Extending pandaroot to run Kalman Filter with all Hypotheses

Walter Ikegami Andersson

Uppsala University on behalf of the $\overline{P}ANDA$ collaboration

PANDA Tracking Workshop September 18-19, 2018 GSI

Kalman filter current design

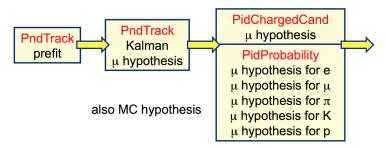


Figure from PANDA Computing Workshop 2017

Kalman filter current design

Following tasks are run during the reco- and PID stages:

- RecoKalmanTask: Perform track fitting (Kalman Filter) with μ hypothesis
- PidCorrelator: Extrapolate tracks to EMC with π hypothesis to distinguish neutrals Cut is made on EMCquality variable (done by user at analysis level instead?)
- PidAlgo: Run all PID algorithms from each subdetector
- McCloner: Filter out MC tracks that are not related to (for 500 $\overline{\Lambda}\Lambda$, 20.000 \rightarrow 2000 tracks)

Pull Distributions (Stretch Functions)

The pull distribution z for an observation i, also known as stretch function, is defined as (calling it MC pull distribution)

$$z_i = \frac{y_{MC,i} - \eta_i}{\sigma(\eta_i)}$$

- Pull distribution used to study performance of kinematic fits
- MC pull used to see performance of Kalman filter

All tests in this presentation on $\overline{p}p \to \overline{\Lambda}\Lambda \to \overline{p}p\pi^+\pi^-$ @ 1.642 GeV/c

Performance of Kalman Filter

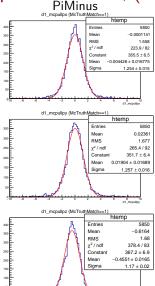
In an ideal fit:

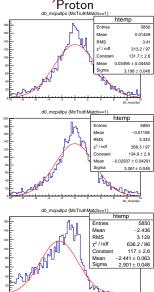
Pull distributions should be normal distributed

There are issues with the fit if:

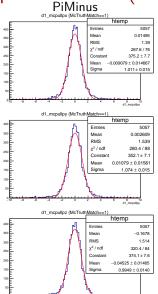
- Skews in pull distributions \implies bias in measurements
- Narrower (broader) normal distributions ⇒ errors over(under)estimated
- Background contamination can also alter pull distributions remove with MC truth info

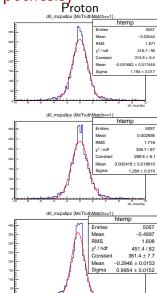
MC pull distribution, (no Ideal Hypothesis) Proton





MC pull distribution (with Ideal Hypothesis)
Proton





Kalman filter new design

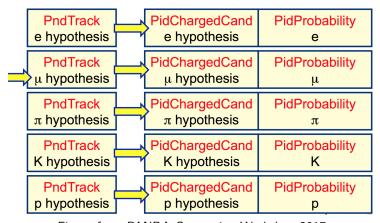


Figure from PANDA Computing Workshop 2017

Reconstruction Stage

- In reco macro, pass option "multikalman", and Kalman filter will run 5 times with all hypotheses
- If e.g. only pions and protons are interesting, pass them in option string

```
// ---- Initial Settings
PndMasterRunAna *fRun= new PndMasterRunAna();
fRun->SetInput(input);
fRun->SetOutput(output);
fRun->SetFriend1(friend1);
fRun->SetFriend2(friend2);
fRun->SetFriend3(friend3);
fRun->SetFriend4(friend4);
fRun->SetParamAsciiFile(parAsciiFile);
fRun->SetUp(prefix);

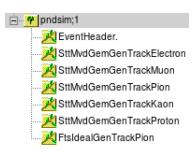
// --- Add tasks
// fRun->SetOptions("multikalman;pion;proton");
fRun->SetOptions("multikalman");
fRun->AddRecoTasks();
```

Reconstruction Stage

PndMultiKalmanTask properties:

- Possible to run with standard pattern recognition as well as ideal PR
- If no hypotheses are specified in the options, all five hypotheses are used
- Currently,
 PndMultiKalmanTask does not run with FTS PR output...
 ⇒ only running pion

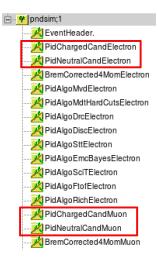
hypothesis for now



PID Stage: PndCorrelator

PndCorrelator - first task PID stage

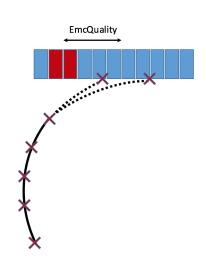
- Produces PndPidCanididates for
 - Track candidates
 - charged
 - EMC bumps
 - neutral
- PndCorrelator run for each track fit hypothesis



PID Stage: PndCorrelator

Why 5 sets of neutral PndPidCandidates?

- A given track can have successful fit with one hypothesis but failed with another
- Neutrals produced if EmcBump distance to track larger than EmcQuality
- Extrapolating track to EMC dependent on particle hypothesis
- Five sets of neutrals produced by extrapolating with different hypothesis



PID Stage: PndPidAlgos, PndMcCloner

Significant changes:

- PndAlgos and PndBremCorrector running for each hypothesis
- New PndMcCloner2 class:
 - Checks if a MC track is related to a track that was successfully fitted with any hypothesis
 (Again, Kalman fit can converge for e.g. pion hypothesis but fail for proton hypothesis).
 - Save MCtrack if that is the case.
- Everything handled in new PndMasterMultiPidTask class
- If no hypotheses specified in options, all five hypotheses are used

```
// --- Add tasks
fRun->SetOptions("multikalman;pion;proton");
// fRun->SetOptions("multikalman");
fRun->AddPidTasks();
```

Analysis Stage

PndAnalysis - main analysis object taking care of file read/write

- Now reads up to five sets of TClonesArrays, one set for each hypothesis
- When filling RhoCandLists, following is specified
 - Particle hypothesis + PID criteria + Charge
 - Name of branch containing subdetector PID info
- Usage same as before

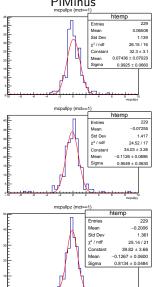
```
// theAnalysis->FillList(piplus, "PionAllPlus");
theAnalysis->FillList(pimlus, "PionAllMinus");
// theAnalysis->FillList(protonplus, "ProtonAllPlus");
theAnalysis->FillList(protonplus, "ProtonAllPlus");

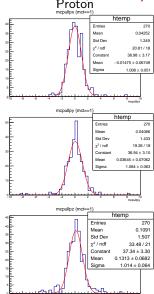
theAnalysis->FillList(pimlus, "PionAllMinus");

theAnalysis->FillList(pimlus, "PionAllMinus", "PidAlgoIdealCharged");
theAnalysis->FillList(protonplus, "ProtonAllPlus", "PidAlgoIdealCharged");
theAnalysis->FillList(protonminus, "ProtonAllMinus", "PidAlgoIdealCharged");

theAnalysis->FillList(pimlus, "PionTightPlus", "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PionTightPlus", "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PionTightPlus", "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PionTightPlus", "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PidAlgoStt;PidAlgoMvd;PidAlgoEmmus, "PidAlgoStt;PidAlgoMvd;PidAlgoMvd;PidAlgoEmmus, "PidAlgoStt;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoMvd;PidAlgoM
```

MC pull distribution (Multikalman, with tight pid cut)





Summary

- Pandaroot simulation and analysis chain modified to run MultiKalman filter
- Preliminary tests shows significant improvements for protons in $\overline{p}p \to \overline{\Lambda}\Lambda$ @ 1.642 GeV/c
- Feature will be pushed to dev branch soon...
- Issues: Ideal FTS pattern recognition not working with all hypotheses...

Outlook

- New implementation runs all tasks 5 times, regardless of physical motivation
 - Running Bremstrahlung corrector for all hypotheses useful for systematic error studies or useless?
- Belle 2 experiment runs π hypothesis for e, μ, π , compromise between accuracy and computing resources (see talk by S. Spataro at CHEP)
- Should cut on EmcQuality variable be available to analysts (instead of a fix cut hidden from the user?)
 - Removing the cut would remove the need of having 5 lists of PndPidCandidates for neutrals

Outlook

- New implementation runs all tasks 5 times, regardless of physical motivation
 - Running Bremstrahlung corrector for all hypotheses useful for systematic error studies or useless?
- Belle 2 experiment runs π hypothesis for e, μ, π , compromise between accuracy and computing resources (see talk by S. Spataro at CHEP)
- Should cut on EmcQuality variable be available to analysts (instead of a fix cut hidden from the user?)
 - Removing the cut would remove the need of having 5 lists of PndPidCandidates for neutrals

Thank you for your attention!

Backup