

ILDPerformance package

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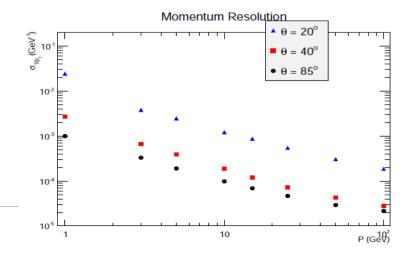
Introduction

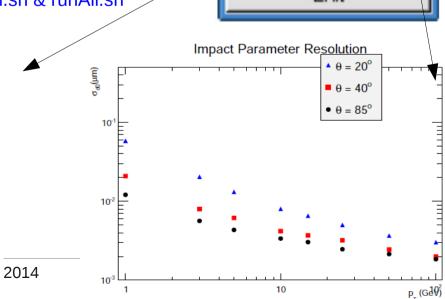
- Scope
 - Allow to evaluate the ILD tracking performance in an easy & fast way
 - > Aid to detector optimisation studies / algorithm development
- Based on Diagnostics processor
 - It's run after the reconstruction
 - Input: studied track collection & relations from RecoMCTruthLinker processor
 - Processor parameters to define the examined MCParticle subsample
 - Processor parameters to define efficiency
 - Default > 75% purity & ≥ 4hits in Si detectors
 - Output: ROOT file
 - ROOT tree with relevant info
 - True reco track parameters & errors, track quality, # of hits, "bad" tracks etc
 - Canvases with track parameters residuals & pulls, finding efficency
- svn co https://svnsrv.desy.de/public/marlinreco/ILDPerformance/trunk
- The user need to build the package and (for time being) export its library to MARLIN_DLL variable

Aid to optimisation studies

- Often it is very useful to study the momentum & impact parameter resolution, and compare them to the std DBD plots
- The user need to
 - source tracking/scripts/runAll.sh
 - It simulates, reconstructs (using the std reconstruction xml file) and analyses 1000 muon tracks for 8 momentum and 4 polar angle values
 - Runs the macros which produce the resolution plot
 - Initiate a ROOT gui
 - One can directly plot the resolutions
- One can do the above procedure step by step by using runMokka.sh, runMarlin.sh and runAll.sh scripts
 - For example, if wants to study the resolution using a different tracking algo. should modify the stdreco_tracking.xml and run runMarlin.sh & runAll.sh







Residuals

Pulls

IPResolution

Momentum Resolution

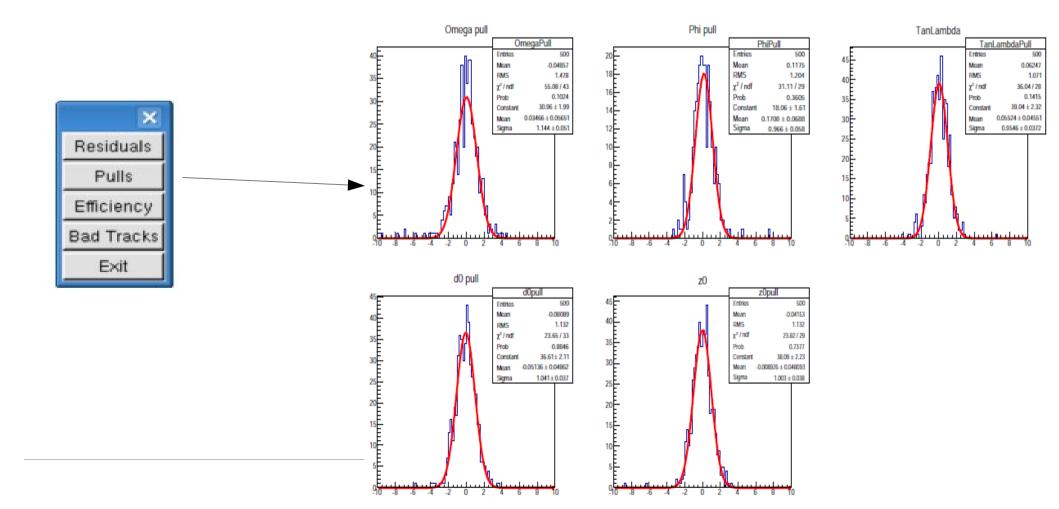
Exit

Pattern recognition

- Evaluation of the efficiency / "bad" track rate of a new tracking algorithm
 - source tracking/scripts/PatRec.sh
 - Edit the script and assign to INFILE variable the relevant reconstructed Icio file
 - Turn the following Diagnostics processor parameters to <u>true</u> (Diagnostics.xml)
 - PhysSampleOn
 - TrkEffOn
 - One will be lead to a similar ROOT gui and plot efficency or "bad" tracks as a function of P
- The script by default examines the full tracks
 - One can change track collection by appropriately editing the RecoMCTruthLinker & Diagnostics processor parameters inside Diagnostics.xml

Track fitting studies

- Plotting of track parameters pulls and residuals
 - tracking/scripts/PatRec.sh
 - Give a reco lcio file with single tracks as INFILE
 - For example for 5 GeV muons at $\theta = 85^{\circ}$



Outlook

- Add a processor which calculates track hit residuals
 - Pattern recognition & track fitting
- Add ROOT files with reference plots corresponding to DBD detector & software
 - Directly compare a new detector configuration or software tool with the standard ones