

Updates on Track Fitting validation

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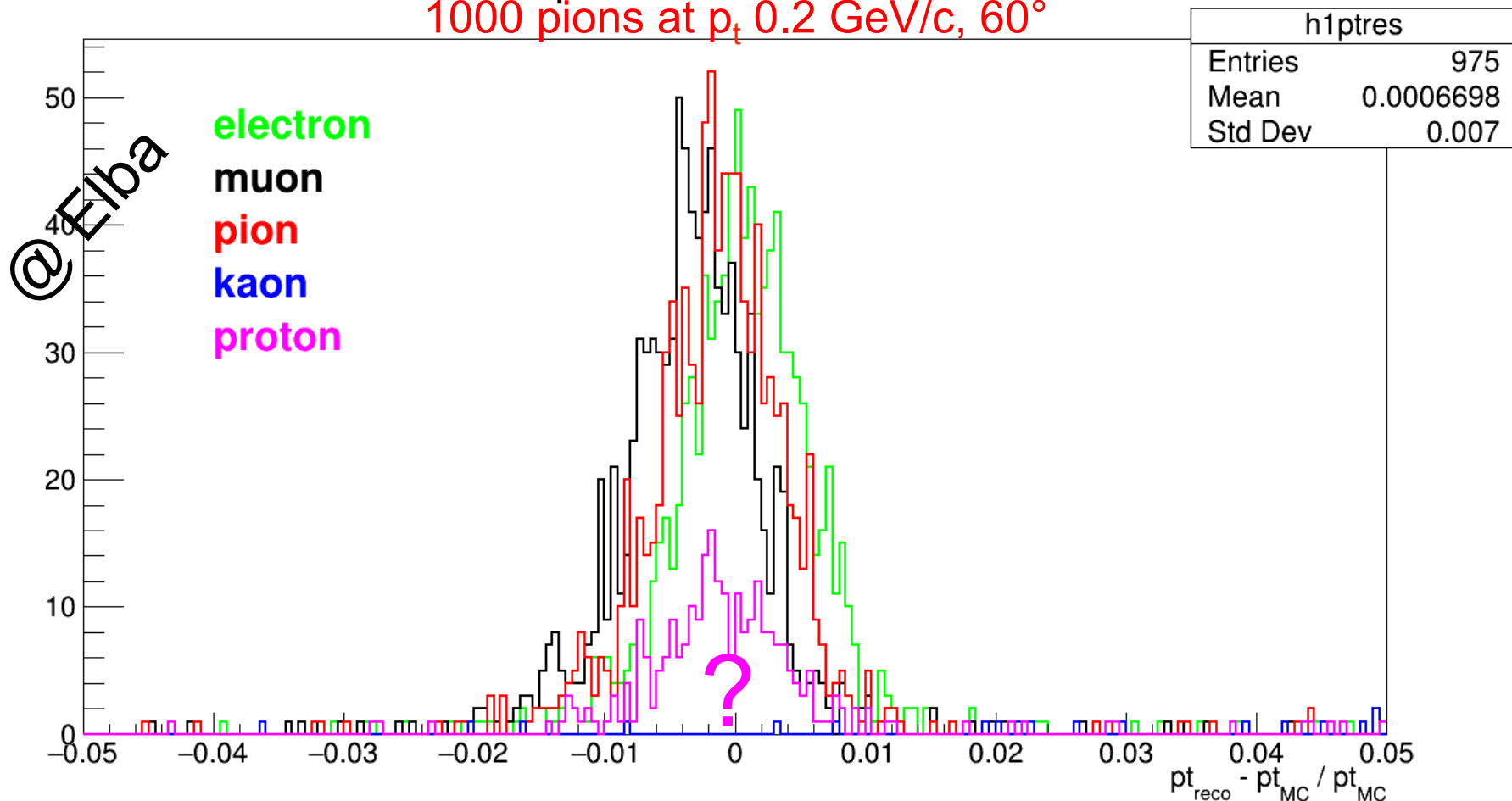
7.5 m

~ 7 m



Friday, 2nd June 2017

pt relative residuals
1000 pions at p_t 0.2 GeV/c, 60°

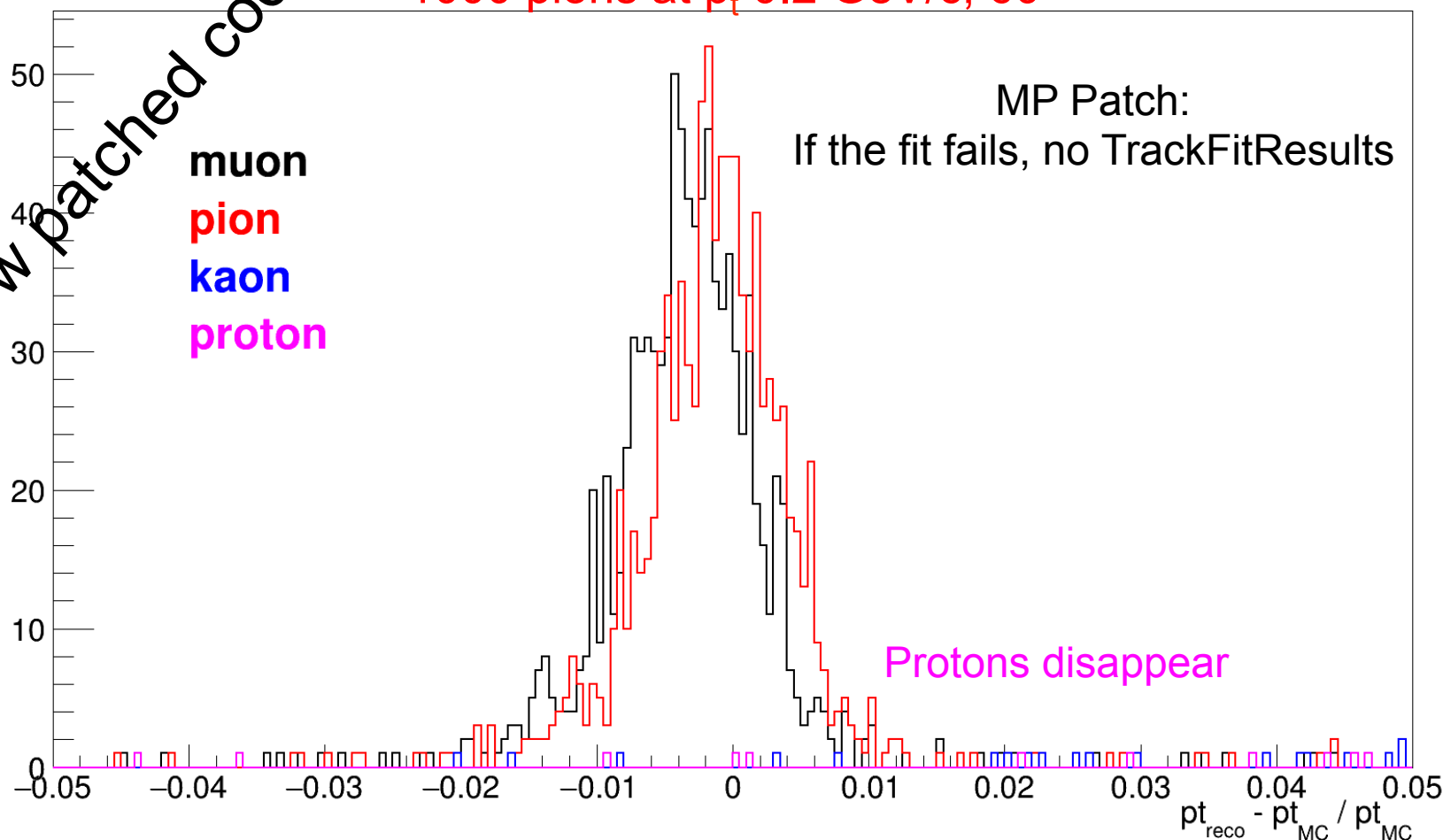


Q: Why proton residuals are peaked at the pion position?

A: If the fit fails, TrackFitResults returns the default pdg (pion) parameters and not the chosen one

@ new patched code

pt relative residuals
1000 pions at p_t 0.2 GeV/c, 60°



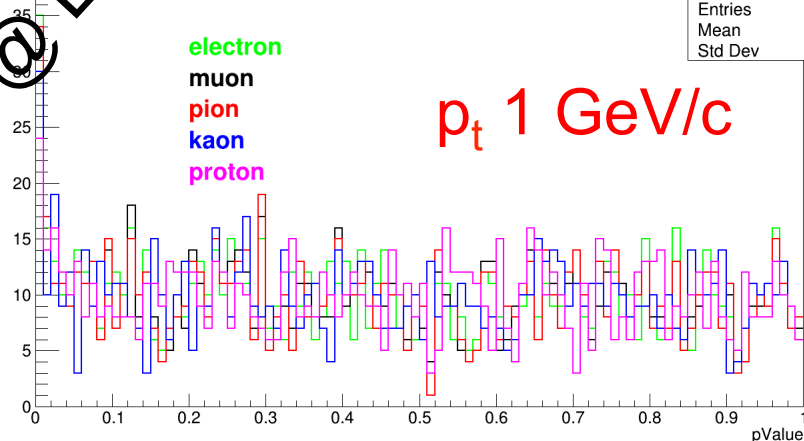
Just to confirm with plots what was said during the workshop

1000 pions at 60° - pValue

@Elba

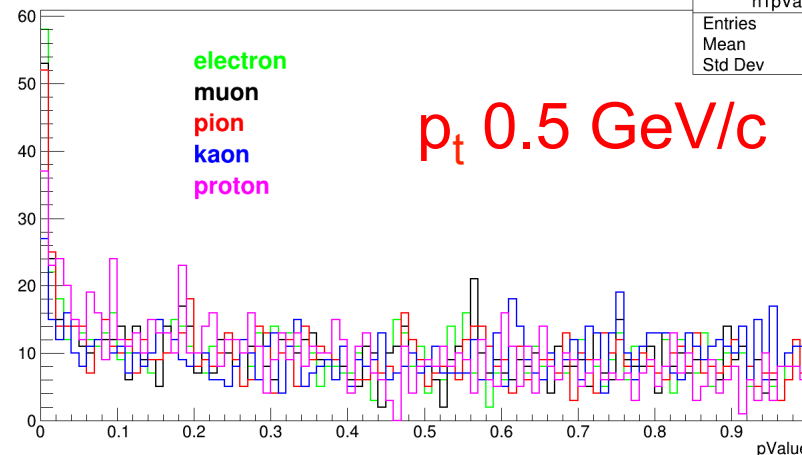
pValue of the fit

h1pValue	
Entries	991
Mean	0.4679
Std Dev	0.2977



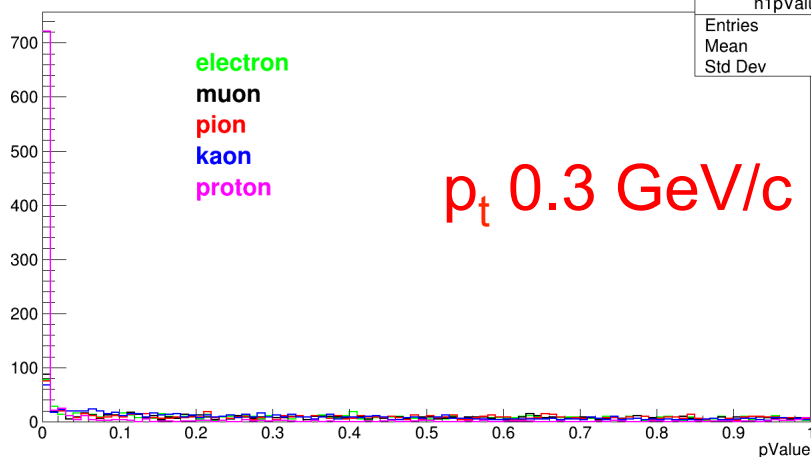
pValue of the fit

h1pValue	
Entries	992
Mean	0.4337
Std Dev	0.303



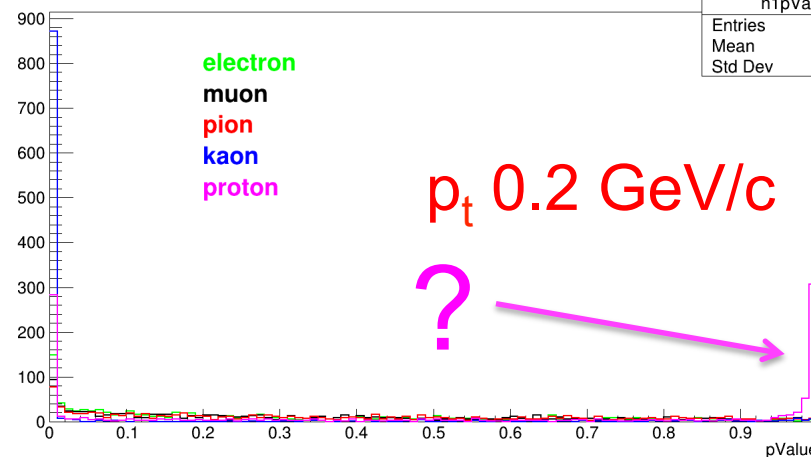
pValue of the fit

h1pValue	
Entries	887
Mean	0.3923
Std Dev	0.3033

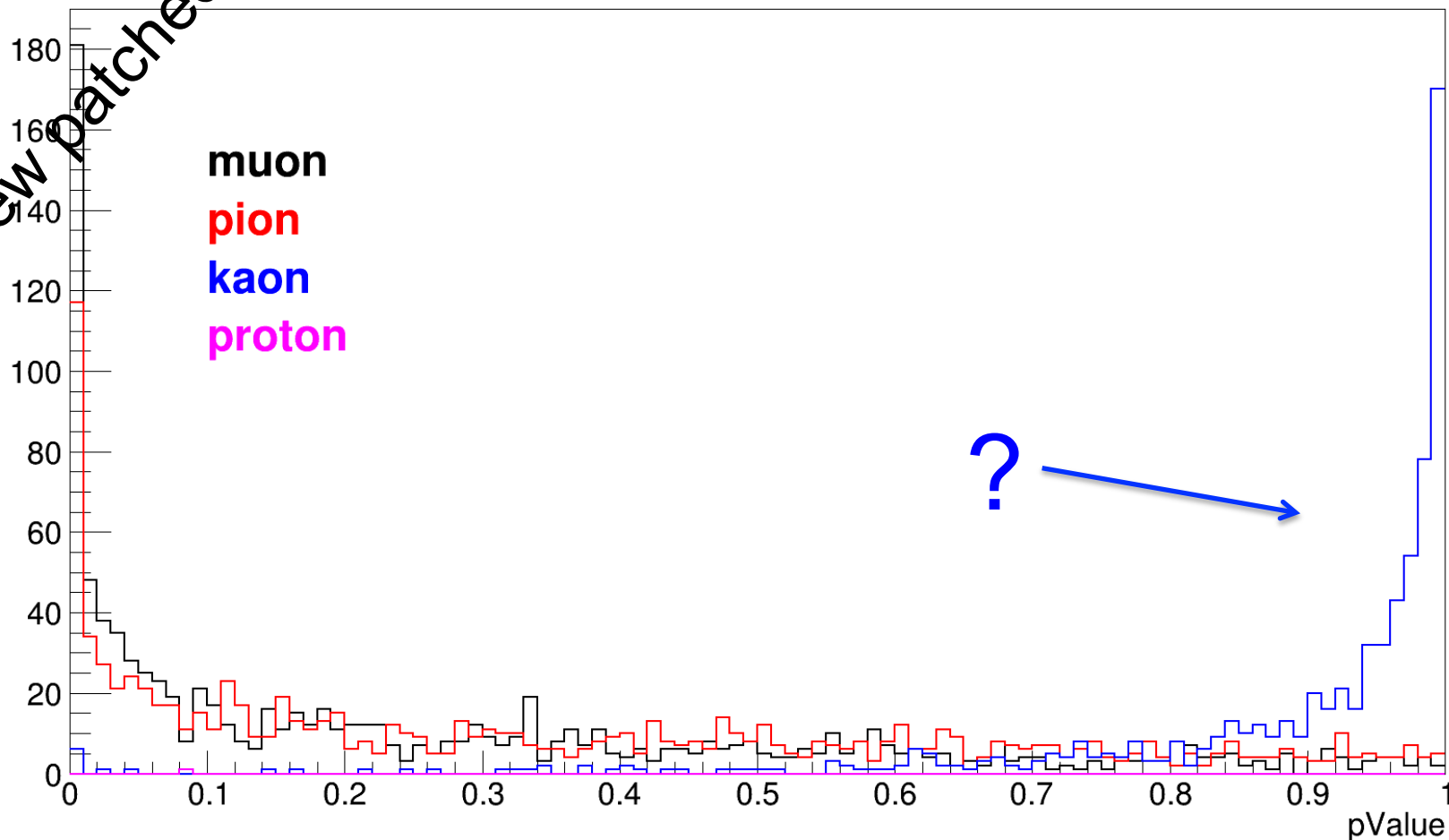


pValue of the fit

h1pValue	
Entries	975
Mean	0.2821
Std Dev	0.2806



@ new patched code 1000 pions at p_t 0.1 GeV/c, 60°
pValue of the fit



Genfit stores inside **FitStatus** object:

- χ^2
- Number of degrees of freedom (NDF)

pValue is calculated from χ^2 and NDF

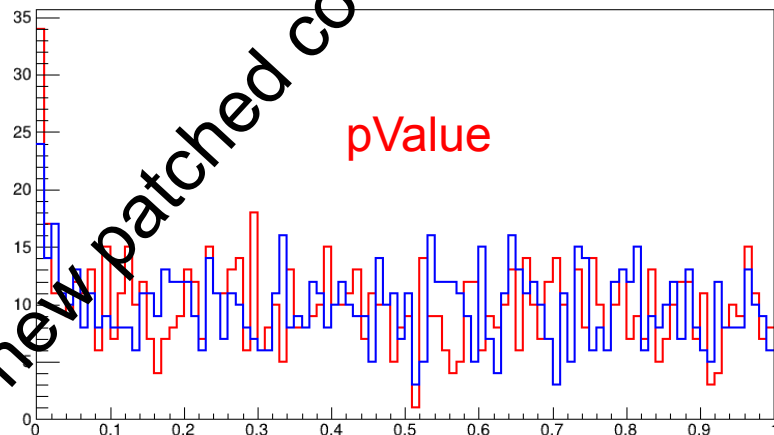
```
double getChi2() const {return chi2_;}  
//! Get the degrees of freedom of the fit.  
double getNdf() const {return ndf_;}  
/**  
 * @brief Get the p value of the fit.  
 *  
 * Virtual, because the fitter may use a different probability distribution.  
 */  
virtual double getPVal() const {return std::max(0.,ROOT::Math::chisquared_cdf_c(chi2_, ndf_));}
```

In **TrackFitResults** we store **ONLY pValue** (TrackBuilder.cc)

I have added few new data members to TrackFitResults as a crosscheck

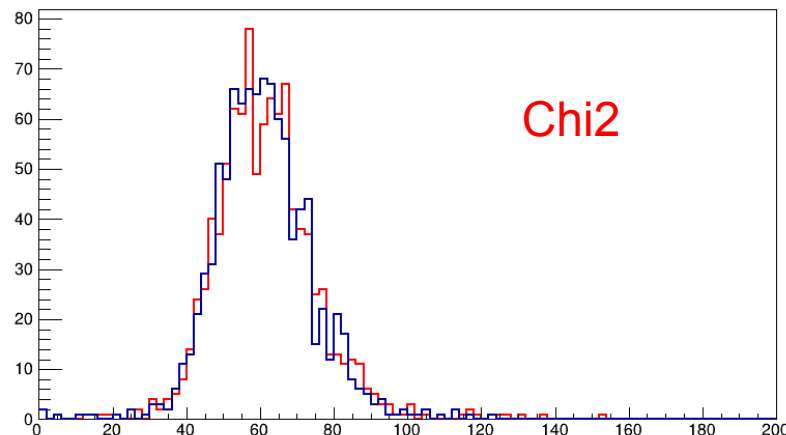
@ new patched code

TrackFitResults.m_pValue {TrackFitResults.m_pdg==211}



pValue

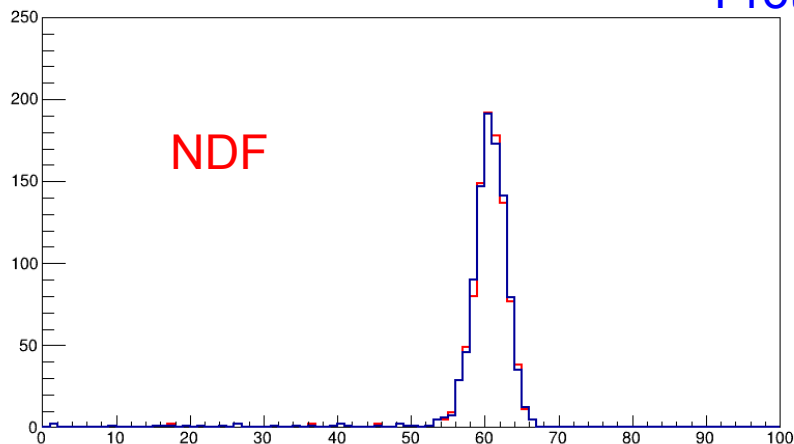
TrackFitResults.m_Chi2 {TrackFitResults.m_pdg==211}



Chi2

Pion hypothesis

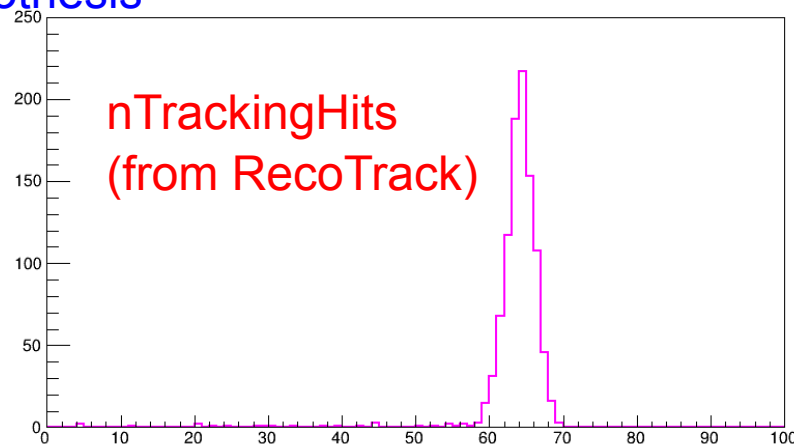
TrackFitResults.m_Ndf {TrackFitResults.m_pdg==211}



NDF

Proton hypothesis

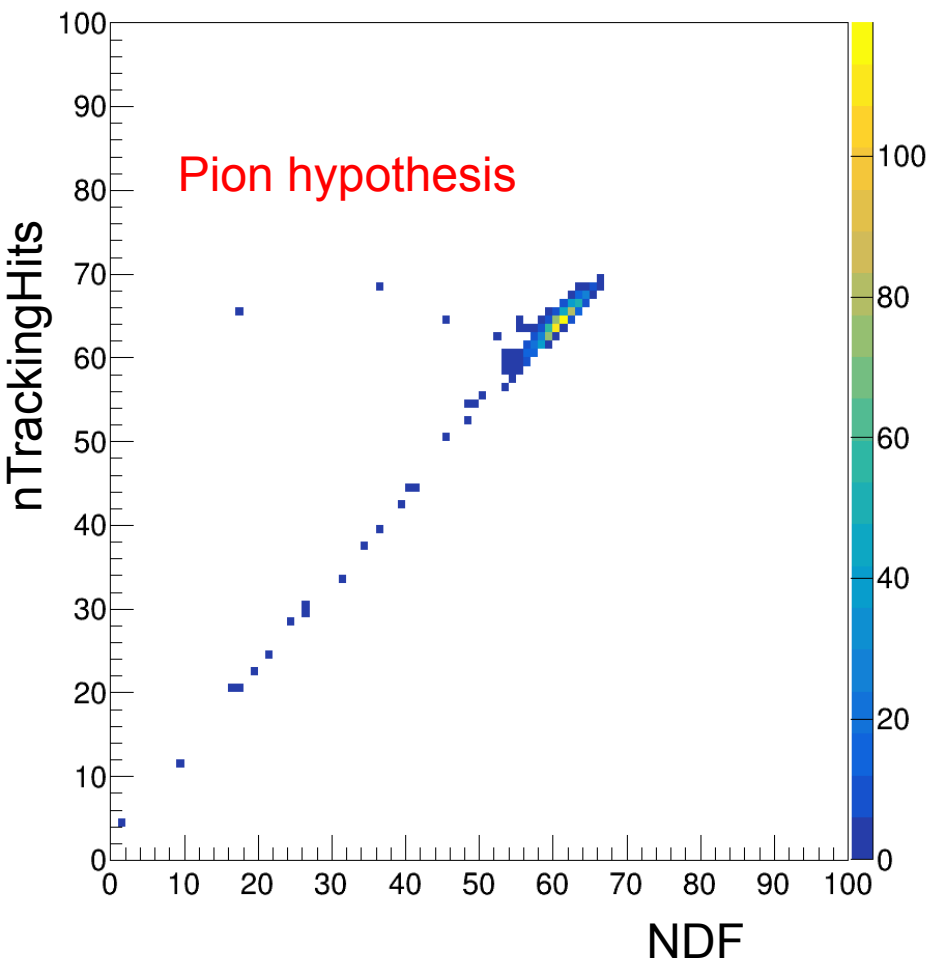
TrackFitResults.m_nTrackingHits {TrackFitResults.m_pdg==211}



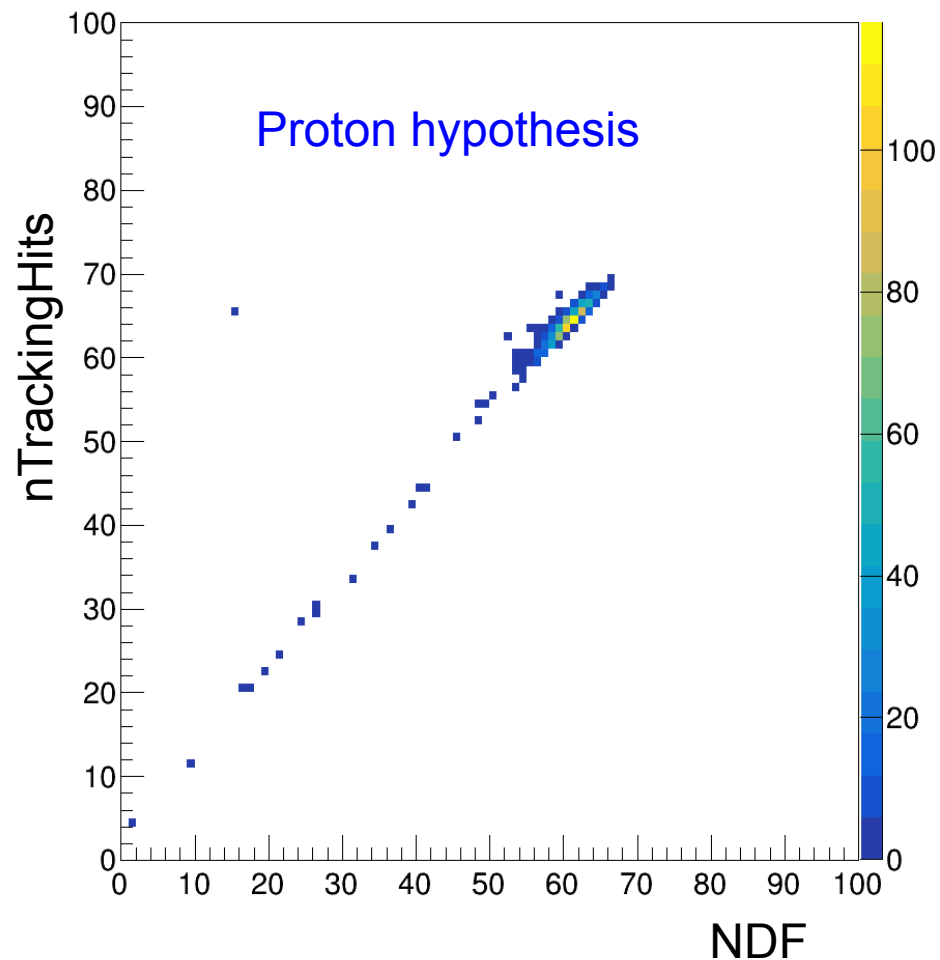
nTrackingHits
(from RecoTrack)

OK, Everything Understood

TrackFitResults.m_nTrackingHits:TrackFitResults.m_Ndf {TrackFitResults.m_pdg==211}

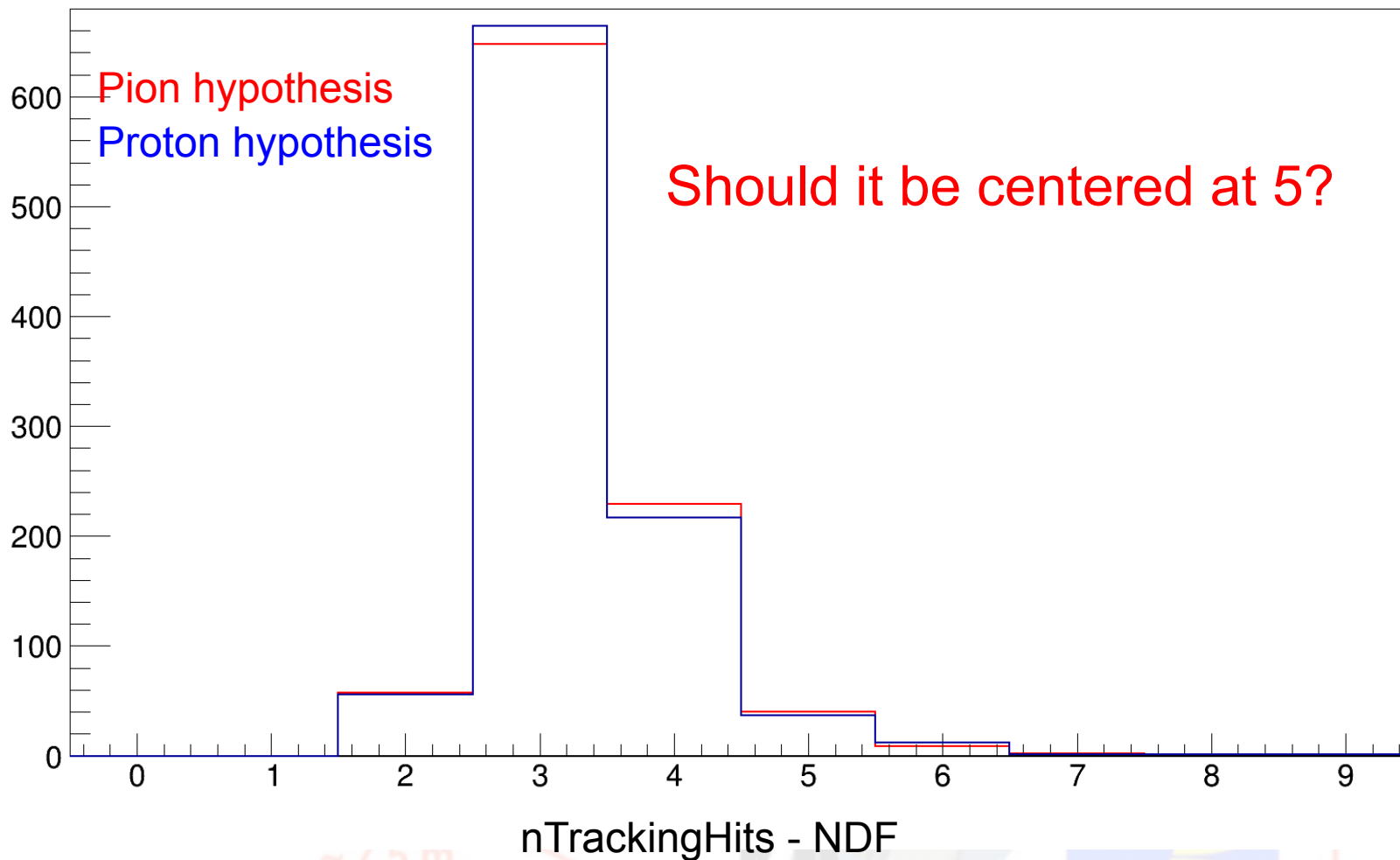


TrackFitResults.m_nTrackingHits:TrackFitResults.m_Ndf {TrackFitResults.m_pdg==2212}



OK, Everything Understood

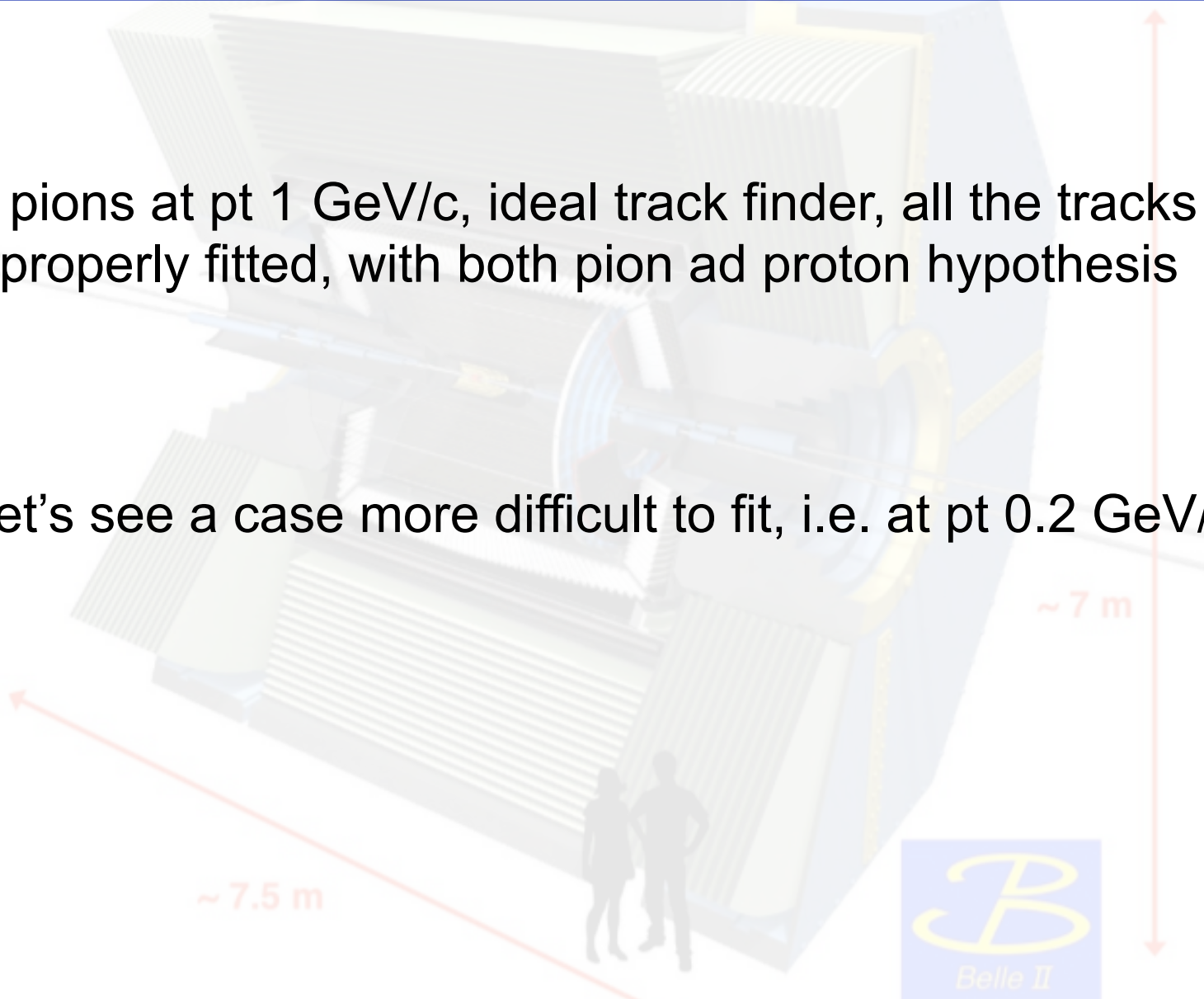
TrackFitResults.m_nTrackingHits-TrackFitResults.m_Ndf {TrackFitResults.m_pdg==211}



Almost Understood

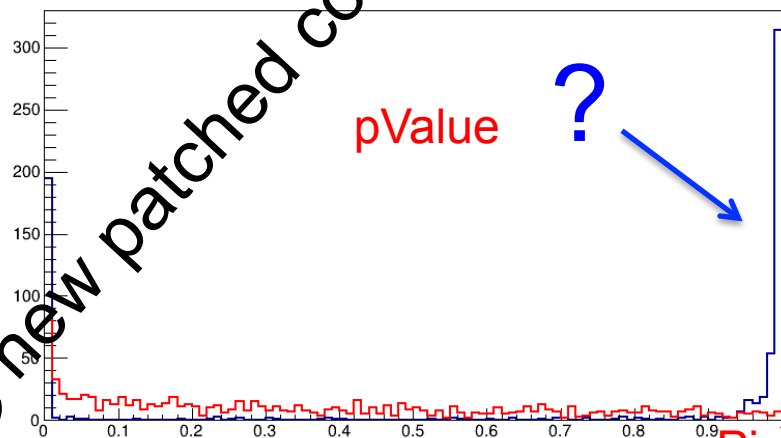
With pions at pt 1 GeV/c, ideal track finder, all the tracks are properly fitted, with both pion and proton hypothesis

Let's see a case more difficult to fit, i.e. at pt 0.2 GeV/c



@ new patched code

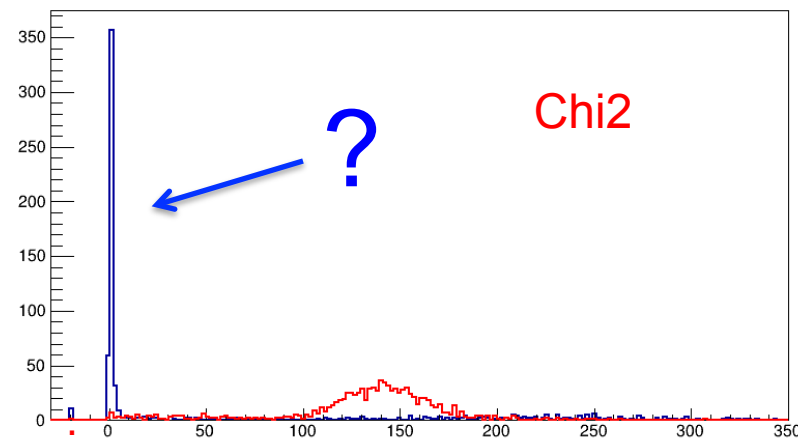
TrackFitResults.m_pValue {TrackFitResults.m_pdg==2212}



pValue

?

TrackFitResults.m_Chi2 {TrackFitResults.m_pdg==2212}



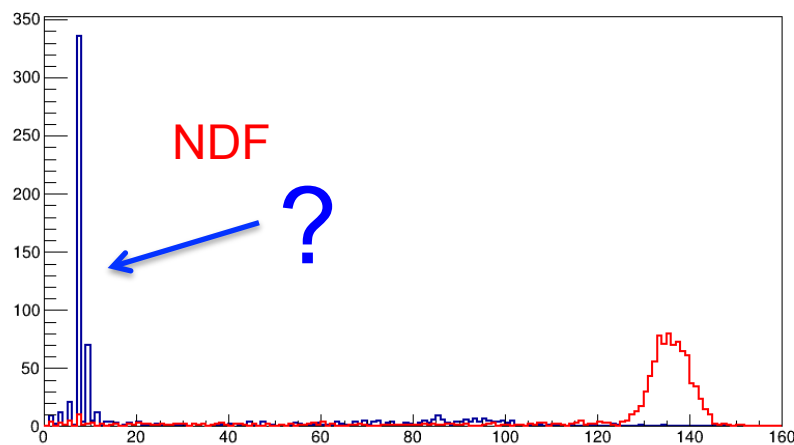
Chi2

?

Pion hypothesis

Proton hypothesis

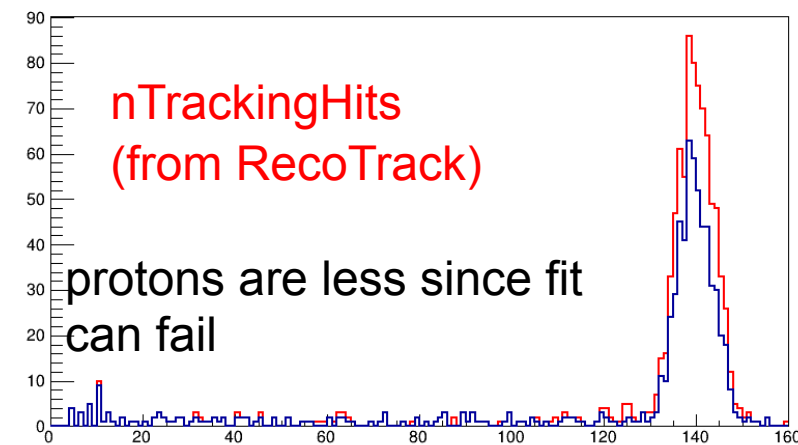
TrackFitResults.m_Ndf {TrackFitResults.m_pdg==2212}



NDF

?

TrackFitResults.m_nTrackingHits {TrackFitResults.m_pdg==211}

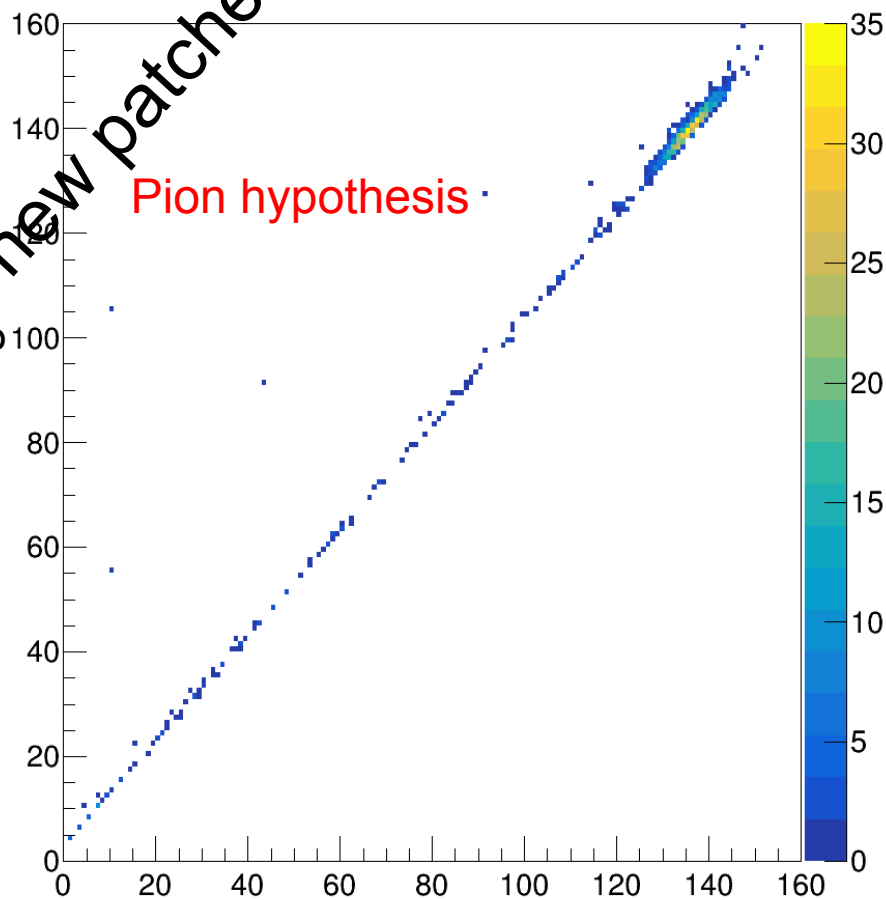


nTrackingHits
(from RecoTrack)

protons are less since fit
can fail

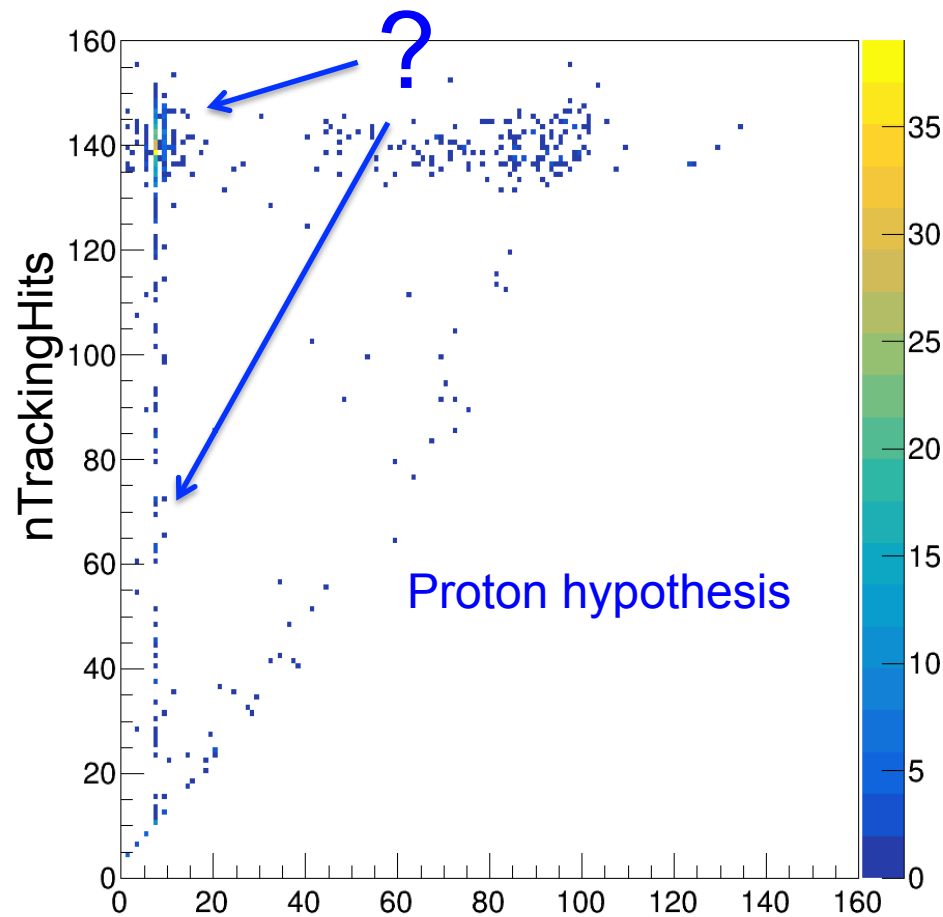
new patched code

TrackFitResults.m_nTrackingHits:TrackFitResults.m_Ndf (TrackFitResults.m_pdg==211)



NDF

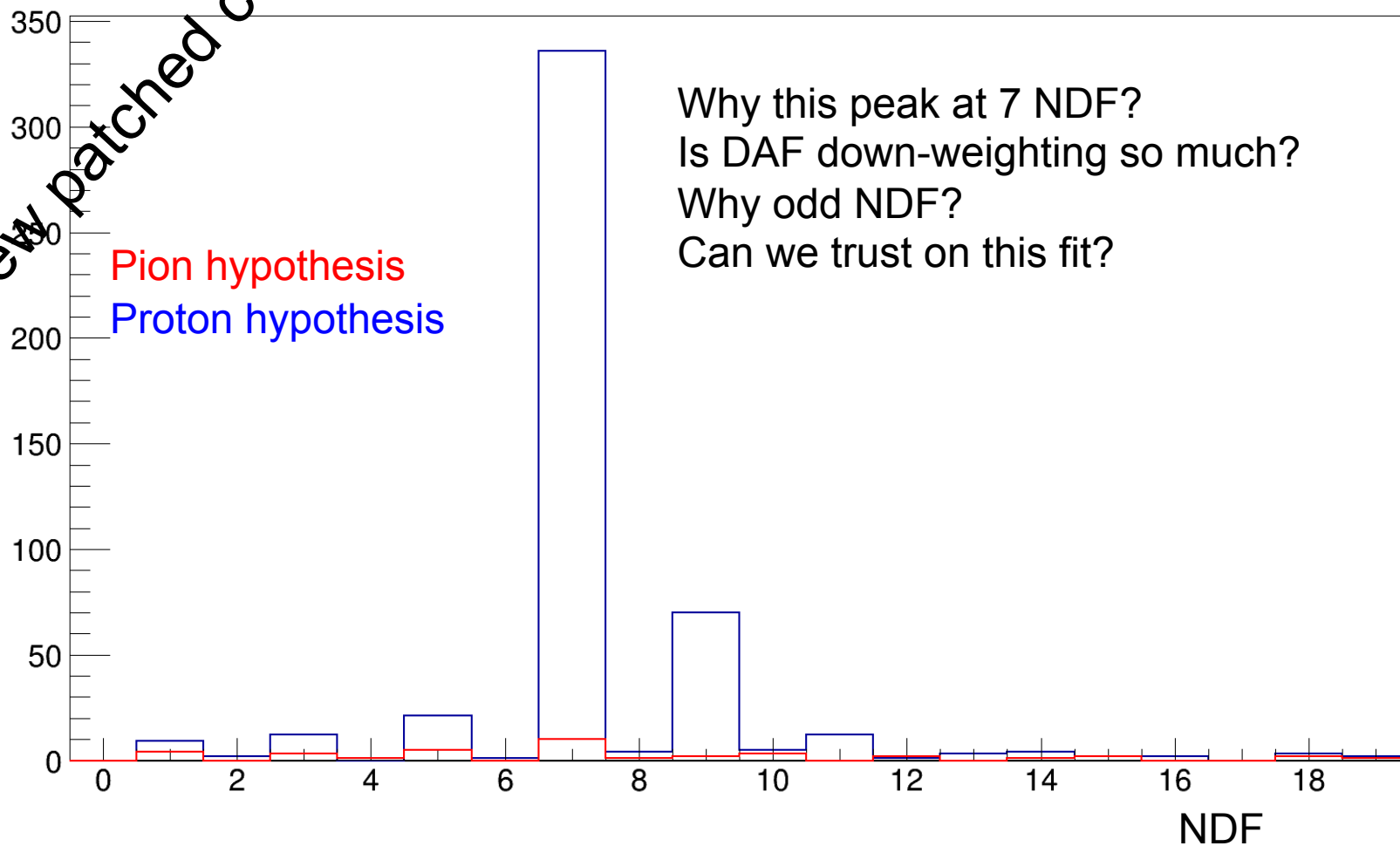
TrackFitResults.m_nTrackingHits:TrackFitResults.m_Ndf (TrackFitResults.m_pdg==2212)



NDF

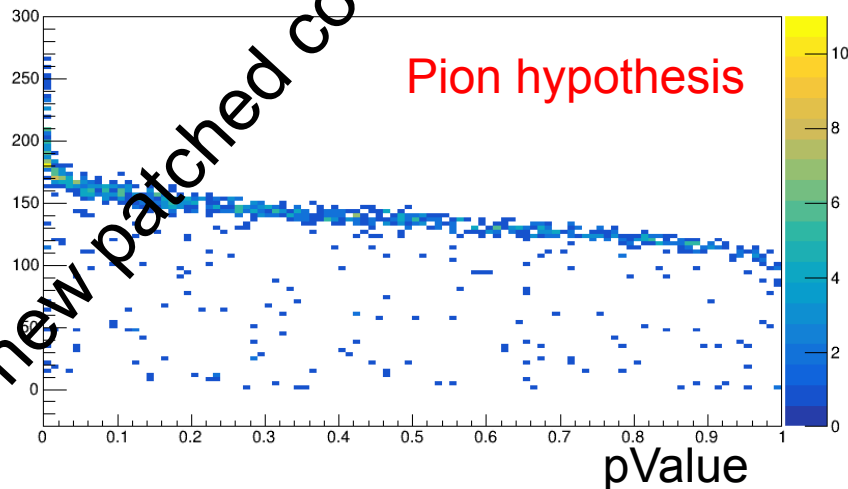
TrackFitResults.m_Ndf {TrackFitResults.m_pdg==2212}

@ new patched code

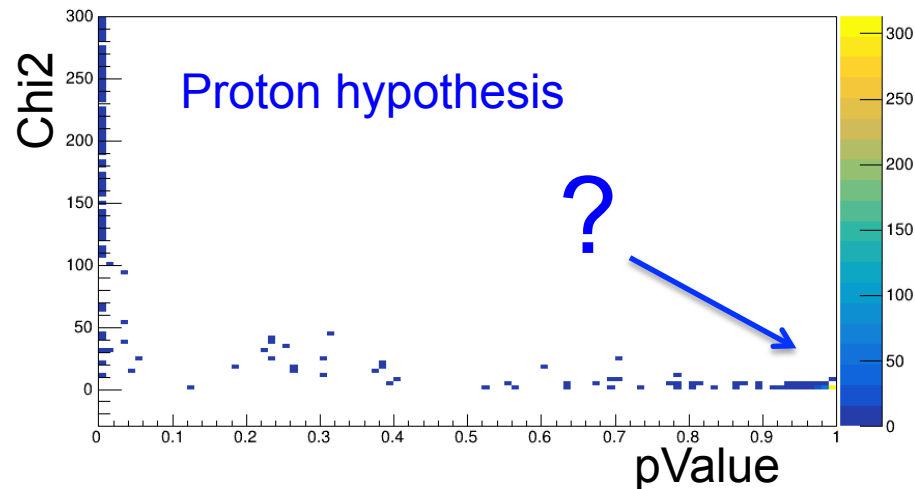


@ new patched code

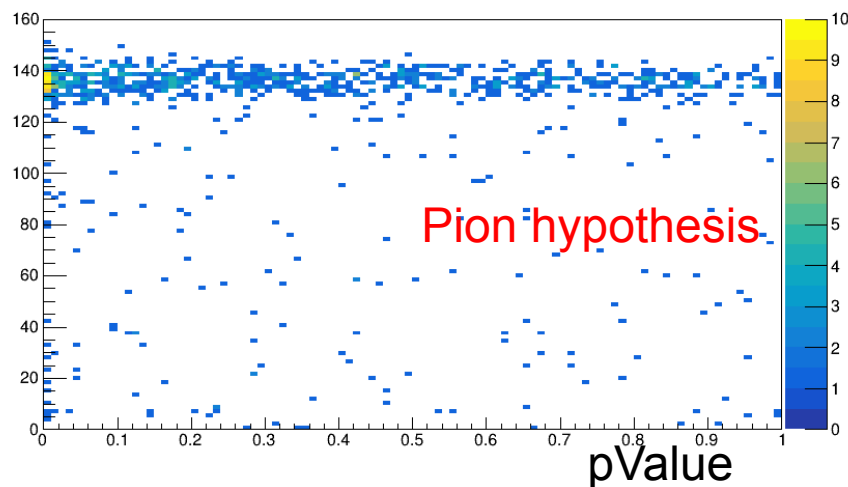
TrackFitResults.m_Chi2:TrackFitResults.m_pValue {TrackFitResults.m_pdg==211}



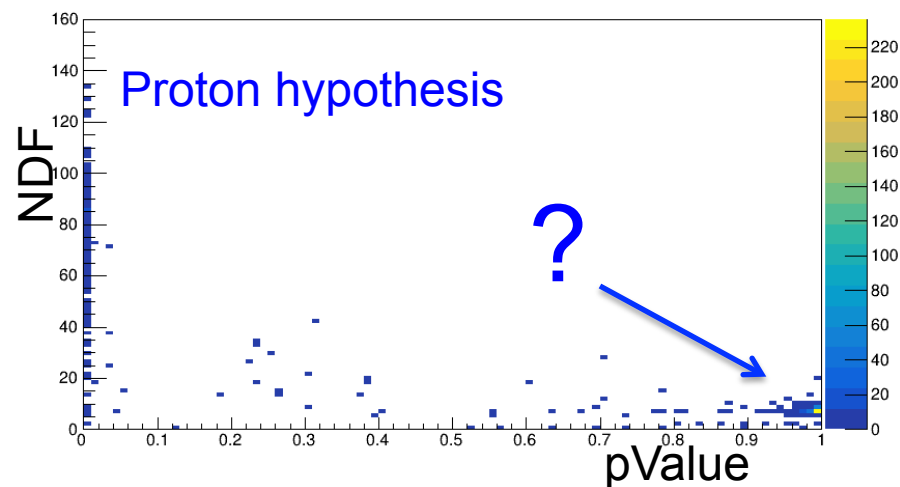
TrackFitResults.m_Chi2:TrackFitResults.m_pValue {TrackFitResults.m_pdg==2212}



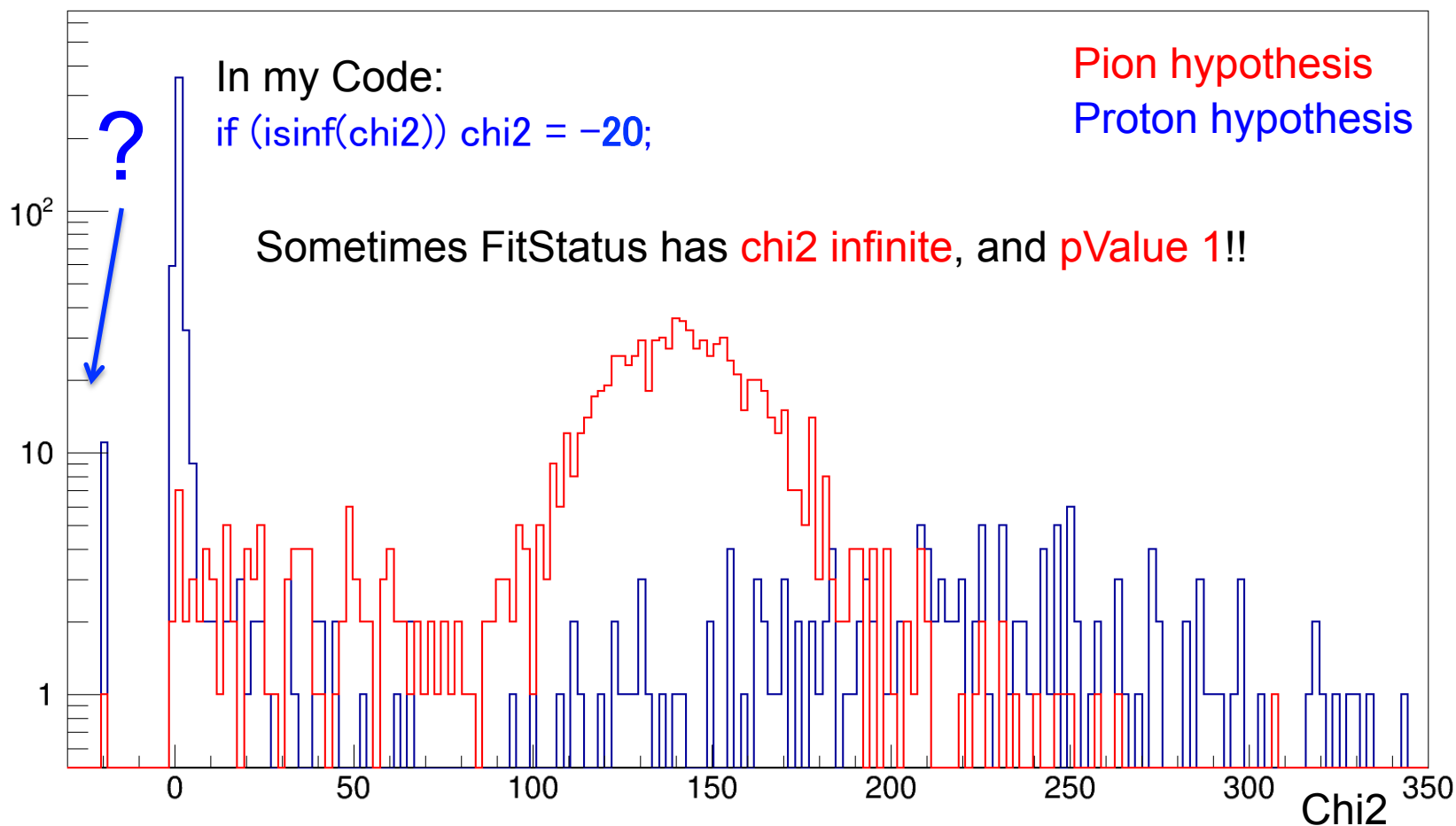
TrackFitResults.m_Ndf:TrackFitResults.m_pValue {TrackFitResults.m_pdg==211}



TrackFitResults.m_Ndf:TrackFitResults.m_pValue {TrackFitResults.m_pdg==2212}



TrackFitResults.m_Chi2 {TrackFitResults.m_pdg==2212}



Sometimes FitStatus returns a “infinite” chi2, which is wrongly associated to pValue 1. This could be easily caught by TrackBuilder.

Very bad fits (i.e. low pt with wrong particle hypothesis) sometimes return a very low number of degrees-of-freedom, I believe not physical, which are associated to pValue = 1. Why NDF 7? Why odd NDF? Maybe one could check nTrackingHits/NDF to estimate the average weight and cut?

It is not clear to me why nTrackingHits-NDF for good fits is peaked at 3.

Maybe storing only pValue in TrackFitResults is not safe enough to catch fit artifacts.

Check real pattern recognition, where tracks can have spurious hits and low pt track candidates could be shorter, without too many curls.

Take a look into eventdisplay, to understand how the bad fits look like.

Fast check of fitting with deuterons.

Any hints?

