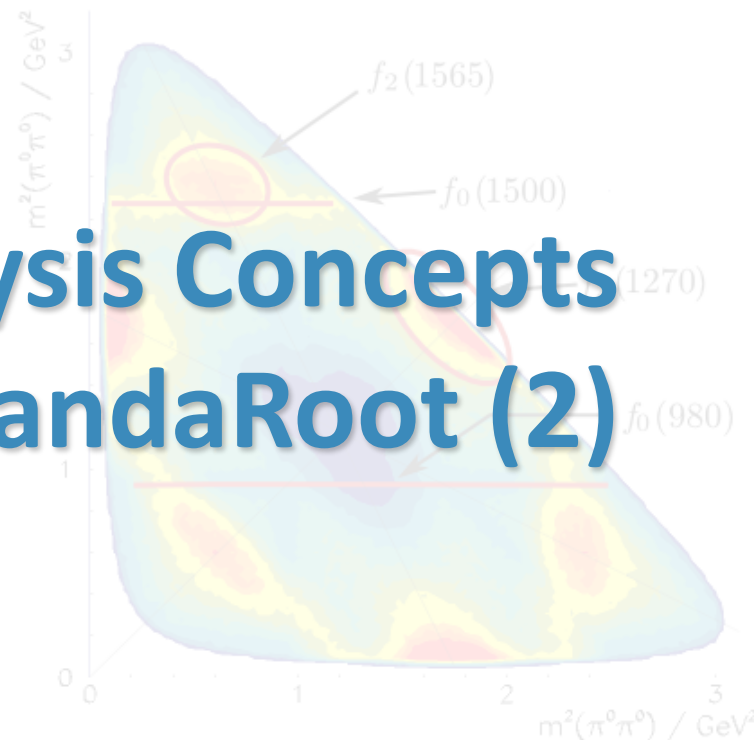


# Physics Analysis Concepts with PandaRoot (2)



*PANDA Lecture Week 2017*

*GSI, Dec 11 - 15, 2017*

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GSI Darmstadt

# Topics

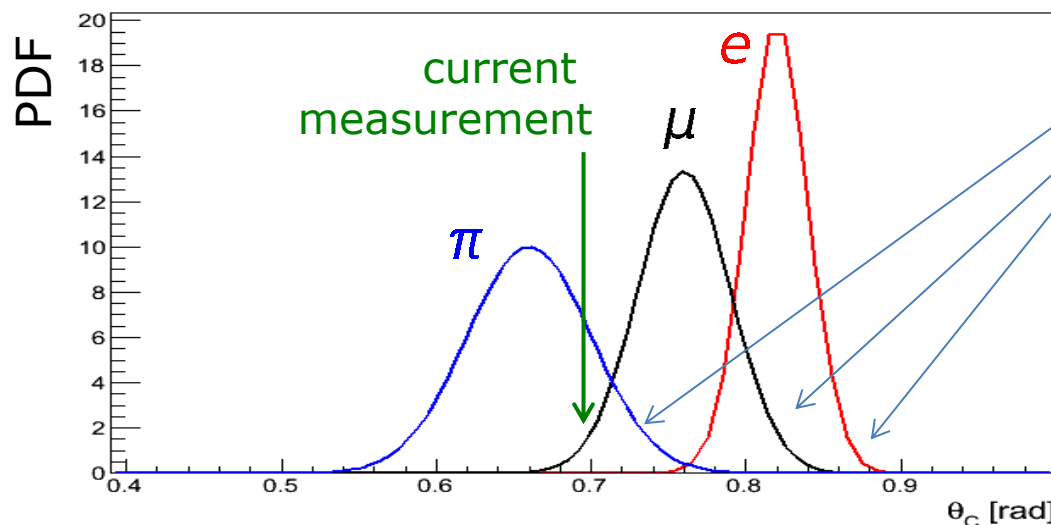
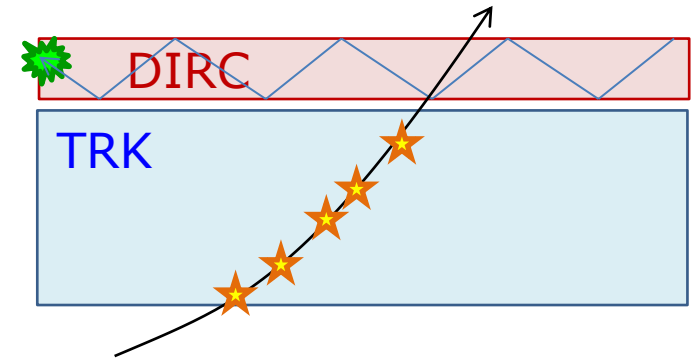
- Particle Identification
- Particle Selectors (*PndPidCombiner*, *PndPidSelector*)
- MC Truth Match (*PndAnalysis::McTruthMatch*)
- Fitting Basics
- Fitting Tools (*PndKinVtxFitter*, *PndKinFitter*)
- Uncertainties

# **PARTICLE IDENTIFICATION**

# Particle Identification Basics

How does PID work in principle?

1. Reconstruct track object
2. Match track with PID detector signal
3. Reconstruct PID information
4. Compute likelihood for various particle types (e.g. based on Likelihood functions)



expected  $\theta_c$  distributions for a certain momentum

# PID Concept in PandaROOT

- Each PID subdetector delivers individual probability/likelihood information
- **PID Combiner**, which combines the likelihood values from different detectors

$$\tilde{P}_{tot,i} = \prod_{Det} P_{Det,i} \quad i \in \{e, \mu, \pi, K, p\}$$

- Normalization via

$$P_{tot,j} = \frac{\tilde{P}_{tot,j}}{\sum_i \tilde{P}_{tot,i}}$$

- **PID Selector**, which requires certain  $P_{tot}$  for positive identification

# PID Concept in PandaROOT

PandaROOT objects: **PndAnaPidCombiner**, **PndAnaPidSelector**

- **PndAnaPidCombiner**
  - combines *on demand* probabilities from **various algorithms** by computing product of all  $P_k$  ( $k=algorithms$ )
  - copies resulting probabilities to RhoCandidate/RhoCandList
- **PndAnaPidSelector**
  - selects particles based on these probabilities
- **PndAnalysis::FillList** is a short-cut to this functionality via

```
pndana.FillList(list, "ElectronLoose", "PidAlgoEmcBayes;PidAlgoDrc");
```
- Predefined selection with keywords (probability based)  
**Electron / Muon / Pion / Kaon / Proton** +  
**All / VeryLoose / Loose / Tight / VeryTight / Best** +  
**Plus / Minus** (*optional*)  
*Simple keywords*  
**Charged / Plus / Minus / Neutral / All**

# PID Concept in PandaROOT

- PandaROOT objects:

**PndAnalysis, PndAnaPidCombiner, PndAnaPidSelector**

```
PndAnalysis *pndana= new PndAnalysis();  
pndana->FillList(eplus, "ElectronLoosePlus", "PidAlgoEmcBayes;PidAlgoMvd");  
pndana->FillList(eminus, "ElectronLooseMinus", "PidAlgoEmcBayes;PidAlgoMvd");
```

Or 'by hand':

```
RhoCandList charged, kaonLoose;  
  
PndAnaPidSelector kaonSel("KaonSelector");  
kaonSel.SetSelection("KaonLoose");           // set selection criterion  
  
PndAnaPidCombiner pidComb("PidCombiner");  
pidComb.SetTcaNames("PidAlgoDrc;PidAlgoMvd"); // set algo's  
  
while (evr->GetEvent())  
{  
    pndana->FillList(charged, "Charged");      // start w/ charged candidates  
  
    pidComb.Apply(charged);                    // copy probab. to candidates  
    kaonLoose.Select(charged, kaonSel);        // select kaons from charged  
}
```

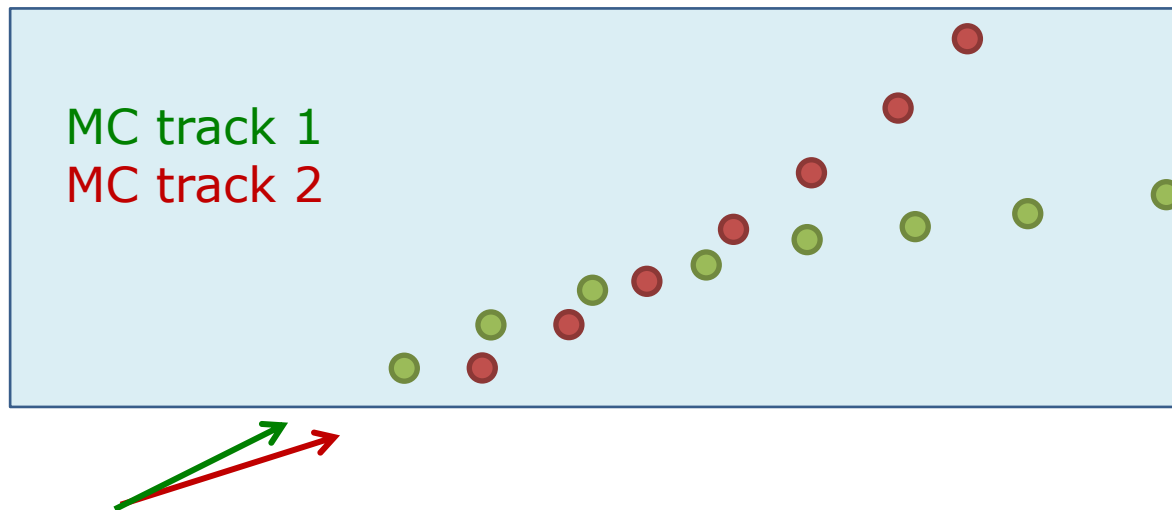
**MONTE CARLO TRUTH**



# Monte Carlo Truth Match

- In simulated reactions  
→ possibility to map reco object to MC object

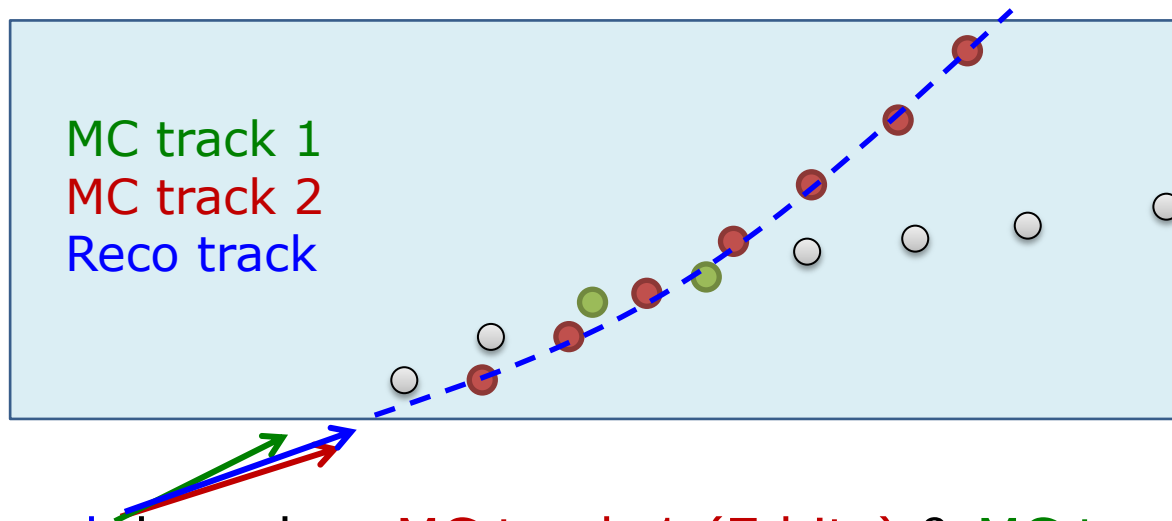
Tracking detector



# Monte Carlo Truth Match

- In simulated reactions  
→ possibility to map reco object to MC object

Tracking detector



- Reco track based on MC track 1 (7 hits) & MC track 2 (2 hits)  
→ Monte Carlo Truth(Reco track) = MC track 2 (max hits)  
[realized by FairLink class]

# Monte Carlo Truth Access

## Use cases:

- a) For typical measurement the **reco efficiency** is needed
  - b) Sometimes the true **signal distribution** is of interest
    - Look in MC, whether you combine the correct candidates
- 
- Concerning MC truth, distinguish between
    1. access MC truth object (or type) for certain reco object (see last slides)
    2. MC truth match for a complete decay tree

# 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

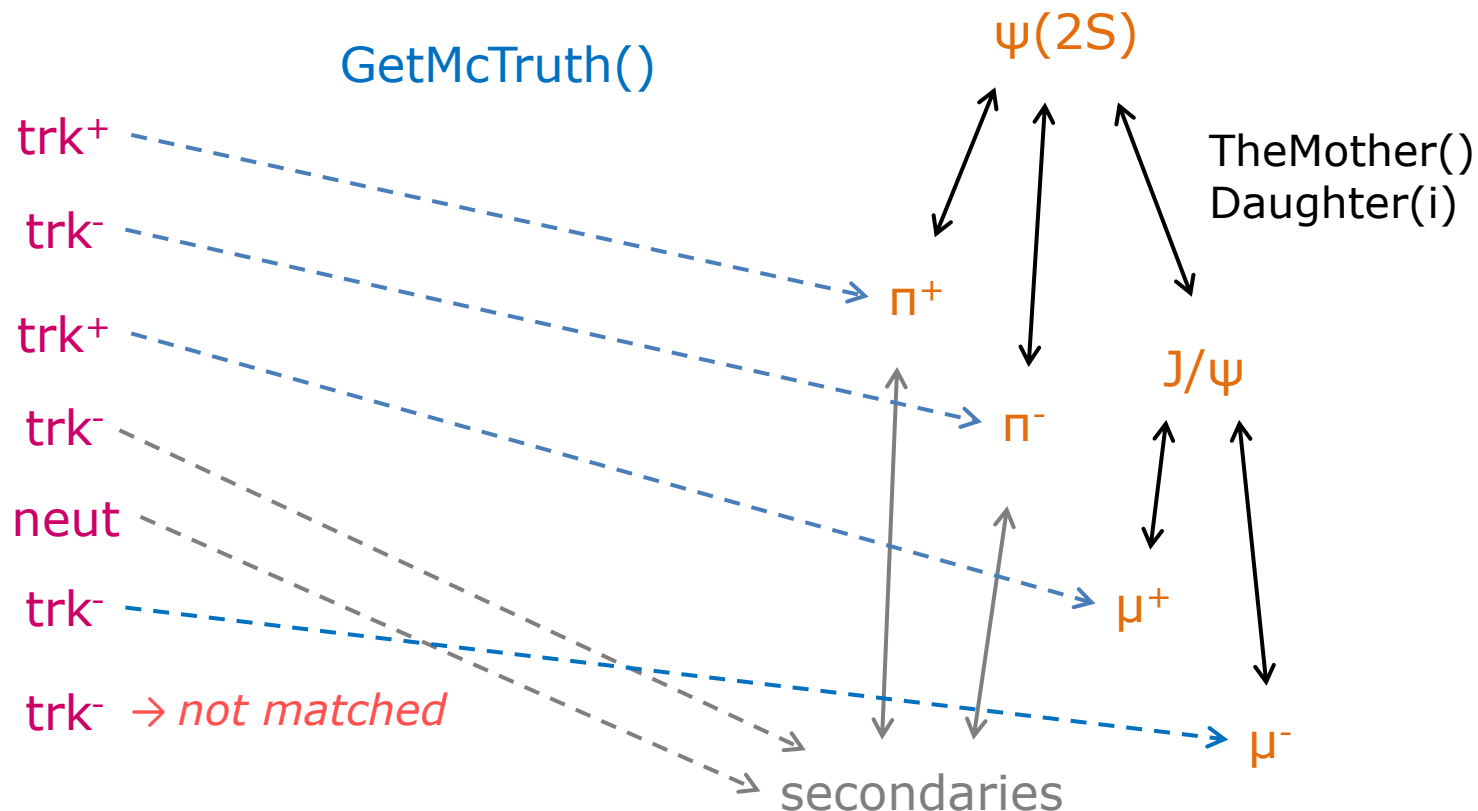
```
ana->FillList( mct, "McTruth" );  
...  
RhoCandidate *firstDaughter = mct[0]->Daughter(0);
```

# Mc Truth Genealogy

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

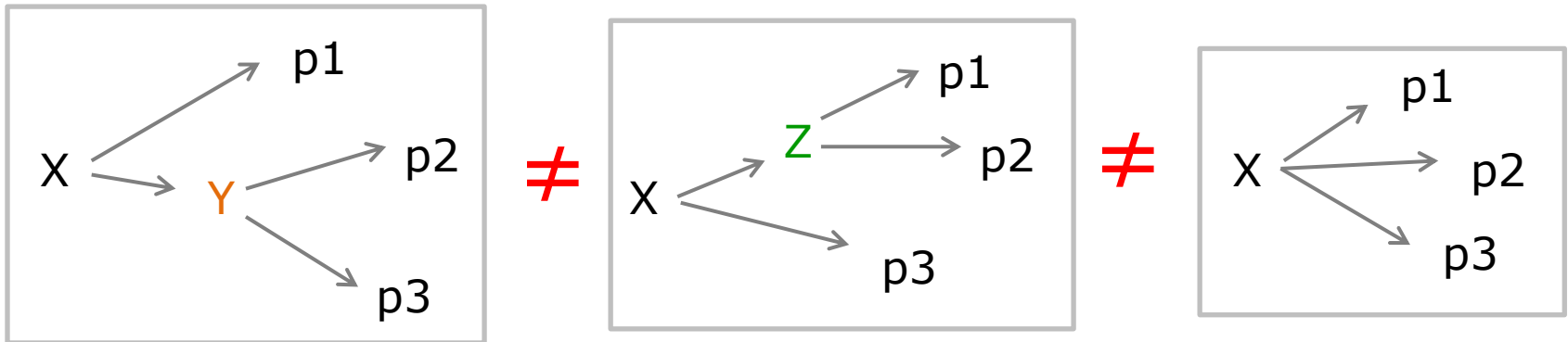
## All Reco

## McTruth



# PndAnalysis: MC Truth Match of Composites

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

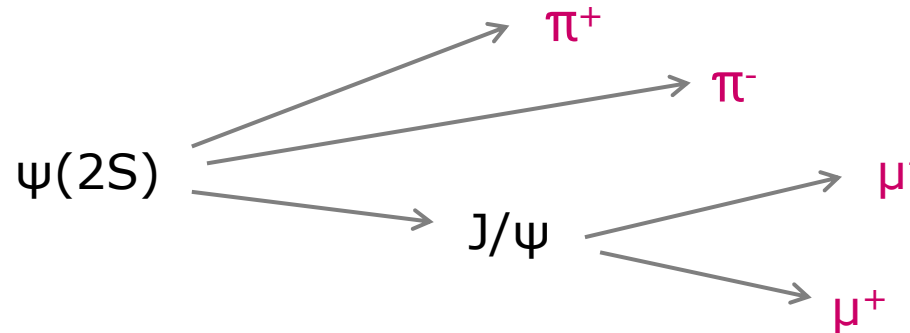


- Functionality in PndAnalysis method

```
PndAnalysis::McTruthMatch( RhoCandidate* );
```

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

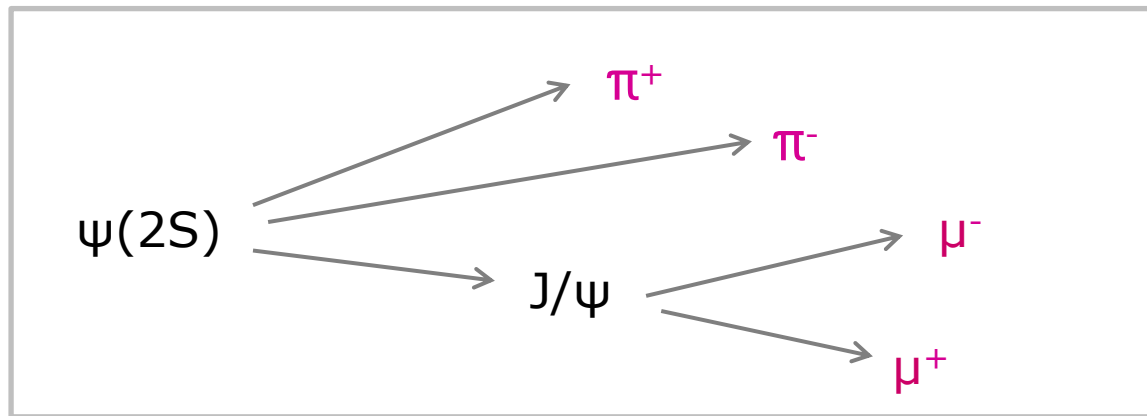
- We want to reconstruct the following decay



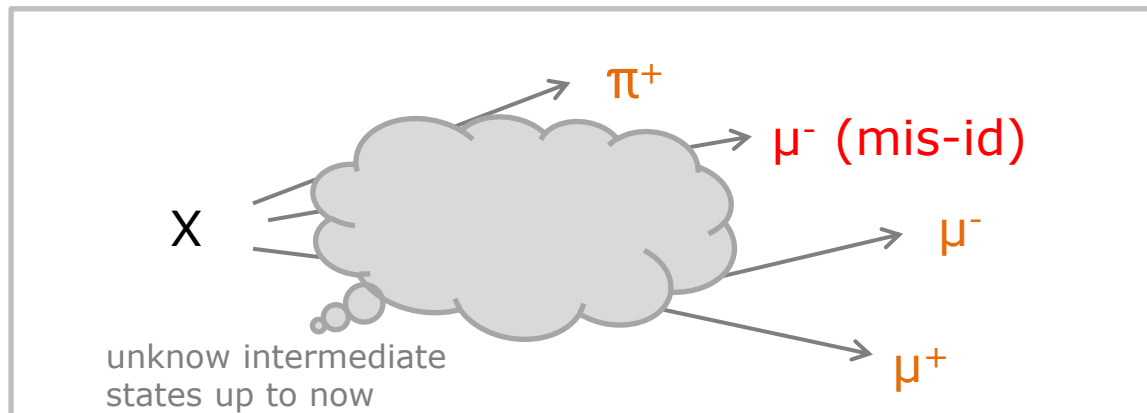
```
RhoCandList muplus, muminus, piplus, piminus, jpsi, psi2s;  
  
ana->FillList( muplus, "MuonLoosePlus", myPidAlgos );  
ana->FillList( muminus, "MuonLooseMinus", myPidAlgos );  
ana->FillList( piplus, "PionLoosePlus", myPidAlgos );  
ana->FillList( piminus, "PionLooseMinus", myPidAlgos );  
  
jpsi.Combine(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

# PndMcTruthMatch

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



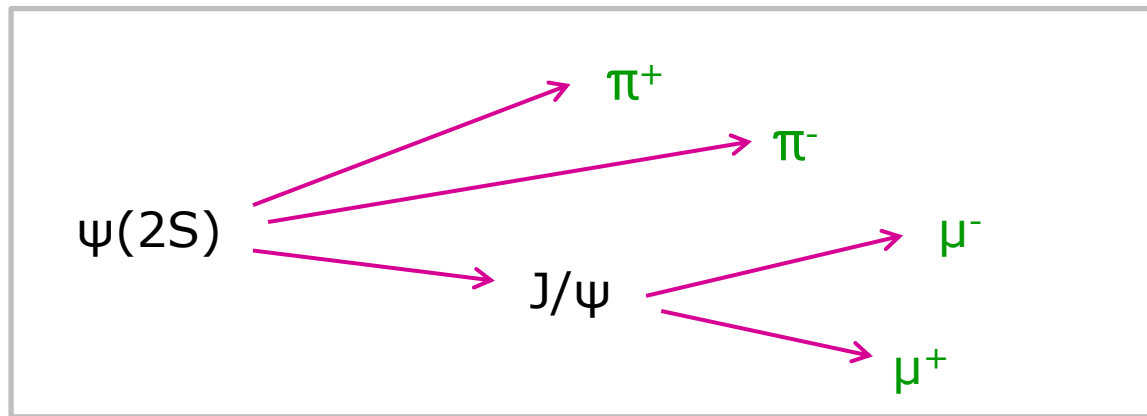
$\neq$



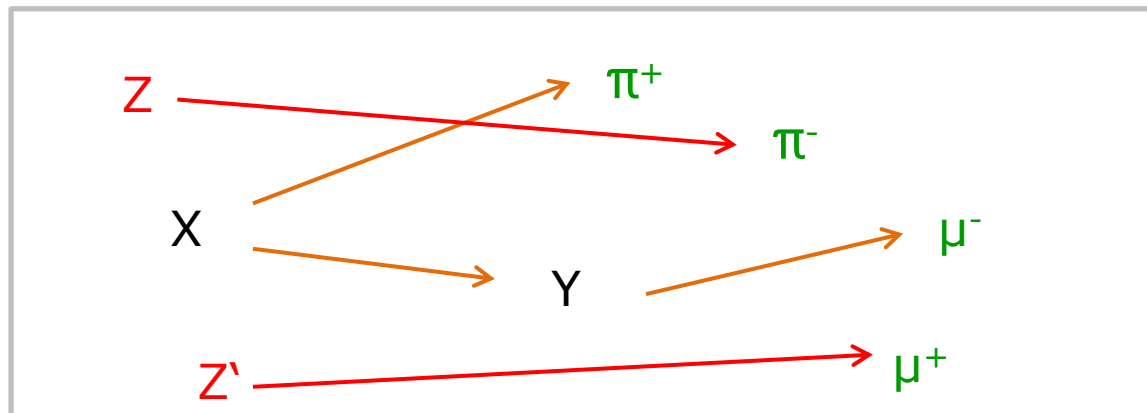


# PndMcTruthMatch

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

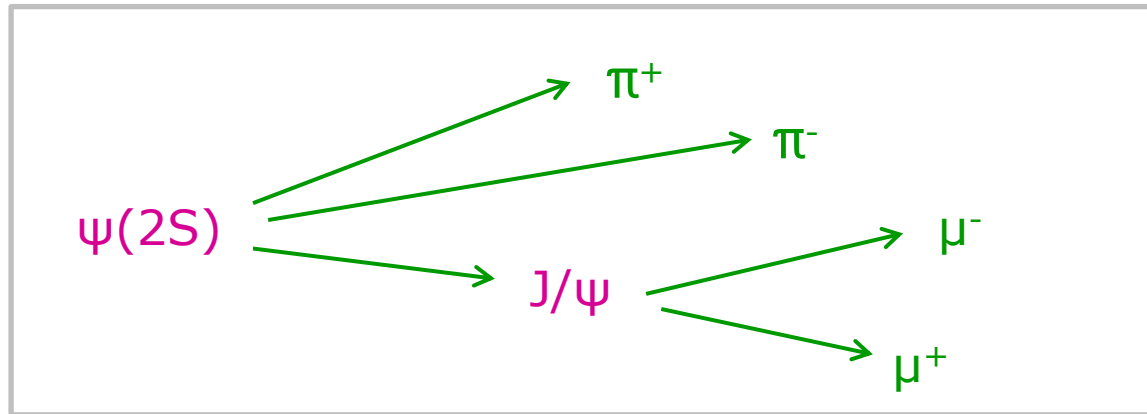


$\neq$

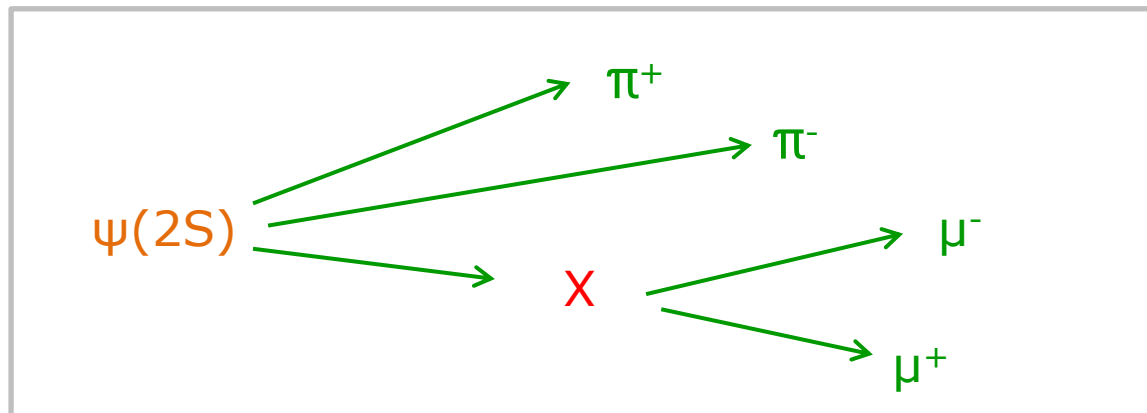


# PndMcTruthMatch

- **Checklist** for full tree match:
  1. mother and daughter have same type
  2. mother and daughter have same charge
  3. mother has required type

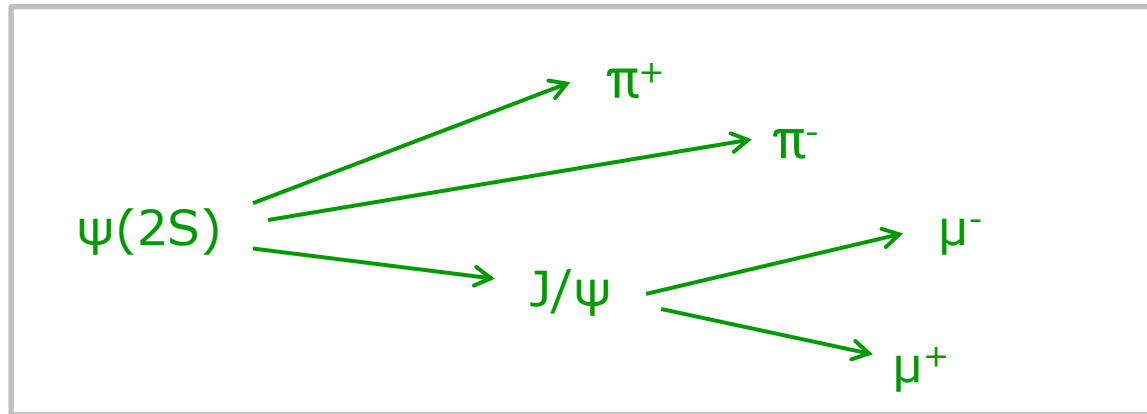


$\neq$

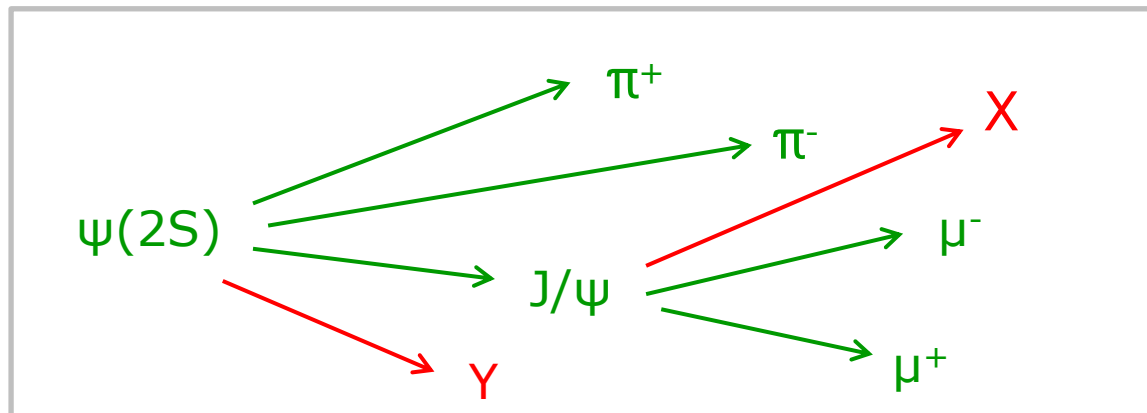


# PndMcTruthMatch

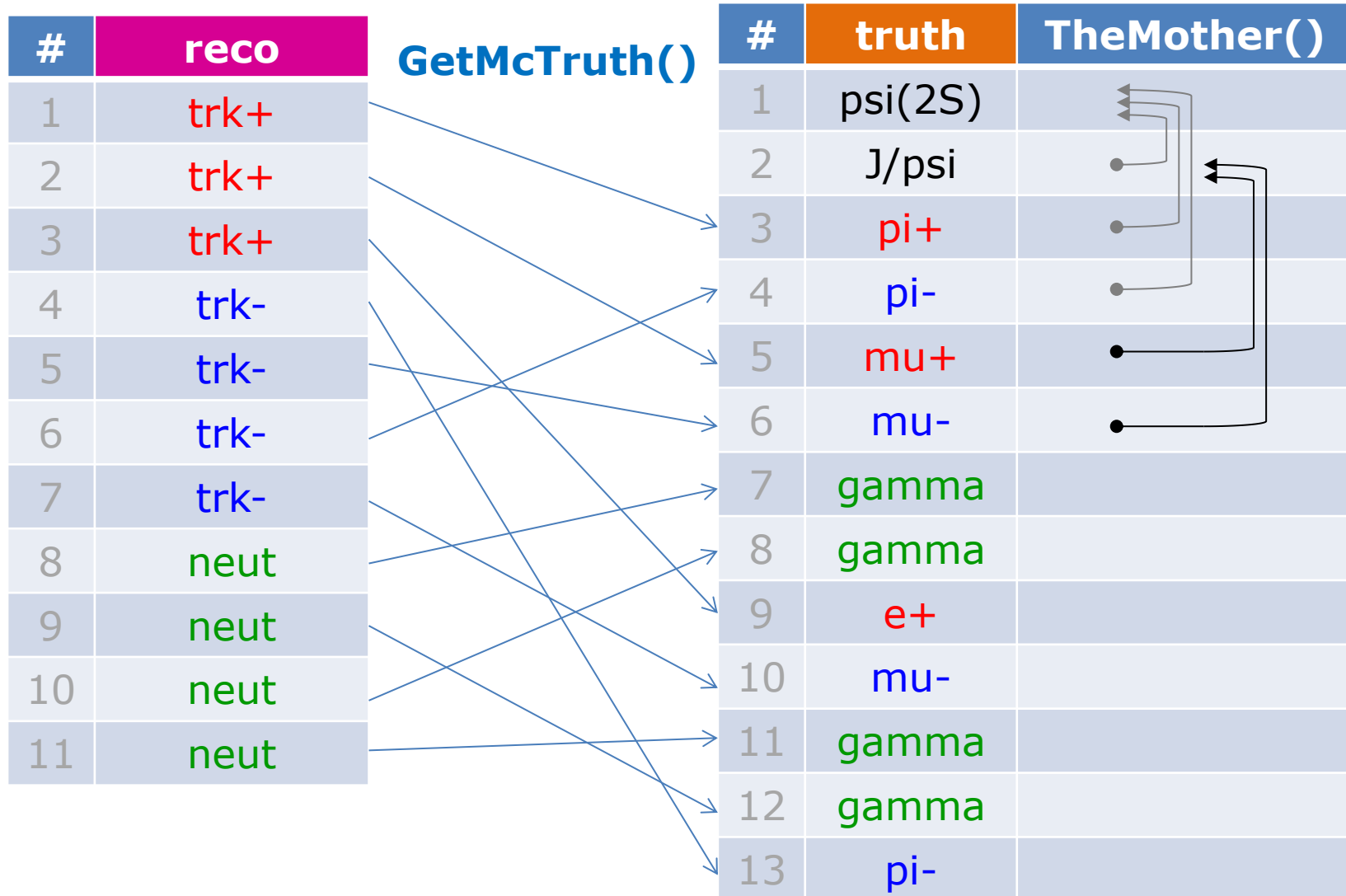
- **Checklist** for full tree match:
  4. mother has correct number of daughters



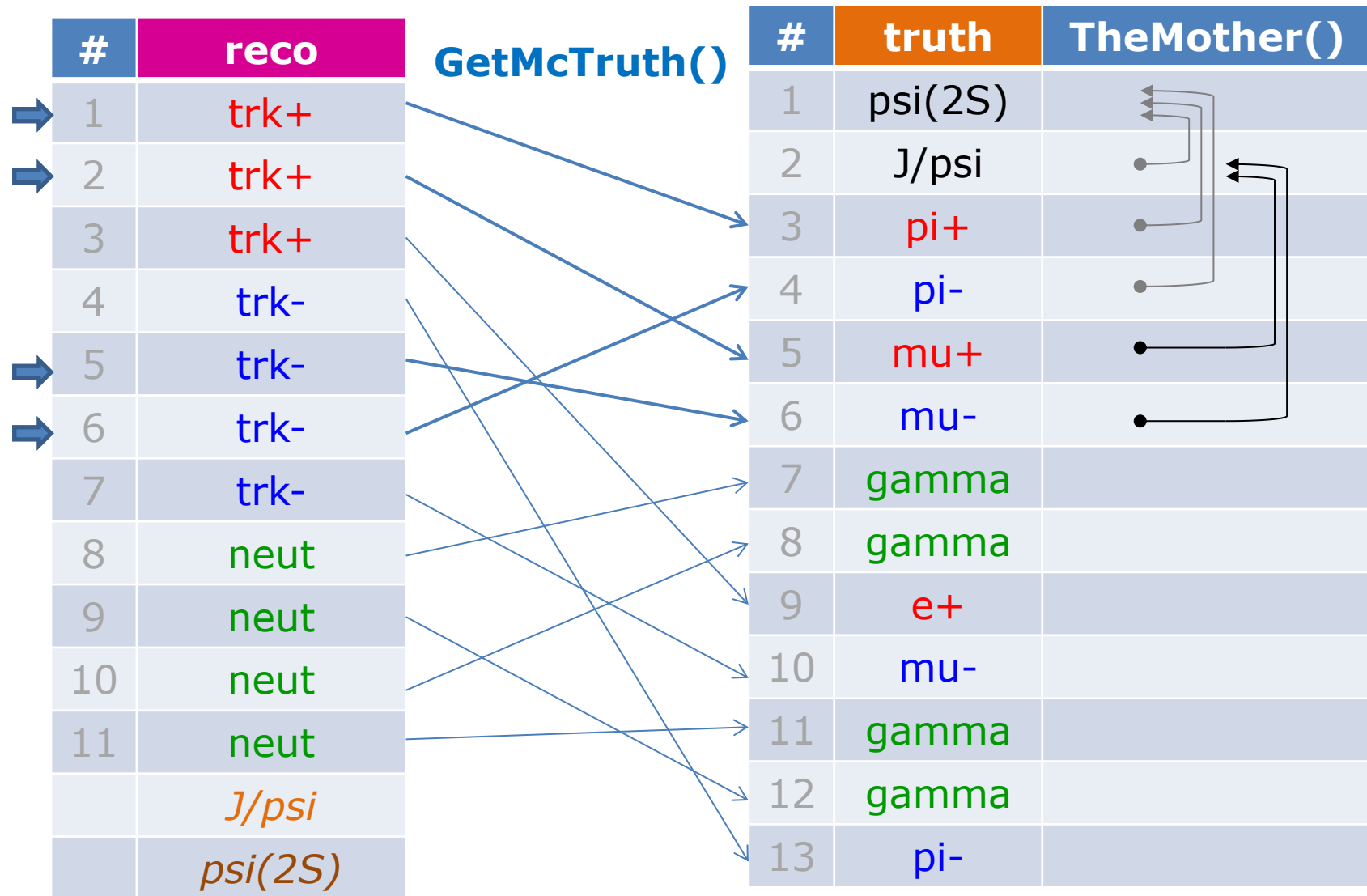
$\neq$



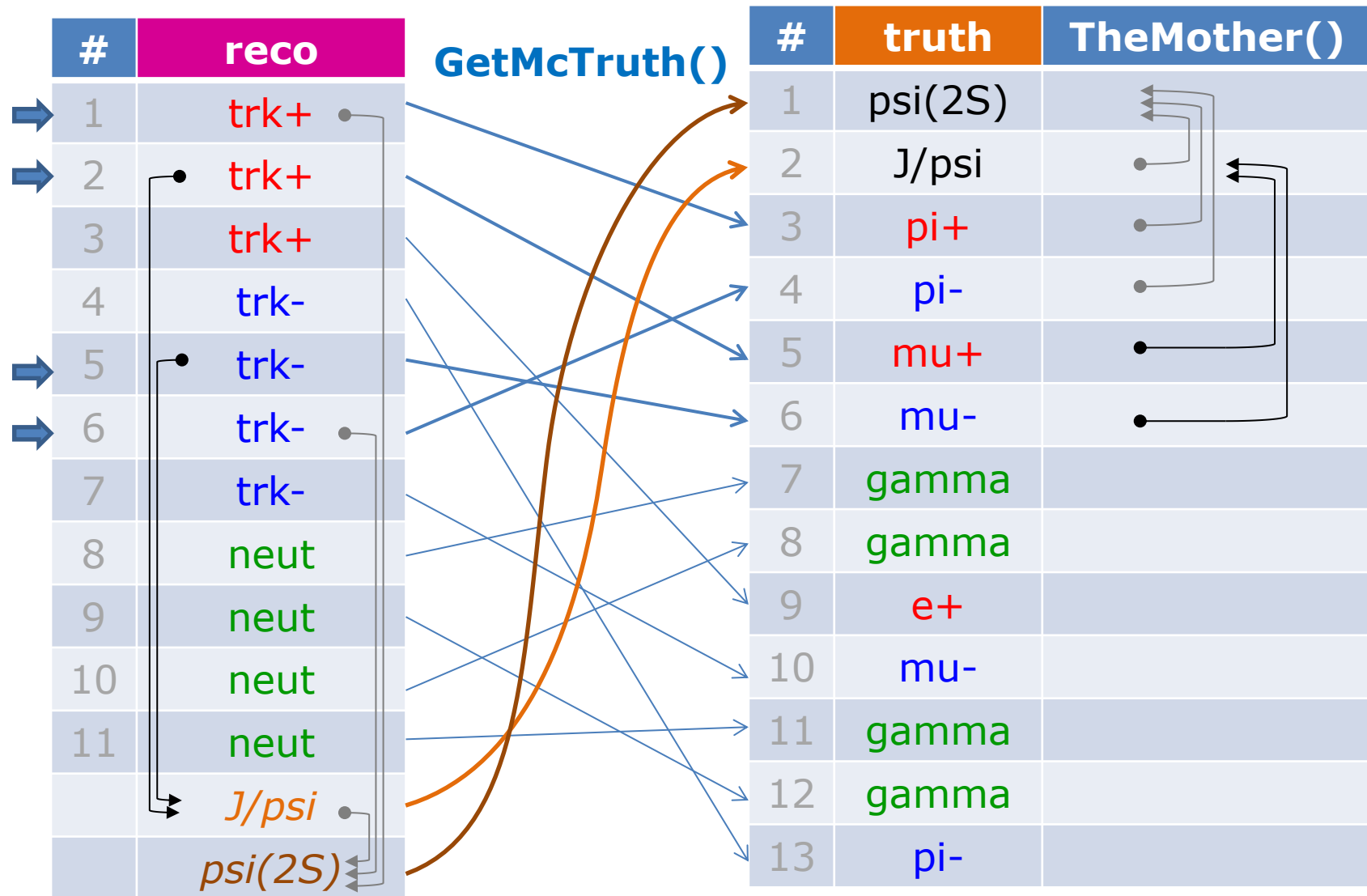
# Example: $\psi(2S) \rightarrow J/\psi (\mu^+\mu^-) \pi^+\pi^-$



# 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, "J/psi");  
    psi2s.Combine(jpsi, piplus, piminus, "pbarpSystem");
```

*although being at psi(2S)  
energy, particle was  
generated as pbarpSystem*

```
    bool match = ana->McTruthMatch( psi2s[0] ); // match for RhoCandidate  
    int nmatch = ana->McTruthMatch( psi2s );    // match complete RhoCandList
```

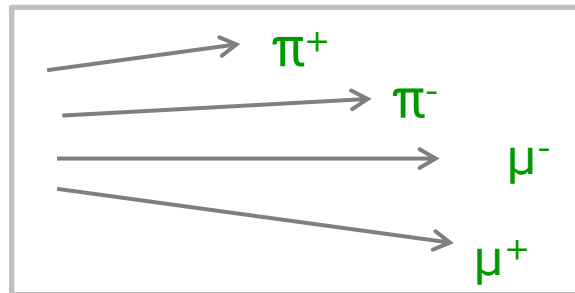
```
    for (int j=0; j<psi2s.GetLength(); ++j)  
    {
```

```
        RhoCandidate *truth = psi2s[j].GetMcTruth(); // access truth of composites  
        ...
```

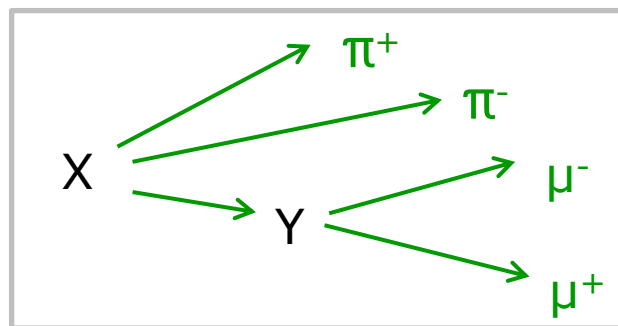
# Different levels of matching

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

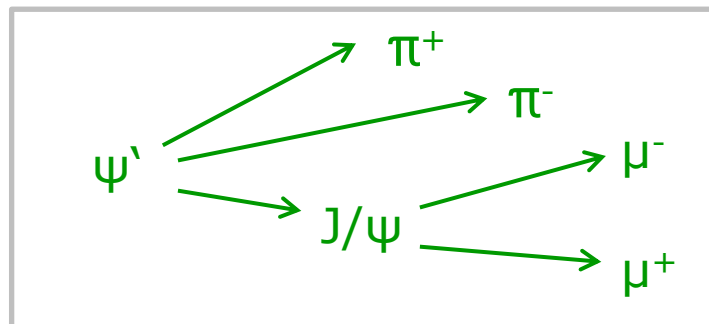
- Level = 0 only looks for correct PID



- Level = 1 matches topology in addition

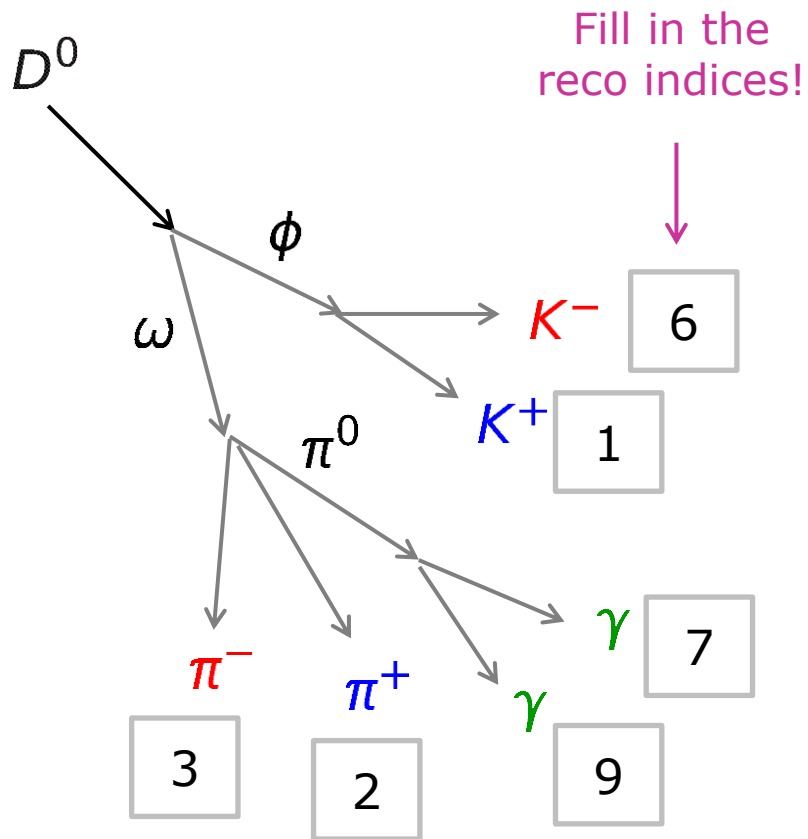


- Level = 2 (default) is full match





# MC Truth Match Exercise



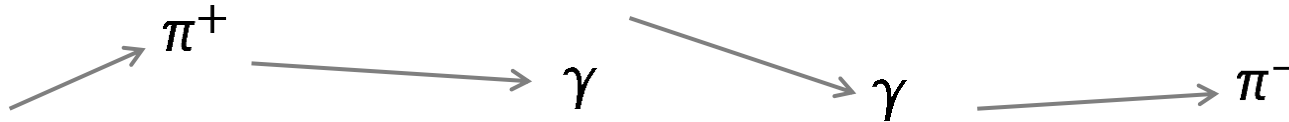
#	reco	mc
1	trk+	7
2	trk+	5
3	trk-	4
4	trk-	11
5	trk-	14
6	trk-	8
7	neut	10
8	neut	12
9	neut	9

#	truth	mum
1	D0	-1
2	omega	1
3	phi	1
4	pi-	2
5	pi+	2
6	pi0	2
7	K+	3
8	K-	3
9	gamma	6
10	gamma	6
11	e-	9
12	gamma	11
13	gamma	11
14	e-	10

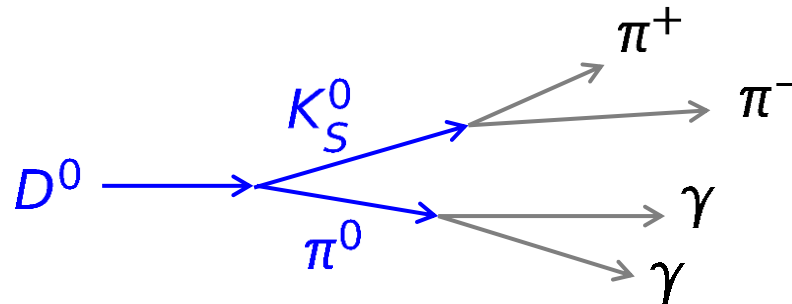
**FITTING**

# Kinematic Fitting

- Let's e.g. consider a set of particle candidates



- You believe what happens is this (hypothesis):

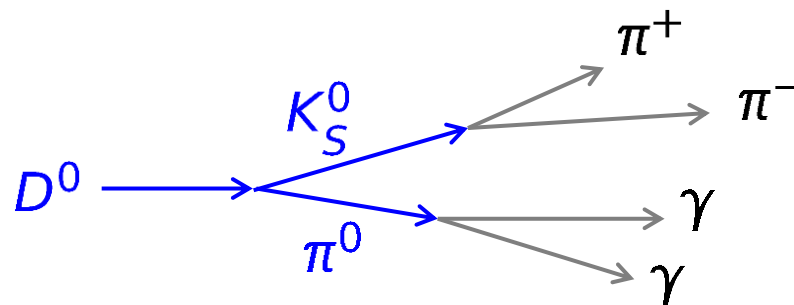


- Kinematic fitting**
  - gives a **measure** for the assumption, **that hypothesis is true**
  - improves the precision/resolution** for kinematic quantities, when hypothesis *actually is* true

More about fitting under <http://www.phys.ufl.edu/~avery/fitting.html>

# Kinematic Fitting: Constraints

- In order to perform fit, you need to apply so-called **constraints**
- Constraints correlate kinematics in the tree



- *Which constraints could be applied here?*
  - Vertex constraint  $K_S$ :  $\pi^+$  &  $\pi^-$  from same origin
  - Mass constraint  $K_S$ : inv.  $\pi^+ \pi^-$  mass =  $K_S$  rest mass
  - Mass constraint  $\pi^0$ : inv.  $2\gamma$  mass =  $\pi^0$  rest mass
  - Mass constraint  $D^0$ : inv.  $K_S \pi^0$  mass =  $D^0$  rest mass

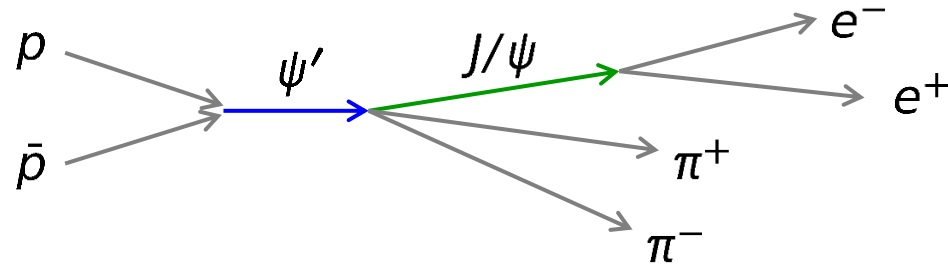
# Kinematic Fitting: Constraints

Most common constraints are

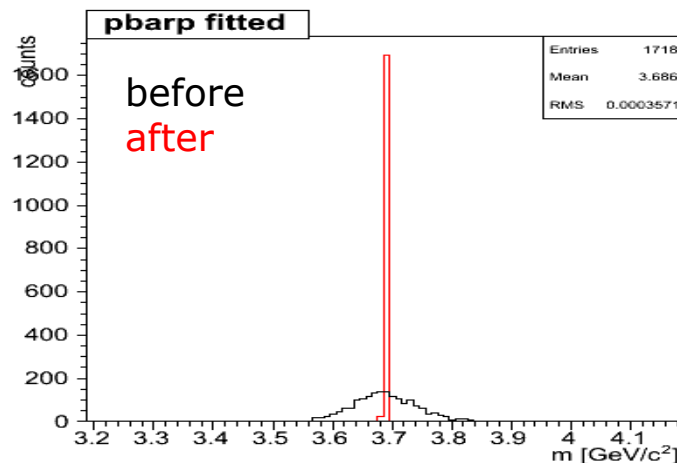
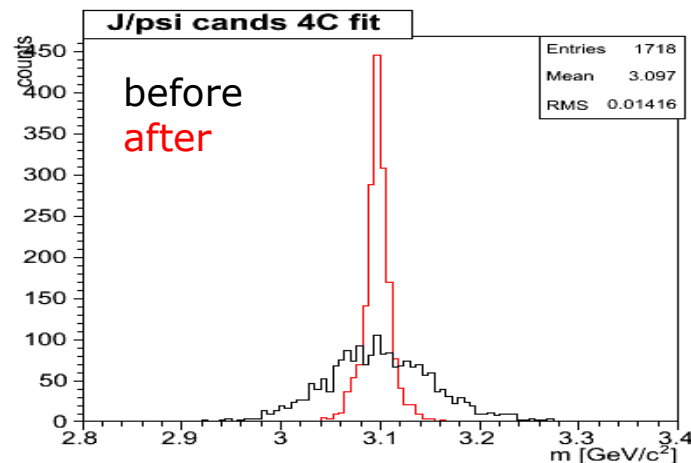
- **Mass Constraint**  
Composite particle has to have a certain invariant mass (only to be applied to quasi stable particles)
- **Vertex Constraint**  
Set of 4-vectors originate from common spatial point (which is determined during the fit!). *At least 2 charged tracks needed!*
- **4-vector-Constraint**  
4-vector of composite particle has to match a certain 4-vector component-wise (used in exclusive reconstruction)
- **Pointing constraint**  
3-vector of composite particle is consistent with originating from a certain point (e.g. the IP)

# Fitting: 4 Constraint Fit

- Can be applied in *exclusive* analysis, where the **initial 4-vector** is fully reconstructed

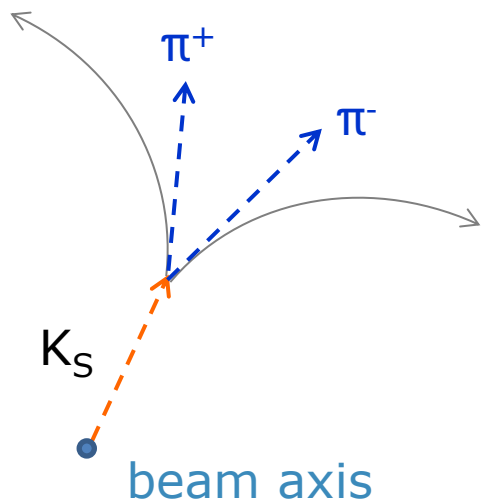


- Fitted total system is not very meaningful (spiked mass), but **intermediate resonances can be improved**
- Can do a **cut on the fit probability to reject background**

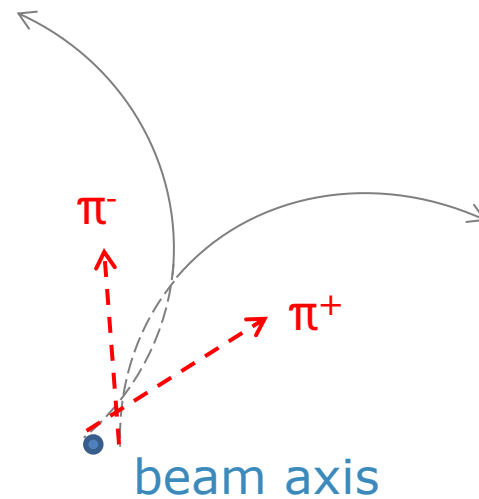


# Vertex Fitting for Long-Living Particles

- For long-living particles (e. g.  $K_S^0$ ,  $\Lambda^0$ ), vertex fitting is essential, since charged particles move on helices in magnet field



The 4-vector sum of daughters is only consistent with resonance 4-vector when evaluated at true points of origin!

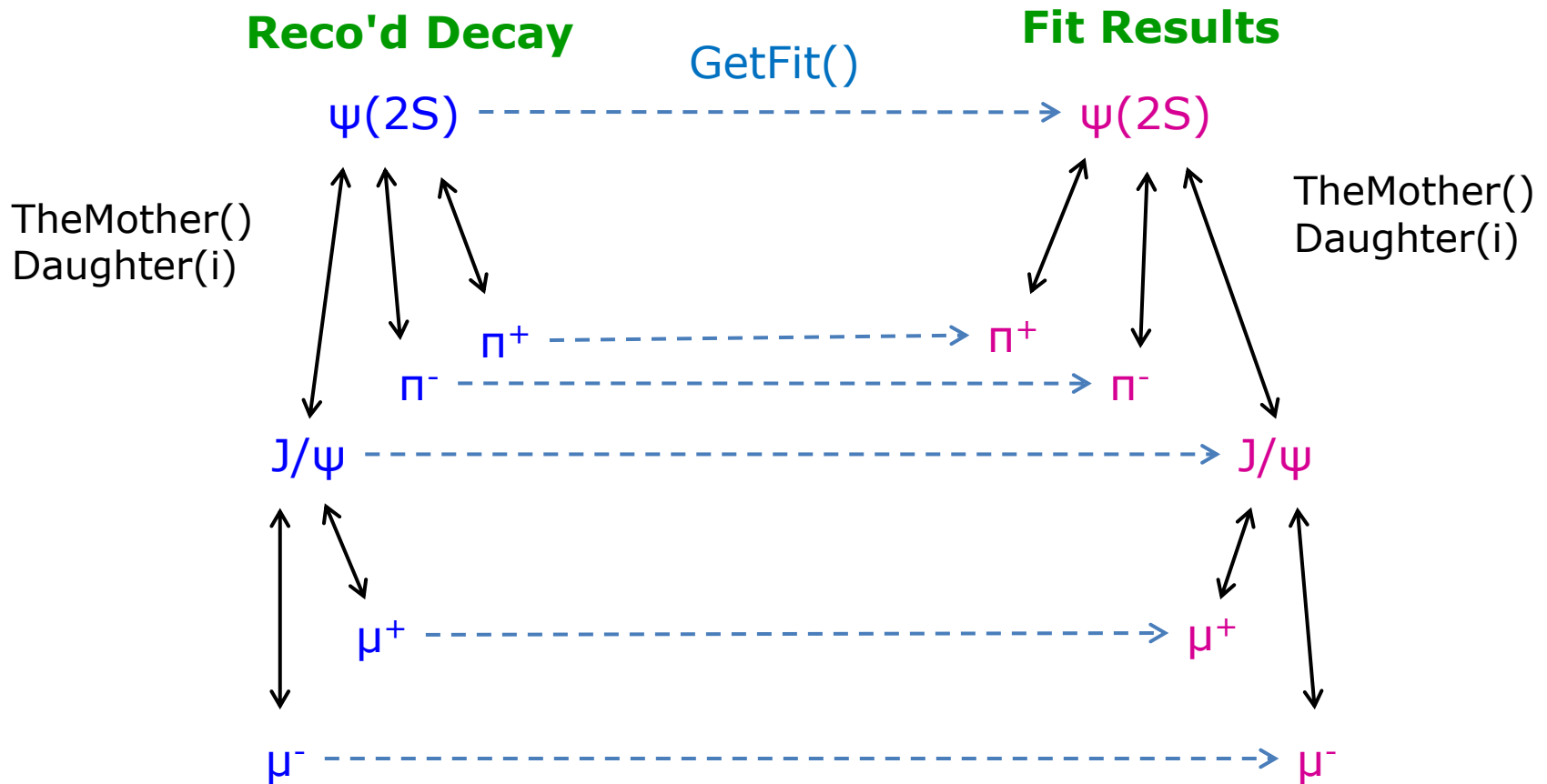


Per default, the track reco computes the 4-vectors at POCA to the IP (or beam axis)  
→ resulting 4-vector is bad!

⇒ Good vertex information necessary for proper resonance reco!

# Fitting: Access to Results

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





# Fitting in PandaROOT

- Fitters basically all have a similar interface
- Fit results are **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.Fit();                               // 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 Example: Vertex-Fit

- PandaROOT object: **PndKinVtxFitter**

```
// ... in event loop ...

for (j=0; j<jpsi.GetLength(); ++j)
{
    PndKinVtxFitter vtxfitter(jpsi[j]);           // instantiate vertex fitter
    vtxfitter.Fit();                             // perform fit

    RhoCandidate *jfit = jpsi[j]->GetFit();       // get fitted candidate

    TVector3 jVtx = jfit->Pos();                  // and the vertex position
    double chi2_vtx = vtxfitter.GetChi2();        // and the chi^2 of the fit

    if ( chi2_vtx<max_chi )                      // if chi2 is good enough
    {                                             // fill some histos
        hjpsim_vf->Fill( jfit->M() );
        hvpos->Fill( jVtx.X(),jVtx.Y() );
    }
}
```

# Fitting Example: 4C-Fit

- PandaROOT object: **PndKinFitter**

```
// the lorentz vector of the initial system; important for the 4C-fit

TLorentzVector ini(0, 0, 6.232, 7.240);

// ... in event loop ...
for (j=0;j<psi2s.GetLength();++j)
{
    PndKinFitter fitter(psi2s[j]);           // instantiate the kinematic fitter
    fitter.Add4MomConstraint(ini);           // set 4 vector constraint
    fitter.Fit();                           // perform fit

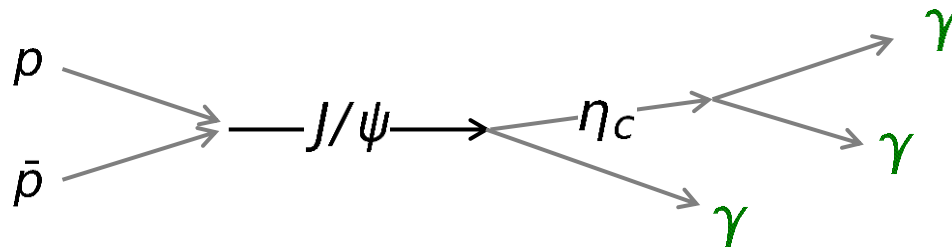
    RhoCandidate *jfit = psi2s[j]->Daughter(0)->GetFit(); // get fitted J/psi

    Double_t chi2 = fitter.GetChi2();        // and the chi^2 of the fit

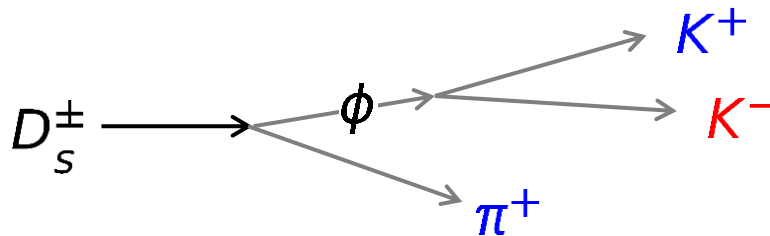
    if ( 0 != jfit )
    {
        hjpsim_4cf->Fill( jfit->M() );      // fill histogram
    }
}
```

# Which constraints do you see?

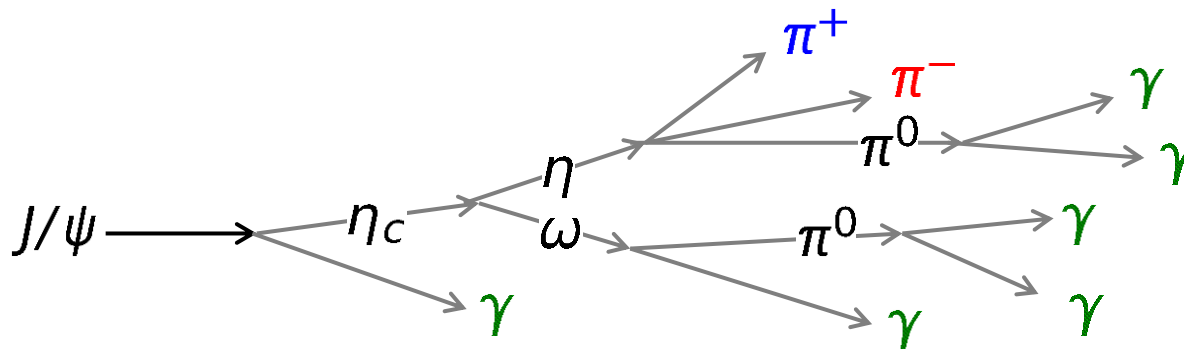
1



2



3

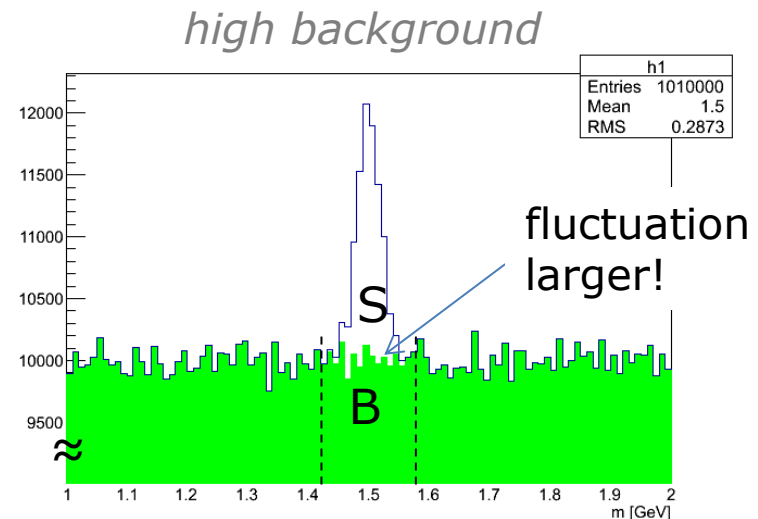
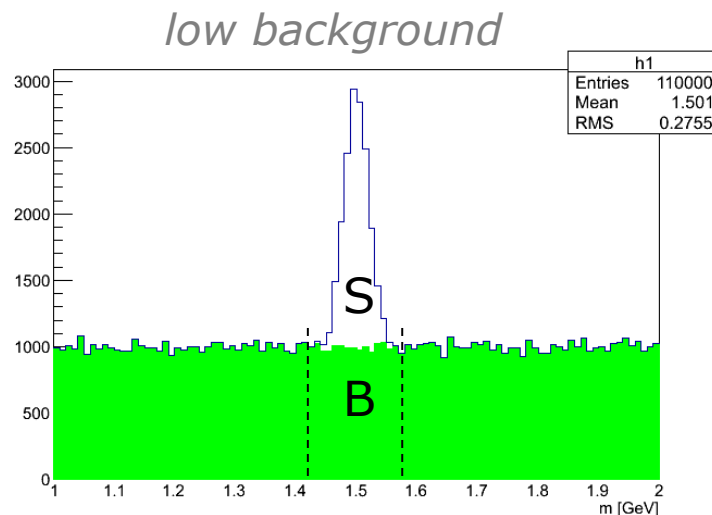


Particle	$\Gamma$ [MeV]
$J/\psi$	0.09
$\eta_c$	28.6
$D_s$	0
$\phi$	4.26
$\omega$	8.49
$\eta$	0.001
$\pi^0$	0

# UNCERTAINTIES

# Uncertainties: Figure of merit

- Figure of Merit for analysis: **Minimization of the relative error!** (*not, as you might think, the measured value itself...*)
- In cut & count analyses, usually maximize significance  $\frac{S}{\sqrt{S+B}}$
- Explanation: Statistical fluctuation of  $S$  is  $\sqrt{S+B}$  (*see below*)  
 $\rightarrow$  relative uncertainty =  $\frac{\sqrt{S+B}}{S} = 1/\text{significance!}$



# Statistic and Systematic Uncertainties

The most common kinds are

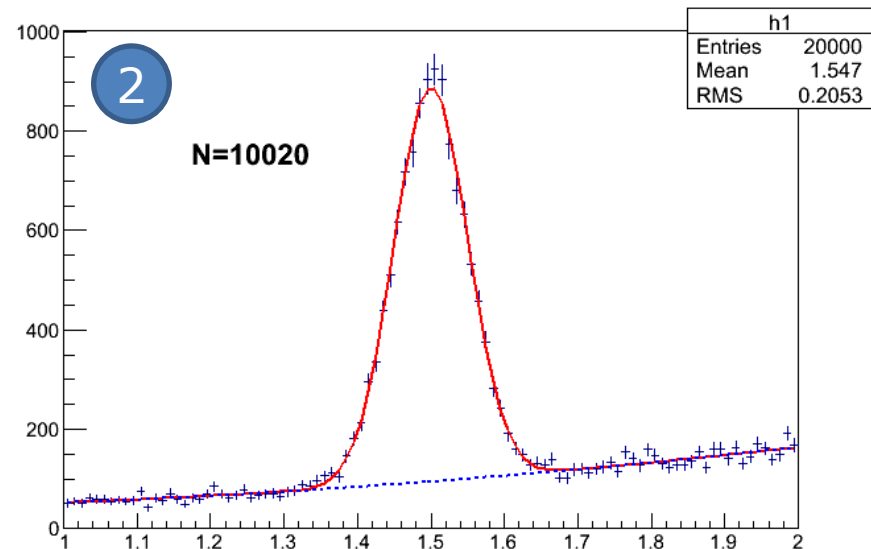
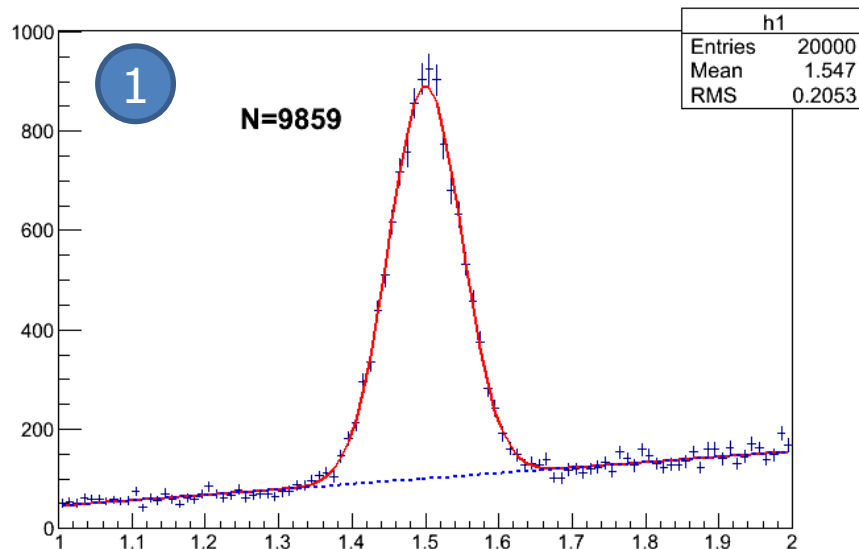
- **Statistic uncertainties**
  - Rule of thumb: Errors, which get smaller when taking more data are statistic errors. (*sometimes also sys. errors get smaller with more statistics*)
- **Systematic uncertainties**
  - Usually do not decrease with more data (when systematics dominate, you can stop measuring longer)
  - *Examples*
    - Track reconstruction efficiency (*BABAR: 1-2%/track*)
    - PID selection efficiency (*BABAR: 1-2%/particle*)
    - Choice of fit model (*e.g. 2<sup>nd</sup> vs. 3<sup>rd</sup> order bkg-polynomial*)

# Systematics Example: Fit model uncertainty

**Task:** Determine number of signals ontop of background

- *Bgk-Model 1:* Linear; integral  $N_1 = 9859$
- *Bgk-Model 2:* Parabolic; integral  $N_2 = 10020$
- Estimate for systematic error (fit model) could be

$$\Delta N_{sys,fit} = \frac{|N_1 - N_2|}{(N_1 + N_2)/2} = 1.6\%$$





# EXERCISES

# Exercises Suggestions

- Rho Tutorial website:  
<http://panda-wiki.gsi.de/cgi-bin/view/Computing/PandaRootRhoTutorial>
  - Take a look to [tutorials/thailand2017/README](#)  
[Exercises: #2 - #6](#)
1. Apply different PID selections and algorithms and compare
  2. Apply MC truth match for your reconstructed channels
  3. Try different fitters/constraints and compare resolution

# Exercises Preparation

Preparation/hints in [tutorials/thailand2017/README](#)

- FIRST setup environment: `source ~/workshop/build/PandaRoot_trunk.../config.sh`
- To have some data for tutorial macros, do one of the following
  - a) `./tut_runall.sh 1000` # sim/reco 1000 events pbarp -> J/psi pi+ pi-
  - b) `cp data/signal_p*root .` # preproduced default data
- Macros named 'tut\_ana...C' are stubs and should be completed by you.
- At places marked with '#### EXERCISE: ...' some code needs to be added;

Run macros (default or different input data) with

```
> root -l tut_ana...C # signal_pid.root, signal_par.root
> root -l 'tut_ana...C(0,"mydata")' # mydata_pid.root, mydata_par.root
```

- If getting stuck, sample solutions are in the subfolder 'solution'

Run solution macros directly (default or different input data) with

```
> root -l solution/tut_ana...C
> root -l 'solution/tut_ana...C(0,"mydata")'
```