

# GENFIT 2

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Belle II F2F Tracking Meeting, Dec 11th to 13th, 2013



Bundesministerium  
für Bildung  
und Forschung





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# New Features



## General:

- More user friendly code (I hope).
- Better data-structure: TrackPoints with Measurements and FitterInfos instead of GFBookkeeping.
- Full Tracks need more memory/disk space, but pruning function allows to strip off unneeded data.
- Easy track merging, adding and removing hits, fast refitting.
- Built-in consistency checks to spot and avoid errors.
- Time of flight calculation.
- Improved visualization.
- Easier Measurement implementation, can use H-Matrices provided by GENFIT.
- H-Matrix for slanted strips.



## Fitters:

- FitStatus objects in track (which fitter was used, number of iterations, fit converged?, NdF,  $\chi^2$ , p-values).
- Kalman filter with reference track, which can also re-sort TrackPoints.
- Possibility to fit tracks partially (Kalman).
- optional square-root formalism (Kalman).
- Possibility to save ThinScatterers in TrackPoints (not yet used, but useful for GBL).

## Configurable Convergence Criteria:

- Minimum and maximum number of iterations.
- Converged if:
  - $\Delta pVal < 1E - 3$
  - relative change in  $\chi^2 < 20 \%$
  - (default values, configurable)



## TrackReps:

- More extrapolation options, possible to stop at material boundaries.
- Get list of materials crossed in last extrapolation.
- Get path / radiation length crossed in last extrapolation.
- Get time of flight of last extrapolation.

## TrackCand:

- TrackCandHits, from which one can inherit and save all necessary information (already in GENFIT 1).
- In the new MeasurementFactory, pointers to the TrackCandHits are provided to the Measurement (fka RecoHit) constructors, so these information can be used to construct the measurements.

This is now used to set the left/right ambiguity of the CDCRecoHits directly in the CDCRecoHit constructor.

# Migration Status

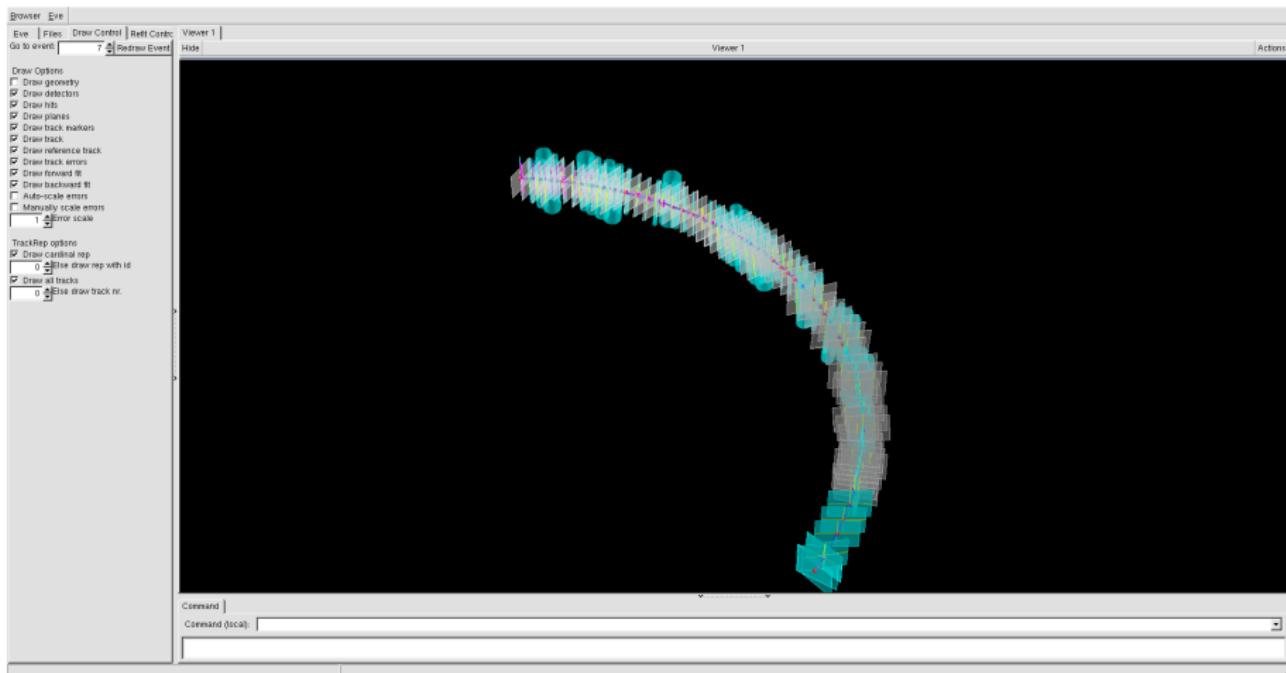


## Migration completed:

- **genfit2** as package in basf2 (via svn:externals).
- New externals v00-05-00 w/o GENFIT.
- Doxygen:  
<https://belle2.cc.kek.jp/internal/software/development/namespacegenfit.html>.
- Problem: svn:externals don't work well with gitsvn, can be worked around.
- Still some problems and crashes: work in progress.

## New Module:

- **GenfitVisModule** passes genfit::Tracks to genfit::EventDisplay.
- Visualize tracks, refit tracks, change fitters and settings, activate debug output, ...



# Validation Status



## Standard Validation:

- Validation scripts to check pull distributions ( $x, y, z, p_x, p_y, p_z$ ), p-value distributions, resolutions and efficiency.
- PXD + SVD + CDC fitting with GENFIT's Kalman filter and RKTrackRep.
- 1.000 tracks with fixed momentum and theta.
- Nightly results at: <https://belle2.cc.kek.jp/internal/validation/index.html>

## Advanced Options:

- In the steering script (`01_steering_genfitStudy.py`), various selections can be made:
  - Turn off/on PXD, SVD, CDC separately.
  - Switch the fitter.
  - Turn on the full test, 100.000 tracks over the whole phase-space.



## GENFIT 2

- Test results look very similar to GENFIT 1 results.
- Pull distributions also look very similar (I showed wrong plots on B2GM).
- Slightly more tracks with p-value  $\simeq 0$ .

## GenFitter module comparison

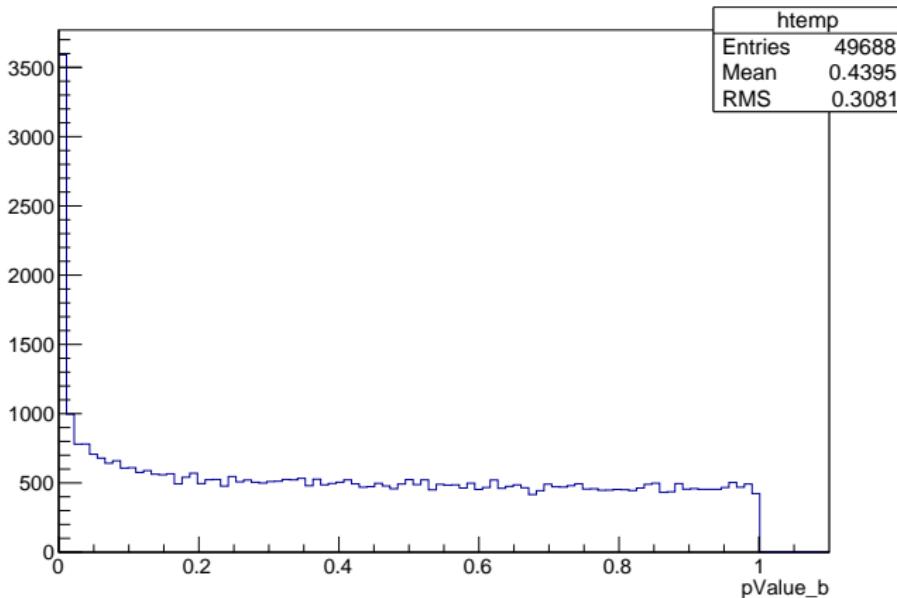
- Pions, 0.4 GeV,  $\theta = 120^\circ$ , pxd + svd + cdc.
- GENFIT 1: 25.168 ms/call (3 iterations)
- GENFIT 2: 25.108 ms/call (3.2 iterations on average)



## GENFIT 1 vs. GENFIT 2: p-Value Distributions



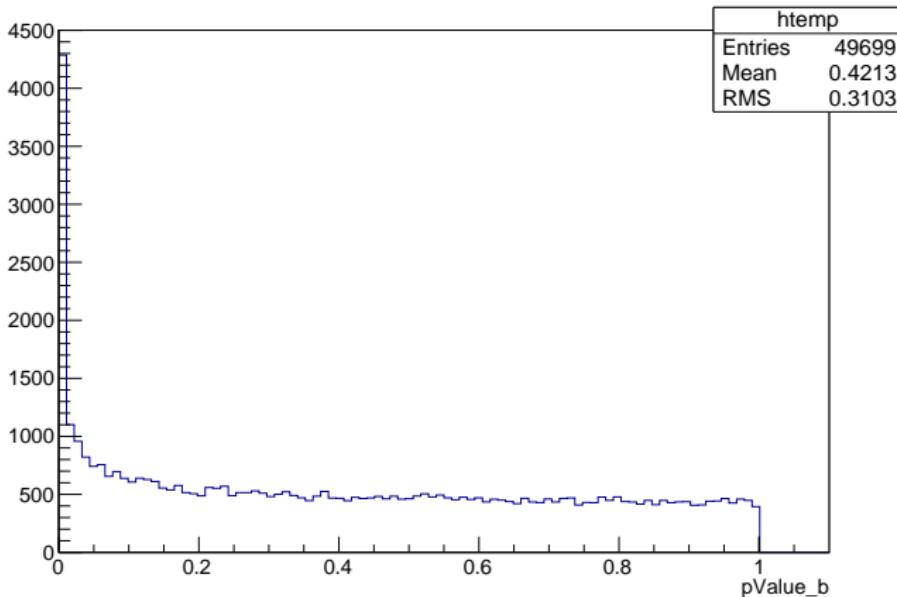
50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: p-Value Distributions

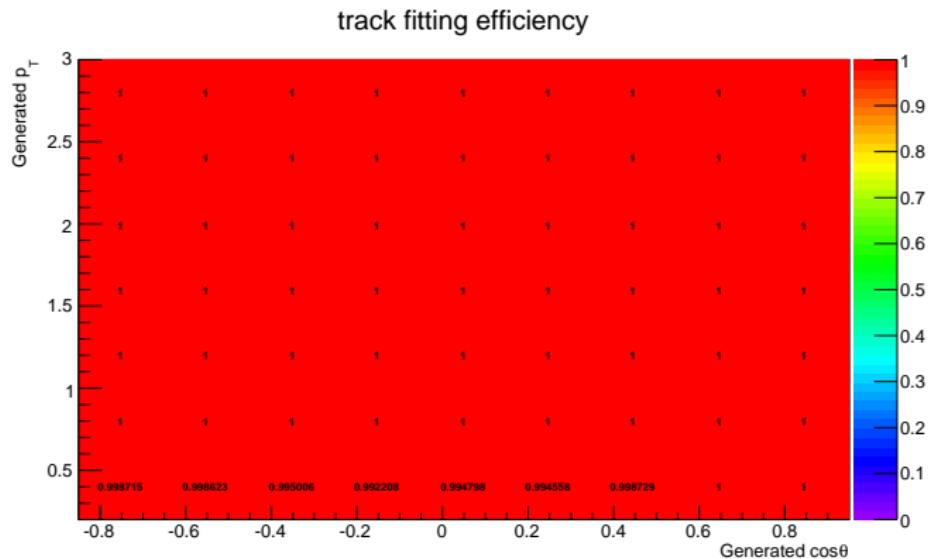
50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

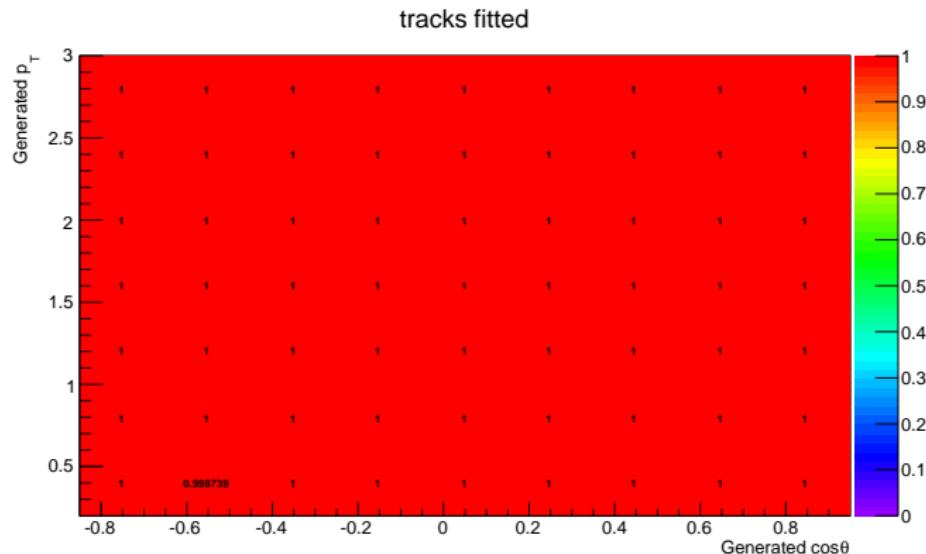


GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



GENFIT 2 with  $\geq 3$  iterations.

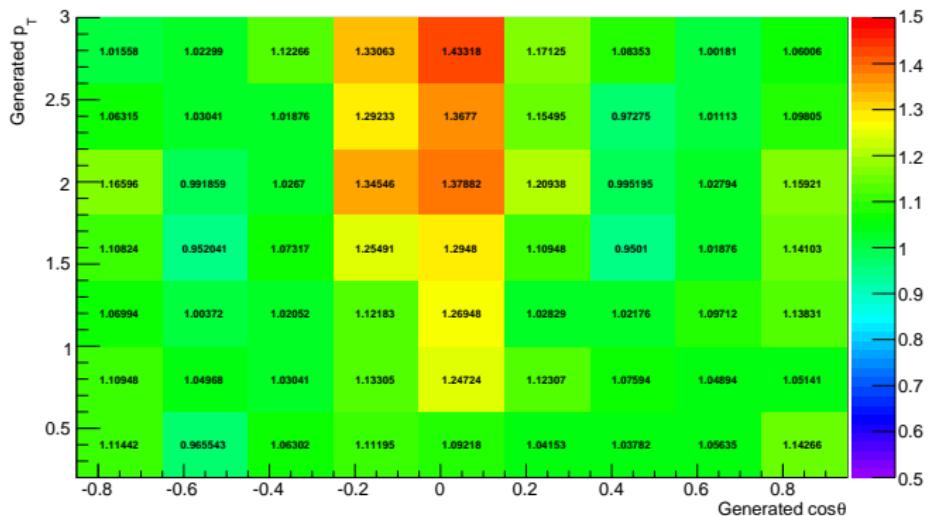


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of z pull distribution



GENFIT 1 with 3 iterations.

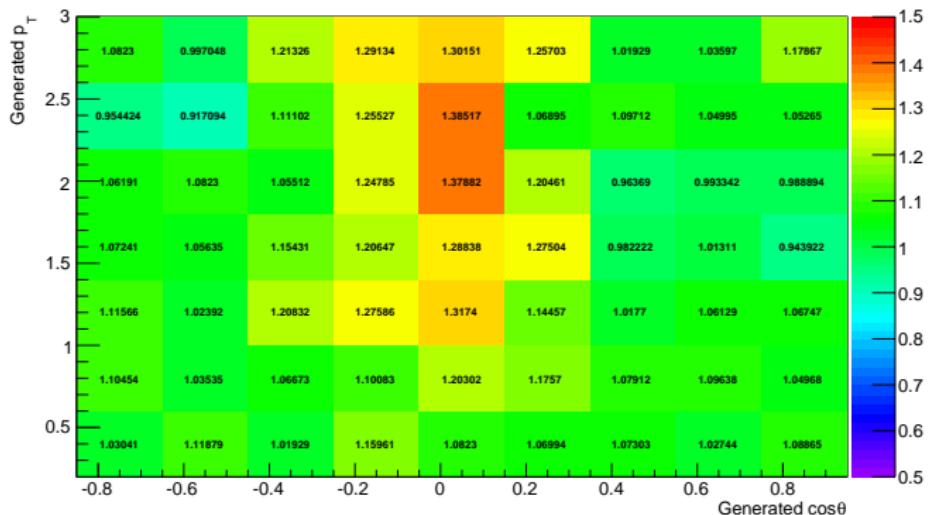


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of z pull distribution



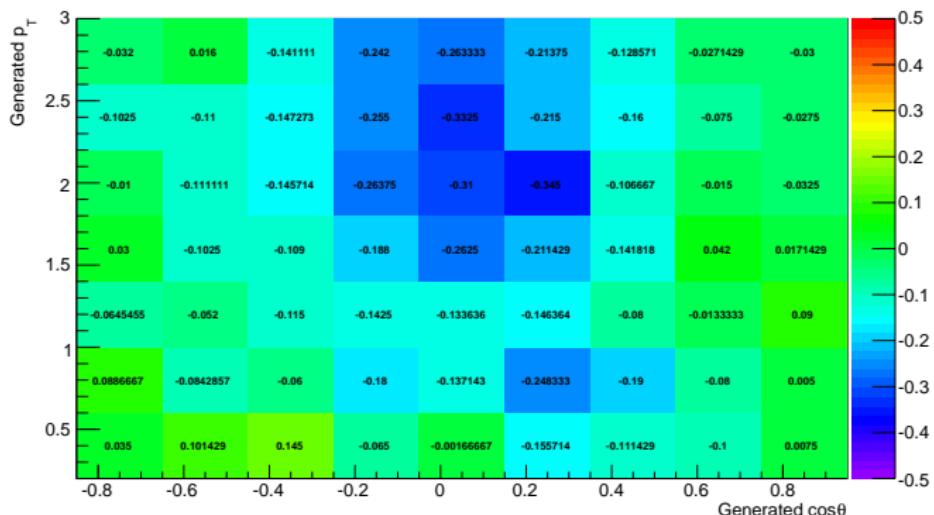
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of pz pull distribution



GENFIT 1 with 3 iterations.

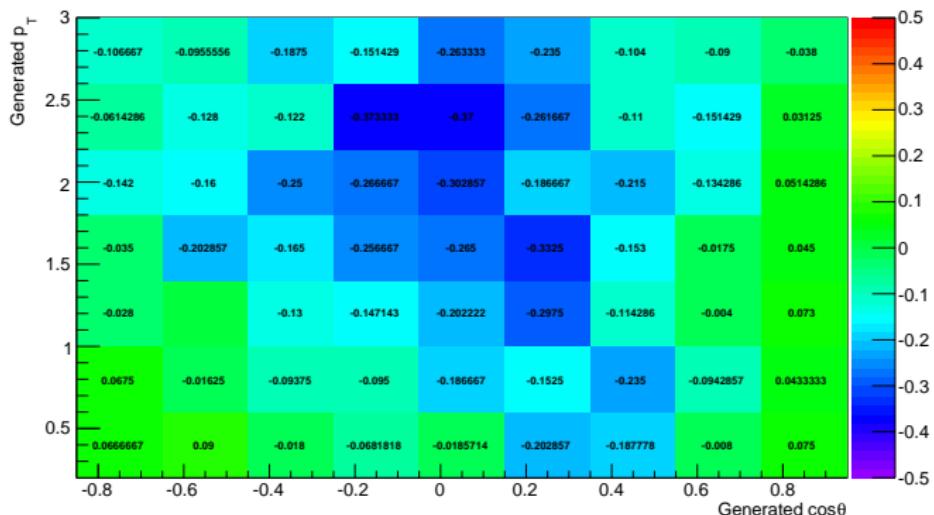


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of pz pull distribution

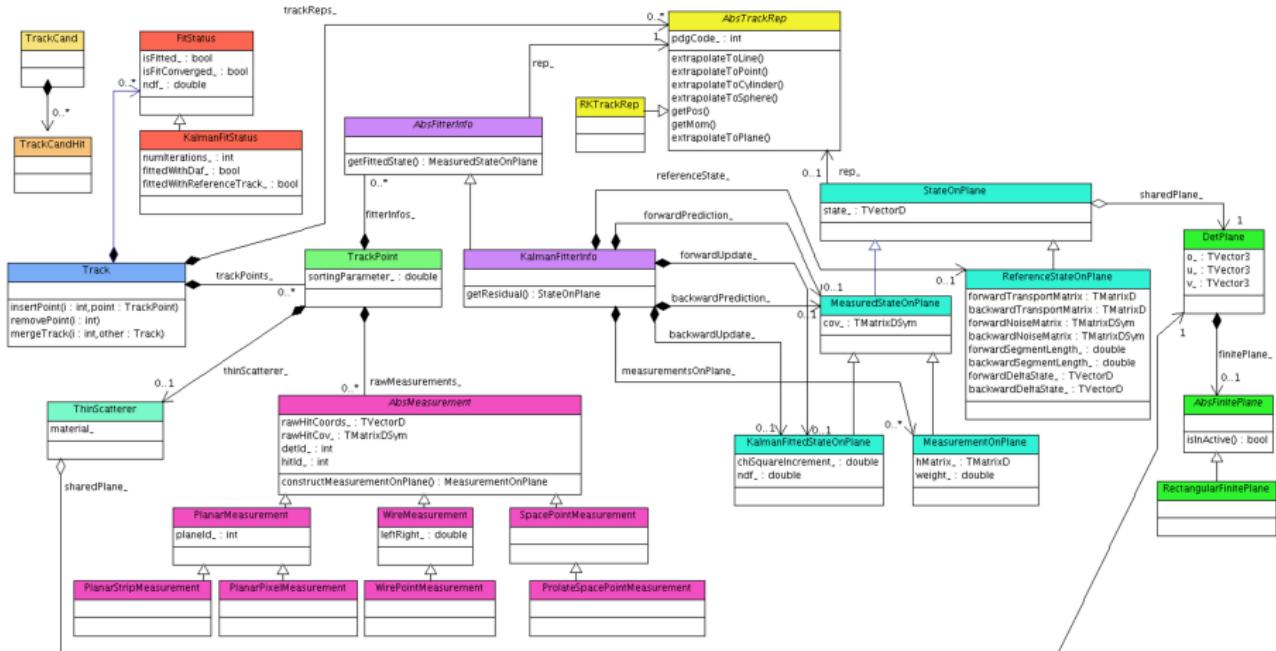


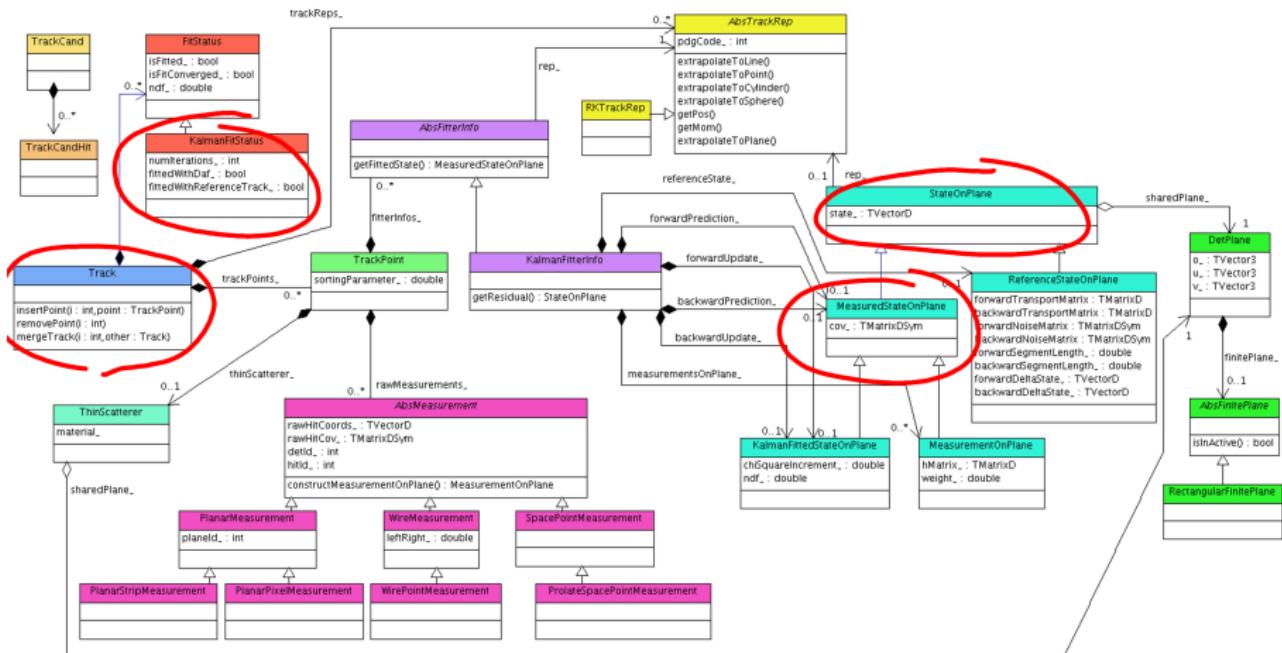
GENFIT 2 with  $\geq 3$  iterations.

# GENFIT 2 Code Examples



# GENFIT Track Data Structure







Getting information about the fit from a fitted track:

## GENFIT 1

```
gfTrack->getCardinalRep()->getStatusFlag() == 0  
  
gfTrack->getCardinalRep()->getChiSqu()  
gfTrack->getCardinalRep()->getNDF()
```

## GENFIT 2

```
gfTrack->getFitStatus()->isFitted()  
gfTrack->getFitStatus()->isFitConverged()  
gfTrack->getFitStatus()->getChi2()  
gfTrack->getFitStatus()->getNdf()  
gfTrack->getFitStatus()->getCharge()
```



# Get Information about the Fit

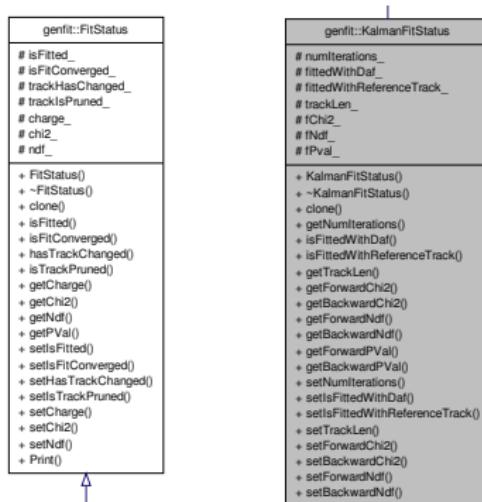
```
FitStatus* genfit::Track::getFitStatus ( const AbsTrackRep * rep = NULL ) const [inline]
```

Get **FitStatus** for a **AbsTrackRep**. Per default, return **FitStatus** for cardinalRep.

Definition at line 136 of file **Track.h**.

References **cardinalRep\_**, and **trackReps\_**.

Referenced by **genfit::GFRaveTrackParameters::getCharge()**, **genfit::AbsKalmanFitter::isTrackFitted()**, **genfit::KalmanFitterRefTrack::prepareTrack()**, **genfit::KalmanFitter::processTrackPartially()**, **genfit::KalmanFitterRefTrack::processTrackWithRep()**, **genfit::KalmanFitter::processTrackWithRep()**, and **genfit::DAF::processTrackWithRep()**.





Biggest difference: State (+ cov) is NOT stored in TrackRep anymore.  
Instead: TrackPoints contain FitterInfos contain StateOnPlane objects.

## GENFIT 1

```
GFAbsTrackRep* rep =  
    gfTrack->getCardinalRep();  
TVector3 pos = rep->getPos();  
TVector3 mom = rep->getMom();  
TMatrixDSym cov = rep->getCov(); // 5x5  
  
TMatrixDSym cov6; // 6x6  
rep->getPosMomCov(pos, mom, cov6);
```

## GENFIT 2

```
const MeasuredStateOnPlane& state =  
    gfTrack->getFittedState();  
TVector3 pos = state.getPos();  
TVector3 mom = state.getMom();  
TMatrixDSym cov = state.getCov(); // 5x5  
  
TMatrixDSym cov6 = state.get6DCov(); // 6x6
```



```
const MeasuredStateOnPlane & genfit::Track::getFittedState ( int id = 0,  
                           const AbsTrackRep * rep = NULL,  
                           bool biased = true  
 ) const
```

Shortcut to get FittedStates.

Uses `getPointWithMeasurementAndFitterInfo(id, rep)`. Per default, the fitted state of the fitterInfo of the first `TrackPoint` with one or more `AbsMeasurement` and `AbsFitterInfo` objects is returned. If no `AbsTrackRep` is specified, the `AbsFitterInfo` of the cardinal rep will be used.

Definition at line 220 of file `Track.cpp`.

References `getCardinalRep()`, `genfit::AbsFitterInfo::getFittedState()`, `genfit::TrackPoint::getFitterInfo()`, `getPointWithMeasurementAndFitterInfo()`, and `genfit::Exception::setFatal()`.



Extrapolate fitted state, e.g. to vertex:

## GENFIT 1

```
TVector3 vertexPos, vertexMom;  
TMatrixDSym vertexCov;  
GFAbsTrackRep* rep =  
    gfTrack->getCardinalRep();  
TVector3 vertex(0,0,0);  
TVector3 poca, dirInPoca;  
rep->extrapolateToPoint(vertex,  
    poca, dirInPoca); // does not alter rep  
GFDetPlane plane(poca, dirInPoca);  
rep->getPosMomCov(plane, vertexPos,  
    vertexMom, vertexCov);
```

## GENFIT 2

```
TVector3 vertexPos, vertexMom;  
TMatrixDSym vertexCov;  
MeasuredStateOnPlane state =  
    gfTrack->getFittedState(); // copy  
TVector3 vertex(0,0,0);  
  
state.extrapolateToPoint(vertex)  
  
state.getPosMomCov(vertexPos,  
    vertexMom, vertexCov);
```



Get time of flight between vertex and cylinder:

## GENFIT 2

```
TVector3 vertex(0,0,0);
double radius = 20;
double tof = gfTrack->getTOF(); // TOF from first to last hit
StateOnPlane state = gfTrack->getFittedState();
state.extrapolateToPoint(vertex);
// TOF is SIGNED!
tof -= state.getRep()->getTOF(); // + TOF from vertex to first hit

state = gfTrack->getFittedState(-1);
state.extrapolateToCylinder(radius);
tof += state.getRep()->getTOF(); // + TOF from last hit to cylinder
```



# Get Track-length and TOF



```
double genfit::Track::getTOF ( AbsTrackRep * rep,
                               int          startId = 0,
                               int          endId = -1
                           )                                const
```

get time of flight in ns between to trackPoints

Definition at line 768 of file [Track.cpp](#).

References [genfit::AbsTrackRep::extrapolateToPlane\(\)](#), [genfit::AbsTrackRep::getMass\(\)](#), [genfit::AbsTrackRep::getMomMag\(\)](#), [genfit::StateOnPlane::getPlane\(\)](#), and [trackPoints\\_](#).

```
double genfit::Track::getTrackLen ( AbsTrackRep * rep,
                                   int          startId = 0,
                                   int          endId = -1
                               )                                const
```

get TrackLength between to trackPoints

Definition at line 727 of file [Track.cpp](#).

References [genfit::AbsTrackRep::extrapolateToPlane\(\)](#), [genfit::StateOnPlane::getPlane\(\)](#), and [trackPoints\\_](#).



```
void genfit::Track::insertPoint ( TrackPoint * point,
                                int           id = -1
                               )
```

Insert [TrackPoint](#) BEFORE [TrackPoint](#) with position id, if id >= 0.

Id -1 means after last [TrackPoint](#). Id -2 means before last [TrackPoint](#). ... Also deletes backwardInfos before new point and forwardInfos after new point. Also sets [Track](#) backpointer of point accordingly.

Definition at line 263 of file [Track.cpp](#).

References [deleteBackwardInfo\(\)](#), [deleteForwardInfo\(\)](#), [deleteReferenceInfo\(\)](#), [fillPointsWithMeasurement\(\)](#), [genfit::TrackPoint::hasRawMeasurements\(\)](#), [genfit::TrackPoint::setTrack\(\)](#), [trackHasChanged\(\)](#), [trackPoints\\_](#), and [trackPointsWithMeasurement\\_](#).

Referenced by [insertPoints\(\)](#), and [Track\(\)](#).

```
void genfit::Track::insertPoints ( std::vector< genfit::TrackPoint * > points,
                                   int                   id = -1
                                  )
```

Definition at line 323 of file [Track.cpp](#).

References [deleteBackwardInfo\(\)](#), [deleteForwardInfo\(\)](#), [deleteReferenceInfo\(\)](#), [fillPointsWithMeasurement\(\)](#), [getNumPoints\(\)](#), [insertPoint\(\)](#), and [trackPoints\\_](#).

Referenced by [mergeTrack\(\)](#).



# Delete Points, Merge Tracks



```
void genfit::Track::deletePoint ( int id )
```

Definition at line 373 of file [Track.cpp](#).

References [deleteBackwardInfo\(\)](#), [deleteForwardInfo\(\)](#), [deleteReferenceInfo\(\)](#), [trackHasChanged\(\)](#), [trackPoints\\_](#), and [trackPointsWithMeasurement\\_](#).

```
void genfit::Track::mergeTrack ( const Track * other,
                                int                  id = -1
                               )
```

Merge two tracks.

The [TrackPoint](#) objects of other will be cloned and inserted after id (per default, they will be appended at the end). The other [Track](#) will not be altered, the [TrackPoint](#) objects will be (deep) copied. Only copies the [TrackPoint](#) objects, NOT the [AbsTrackRep](#), [FitStatus](#), seed state and other objects of the other track.

Definition at line 406 of file [Track.cpp](#).

References [getNumPoints\(\)](#), [insertPoints\(\)](#), [genfit::Exception::setFatal\(\)](#), [trackPoints\\_](#), and [trackReps\\_](#).

# TODOs



## Urgent:

- Fix remaining bugs from migration.

## New GENFIT features which should be used by basf2:

- Use GENFIT's time-of-flight calculation for CDC drift-time corrections.
- Use HMatrixPhi for slanted SVD sensor-planes.
- Re-integrate GENFIT visualization into basf2 display.

## GenFitterModule:

- Fitted vs. fitted and converged?

Support from TUM



## My Status

- I will switch over to Belle I analysis soon.
- I can still do maintenance, but no more big new developments.

## E18 Status:

- E18 wants to hire another Belle (II) PhD student.
- Two candidates, start around Feb/Mar 2014.
- Will also work on GENFIT as service task.
- “We will take care of maintenance of GENFIT.”
- Active development of new features will be reduced for a while.

Thanks for your attention!

# Backup Slides



The following descriptions and diagrams are taken from GENFIT's doxygen documentation.

```
void genfit::AbsFitter::processTrack ( Track * tr,
                                      bool   resortHits = true
                                    )
```

Process all reps. Start with the cardinalRep and resort the hits if necessary (and supported by the fitter)

Definition at line 25 of file [AbsFitter.cpp](#).

References [genfit::Track::checkConsistency\(\)](#), [genfit::Track::getCardinalRep\(\)](#), [genfit::Track::getNumReps\(\)](#), [genfit::Track::getTrackRep\(\)](#), and [processTrackWithRep\(\)](#).

```
virtual void genfit::AbsFitter::processTrackWithRep ( Track * ,
                                                      const AbsTrackRep * ,
                                                      bool             resortHits = false
                                                    )                                     [pure virtual]
```

Process [Track](#) with one [AbsTrackRep](#) of the [Track](#). Optionally resort the hits if necessary (and supported by the fitter)

Implemented in [genfit::DAF](#), [genfit::KalmanFitter](#), and [genfit::KalmanFitterRefTrack](#).

Referenced by [processTrack\(\)](#).



```
const MeasuredStateOnPlane & genfit::Track::getFittedState ( int id = 0,  
                           const AbsTrackRep * rep = NULL,  
                           bool biased = true  
 ) const
```

Shortcut to get FittedStates.

Uses `getPointWithMeasurementAndFitterInfo(id, rep)`. Per default, the fitted state of the fitterInfo of the first `TrackPoint` with one or more `AbsMeasurement` and `AbsFitterInfo` objects is returned. If no `AbsTrackRep` is specified, the `AbsFitterInfo` of the cardinal rep will be used.

Definition at line 220 of file `Track.cpp`.

References `getCardinalRep()`, `genfit::AbsFitterInfo::getFittedState()`, `genfit::TrackPoint::getFitterInfo()`, `getPointWithMeasurementAndFitterInfo()`, and `genfit::Exception::setFatal()`.

Important: The TrackReps do NOT contain the state anymore.  
Instead, all states are stored in the FitterInfos in the TrackPoints.



```
const SharedPlanePtr& genfit::StateOnPlane::getPlane( ) const [inline]
```

Definition at line 63 of file [StateOnPlane.h](#).

References [sharedPlane\\_](#).

Referenced by [genfit::KalmanFitterInfo::addMeasurementOnPlane\(\)](#), [genfit::RKTrackRep::checkCache\(\)](#), [genfit::RKTrackRep::extrapolateBy\(\)](#), [genfit::RKTrackRep::extrapolateToCylinder\(\)](#), [genfit::RKTrackRep::extrapolateToLine\(\)](#), [genfit::RKTrackRep::extrapolateToPlane\(\)](#), [genfit::RKTrackRep::extrapolateToPoint\(\)](#), [genfit::RKTrackRep::extrapolateToSphere\(\)](#), [genfit::KalmanFitterInfo::getResidual\(\)](#), [genfit::RKTrackRep::getState5\(\)](#), [genfit::RKTrackRep::getState7\(\)](#), [genfit::Track::getTOF\(\)](#), [genfit::Track::getTrackLen\(\)](#), [genfit::KalmanFitterRefTrack::prepareTrack\(\)](#), [genfit::KalmanFitterRefTrack::processTrackPoint\(\)](#), [genfit::RKTrackRep::setPosMom\(\)](#), [genfit::RKTrackRep::setPosMomErr\(\)](#), [genfit::RKTrackRep::transformM6P\(\)](#), [genfit::RKTrackRep::transformM7P\(\)](#), [genfit::RKTrackRep::transformPM6\(\)](#), and [genfit::RKTrackRep::transformPM7\(\)](#).



```
const TVectorD& genfit::StateOnPlane::getState() const [inline]
```

```
TVectorD& genfit::StateOnPlane::getState() [inline]
```

```
const TMatrixDSym& genfit::MeasuredStateOnPlane::getCov() const [inline]
```

Get the state (and covariance) in the parametrization of the TrackRep.  
For the RKTrackRep, this is  $(q/p, u', v', u, v)$ .



```
void genfit::MeasuredStateOnPlane::getPosMomCov ( TVector3 & pos,  
                                                 TVector3 & mom,  
                                                 TMatrixDSym & cov  
) const [inline]
```

Definition at line 63 of file [MeasuredStateOnPlane.h](#).

References [genfit::AbsTrackRep::getPosMomCov\(\)](#), and [genfit::StateOnPlane::getRep\(\)](#).

These and the following functions call the corresponding functions of the TrackRep connected to the StateOnPlane/MeasuredStateOnPlane.



```
TVector3 genfit::StateOnPlane::getPos ( ) const [inline]
```

Definition at line 91 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getPos\(\)](#), and [rep\\_](#).

```
void genfit::StateOnPlane::getPosDir ( TVector3 & pos,  
                                      TVector3 & dir  
                                     ) const [inline]
```

Definition at line 95 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getPosDir\(\)](#), and [rep\\_](#).

```
void genfit::StateOnPlane::getPosMom ( TVector3 & pos,  
                                       TVector3 & mom  
                                      ) const [inline]
```

Definition at line 94 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getPosMom\(\)](#), and [rep\\_](#).

```
double genfit::StateOnPlane::getQop ( ) const [inline]
```

Definition at line 100 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getQop\(\)](#), and [rep\\_](#).



```
TVector3 genfit::StateOnPlane::getMom( ) const [inline]
```

Definition at line 92 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getMom\(\)](#), and [rep\\_](#).

```
double genfit::StateOnPlane::getMomMag( ) const [inline]
```

Definition at line 97 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getMomMag\(\)](#), and [rep\\_](#).

```
int genfit::StateOnPlane::getPDG( ) const [inline]
```

Definition at line 98 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getPDG\(\)](#), and [rep\\_](#).



```
double genfit::StateOnPlane::getCharge( ) const [inline]
```

Definition at line 99 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getCharge\(\)](#), and [rep\\_](#).

```
TVector3 genfit::StateOnPlane::getDir( ) const [inline]
```

Definition at line 93 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getDir\(\)](#), and [rep\\_](#).

```
double genfit::StateOnPlane::getMass( ) const [inline]
```

Definition at line 101 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::getMass\(\)](#), and [rep\\_](#).



```
double genfit::StateOnPlane::extrapolateBy ( double step,
                                             bool    stopAtBoundary = false
                                         )                                     [inline]
```

Definition at line 88 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::extrapolateBy\(\)](#), and [rep\\_](#).

```
double genfit::StateOnPlane::extrapolateToCylinder ( double           radius,
                                                   const TVector3 & linePoint = TVector3(0.,0.,0.),
                                                   const TVector3 & lineDirection = TVector3(0.,0.,1.),
                                                   bool            stopAtBoundary = false
                                         )                                     [inline]
```

Definition at line 81 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::extrapolateToCylinder\(\)](#), and [rep\\_](#).

```
double genfit::StateOnPlane::extrapolateToLine ( const TVector3 & linePoint,
                                                const TVector3 & lineDirection,
                                                bool            stopAtBoundary = false
                                         )                                     [inline]
```

Definition at line 76 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::extrapolateToLine\(\)](#), and [rep\\_](#).



```
double genfit::StateOnPlane::extrapolateToPlane ( const SharedPlanePtr & plane,
                                                 bool                  stopAtBoundary = false,
                                                 bool                  calcJacobianNoise = false
)
                                                [inline]
```

Definition at line 73 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::extrapolateToPlane\(\)](#), and [rep...](#).

```
double genfit::StateOnPlane::extrapolateToPoint ( const TVector3 & point,
                                                 bool                  stopAtBoundary = false
)
                                                [inline]
```

Definition at line 79 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::extrapolateToPoint\(\)](#), and [rep...](#).

```
double genfit::StateOnPlane::extrapolateToSphere ( double           radius,
                                                 const TVector3 & point = TVector3(0.,0.,0.),
                                                 bool            stopAtBoundary = false
)
                                                [inline]
```

Definition at line 85 of file [StateOnPlane.h](#).

References [genfit::AbsTrackRep::extrapolateToSphere\(\)](#), and [rep...](#).



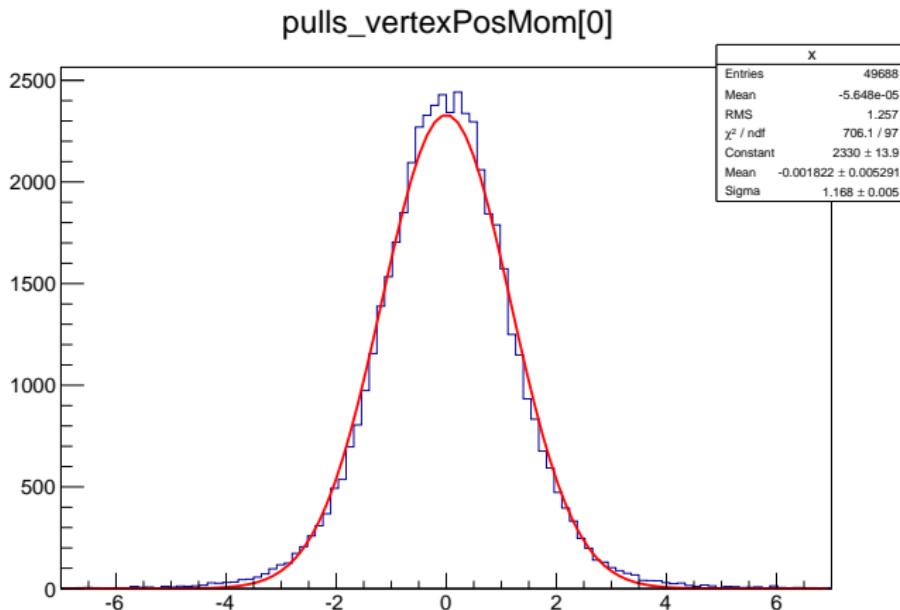
```
void genfit::StateOnPlane::setPosMom ( const TVector3 & pos,  
                                      const TVector3 & mom  
                                     ) [inline]
```

Definition at line 103 of file [StateOnPlane.h](#).

References [rep\\_](#), and [genfit::AbsTrackRep::setPosMom\(\)](#)



## GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

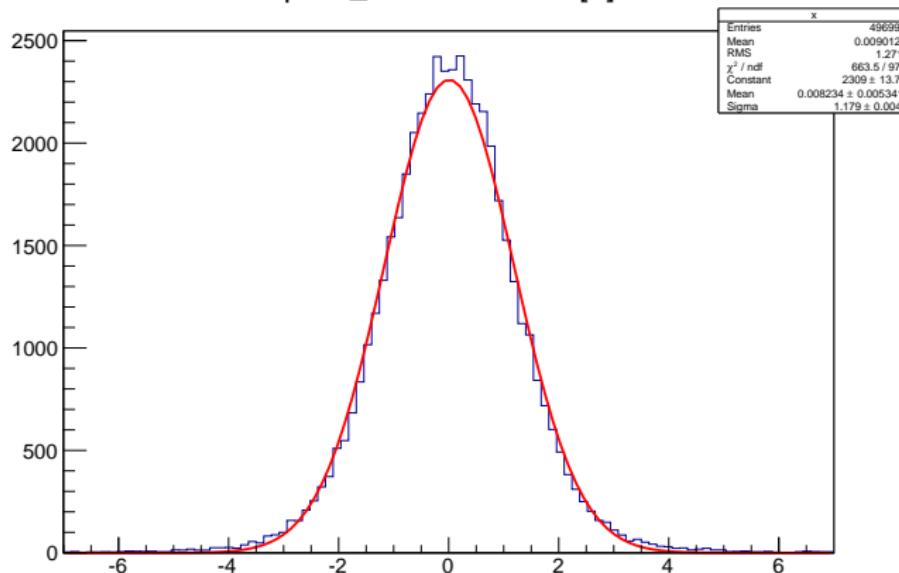
GENFIT 1 with 3 iterations.



# GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[0]



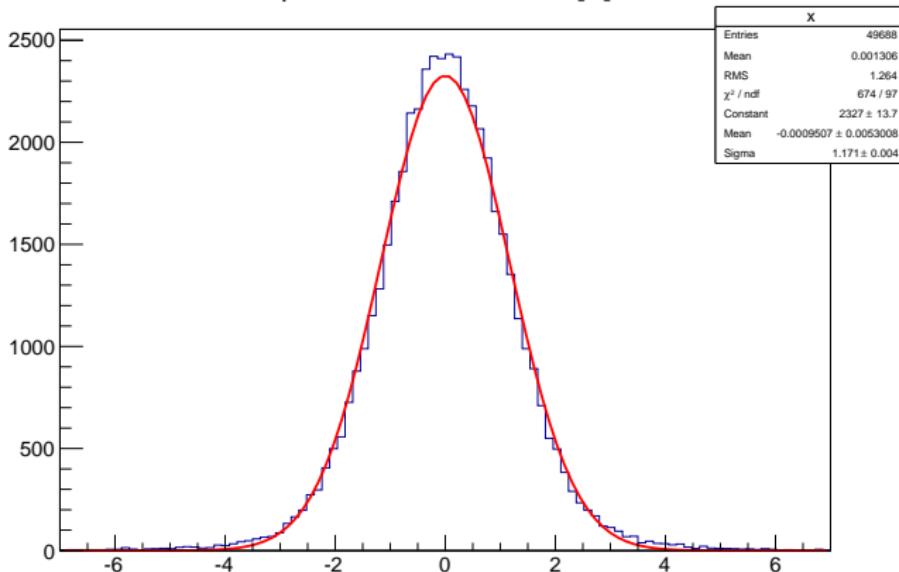
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[1]



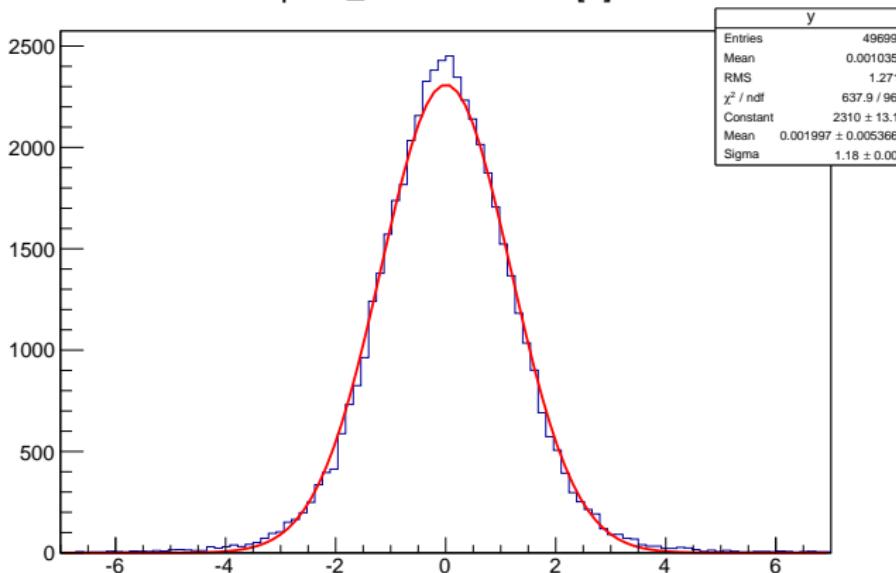
GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[1]

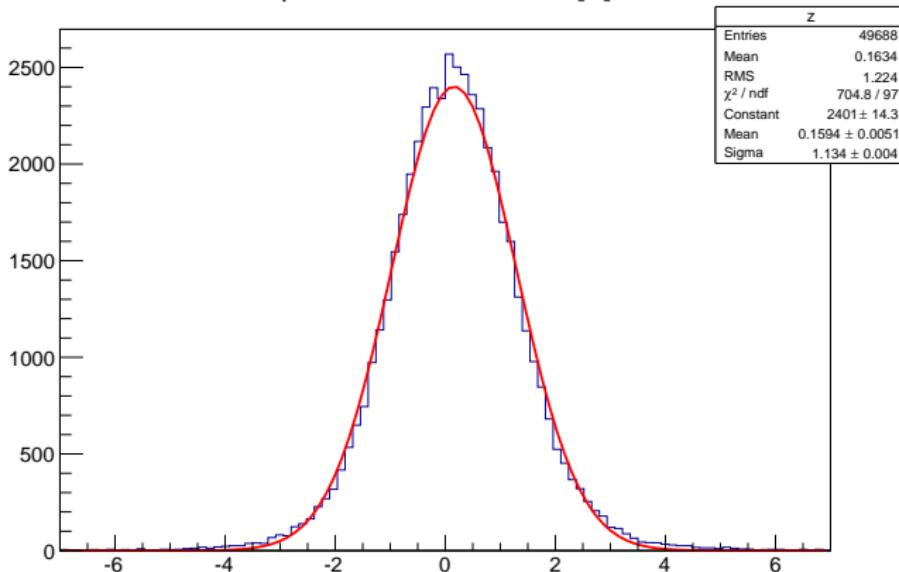
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[2]



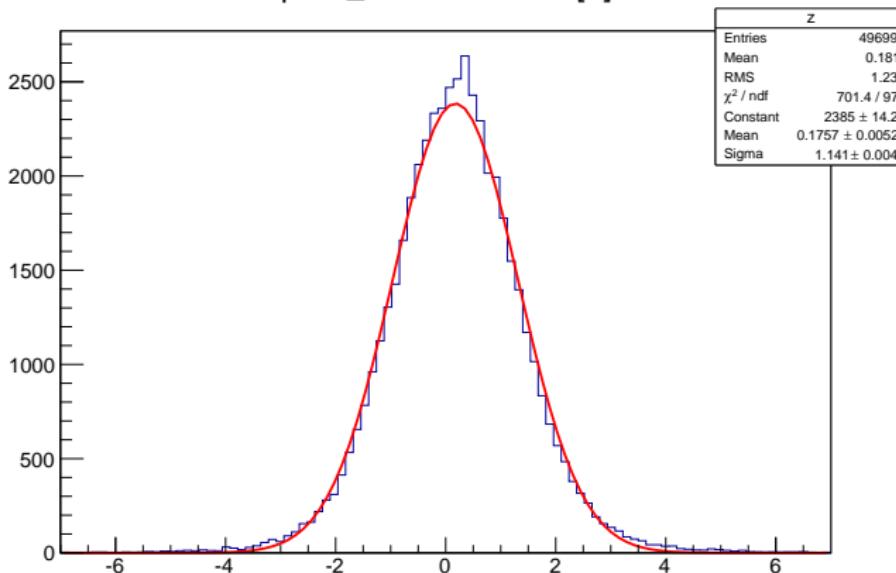
GENFIT 1 with 3 iterations.



# GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[2]



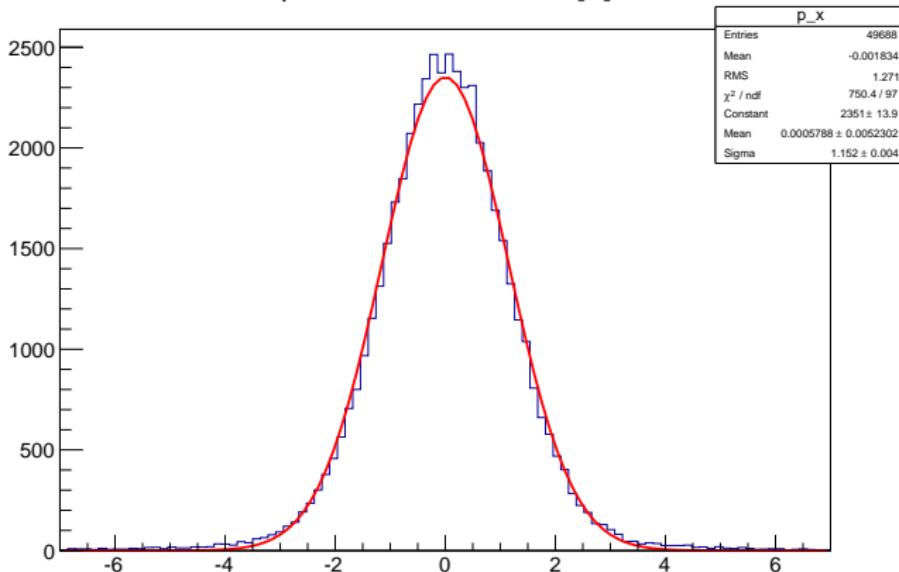
GENFIT 2 with  $\geq 3$  iterations.



# GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[3]



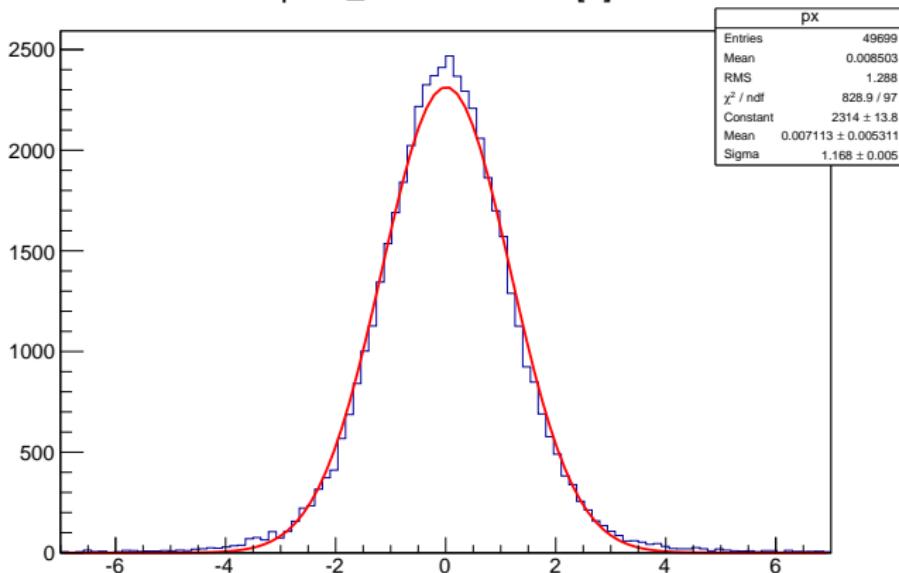
GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[3]

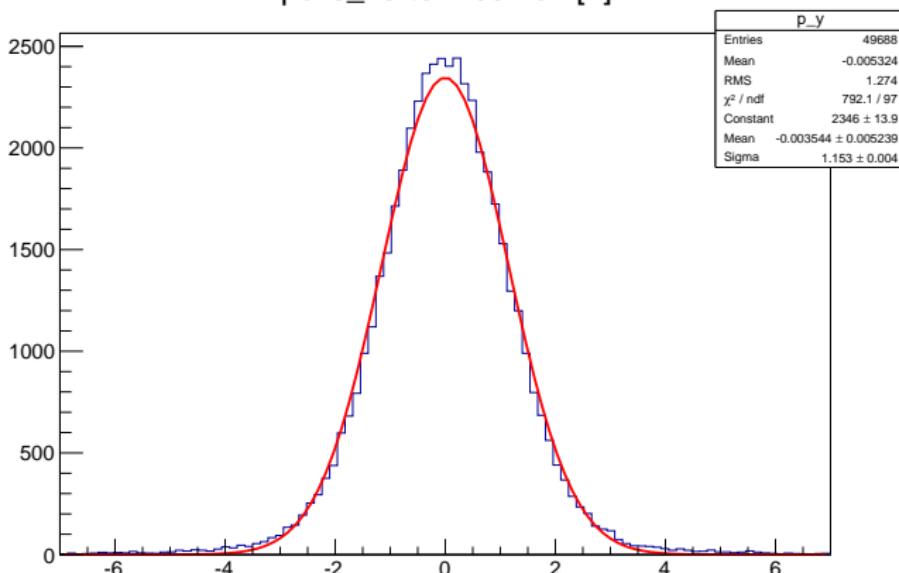
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[4]



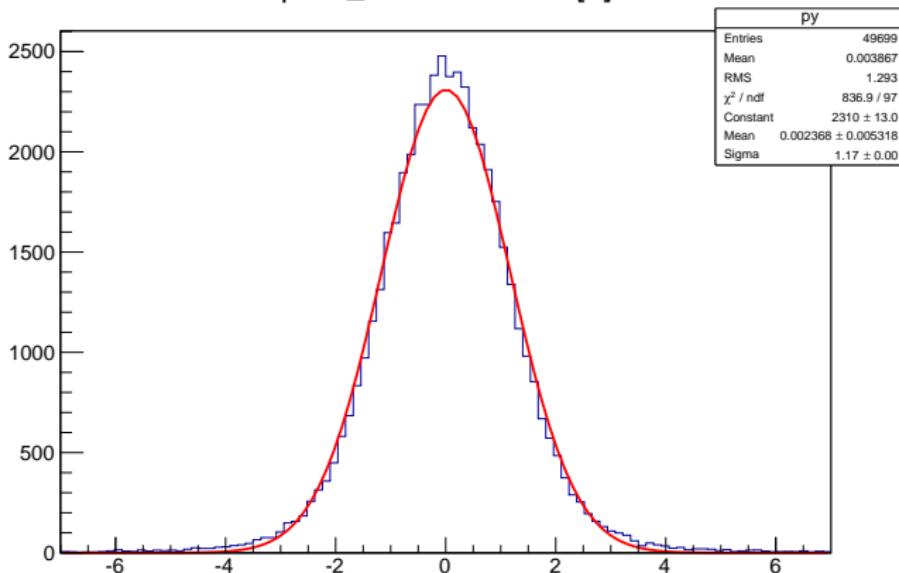
GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[4]

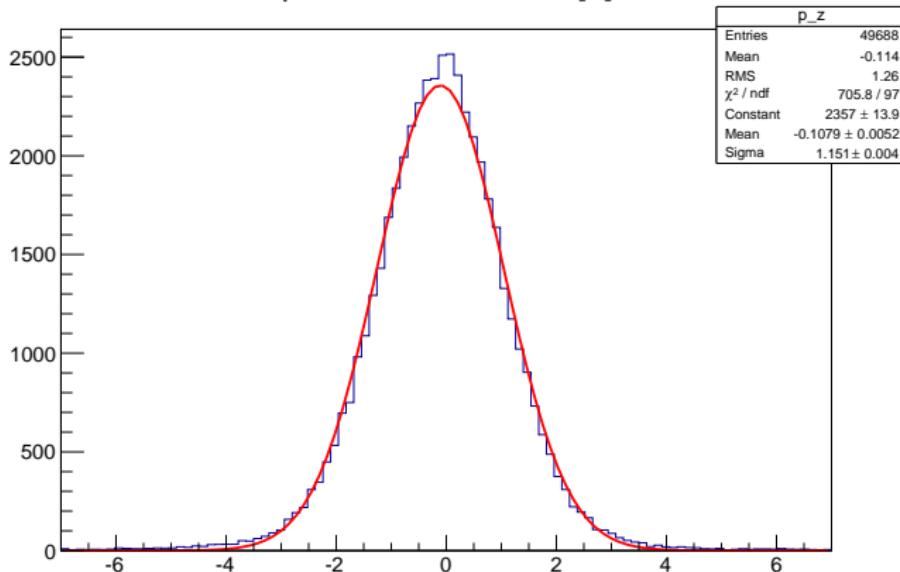
GENFIT 2 with  $\geq 3$  iterations.



# GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

pulls\_vertexPosMom[5]



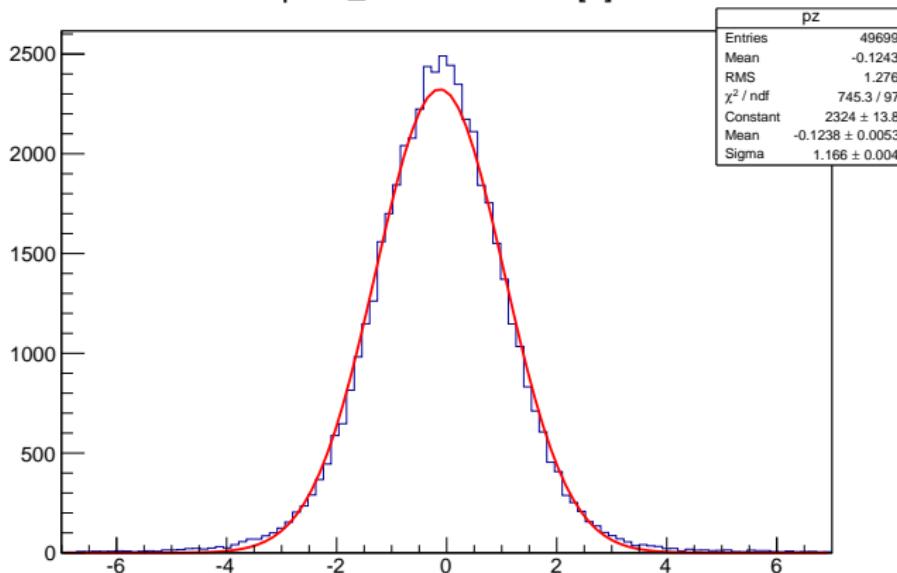
GENFIT 1 with 3 iterations.



# GENFIT 1 vs. GENFIT 2: Pull Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

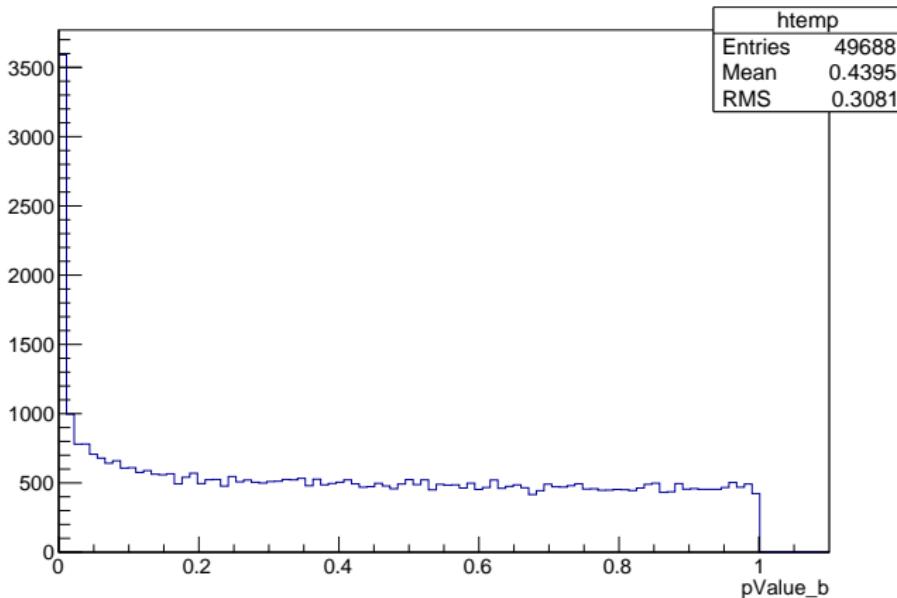
pulls\_vertexPosMom[5]



GENFIT 2 with  $\geq 3$  iterations.



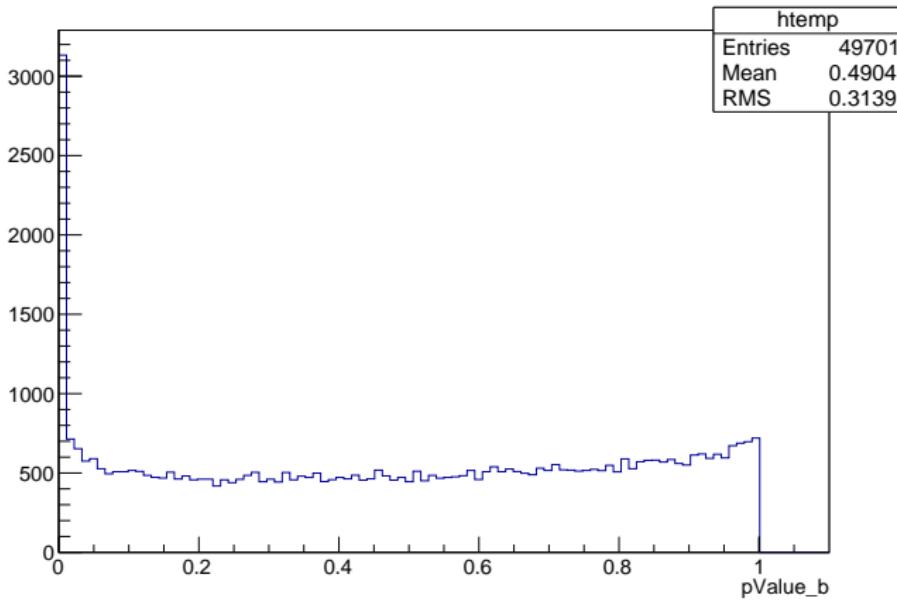
## GENFIT 1 vs. GENFIT 2: P-Value Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

GENFIT 1 with 3 iterations.

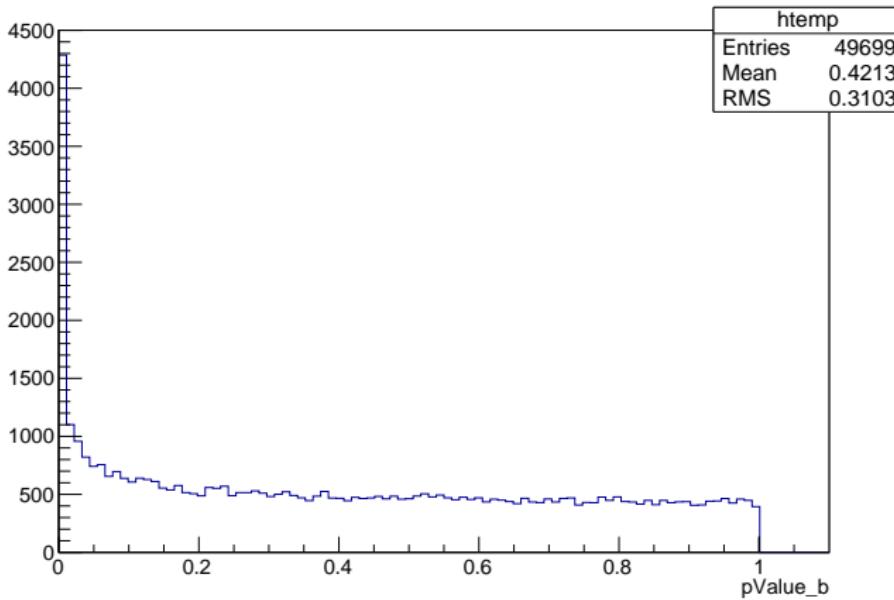


## GENFIT 1 vs. GENFIT 2: P-Value Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.GENFIT 2 with  $\geq 2$  iterations.

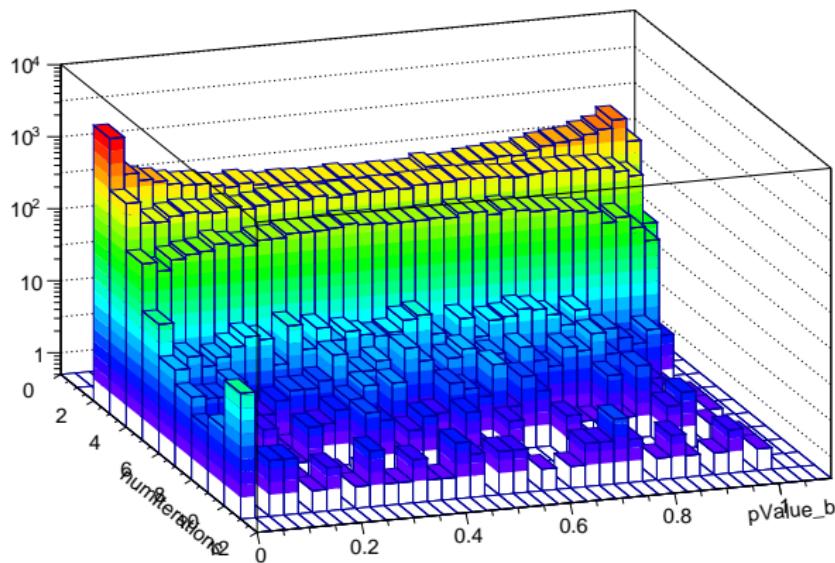


## GENFIT 1 vs. GENFIT 2: P-Value Distributions

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.GENFIT 2 with  $\geq 3$  iterations.



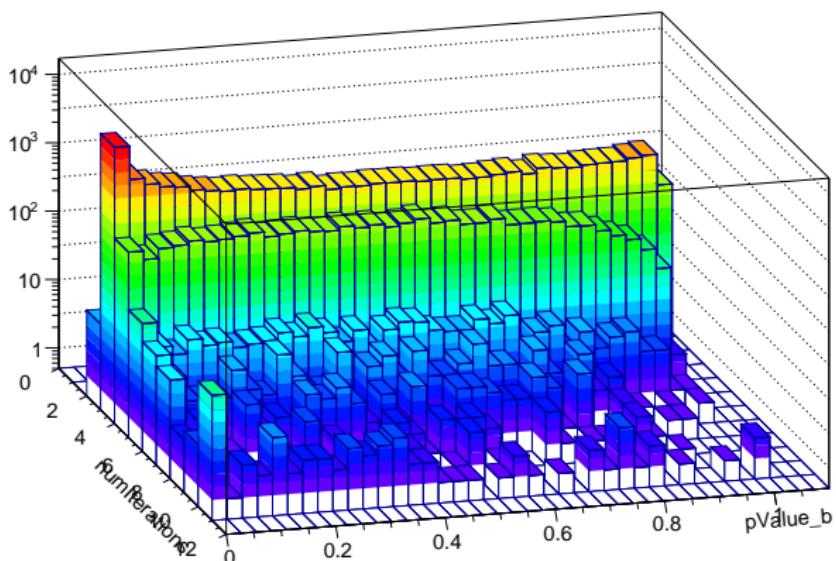
## GENFIT 2: P-Values vs. Number of Iterations

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.GENFIT 2 with  $\geq 2$  iterations.



## GENFIT 2: P-Values vs. Number of Iterations

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



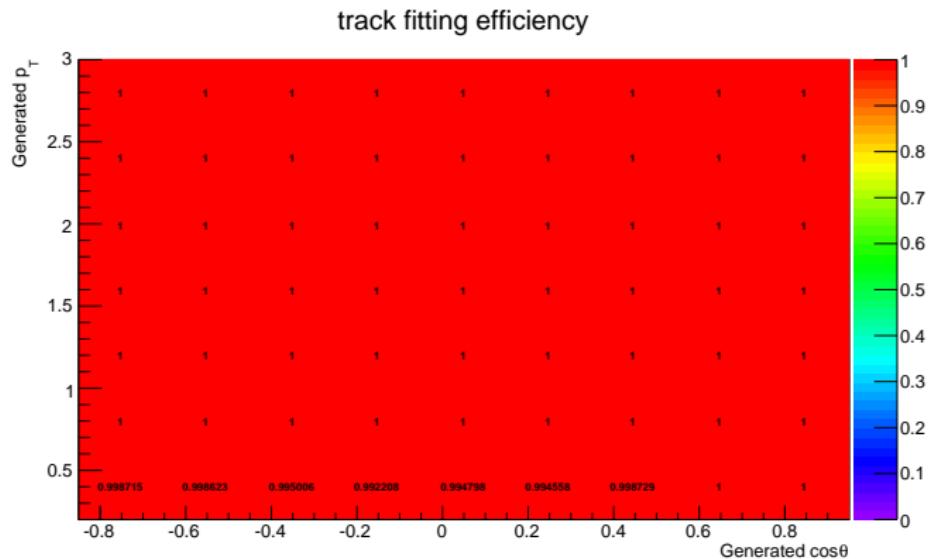
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



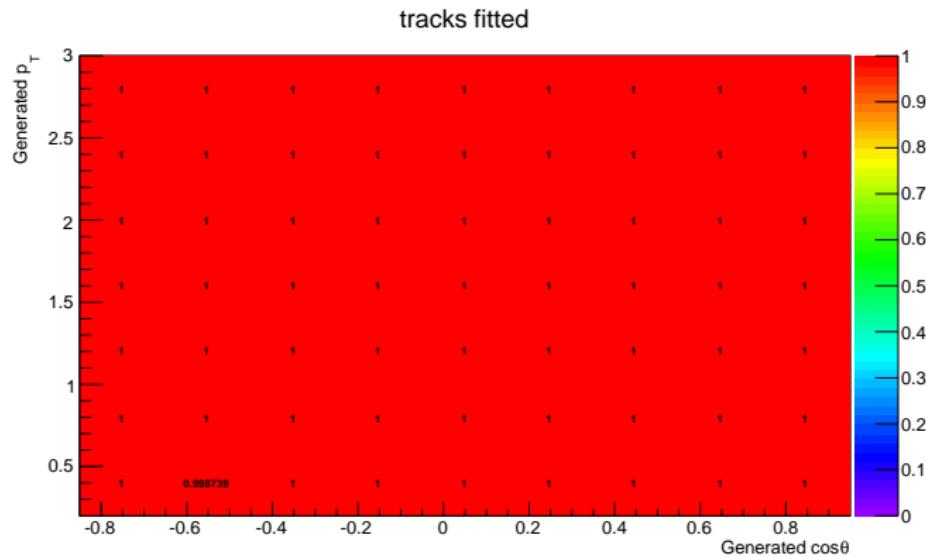
GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



GENFIT 2 with  $\geq 2$  iterations.



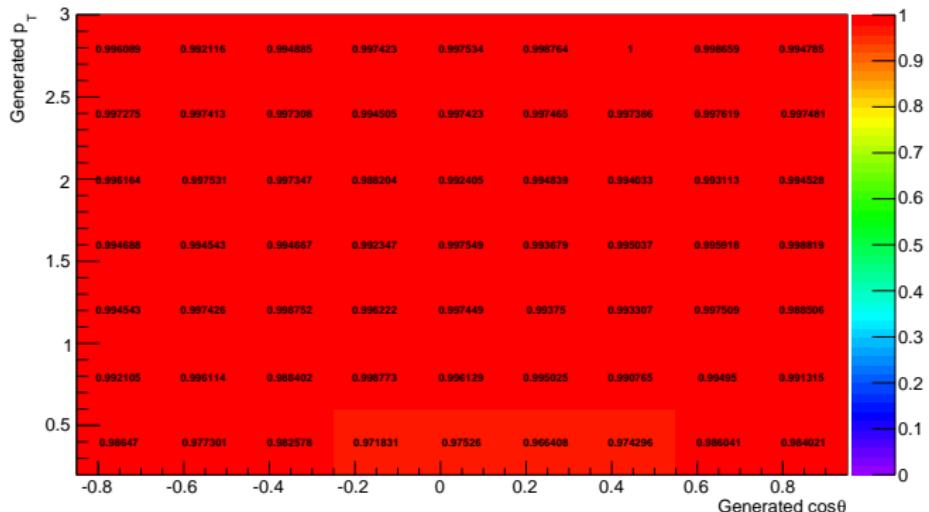
## GENFIT 1 vs. GENFIT 2: 2D QA Plots



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50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

tracks fitted and converged

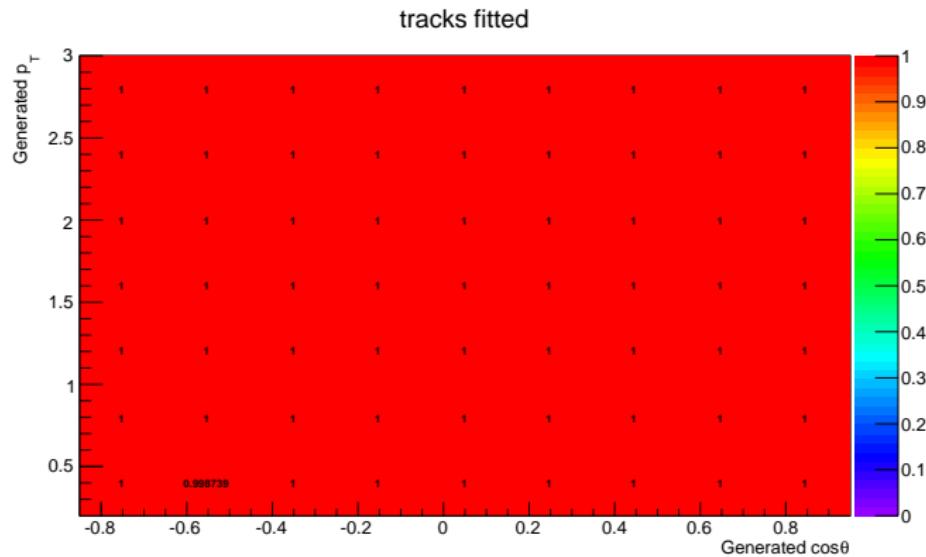
GENFIT 2 with  $\geq 2$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



GENFIT 2 with  $\geq 3$  iterations.

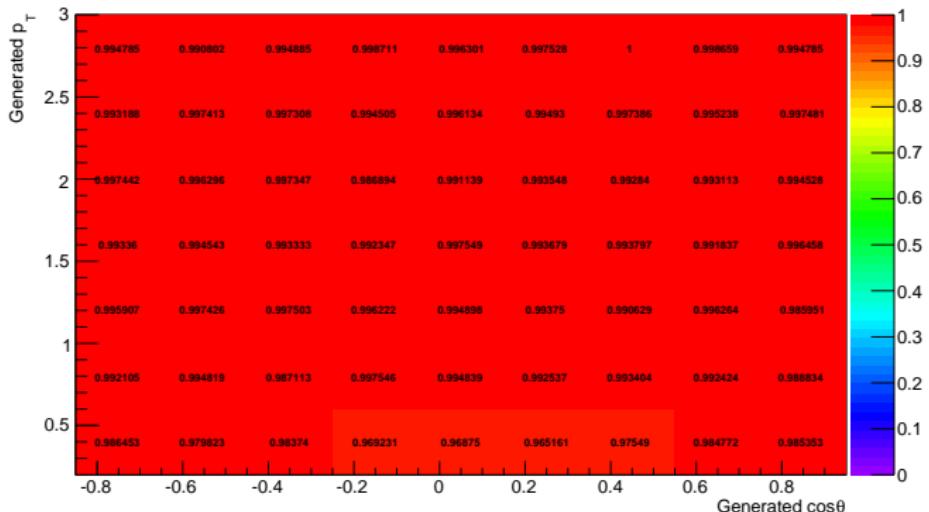


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

tracks fitted and converged



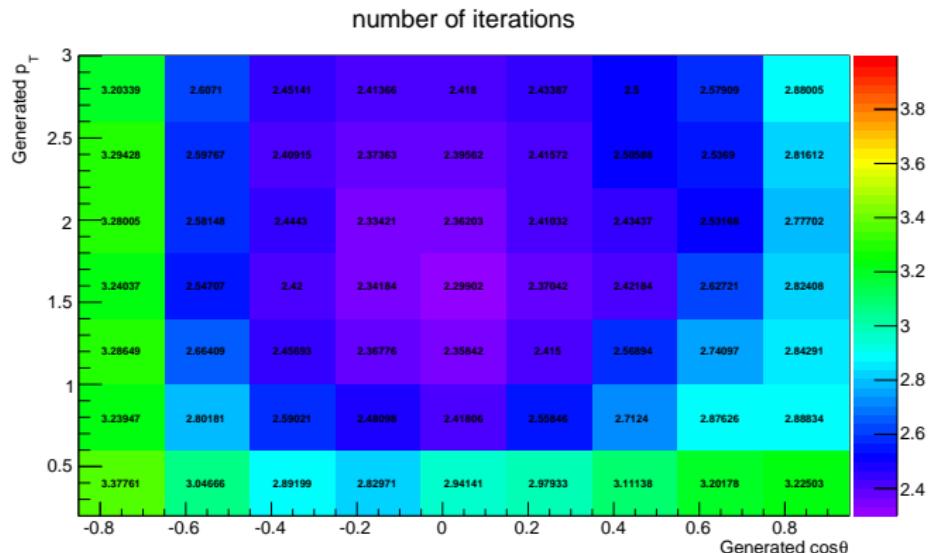
GENFIT 2 with  $\geq 3$  iterations.



# GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



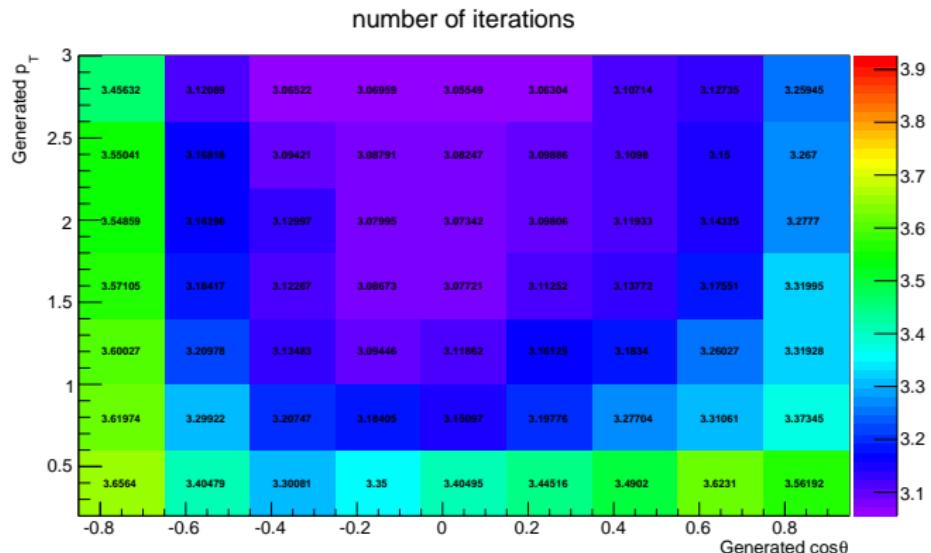
GENFIT 2 with  $\geq 2$  iterations.



# GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



GENFIT 2 with  $\geq 3$  iterations.

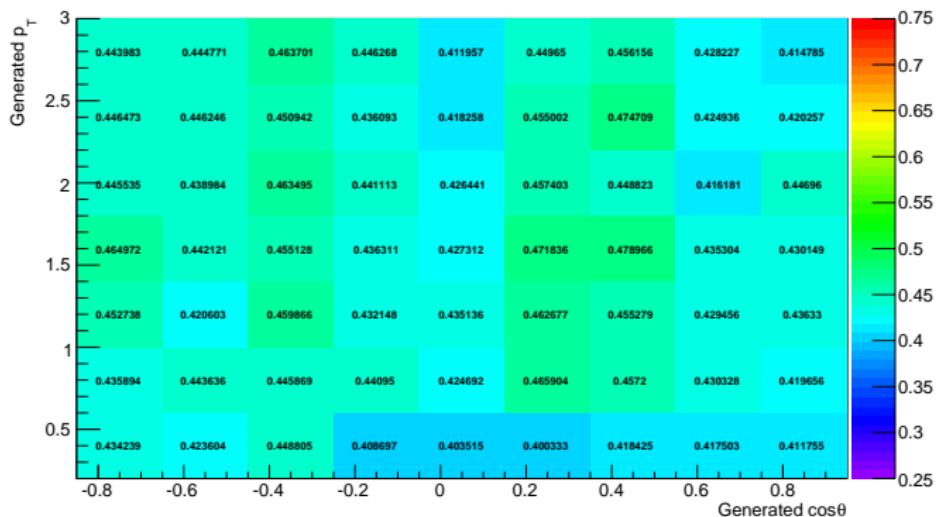


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

mean of p-value distribution from the backward filter



GENFIT 1 with 3 iterations.

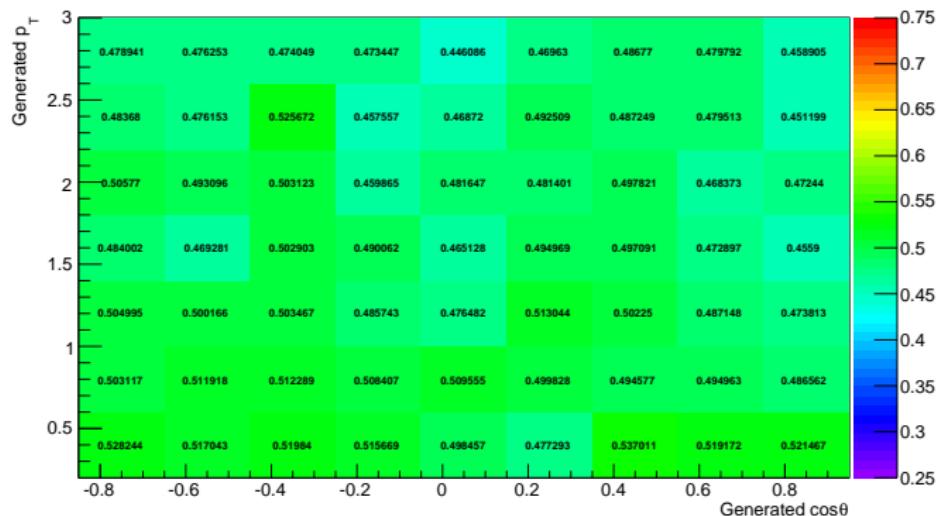


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

mean of p-value distribution from the backward filter



GENFIT 2 with  $\geq 2$  iterations.

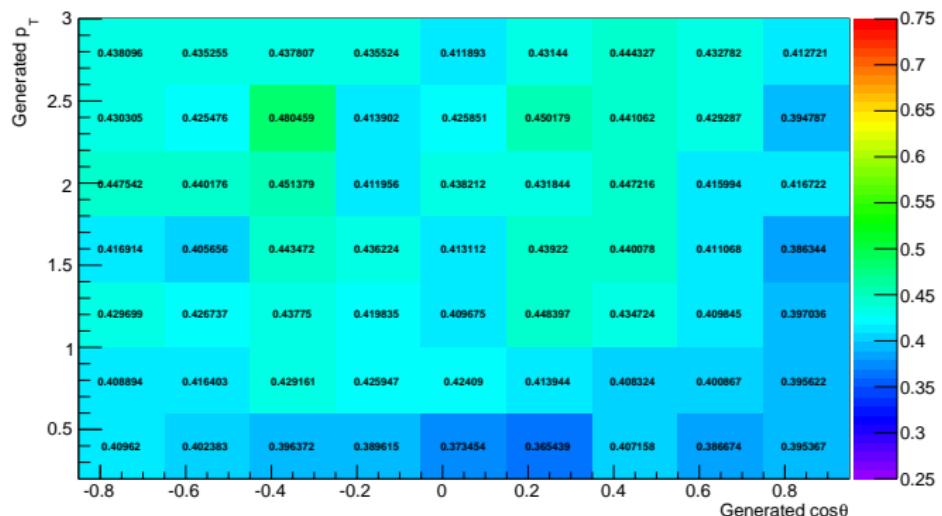


# GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

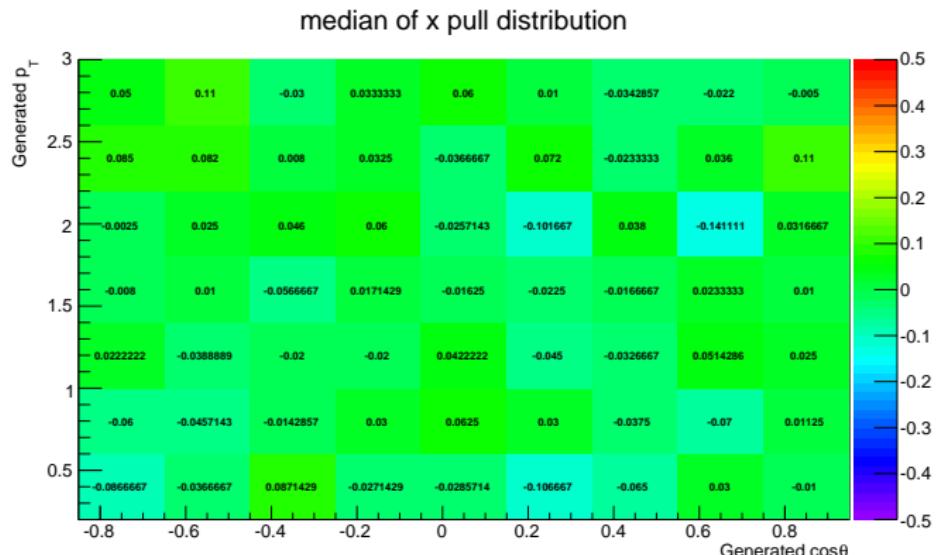
mean of p-value distribution from the backward filter



GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

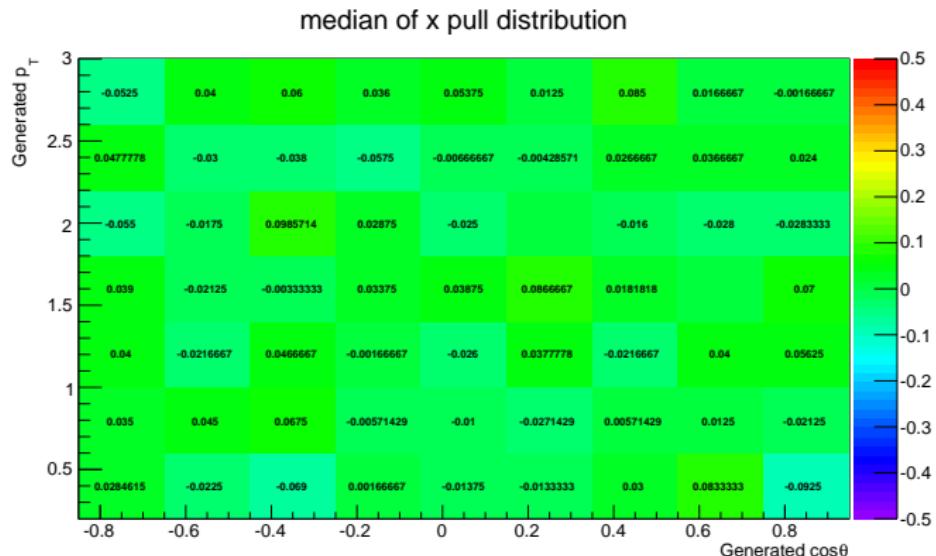
GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



GENFIT 2 with  $\geq 2$  iterations.



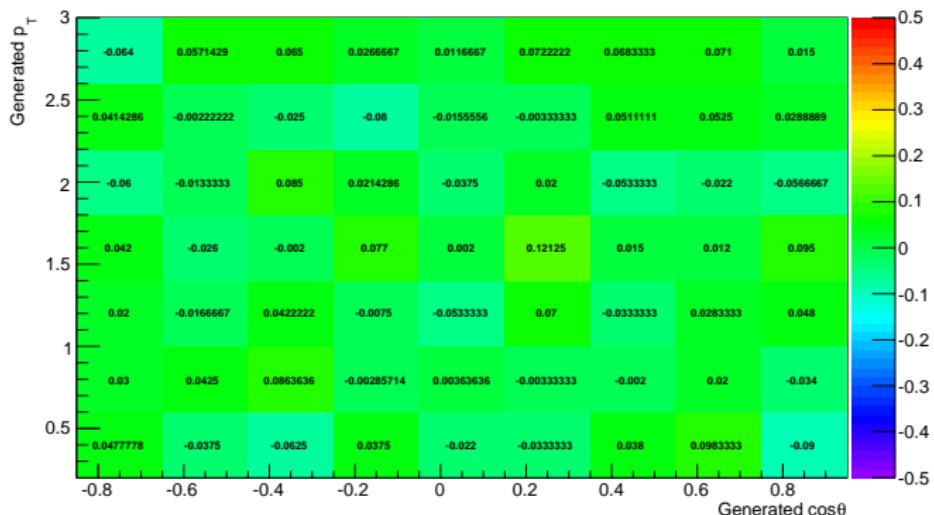
## GENFIT 1 vs. GENFIT 2: 2D QA Plots



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50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of x pull distribution

GENFIT 2 with  $\geq 3$  iterations.

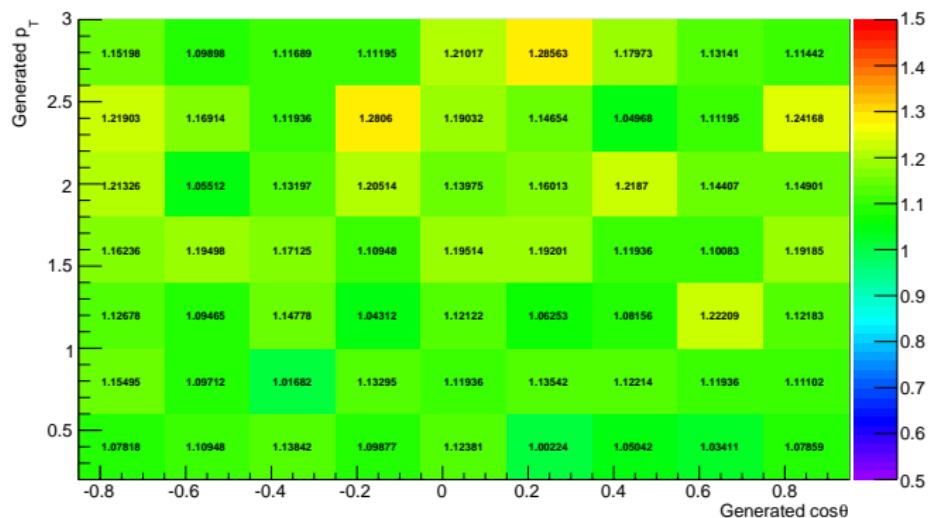


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of x pull distribution



GENFIT 1 with 3 iterations.

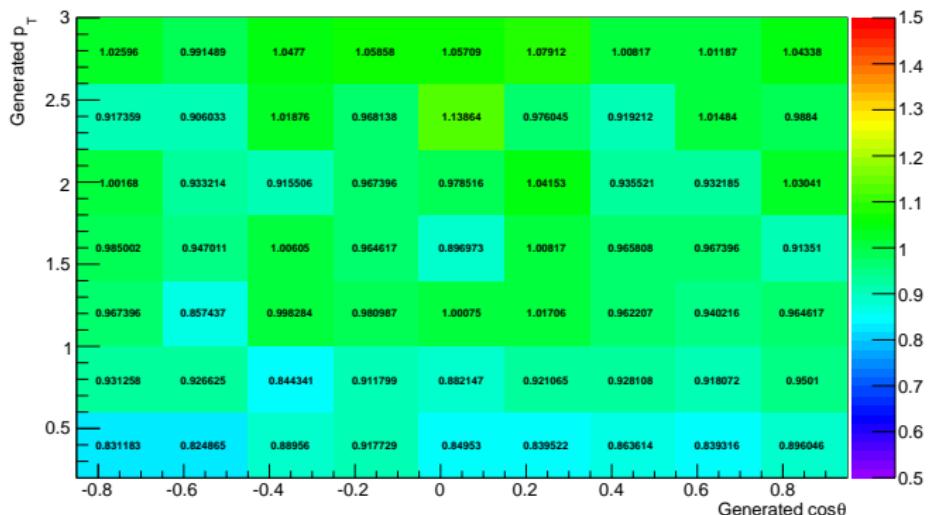


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of x pull distribution



GENFIT 2 with  $\geq 2$  iterations.

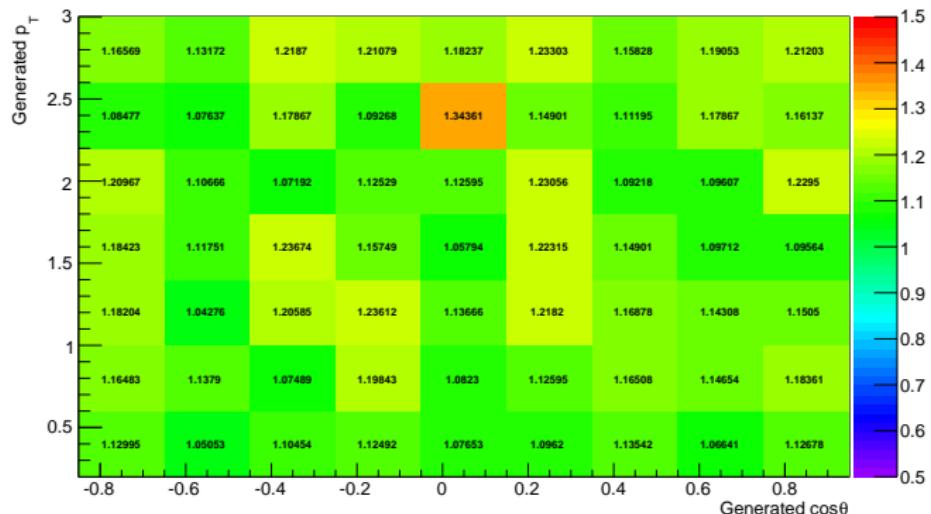


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of x pull distribution



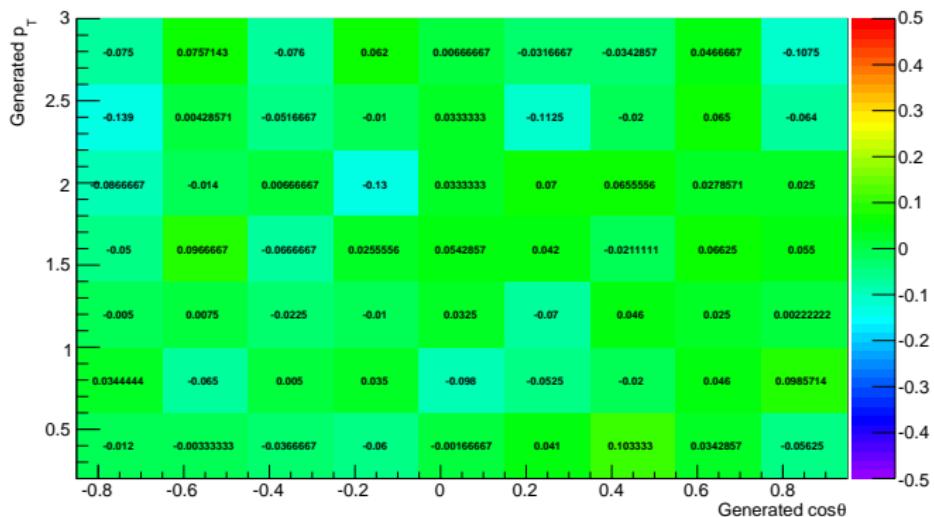
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of y pull distribution



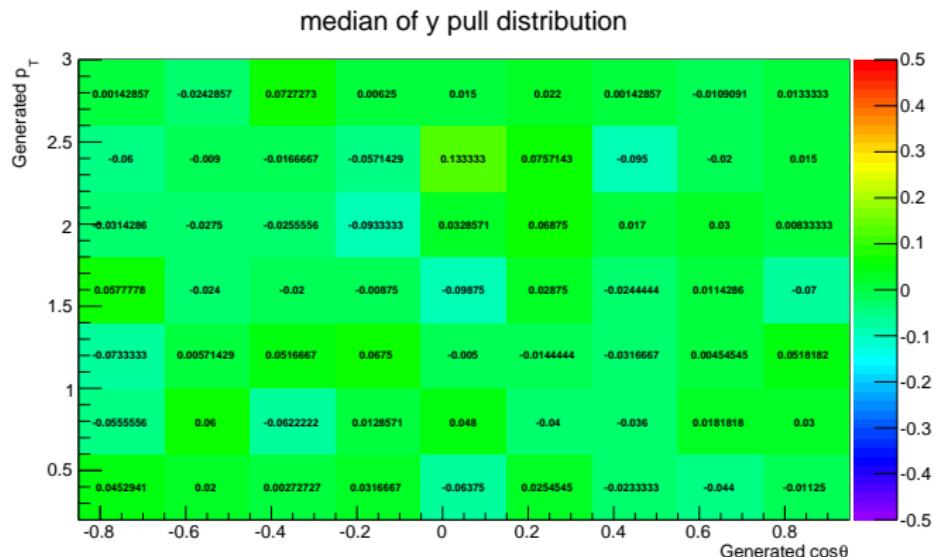
GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.



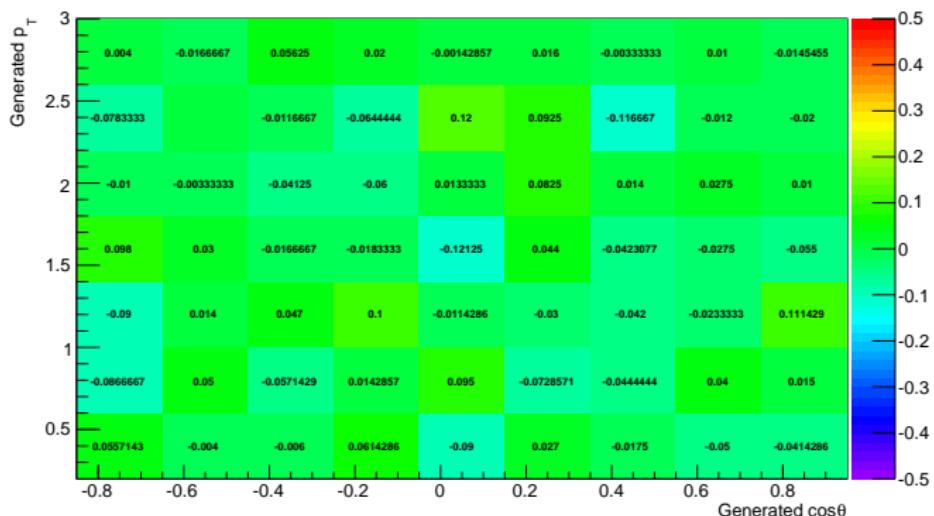


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of y pull distribution



GENFIT 2 with  $\geq 3$  iterations.

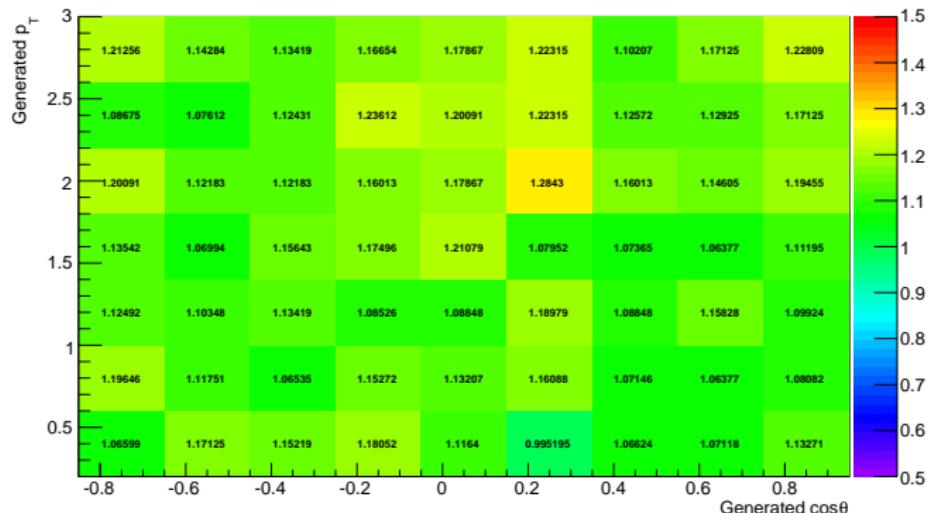


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of y pull distribution



GENFIT 1 with 3 iterations.

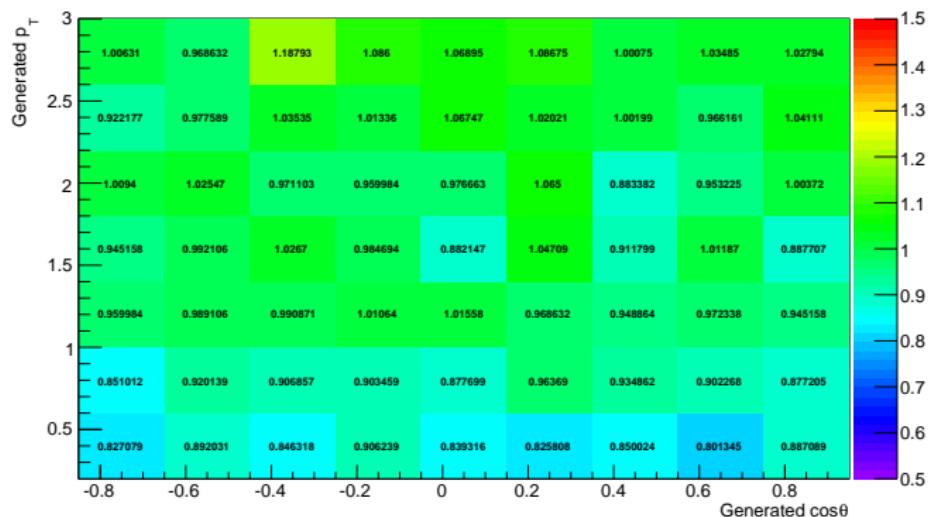


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of y pull distribution



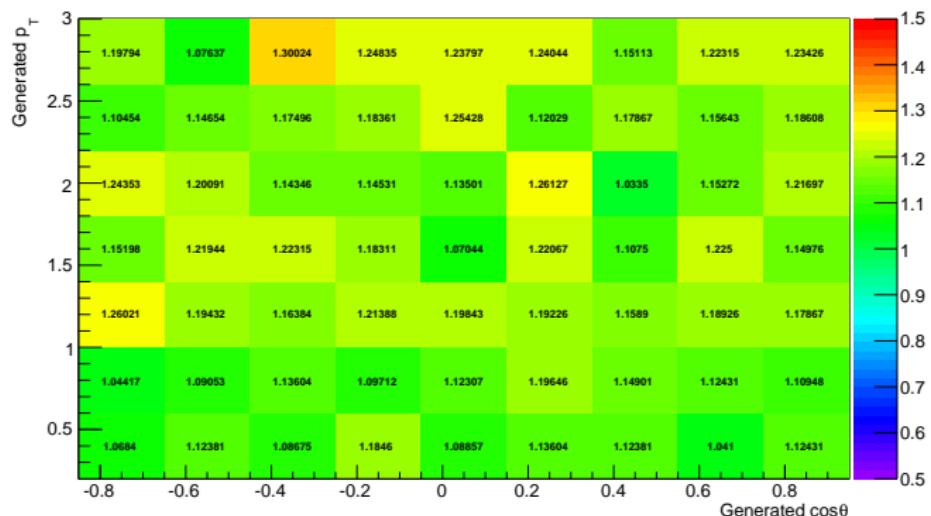
GENFIT 2 with  $\geq 2$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of y pull distribution



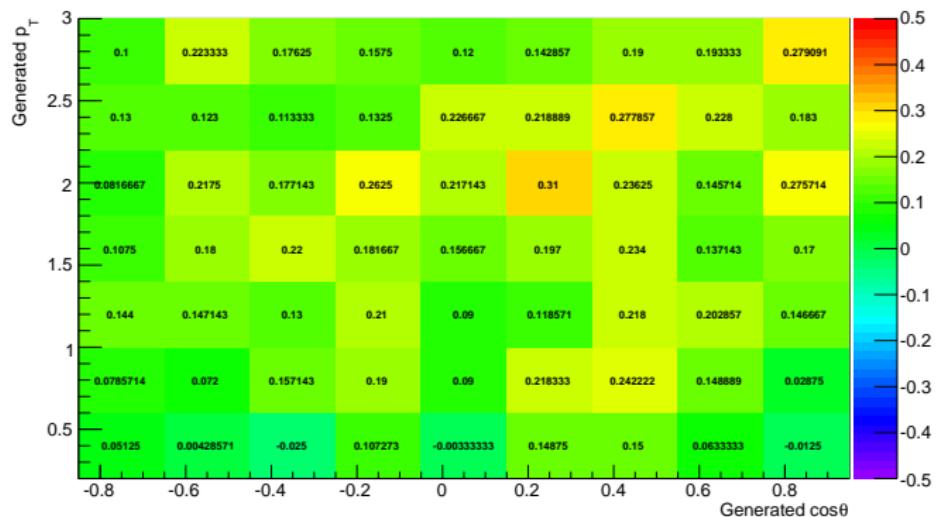
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of z pull distribution



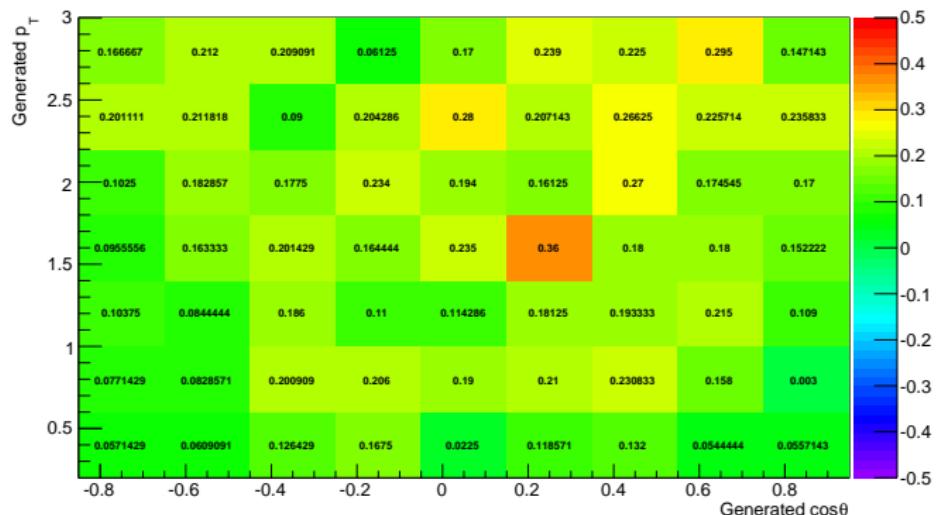


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of z pull distribution



GENFIT 2 with  $\geq 2$  iterations.

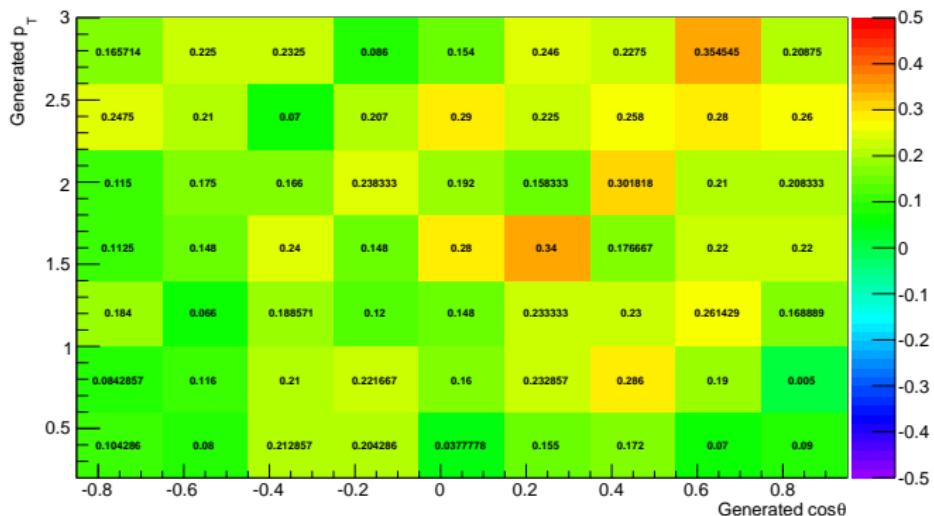


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of z pull distribution



GENFIT 2 with  $\geq 3$  iterations.

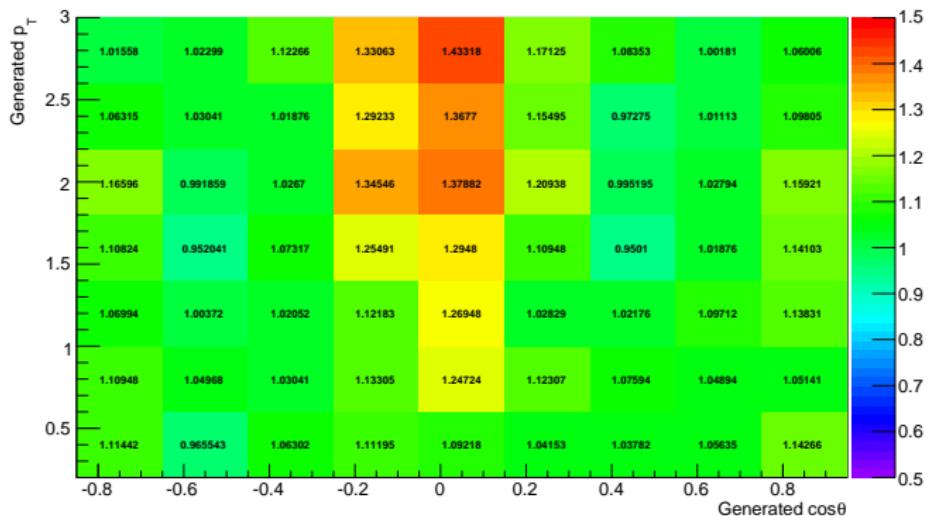


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of z pull distribution



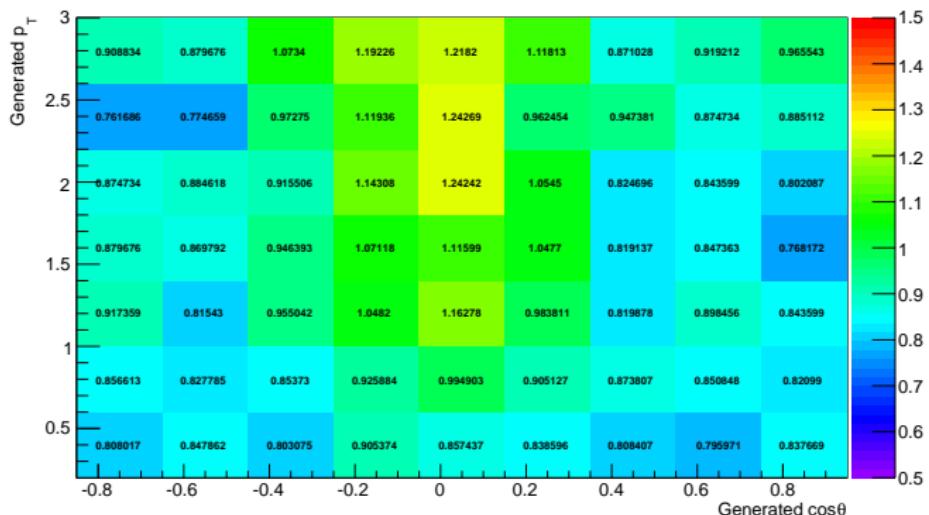
GENFIT 1 with 3 iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of z pull distribution

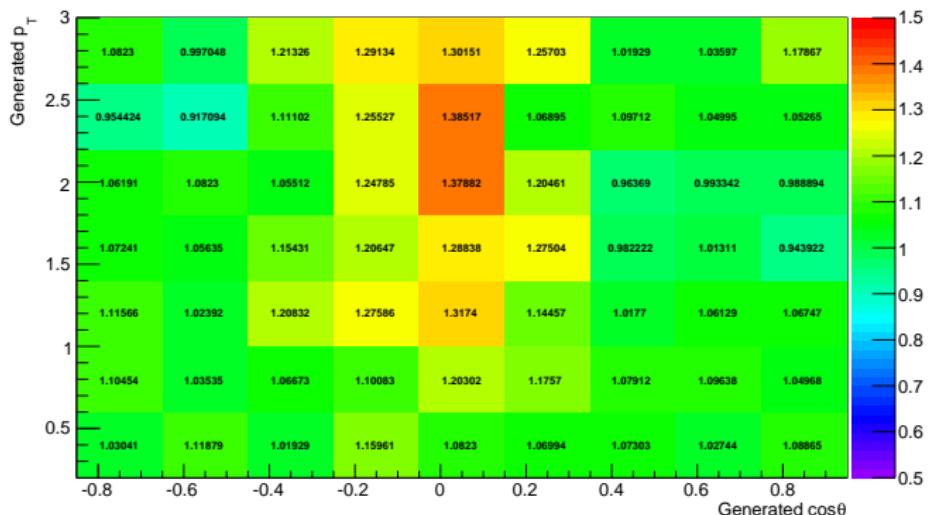
GENFIT 2 with  $\geq 2$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of z pull distribution

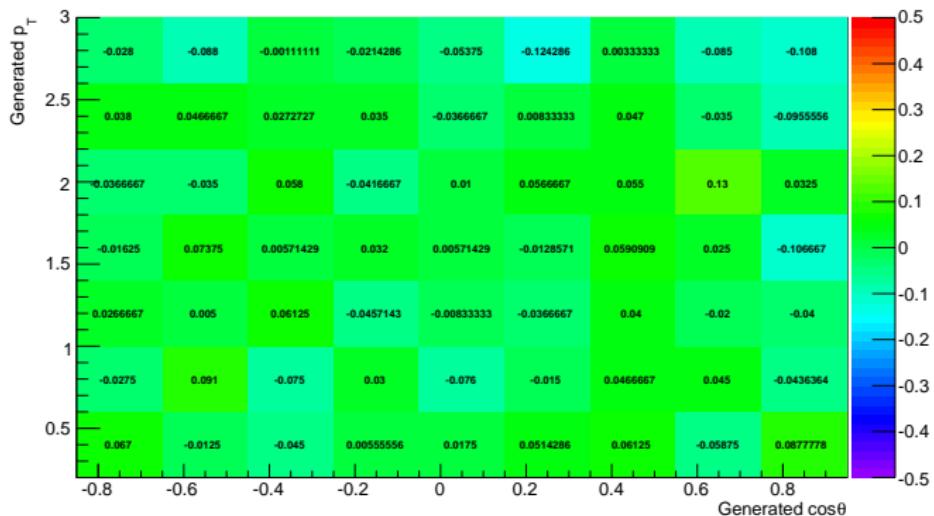
GENFIT 2 with  $\geq 3$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of px pull distribution



GENFIT 1 with 3 iterations.

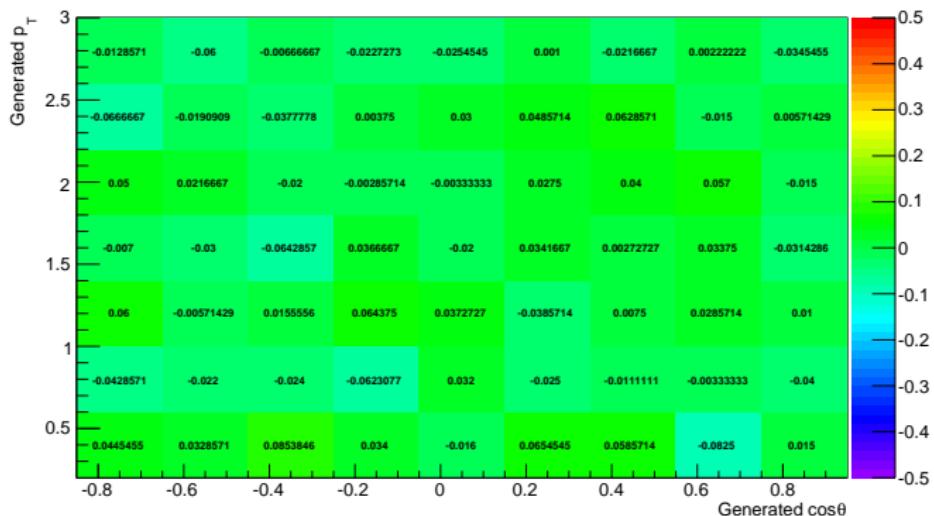


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of px pull distribution



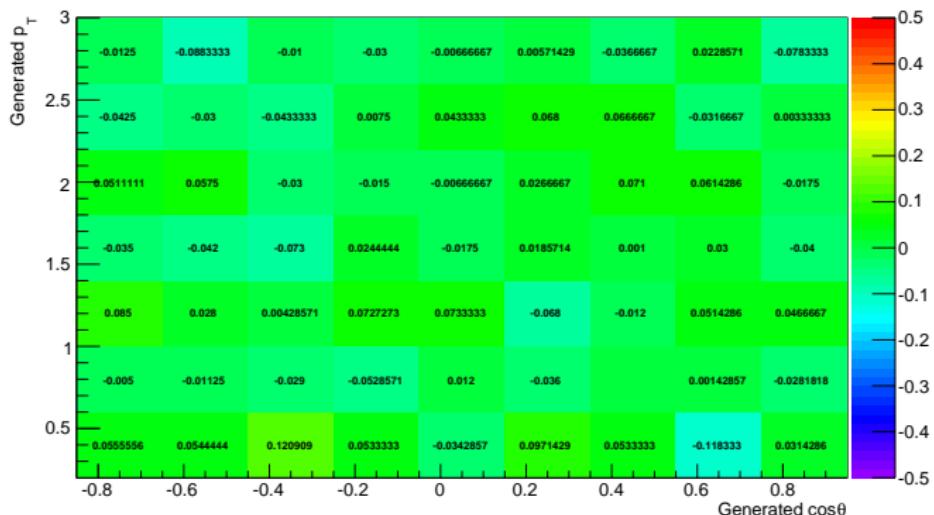
GENFIT 2 with  $\geq 2$  iterations.



## GENFIT 1 vs. GENFIT 2: 2D QA Plots

50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of px pull distribution

GENFIT 2 with  $\geq 3$  iterations.

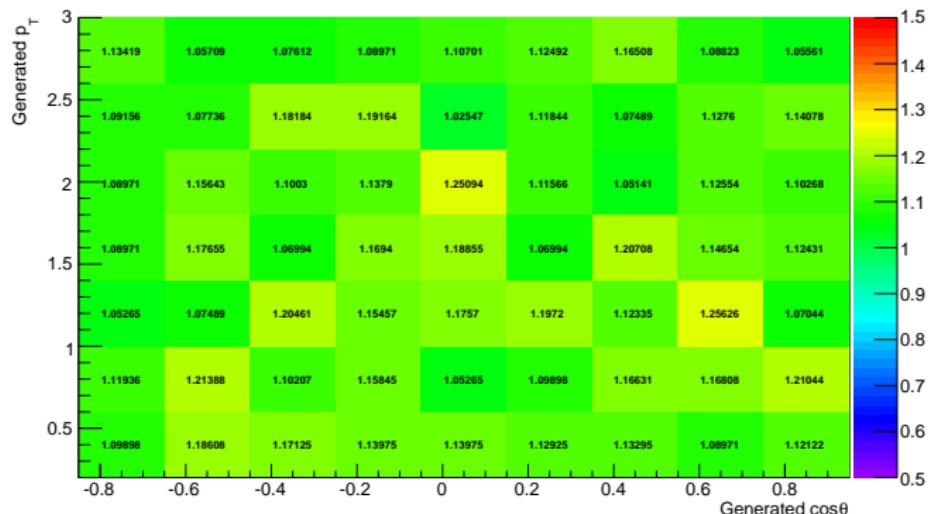


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of px pull distribution



GENFIT 1 with 3 iterations.

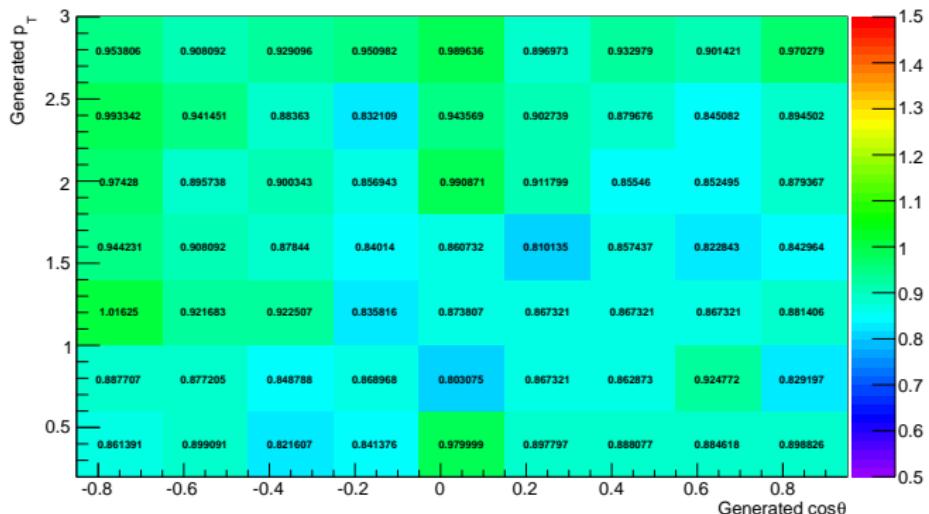


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50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

MAD std of px pull distribution



GENFIT 2 with  $\geq 2$  iterations.

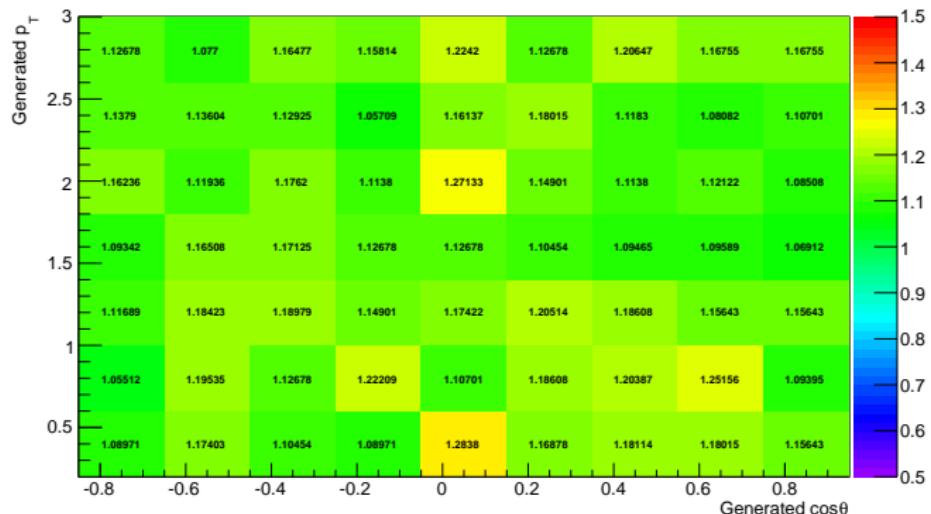


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MAD std of px pull distribution



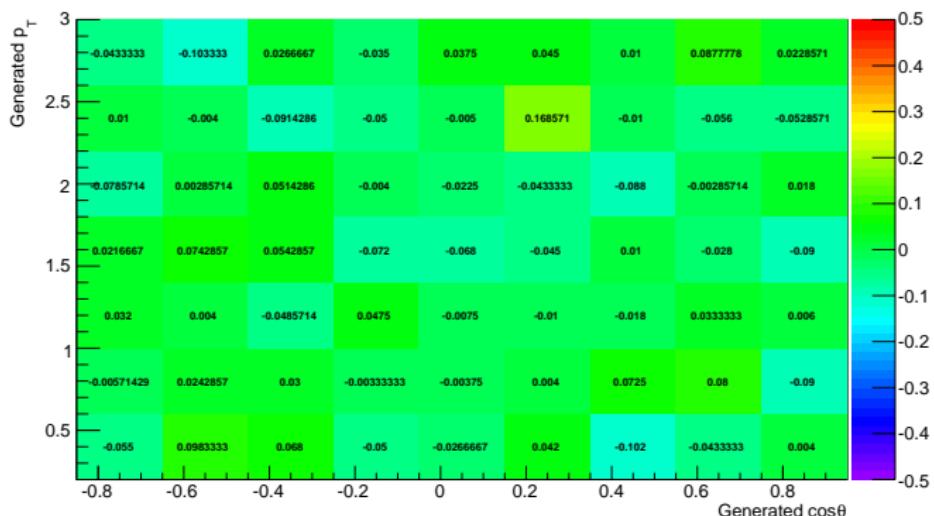
GENFIT 2 with  $\geq 3$  iterations.



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median of py pull distribution



GENFIT 1 with 3 iterations.

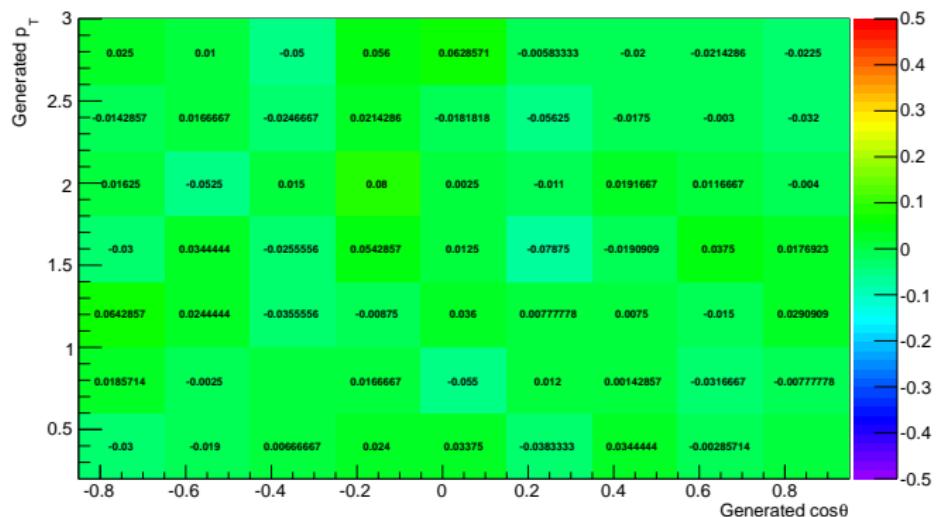


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median of py pull distribution



GENFIT 2 with  $\geq 2$  iterations.

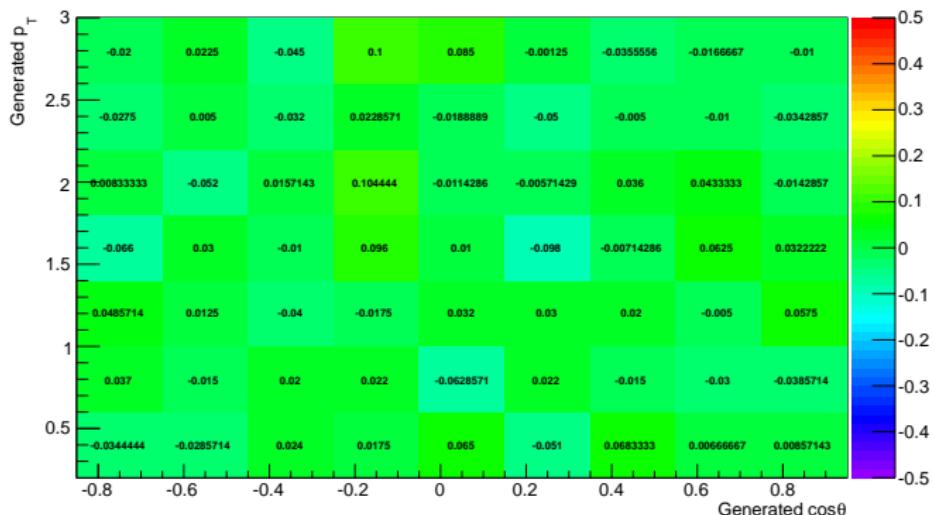


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median of py pull distribution



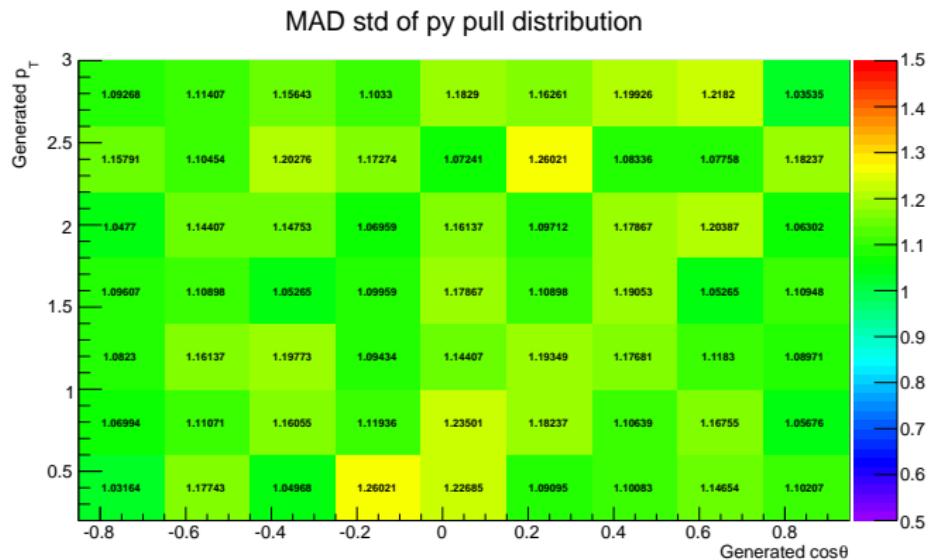
GENFIT 2 with  $\geq 3$  iterations.



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GENFIT 1 with 3 iterations.

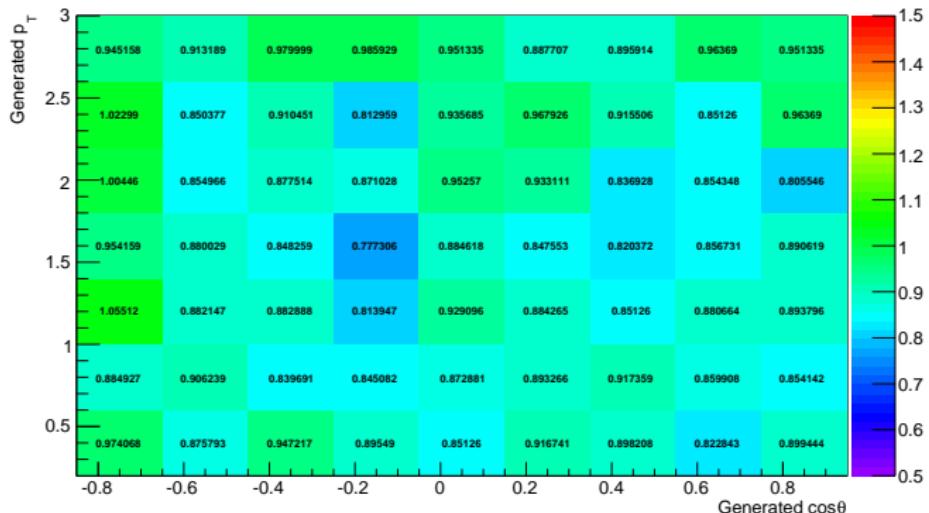


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MAD std of py pull distribution



GENFIT 2 with  $\geq 2$  iterations.

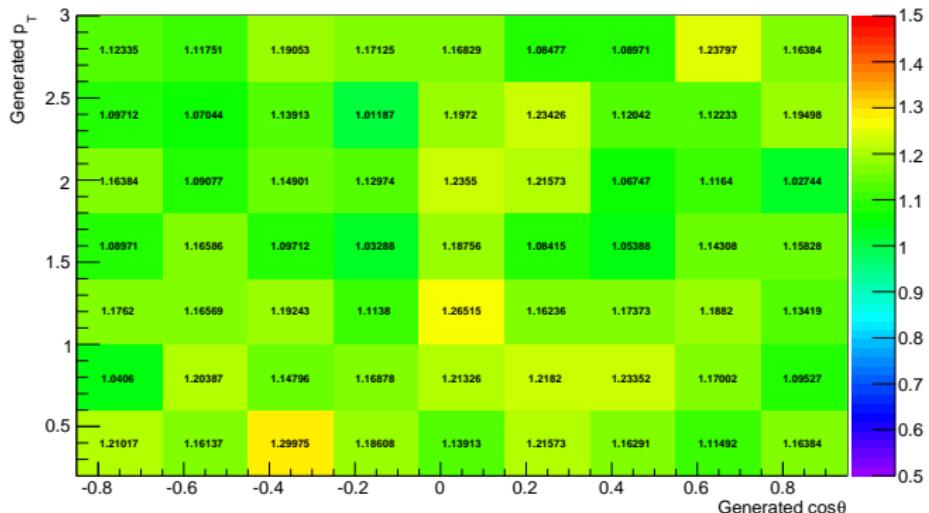


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MAD std of py pull distribution



GENFIT 2 with  $\geq 3$  iterations.

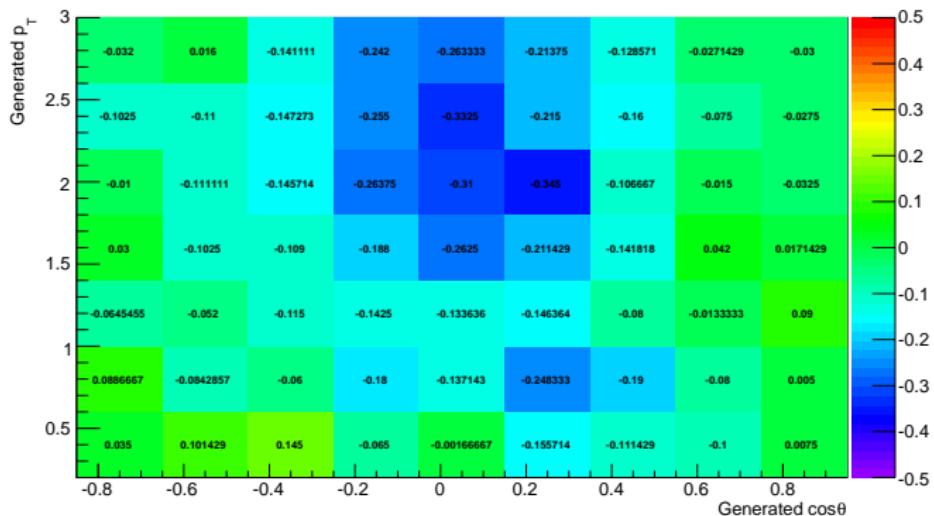


## GENFIT 1 vs. GENFIT 2: 2D QA Plots



50000 Pions, (0.2 to 3.0) GeV,  $\theta = (17 \text{ to } 150)^\circ$ , pxd + svd + cdc.

median of pz pull distribution



GENFIT 1 with 3 iterations.

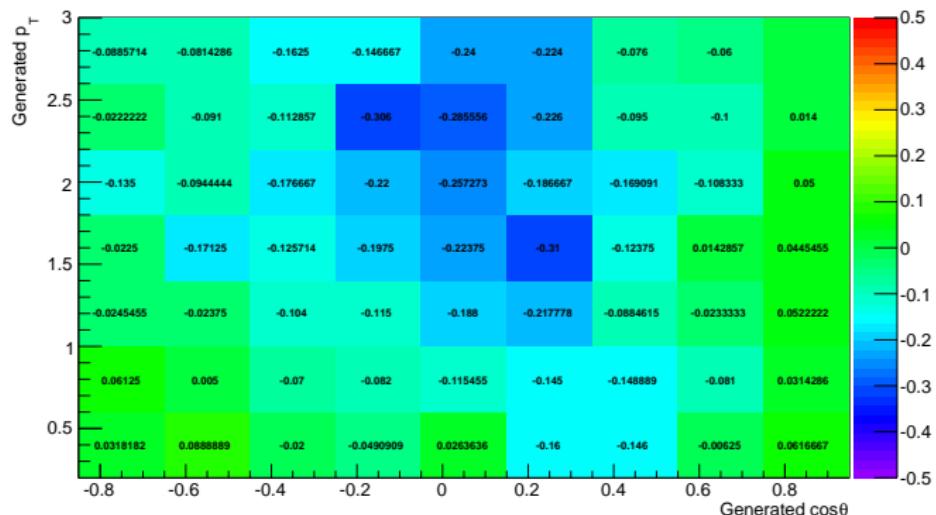


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GENFIT 2 with  $\geq 2$  iterations.

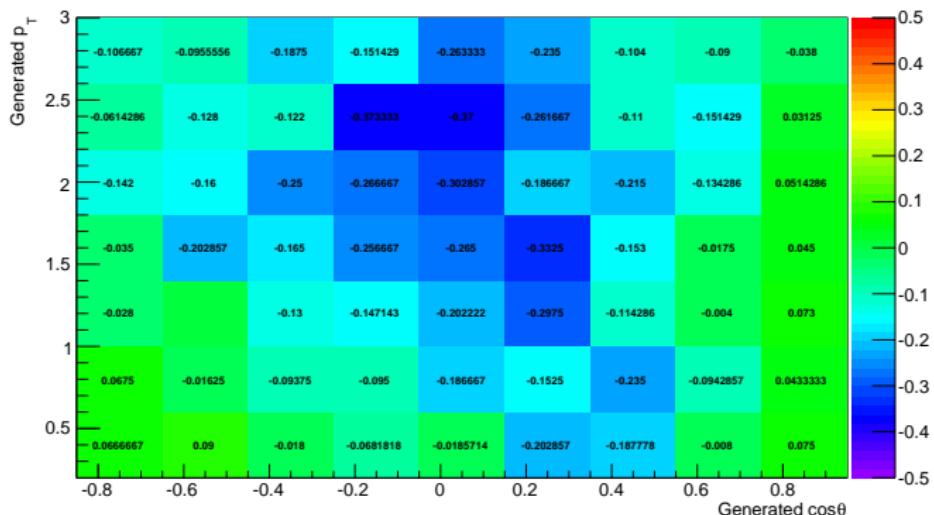


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median of pz pull distribution

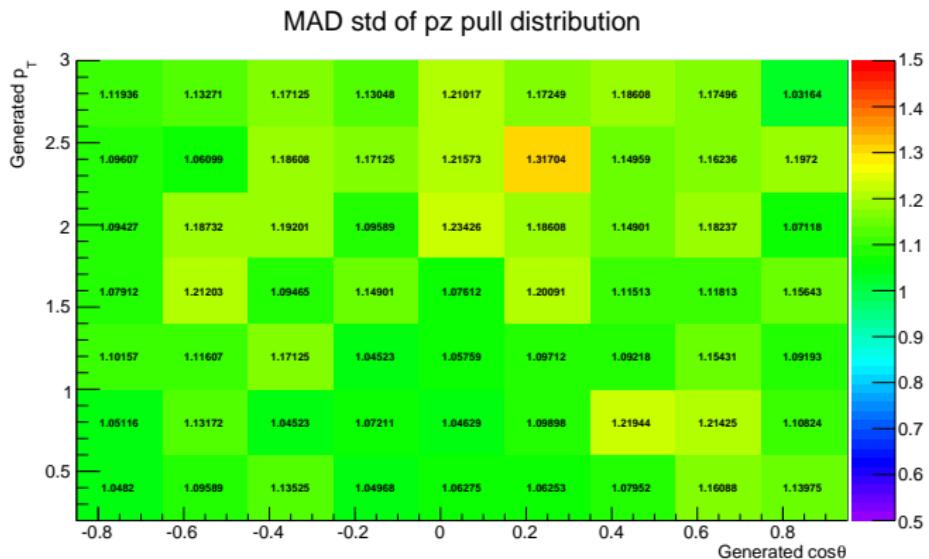


GENFIT 2 with  $\geq 3$  iterations.



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## GENFIT 1 with 3 iterations.

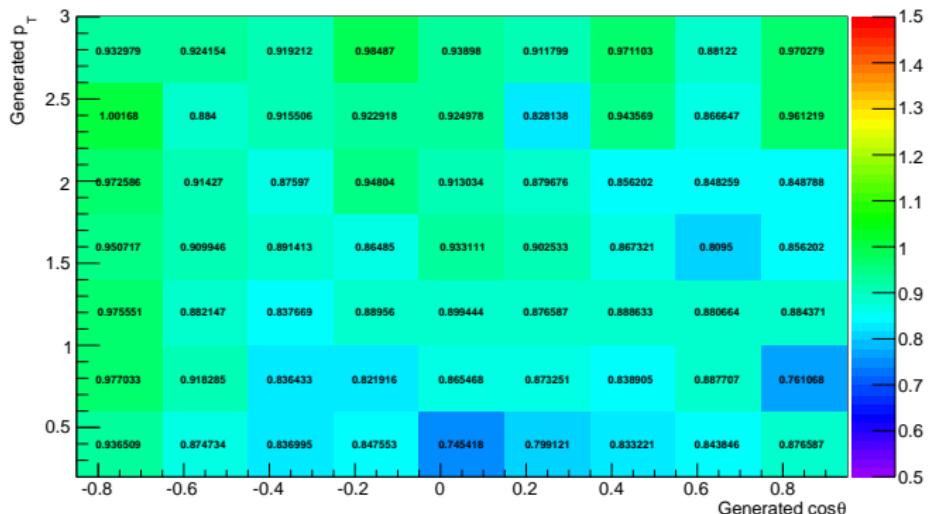


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MAD std of pz pull distribution



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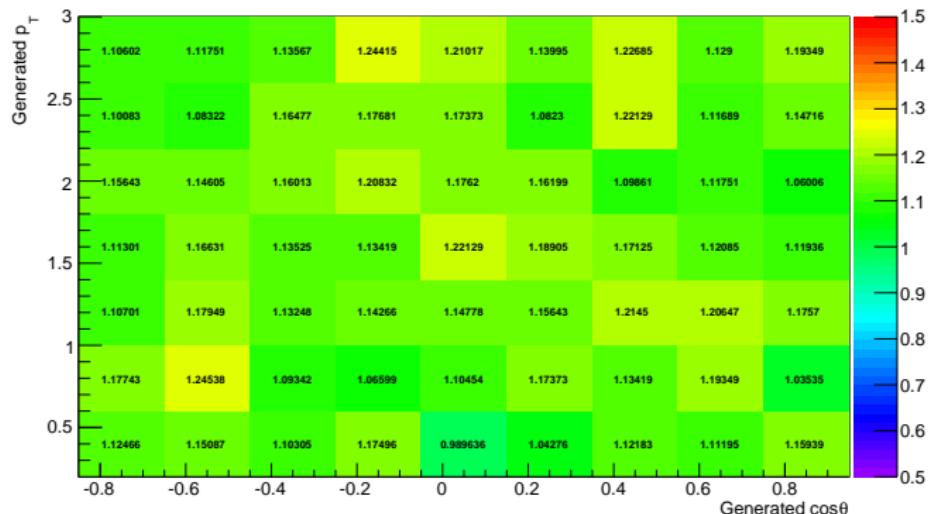


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