Status of Ke2/Kµ2

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Status of the software

- BUG on recovery corrected + prompt γs for Ke2g and ke3 added
- Critical issue rare decays need several days to optimize each cut
- Move from PC to Farm
 - Modified HBOOK open Gzipped ntuples directly (save factor ~2 disk space)
- All chain tested and ready

The way to the 1% accuracy: statistical error

- More MC for background
 - Analysis of new production well under way ('04+'05 0.5 scale factor almost ready)
- Reduce background contamination
 - Use kinematics
 - Redundant measurement of K momentum + tighter cut on error a priori on missing mass
 - Better particle ID using calorimeter info:
 - Recover all cells with one only PMT fired
 - New definition of shower direction and impact point
 - New variables: higher momenta of shower distribution

Kinematics



Kinematics: error on lepton mass

 Error on lepton mass still not fully exploited



Cells recovery

- Use incomplete cells (1 pmt only fired) to determine all cluster properties*
- Coordinate along fiber from centroid position



*only for energy evaluation all cells are used in the standard reconstruction

Cluster direction & impact point position



New variables

- All variables benefits from new procedure
- Skewness brings additional separation



Effect

Selection still to be tuned: **DATA(NEW)** 3000 factor 3 more rejection DATA(OLD) 2500 can be obtained, paying 25% on efficiency 2000 1500 Nsig = 5364.4800 Nsig = 7303.9350 700 300 600 250 500 200 400 -4000 -2000 -6000 6000 150 0 2000 4000 300 MeV^2 M 100 200 50 100 -10000 -10000 -8000 -6000 -4000 -2000 2000 4000 -8000 -6000 -4000 -2000 2000 4000 0 0

M_{lcp}, MeV²

M_{lcp}, MeV²

The way to the 1% accuracy: systematic error

- Dominated by PID selection efficiency
 - Adjusting MC response to data
 - Energy scale
 - Threshold
 - Energy response saturation
 - Control samples
 - Increased statistics for Ke3 CS
 - Trying to use Bhabha events
 - Ke3 from charged kaons

Threshold

- Cell energy from MC distribution shows an excess wrt data at low energy
- <# of cell> from MC slightly higher wrt data
- Apply additional threshold at 0.8 pe



Energy scale and resolution

 Resolution in MC worse that in data Scale almost OK Rescale MC cell energies using E tru democratically: $E_i^{new} = E_i a + E_i (1-a) \frac{P}{\sum E_i}$ a=0.92



Saturation?

- E_{Cell} distribution from MC has a longer tail wrt data
- Hints of saturation?
 - Physics (Mario)
 - PMT (Alex,Tom)
- Practically:
 - Apply saturation with Fermi-Dirac
 - New scale ~1.15: almost able to recover e/µ energy response ratio



Conclusion

- New analysis chain is ready
- Refined kinematic cuts
- Improvement in particle ID:
 - New cluster depth
 - Skewness
 - Threshold refinement
 - Cell calibration
- Almost ready to finilize, hope to do so for late winter conferences