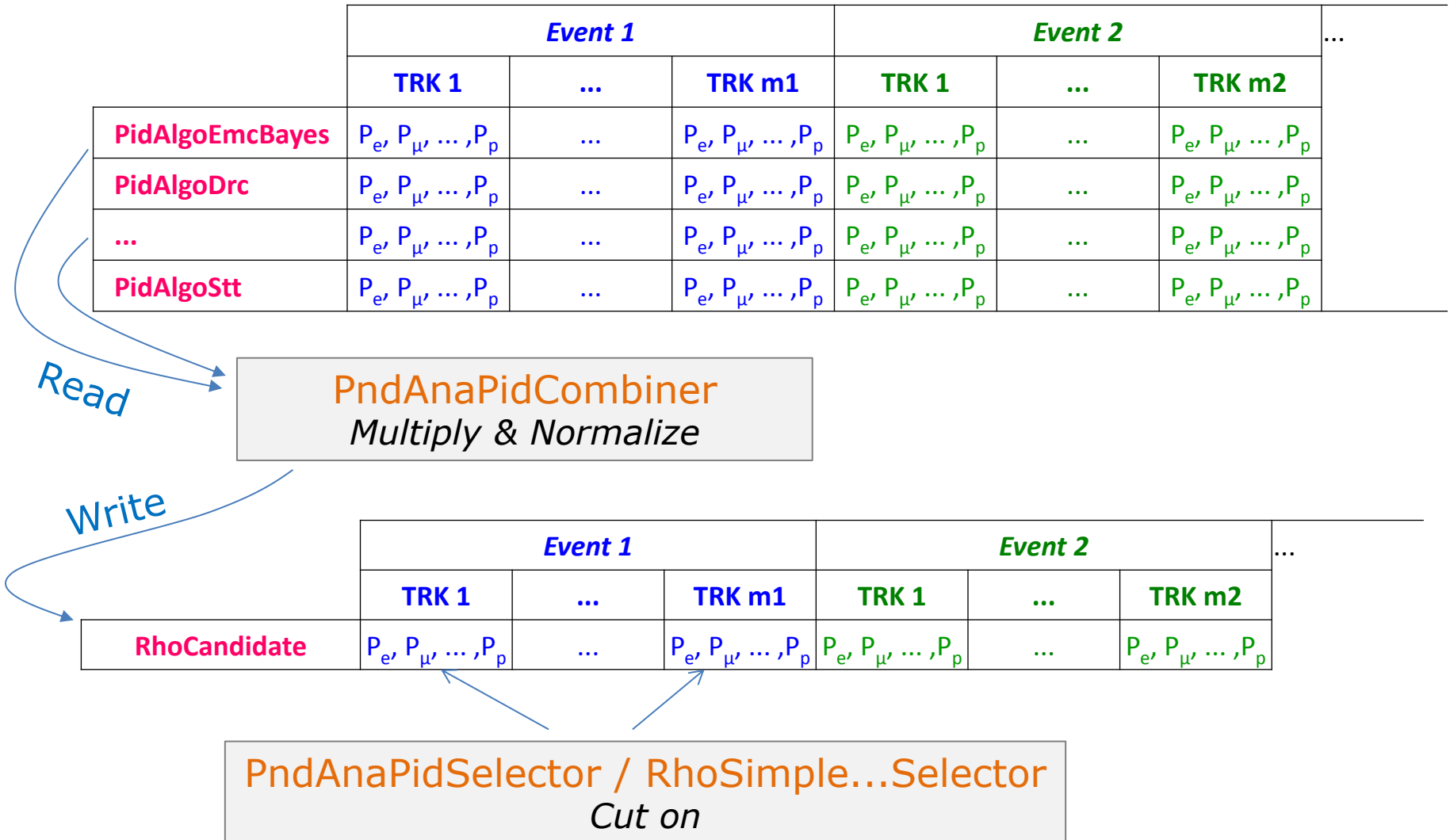
- Simple access to reco candidates and McTruth objects
- Various PID algorithms directly accessible

# Particle Identification Concept

- PID probabilities are computed by various algorithms  
*PidAlgoEmcBayes*, *PidAlgoDrc*, *PidAlgoMvd*, ...
- *PndAnaPidCombiner*
  - combines *on demand* probabilities from various algo's by computing product of all  $P_i$  ( $i=algo's$ )
  - copy probabilities to RhoCandidate/RhoCandList
- *PndAnaPidSelector* / *RhoSimple...Selector*
  - selects particles based on these probabilities
- *PndAnalysis::FillList* is a short-cut to this functionality via

```
ana->FillList(..., "PidAlgoEmcBayes;PidAlgoDrc");
```

# Particle Identification Concept



# Stand-alone usage of PID-Selector

- Selector can be used independent of PndAnalysis::FillList
- Say, you have a list of charged particles and want to identify **loose kaons** with **PidAlgoDrc & PidAlgoMvd**

```
...
PndAnalysis *ana = new PndAnalysis();

RhoCandList charged, kaonLoose;

PndAnaPidSelector kaonSel("KaonSelector");
kaonSel.SetSelection("KaonLoose"); // set selection criterion

PndAnaPidCombiner pidComb("PidCombiner");
pidComb.SetTcaNames("PidAlgoDrc;PidAlgoMvd"); // set algo's

while ( ana->GetEvent() )
{
    ana->FillList(charged, "Charged"); // start w/ charged candidates

    pidComb.Apply(charged); // copy P to candidates
    kaonSelector.Select(charged, kaonLoose); // select kaons from charged
}
...
```

*Possible selection keywords are described in Rho class documentation wiki!*

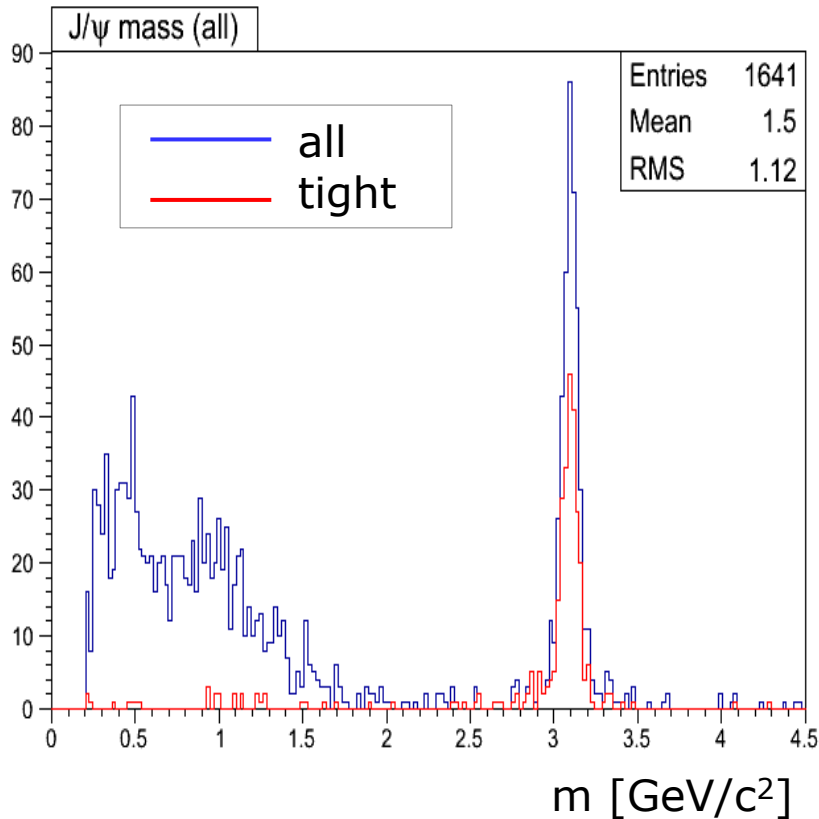


# PID: Additional Notes

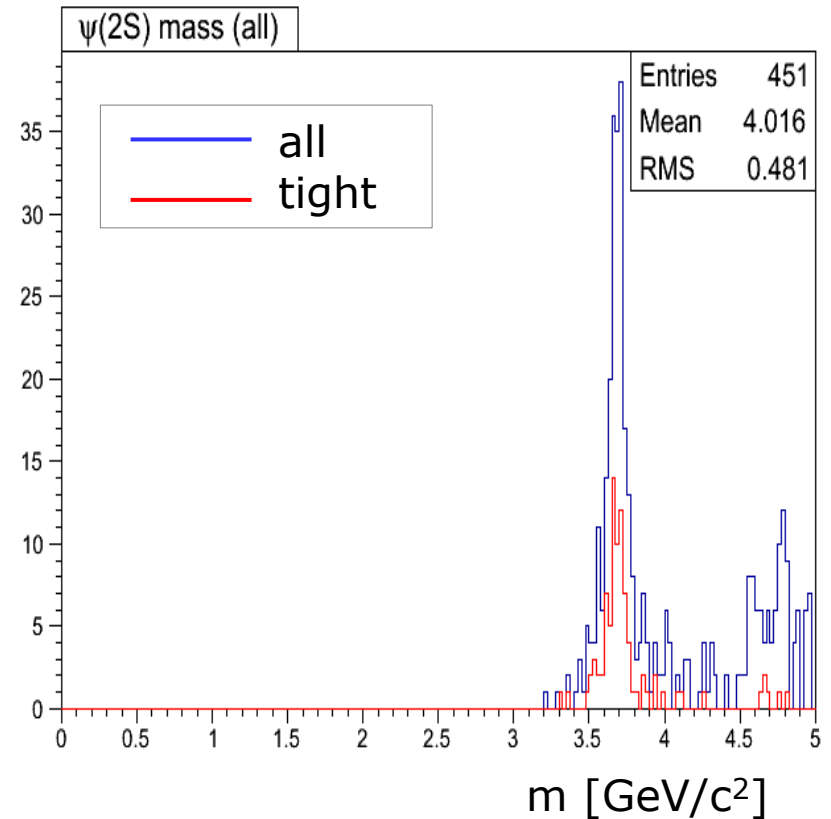
- There are some **default PID-criteria** (*[ANAPidSelections]* set in `ana.par` in `macro/params/`)
  - "VeryLoose" :  $P_i \geq 0$
  - "Loose" :  $P_i \geq 0.25$  (*was previously 0.2*)
  - "Tight" :  $P_i \geq 0.5$
  - "VeryTight" :  $P_i \geq 0.9$
  - "Variable" :  $P_i \geq 0.5$  (user value)
  - "Best" :  $P_i \geq P_j, \forall j \neq i$
- **Ideal PID algorithm:** `PidAlgoIdealCharged`
  - During reconstruction looks up true MC particle
  - Sets  $P = 1$  for correct species, for all other  $P = 0$
  - *Not equivalent to MC truth match!*

# Example: PID

$$J/\psi \rightarrow \mu^+\mu^-$$



$$\psi(2S) \rightarrow J/\psi \pi^+\pi^-$$



$\mu^\pm$ : "PidAlgoMdtHardCuts"

$\pi^\pm$ : "PidAlgoMvd;PidAlgoStt;PidAlgoDrc"

# MC Truth Access

- For final state particles via `RhoCandidate::GetMcTruth`

```
ana->FillList( eplus, "ElectronTightPlus", "PidAlgoEmcBayes" );  
RhoCandidate *truth = eplus[0]->GetMcTruth();
```

- The MC truth particles have the complete genealogy

```
if ( truth != 0x0 ) {  
    RhoCandidate *mother = truth->TheMother();  
    if ( truth->NDaughters() > 1 ) {  
        RhoCandidate *firstDaughter = truth->Daughter(0);  
        RhoCandidate *secondDaughter = truth->Daughter(1);  
    }  
}
```

- Also complete MC truth list available

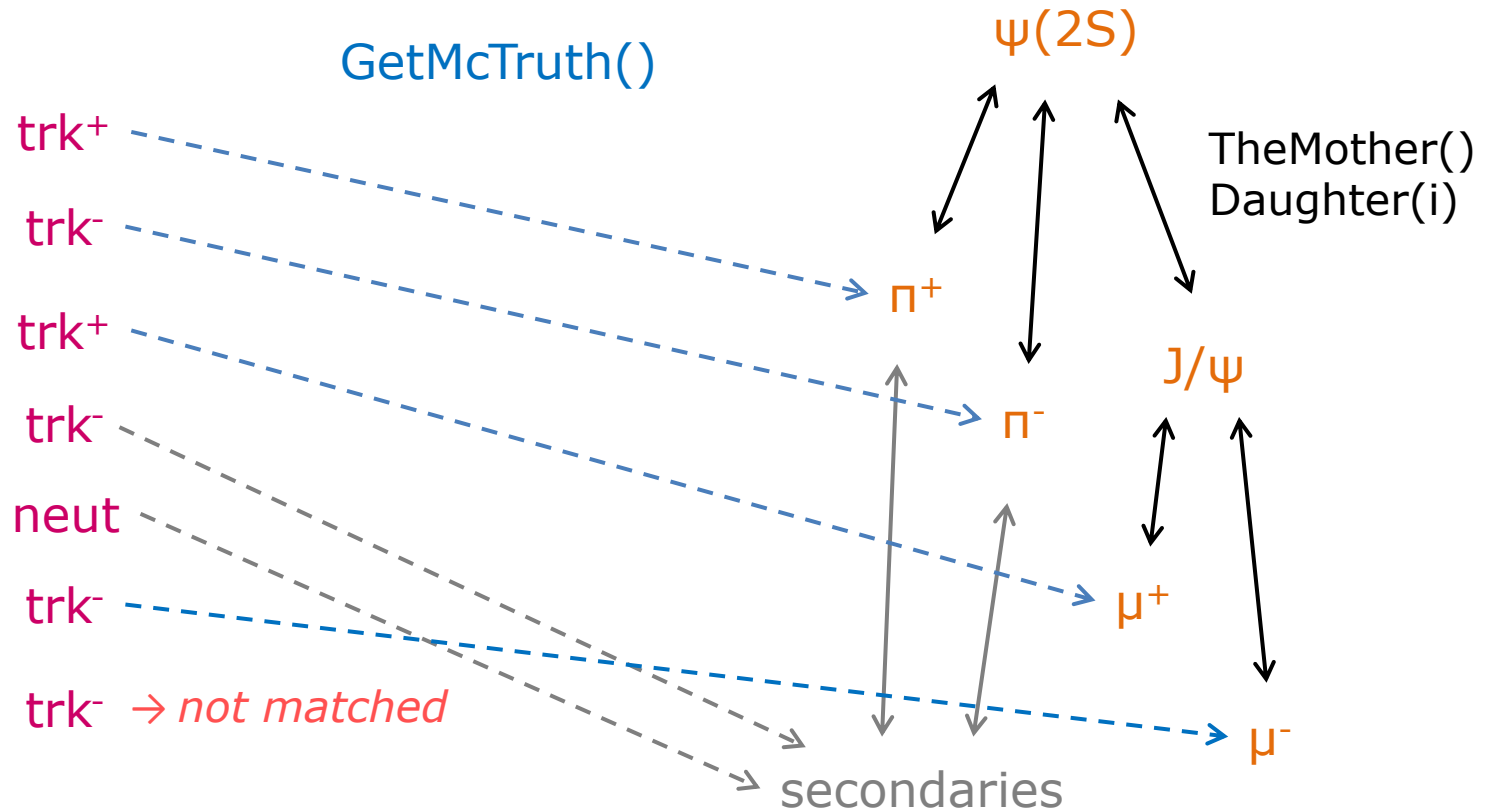
```
ana->FillList( mct, "McTruth" );  
...  
RhoCandidate *firstDaughter = mct[0]->Daughter(0);
```

# Mc Truth Genealogy

- Idea:** Each **Reco** points to an **McTruth** object, from which the full true tree can be accessed

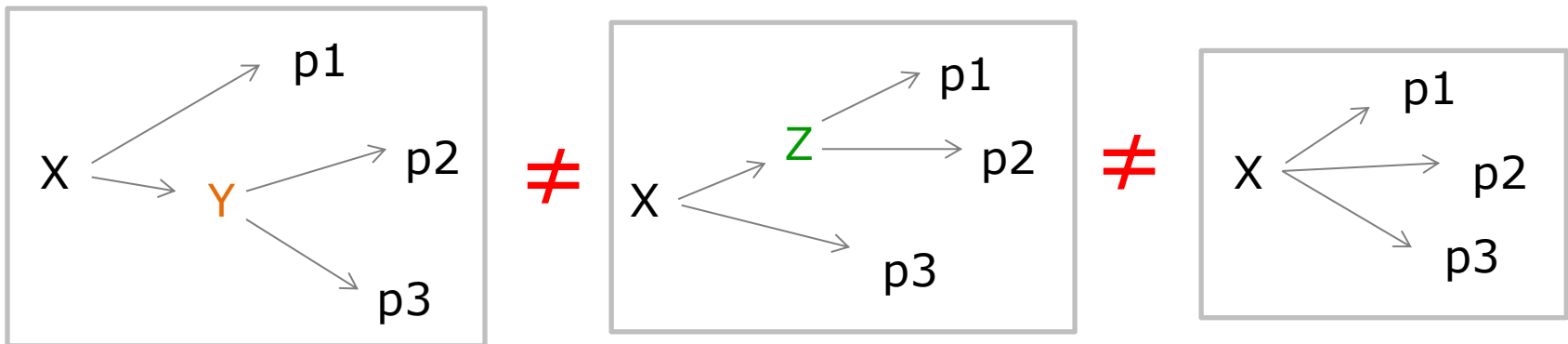
**All Reco**

**McTruth**



# PndAnalysis: MC Truth Match of Composites

- Physics analysis might require a full mc truth match for composite particles for efficiency calculation, because



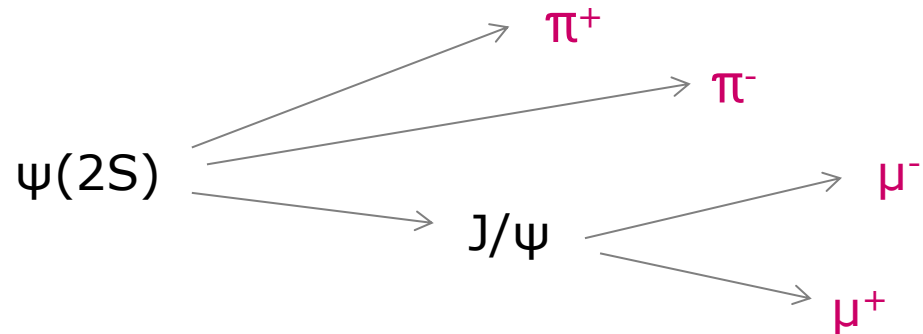
- Functionality, formerly hosted in PndMcTruthMatch now in

`PndAnalysis::McTruthMatch( RhoCandidate* );`

*Requires PndEvtGenDirect with SetStoreTree!*

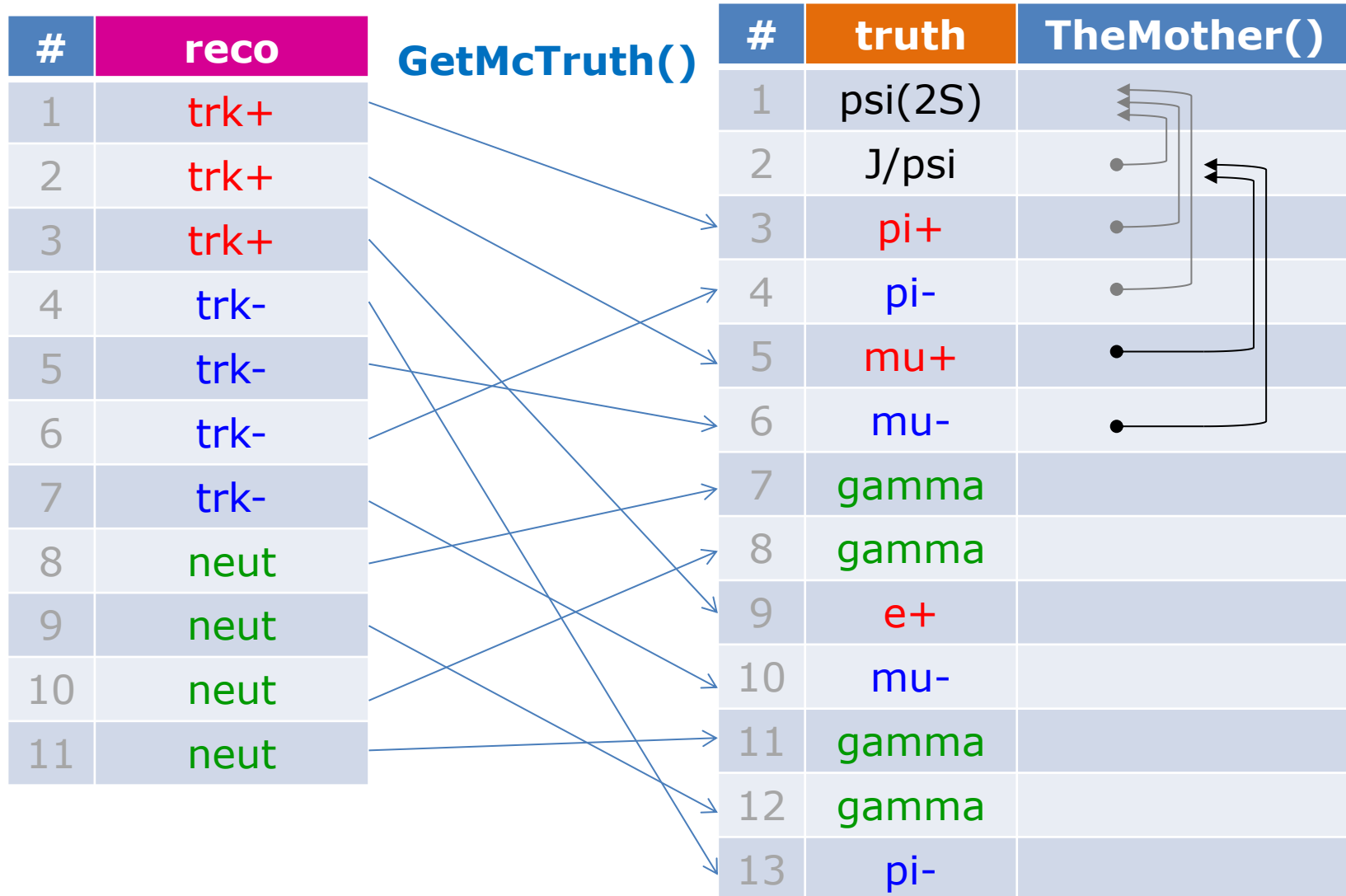
# Example: $\psi(2S) \rightarrow J/\psi (\mu^+\mu^-) \pi^+\pi^-$

- We want to reconstruct the following decay

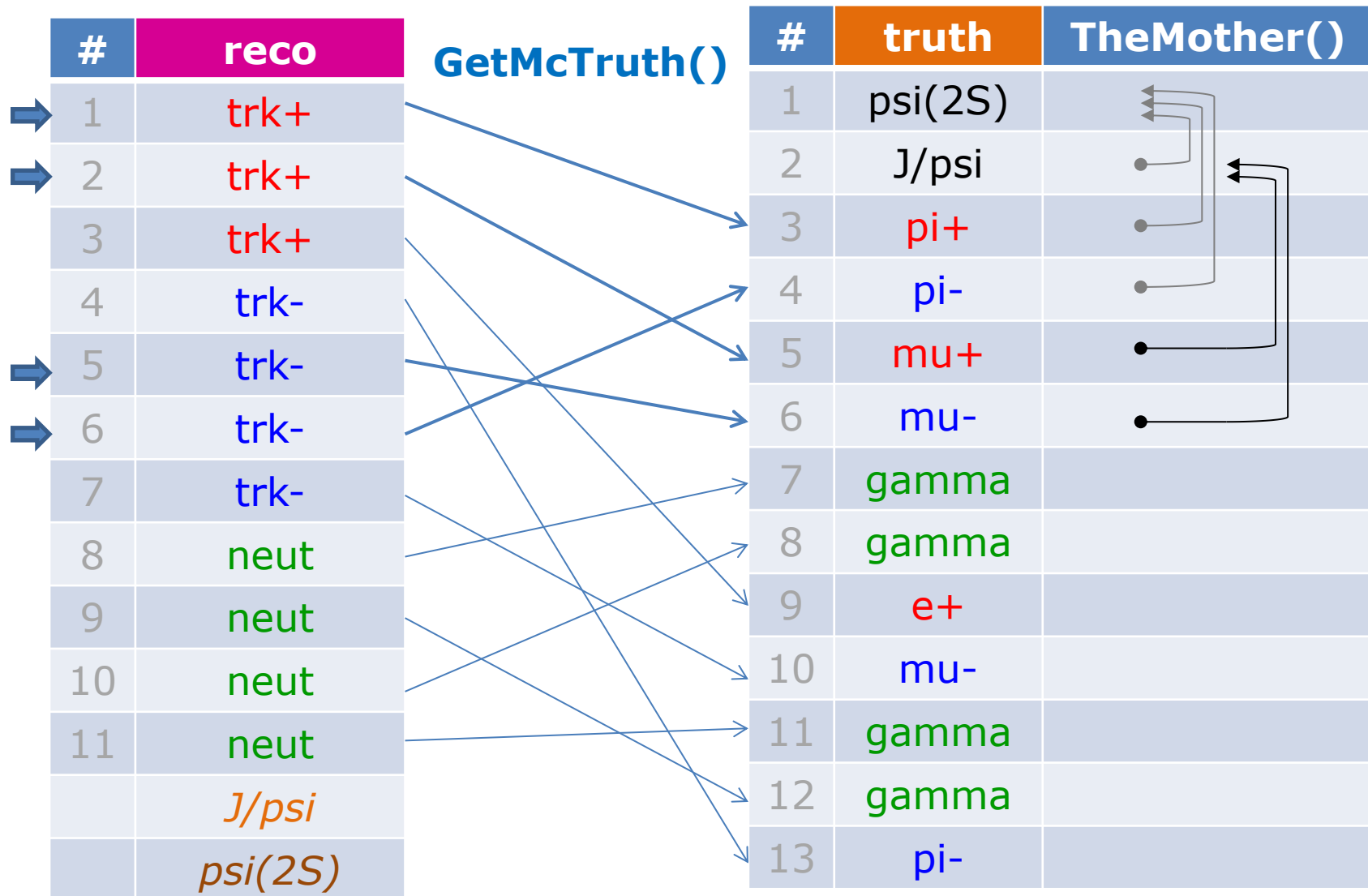


```
RhoCandList muplus, muminus, piplus, piminus, jpsi, psi2s;  
  
ana->FillList( muplus, "MuonLoosePlus", myPidAlgos );  
ana->FillList( muminus, "MuonLooseMinus", myPidAlgos );  
ana->FillList( piplus, "PionLoosePlus", myPidAlgos );  
ana->FillList( piminus, "PionLooseMinus", myPidAlgos );  
  
jpsi.Combine( muplus, muminus );  
psi2s.Combine( jpsi, piplus, piminus );
```

# Example: $\psi(2S) \rightarrow J/\psi (\mu^+\mu^-) \pi^+\pi^-$

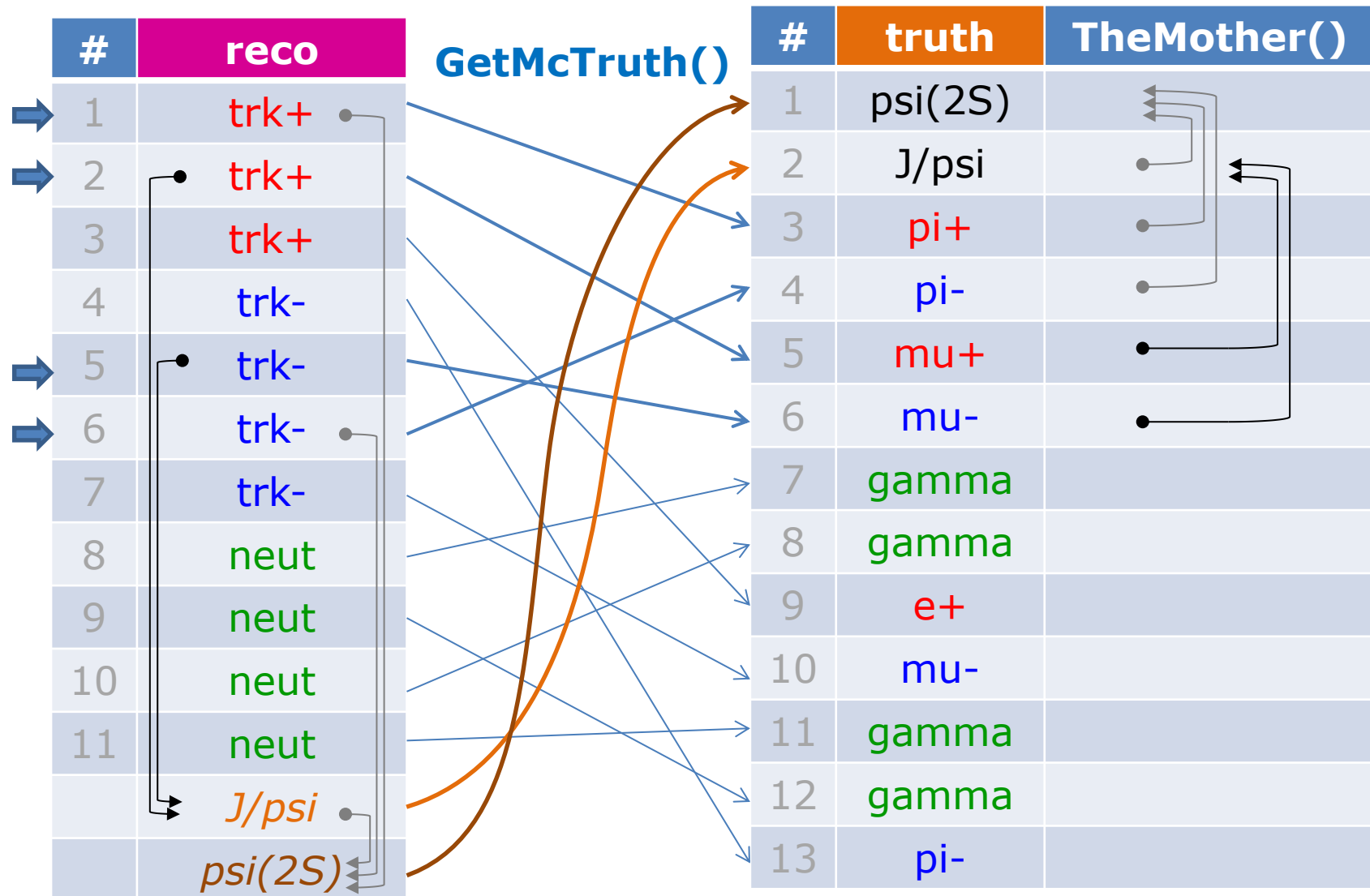


# Example: $\psi(2S) \rightarrow J/\psi (\mu^+\mu^-) \pi^+\pi^-$





# Example: $\psi(2S) \rightarrow J/\psi (\mu^+\mu^-) \pi^+\pi^-$



# Usage of PndAnalysis::McTruthMatch

- For matching, **composite** candidates have to have **type set**

```
PndAnalysis *ana = new PndAnalysis();

while ( ana->GetEvent() )
{
    ana->FillList(muplus, "MuonPlus");
    ...
    jpsi.Combine(muplus, muminus);
    jpsi.SetType( "J/psi" );    // set type for J/psi (names like TDatabasePDG)

    psi2s.Combine(jpsi, piplus, piminus);
    psi2s.SetType( "psi(2S)" ); // set type for psi(2S)

    bool match = ana->McTruthMatch( psi2s[0] ); // match for RhoCandidate

    int nmatch = ana->McTruthMatch( psi2s );    // match complete RhoCandList

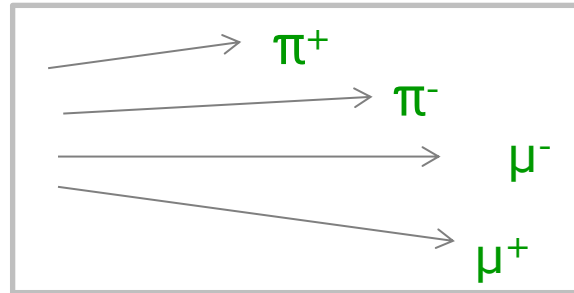
    for (int j=0; j<psi2s.GetLength(); ++j)
    {
        RhoCandidate *truth = psi2s[j].GetMcTruth(); // access truth of composites
        ...
    }
}
```

*still some inconsistency  
with TDatabasePDG names*  
*psi(2S) <-> psi'*  
*30443 <-> 100443*

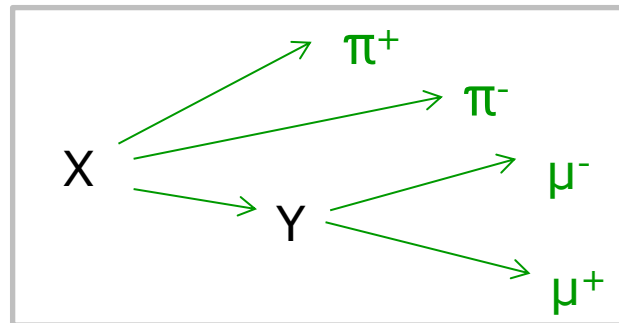
# Different levels of matching

- Three different match levels are available via  
`ana->McTruthMatch( psi[0], level );`

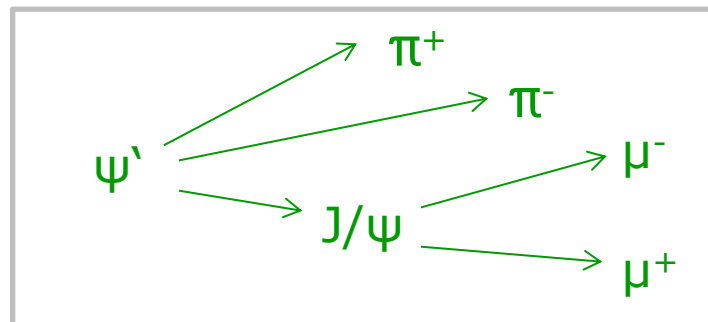
- Level = 0 only looks for correct PID



- Level = 1 matches topology in addition

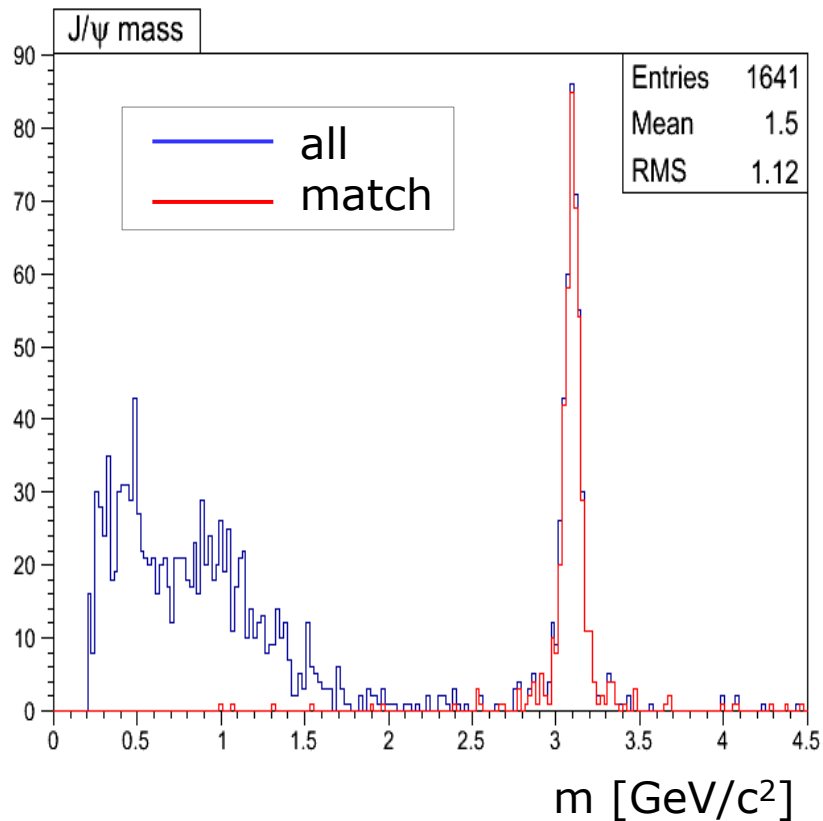


- Level = 2 (default) is full match

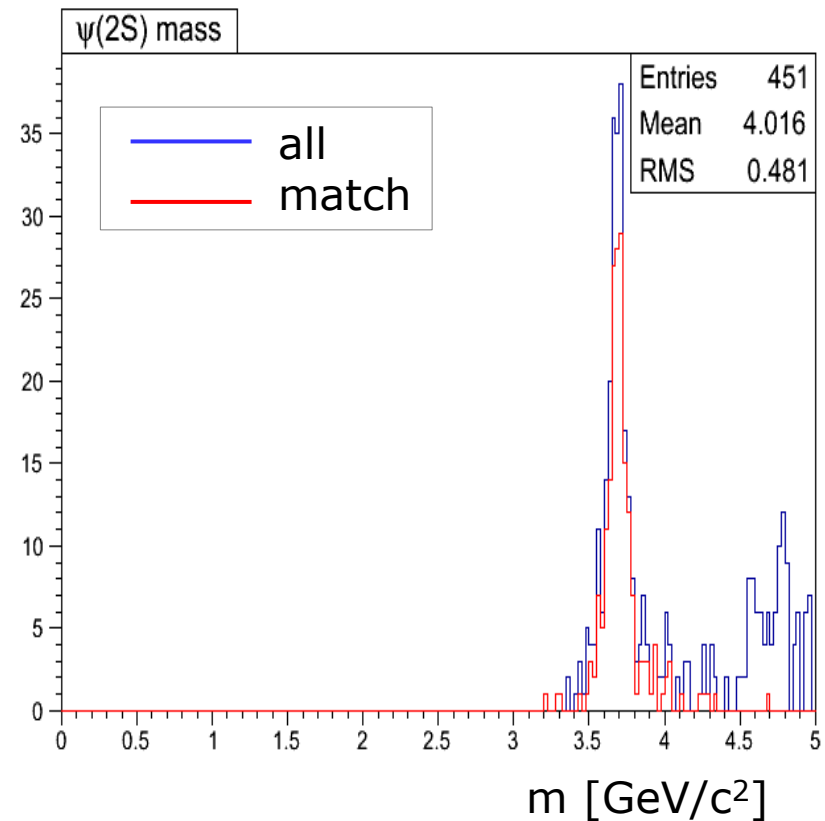


# Example: Mc Truth Match

$J/\psi \rightarrow \mu^+\mu^-$



$\psi(2S) \rightarrow J/\psi \pi^+\pi^-$



# Fitting

- Fitters basically all have a similar interface
- Fit results are now **attached to RhoCandidates**
- Can be accessed as full tree, **allows cascaded fitting!**
- E.g. vertex fitting + mass fitting might look like this:

```
RhoCandidate *lambda = pplus->Combine( pminus );
TVector3 IP( 0,0,0 );
...
PndKinVtxFitter fitvtx( lambda );           // setup vertex fitter
fitvtx.AddPointingConstraint( IP );         // add pointing constraint
fitvtx.FitAll();                           // perform fit

RhoCandidate *lambda_vtx = lambda->GetFit(); // access fit results

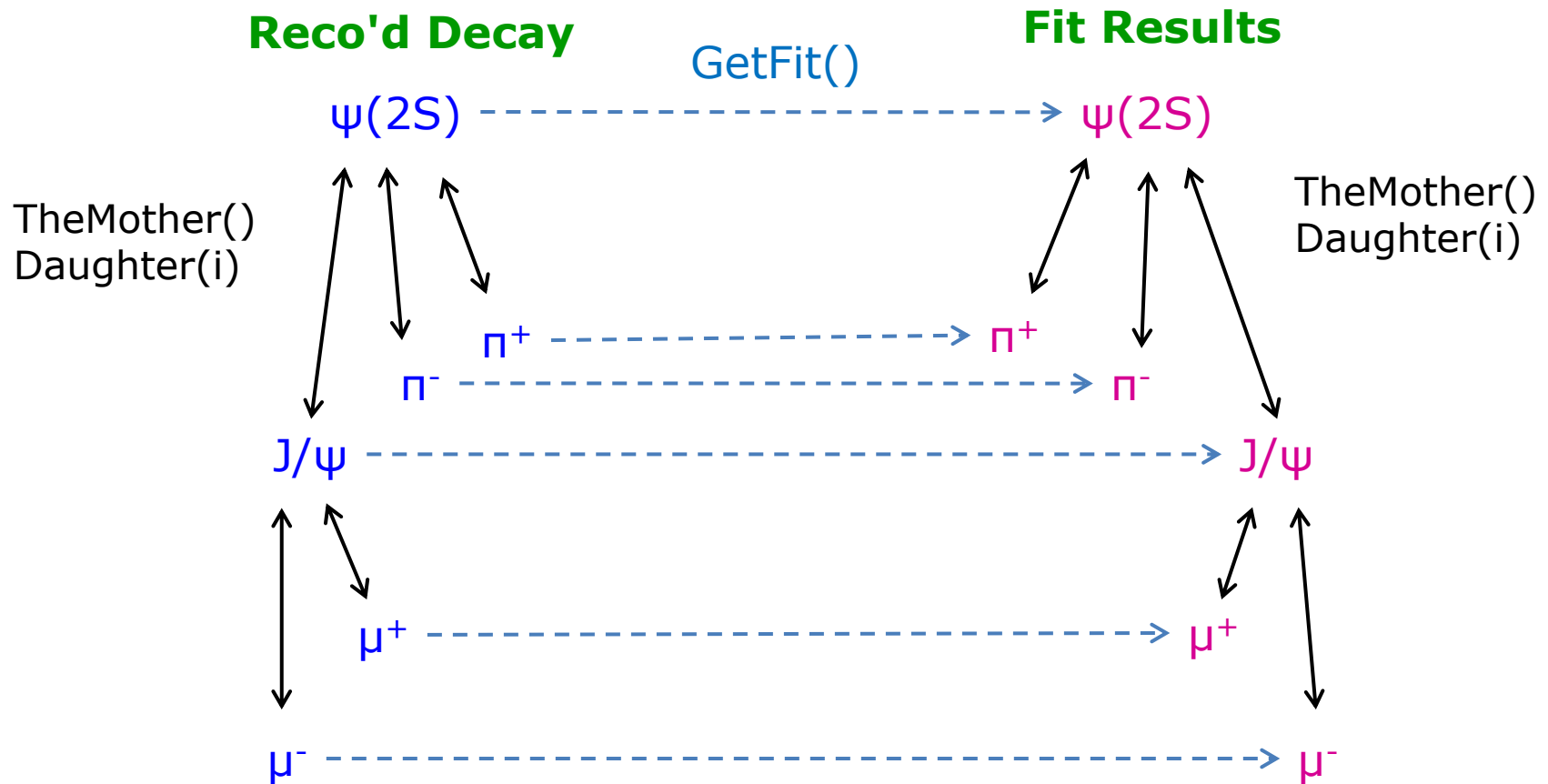
PndKinFitter fitmass( lambda_vtx );         // setup mass fitter
fitmass.SetMassConstraint( 1.115 );        // set mass constraint
fitmass.Fit();                             // perform fit

RhoCandidate *lambda_mass = lambda_vtx->GetFit(); // access cascaded fit results

RhoCandidate *fit_pplus  = lambda_mass->Daughter( 0 );
RhoCandidate *fit_pminus = lambda_mass->Daughter( 1 );
```

# Fitting: Access to Results

- After fit → full fitted tree is attached to the reco object



# Analysis Code in a Task

- **Problem:**
  - Interpreted code is harder to be debugged
  - It runs slower when using many loops
  - Beyond certain macro-complexity
    - *CINT starts getting slower from event to event*
- **Solution:**
  - Transformation of the macro code into task
    - Code is compiled and thus more stable/reliable
    - Analysis runs (usually) much faster
    - Doesn't have the problem from above
- Much more simple to achieve than you might think
- How-to is given in the new tutorial wiki!

# Analysis Code in a Task

## AnalysisMacro.C

```
void AnalysisMacro()  
{  
    FAIRROOT Initialization  
  
    HISTOGRAM Declaration  
    HISTOGRAM Initialization  
  
    GLOBALS Declaration  
    GLOBALS Initialization  
  
    while ( fAna->GetEvent() )  
    {  
        ANALYSIS CODE  
    }  
  
    SAVING HISTOGRAMS  
}  
  
ROUTINES Definition  
ROUTINES Implementation
```

## AnalysisMacroTask.C

```
void AnalysisMacroTask()  
{  
    FAIRROOT Initialization  
    PndMyAnaTask *mytask = new PndMyAnaTask()  
    fRunAna->AddTask(mytask);  
}
```

## PndMyAnaTask.h

```
class PndMyAnaTask : public FairTask  
{  
private:  
    GLOBALS Declaration  
    HISTOGRAM Declaration  
    ROUTINES Definition  
};
```

## PndMyAnaTask.cxx

```
ROUTINES Implementation  
  
PndMyAnaTask::Init()  
HISTOGRAM/GLOBALS Initialization  
  
PndMyAnaTask::Exec()  
ANALYSIS CODE  
  
PndMyAnaTask::Finish()  
SAVING HISTOGRAMS
```



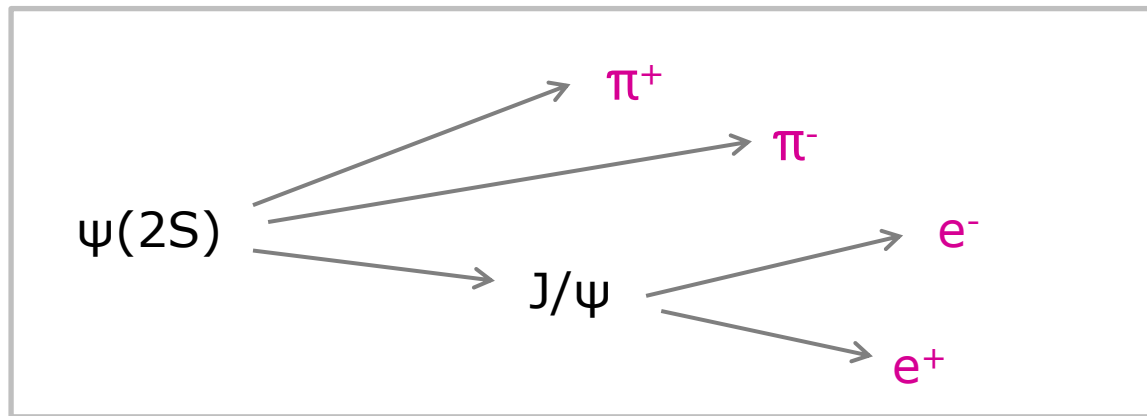
# Summary

- **New Tutorial is available**
  - Plan: Keep functioning (+ extend it) as standard tutorial
  - Rho Class Docu available in addition
- **Extended PID concept**
  - Easy use with `PndAnalysis::FillList`
  - Stand-alone use possible
- **Monte Carlo Truth Match**
  - Should work for all kinds of complicated decay trees
  - Allows access to non-final-state MC truth resonances
- **Fitting**
  - Full fitted tree created and attached to reco object
  - Cascaded fitting possible on fit result
- **Analysis in Task**
  - Documented in Tutorial → more stable running

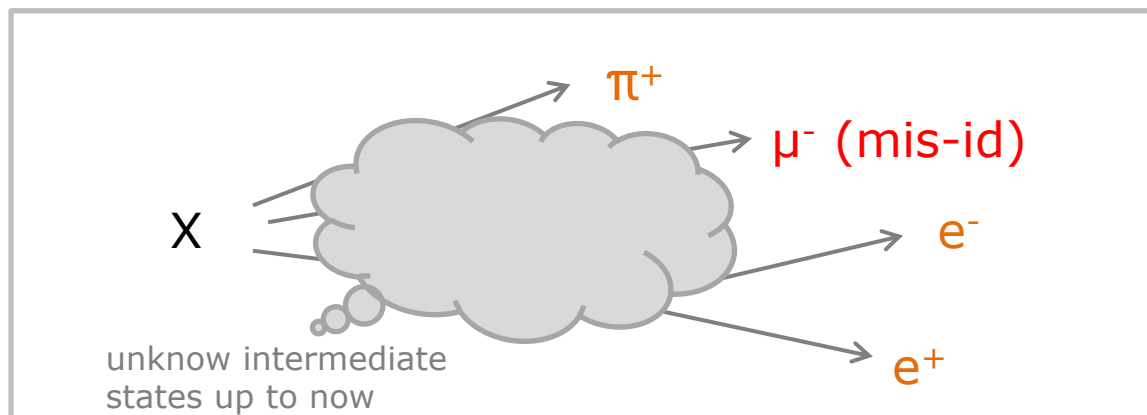
# BACKUP

# PndMcTruthMatch

- **Checklist** for full tree match:
  1. truth objects of final states have the correct PID types

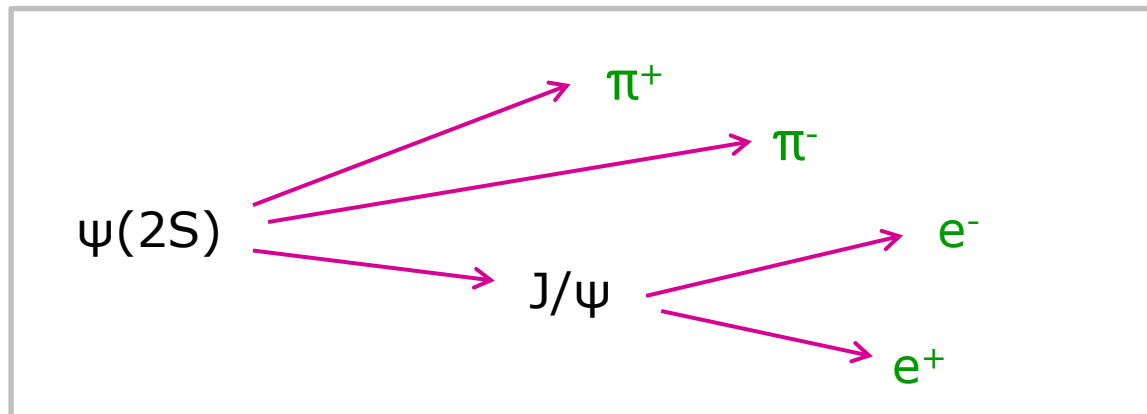


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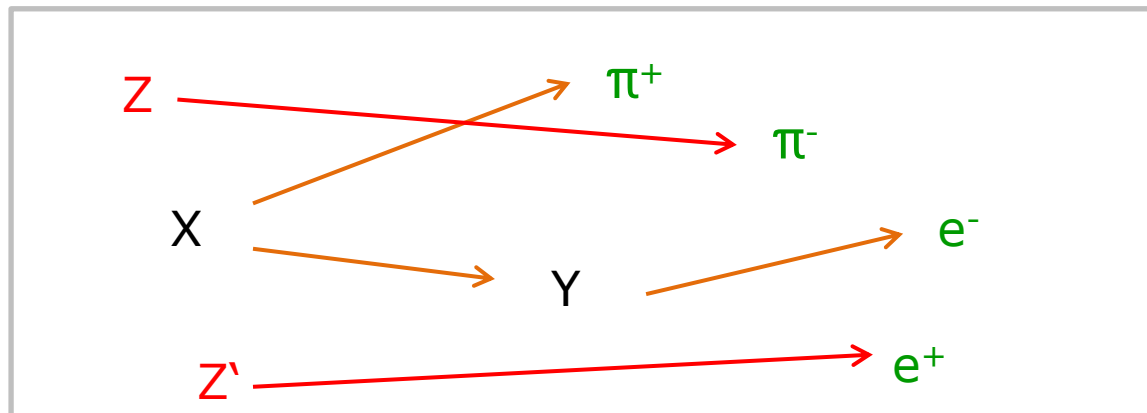


# PndMcTruthMatch

- **Checklist** for full tree match:
  1. truth object of final states have the same mother

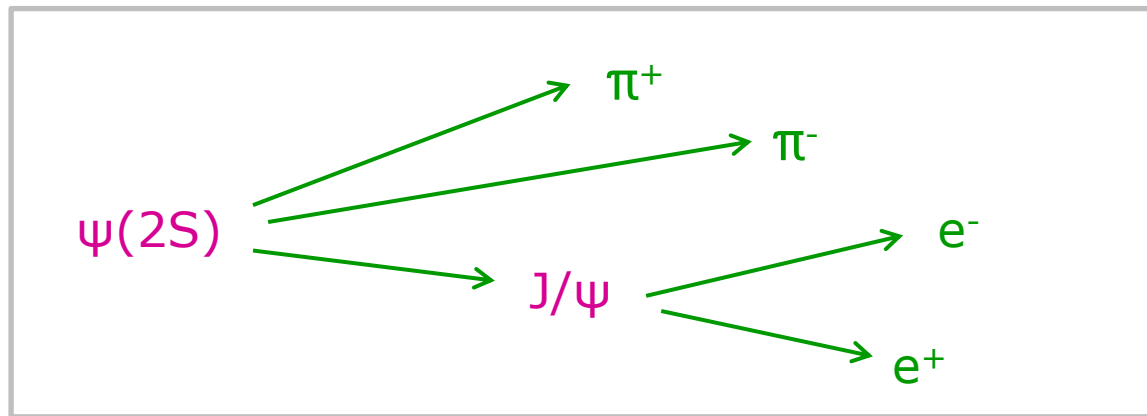


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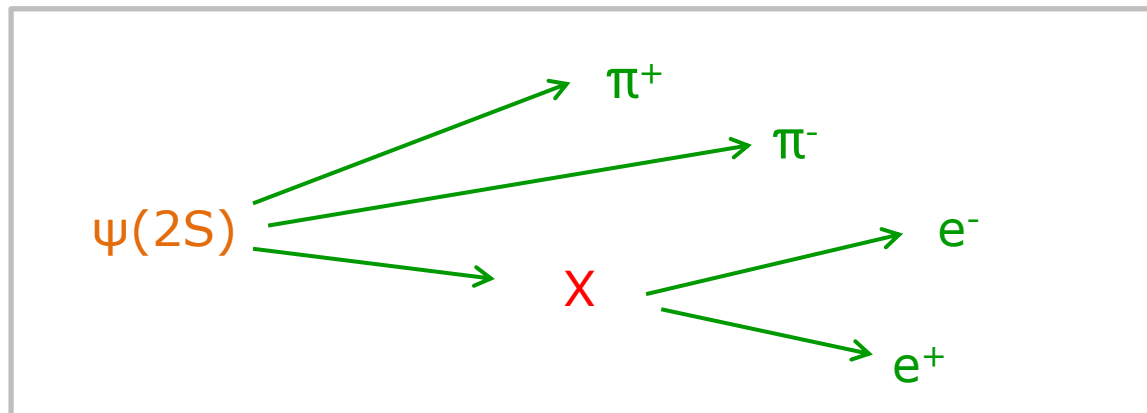


# PndMcTruthMatch

- **Checklist** for full tree match:
  1. mother has required type
  2. mother has required daughters
  3. mother has required type

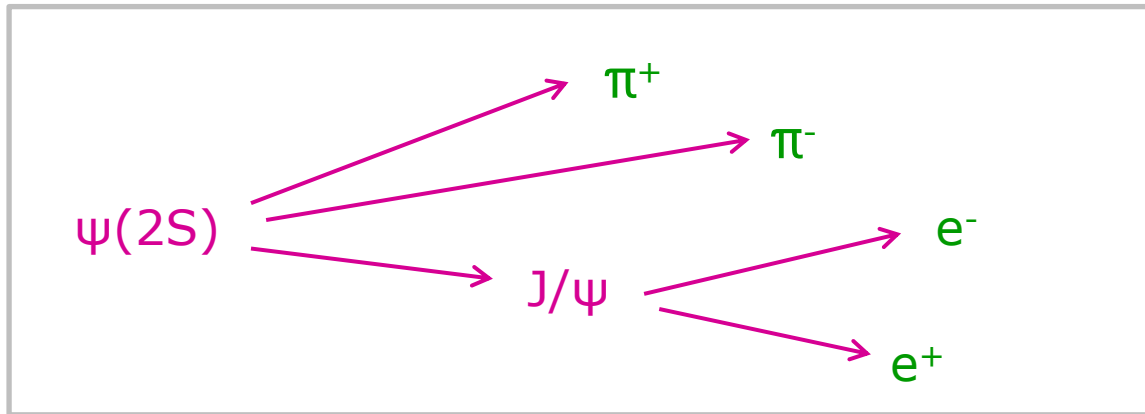


$\neq$



# PndMcTruthMatch

- **Checklist** for full tree match:
  4. mother has correct number of daughters



$\neq$

