

Preliminary analysis of LMD in RGA

Finding light mesons and the code that does it

Tyler Viducic

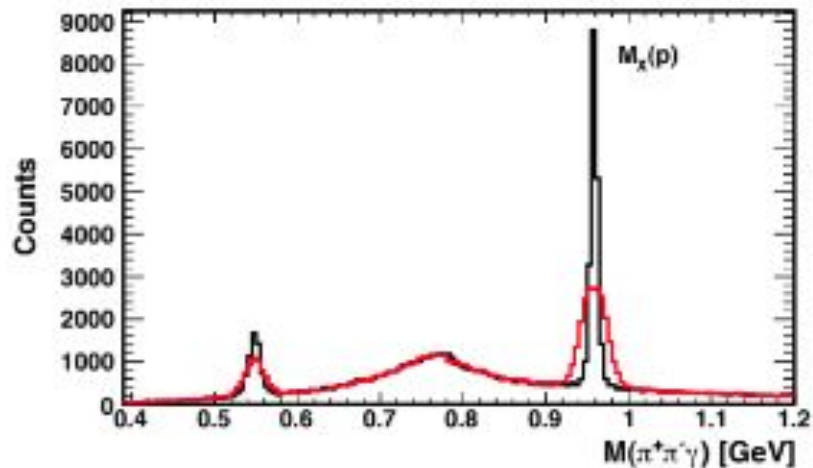
Thursday, June 13, 2019

```
354 //get particle data
355
356 PhysicsEvent physEvent = setPhysicsEvent(beam, event);
357 if (filter.isValid(physEvent)) {
358     //System.out.println(physEvent.toLundString());
359     Particle mx_P = physEvent.getParticle( operator: "[b] + [t] - [2212]");
360     Particle mx_PPipPim = physEvent.getParticle( operator: "[b] + [t] - [2212] - [211] - [-211]");
361     Particle mx_PPipPimGam = physEvent.getParticle( operator: "[b] + [t] - [2212] - [211] - [-211] - [221]");
362     Particle pgam = physEvent.getParticle( operator: "[22]");
363     Particle me_PPipPim = physEvent.getParticle( operator: "[b] + [t] - [2212] - [211] - [-211]");
364     Particle im_PipPim = physEvent.getParticle( operator: "[211] + [-211]");
365     Particle pP = physEvent.getParticle( operator: "[2212]");
366
367 //Fill Histograms
368
369 if (pgam.e() > cutPgam && me_PPipPim.e() > cutMe_PPipPim && Math.abs(mx_PPipPim.mass2()) < cutMx_PPipPim
370     && Math.abs(me_PPipPim.e() - pgam.e()) < cutMe_PPipPimPgamSubtract
371     && Math.abs(mx_PPipPimGam.mass2()) < cutMx_PPipPimGam) {
372     hMxPuncut.fill(mx_P.mass());
373 }
374 if (Math.abs(mx_PPipPimGam.mass2()) < 0.01 && Math.abs(mx_PPipPim.mass2()) < 0.005) {
375     hMxP.fill(mx_P.mass());
376 }
377 if (Math.abs(mx_P.mass() - mRho) < cutRhoRegion && me_PPipPim.e() > cutMe_PPipPim) {
378     hPGam.fill(pgam.e());
379 }
380 if (Math.abs(mx_P.mass() - mRho) < cutRhoRegion && pgam.e() > cutPgam) {
381     hMe_PPipPim.fill(me_PPipPim.e());
382 }
383 if (pgam.e() > cutPgam && me_PPipPim.e() > cutMe_PPipPim && Math.abs(mx_P.mass() - mRho) < cutRhoRegion
384     && Math.abs(mx_PPipPimGam.mass2()) < cutMx_PPipPimGam && Math.abs(mx_PPipPim.mass2()) < cutMx_PPipPim) {
385     hMe_PPipPimPgam.fill(me_PPipPim.e() - pgam.e());
386 }
387 if (pgam.e() > cutPgam && me_PPipPim.e() > cutMe_PPipPim && Math.abs(mx_P.mass() - mRho) < cutRhoRegion
388     && Math.abs(mx_PPipPim.e() - pgam.e()) < cutMe_PPipPimPgamSubtract
389     && Math.abs(mx_PPipPim.mass2()) < cutMx_PPipPim) {
390     hMxPPipPimGam.fill(mx_PPipPimGam.mass2());
391 }
392 if (pgam.e() > cutPgam && me_PPipPim.e() > cutMe_PPipPim && Math.abs(mx_P.mass() - mRho) < cutRhoRegion
393     && Math.abs(me_PPipPim.e() - pgam.e()) < cutMe_PPipPimPgamSubtract
394     && Math.abs(mx_PPipPim.mass2()) < cutMx_PPipPim) {
395     hMxPPipPimGam.fill(mx_PPipPimGam.mass2());
396 }
397 if (pgam.e() > cutPgam && me_PPipPim.e() > cutMe_PPipPim && Math.abs(mx_P.mass() - mRho) < cutRhoRegion
398     && Math.abs(me_PPipPim.e() - pgam.e()) < cutMe_PPipPimPgamSubtract
399     && Math.abs(mx_PPipPimGam.mass2()) < cutMx_PPipPimGam) {
400     hMxPPipPimGam.fill(mx_PPipPimGam.mass2());
401 }
402 if (pgam.e() > cutPgam && me_PPipPim.e() > cutMe_PPipPim
403     && Math.abs(me_PPipPim.e() - pgam.e()) < cutMe_PPipPimPgamSubtract
404     && Math.abs(mx_P.mass() - mRho) < cutRhoRegion && Math.abs(mx_PPipPim.mass2()) < cutMx_PPipPim
```

Goals of physics analysis

- Extract physics events with final state $ep\pi^+\pi^-\gamma$
- Identify events where parent particle for the $\pi^+\pi^-\gamma$ is a light unflavored meson (η, ρ, η')

-Pictured at right is what we hope to see



A look at the code

- Code written using Groovy and JAVA analysis tools
- An example of how to do physics analysis using the JAVA analysis framework
- Please feel free to ask questions about the script as we go

A look at the code

-First step:

-Filter code for desired final state

-Attempted with exclusive final state of $e p \pi^+ \pi^- \gamma$

-Not enough statistics yet

-Tried semi-inclusive $e p \pi^+ \pi^- X_n$

-Much better statistics

-Runs used (10 files/run)

```
viducic@ifarm1801> cd /work/clas12/rg-a/production/recon/pass0/v5/mon/  
005030/ 005040/ 005053/ 005126/ 005153/ 005165/ 005181/ 005195/ 005201/ 005208/ 005258/  
005031/ 005041/ 005116/ 005128/ 005158/ 005166/ 005182/ 005196/ 005202/ 005211/ 005259/  
005032/ 005043/ 005117/ 005129/ 005159/ 005167/ 005183/ 005197/ 005203/ 005212/ 005261/  
005036/ 005046/ 005120/ 005130/ 005160/ 005168/ 005190/ 005198/ 005204/ 005252/  
005038/ 005051/ 005124/ 005138/ 005162/ 005169/ 005191/ 005199/ 005205/ 005253/  
005039/ 005052/ 005125/ 005139/ 005163/ 005180/ 005193/ 005200/ 005206/ 005257/
```

A look at the code

-Second step:

- Filter file for wanted banks (REC::Particle, REC::Event)

- This dramatically speeds up read time

-How-to:

- On farm:

- ```
~/home/viducic/jaw-0.9/bin/hipoutils.sh -filter -b
REC::Particle,REC::Event -o output_file_name.hipo
input_file_name.hipo
```

# Time for physics

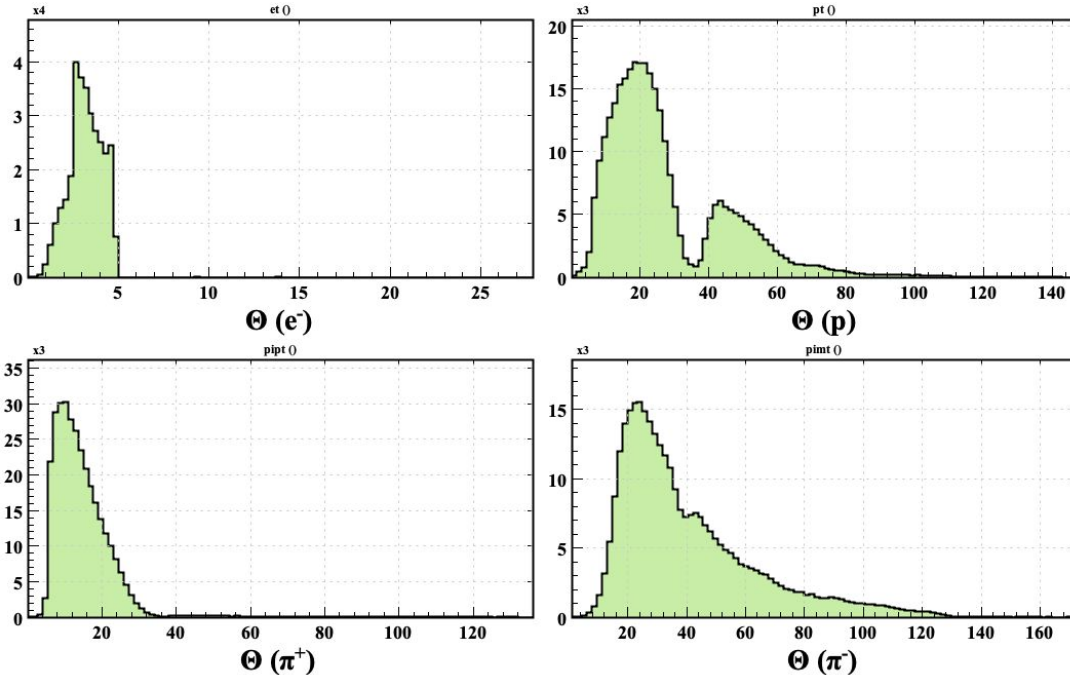
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-First a look at the code:

# Time for physics

-Cut selection

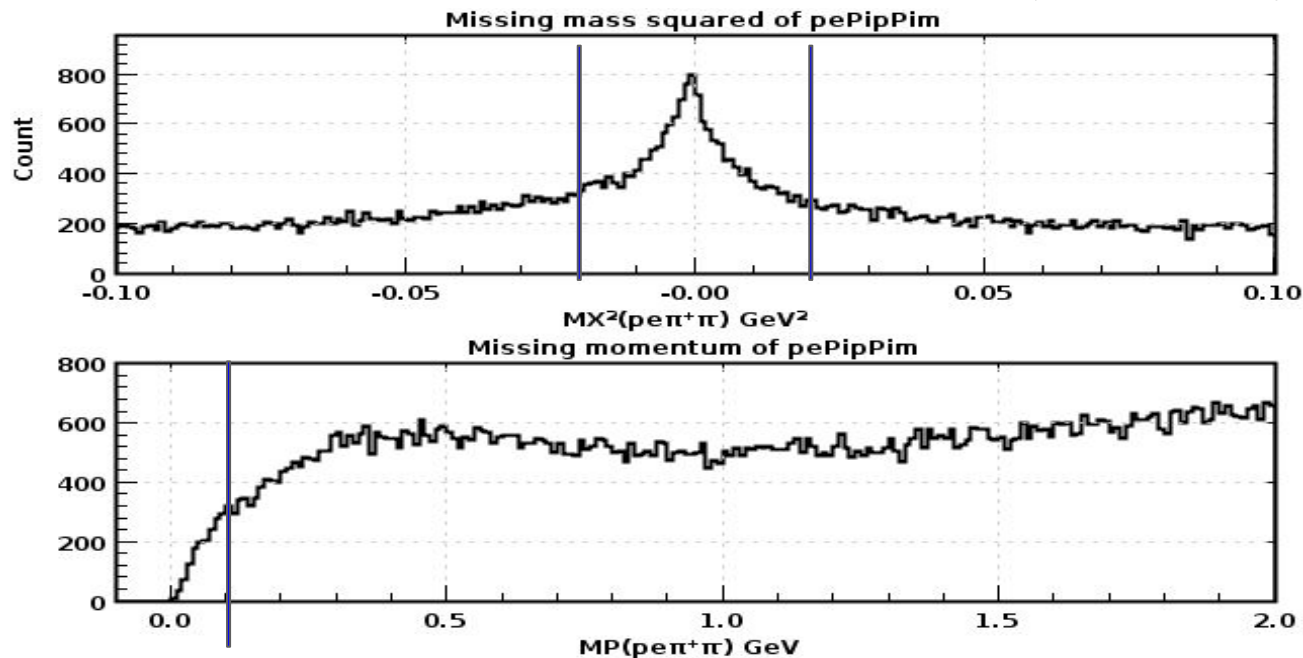
-Electron in FT -> decay particles in FD



# Time for physics

-Cut selection

$$-|MX^2(pe\pi^+\pi^-)| < 0.02 \text{ GeV}^2 \ \&\& \ MP(pe\pi^+\pi^-) > 0.1 \text{ GeV}$$





# Time for physics

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-And now a look at the results.

-How to open saved histograms:

```
-/home/gavalian/coatjava/bin/browser my_file.hipo
```

# What's Next

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- In conclusion: Not great!
- Need more statistics and better calibration
- Evaluate viability of analysis with FastMonteCarlo
  - FastMC gives acceptance within minutes to single digit%
  - Cuts down on need to fully simulate with GEMC

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```
if(seminar.hasQuestions()){
 String question = seminar.ask();
 tyler.answer(question);
}else{return;}
```