

Studies with Matrix Element Method

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May 28, 2013

Outline

MEM and Normalization

Likelihood function and normalization of Matrix Element Method outputs

Validation

Basic validation and application studies to a search for heavy di-muon resonances.

MEM and Normalization of Outputs

Analysis Level Selections

In an experimental set-up one usually measures the physics objects with **detection efficiencies** after some online/offline **selections**.

X : full phase space of an experimental event $\rightarrow X'$: reduced phase space of the detected object.

Y : full phase space at parton level $\rightarrow Y'$: sub-phase space

It can be proven that likelihood function of underlying model parameters M , for one experimentally observed events x in the reduced phase space X' of selectable events (analysis-driven) is given by;

Likelihood Function :

$$L(\alpha) = pdf(x|\alpha \text{ AND } x \text{ in } X') = \frac{1}{\epsilon' \sigma'} \int_{Y'} \frac{d\sigma_\alpha}{dy} w(x|y) dy$$

Likelihood Function with MEM

$$L(\alpha) = pdf(x|\alpha \text{ AND } x \text{ in } X') = \frac{1}{\epsilon' \sigma'} \int_{Y'} \frac{d\sigma_\alpha}{dy} w(x|y) dy$$

Where;

- $\frac{d\sigma_\alpha}{dy}$: is parton level cross-section for a given model
- $w(x|y)$: "transfer functions" (Note to remark on next slide)
- Y' : reduced parton level phase-space in which $w(x|y) = 0$ for any x in X' and y outside Y'
- σ' : reduced cross-section at the parton level of Y'
- ϵ' : efficiency of events in Y' to end up in reduced phase space X'
- σ' : the reduced cross-section for parton level events in Y'

Remarks on Normalisation of Outputs

Programs like MadWeight returns only the so-called "weight"

$$W = \int_{Y'} \frac{d\sigma_\alpha}{dy} w(x|y) dy$$

- Where $w(x|y)$ is what the user has defined, i.e. often unit normalized function of x and y .
- Factorization of single object is assumed, so y and x will indicate in the following, for simplicity, a single object (parton and reco-level)

$w(x|y)$ must satisfy $\int_X w(x|y) dx = 1$ for any y in Y

- Where X is full experimental event space, which includes non-reconstructed objects.
- In other words if $u(x|y)$ is unit-normalized function of x for given y , then one should use $w(x|y) = \varepsilon(y)u(x|y)$ where $\varepsilon(y)$ (usually < 1 for leptons) is the efficiency of the reconstructed object

Likelihood Function with MEM

$$L(M) = \frac{1}{\epsilon' \sigma'} \int_{Y'} \frac{d\sigma_\alpha}{dy} \epsilon(y) u(x|y) dy$$

Overall Normalization :

- Fundamental for parameter estimation:

$$L(\{x_i\}|\mu, M) = \prod_i \frac{\mu}{\mu\sigma_{SES} + \sigma_{BEB}} W_S(x_i, M) + \frac{1}{\mu\sigma_{SES} + \sigma_{BEB}} W_B(x_i|B)$$

- For simplified hypothesis testing there is no problem because one usually ends up with a monotonous function of the likelihood ratio. Still it has an impact for significance computation using PLR test statistics.

$$K_D = \frac{W_S}{W_S + W_B} = \frac{1}{1 + \frac{\epsilon_B \sigma_B L_B}{\epsilon_S \sigma_S L_S}}$$

Likelihood Function with MEM

$$L(M) = \frac{1}{\epsilon' \sigma'} \int_{Y'} \frac{d\sigma_{\alpha}}{dy} \epsilon(y) u(x|y) dy$$

Normalization of transfer function :

- Only matters if $\epsilon(y)$ varies over an interval centered around the experimentally measured value x and as large as few times the resolution. If this is not the case, it goes out of integration and:
 - In Hypothesis testing it cancels out event-by-event in the likelihood ratio.
 - In parameter estimation ends up a overall multiplicative factor of a likelihood function.
- Special relevant case is $H \rightarrow ZZ \rightarrow 4l$, where it would require attention : Higgs properties with the same channel, if one pulls down the lepton P_T very low

Hypothesis Testing and Significance

Likelihood function can be used to compute the significance of a discovery or excluded cross-sections.

We use the asymptotic formulae with likelihood ratio test statistics to reduce computation time (G.Cowan, K.Cranmer, E. Gross,O.Vitells, arxiv:1007.1727)

Profile Likelihood Ratio for μ signal strength : $\lambda(\mu) = \frac{L(\mu, \hat{\hat{\theta}})}{L(\hat{\mu}, \hat{\theta})}$

Test statistics for discovery of a signal :

$$q_0 = \begin{cases} -2 \ln \lambda(0) & \hat{\mu} \geq 0 \\ 0 & \hat{\mu} < 0 \end{cases}$$

$\lambda(0)$ is the profile likelihood ratio for $\mu = 0$

Significance of the test is $Z = \sqrt{q_0}$

Validation

Validation of the method with
A search for Z' high mass resonance

Overview of the Validation

- MadGraph generator used for exclusive signal and background event generation.
 - Signal Events : $Z' \rightarrow \mu^- \mu^+$ (Seq. Standard Model)
 - Background Events : $DY \rightarrow \mu^- \mu^+$

Generator-level inv-mass range (Y') : $600 \text{ GeV} \leq M \leq 1400 \text{ GeV}$
Selection-level inv-mass range (X') : $800 \text{ GeV} \leq M \leq 1200 \text{ GeV}$
- Estimation of parameters: resonance mass, signal fraction.
- Significance of signal hypothesis
- Comparison with CMS Z' shape analysis was performed (with RooFit/RooStats)

Generation, Selection, Normalization

Signal mass = $1 \text{ TeV}/c^2$

Signal width = 3%

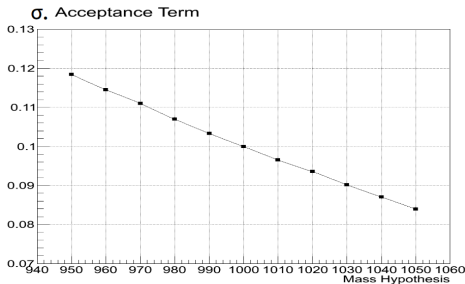
5% detector resolution effect
via Gaussian smearing on muon
energy

Analysis Selections :

- $800 \text{ GeV} \leq M \leq 1200 \text{ GeV}$
- $P_t^\mu > 45 \text{ GeV}/c$
- $|\eta| < 2.4$

$\sigma_s \cdot \epsilon_s$

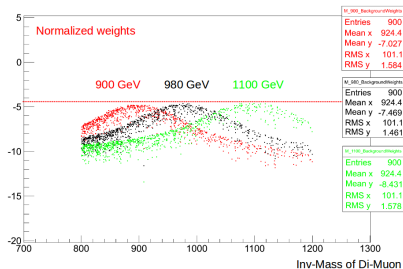
for different mass hypotheses.



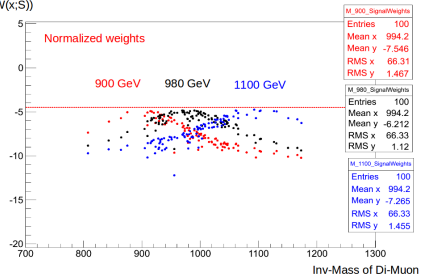
Likelihood Function Validation

$\log\text{-L}(M)$ for DY and Signal events under signal hypothesis of 900, 980, 1100 GeV

$\log(W(x;S))$

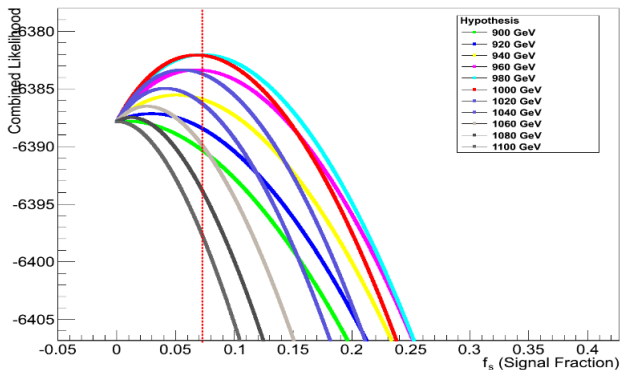


$\log(W(x;S))$



Parameter Estimation

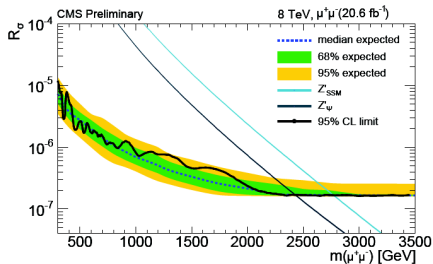
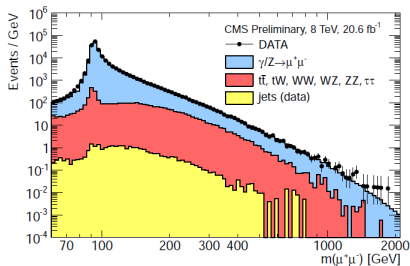
log-likelihood for Signal + Background events for one toy experiment



Signal fraction (f_s) and the resonance mass M_Z' are estimated from maximization of the combined likelihood function.

Comparison with CMS classical shape analysis

Classical analysis: search in the di-muon invariant mass distribution a peak compatible with a "narrow" resonance on top of a smooth background.



Comparison with shape analysis

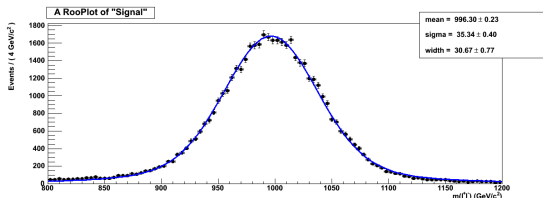
- Fit for signal fraction for a fixed mass hypothesis
- Simultaneous fit for signal fraction and mass.

Classical
shape based
analysis

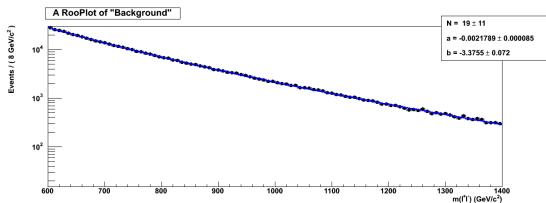


MEM with
MAD-Weight

Shapes for Classical Analysis



Signal Model :
 Breit-Wigner \otimes Gaussian

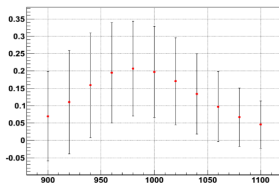


Background Model :
 $N \exp^{am} mb$

Comparison : Signal fraction and significance

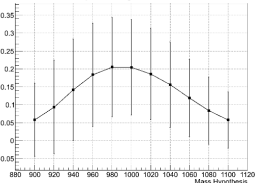
Pseudo experiments with 20B+5S events for fixed mass points

Classical Shape Analysis

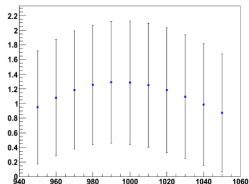
Mean Signal Fraction vs. Z' mass hypothesis

Matrix Element Method

Expected Signal Fraction

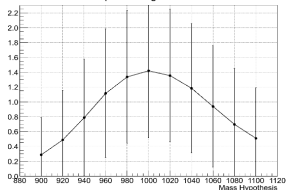


Classical Shape Analysis

Mean Significance vs. hypothesized Z' mass

Matrix Element Method

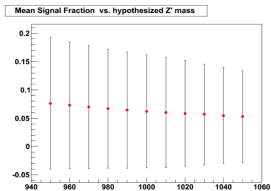
Expected Significance



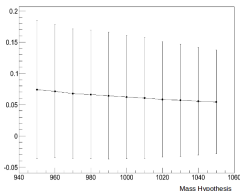
Comparison : Background-only experiments

Signal fraction and significance for 20 background-only experiments

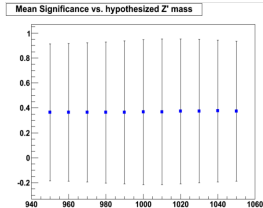
Classical Shape Analysis



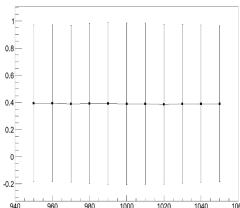
Matrix Element Method



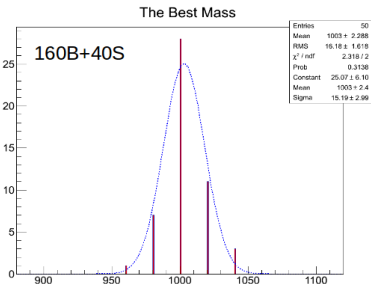
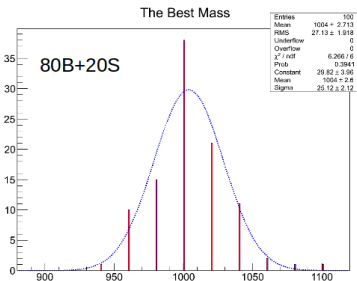
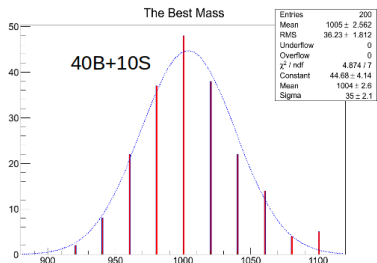
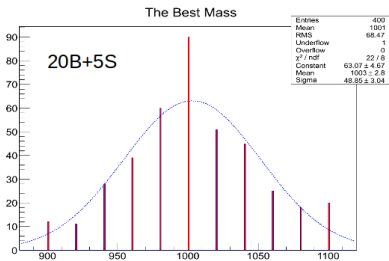
Classical Shape Analysis



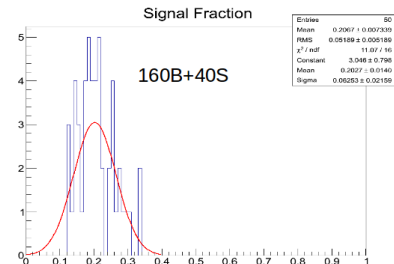
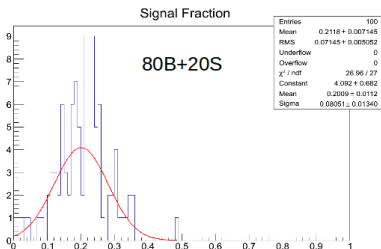
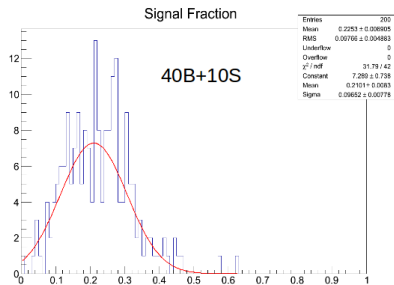
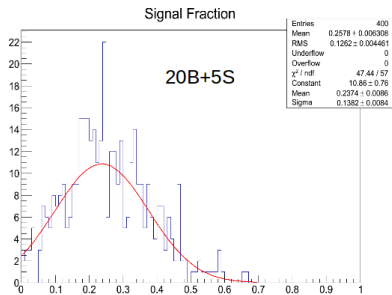
Matrix Element Method



MEM - simultaneous fit - mass

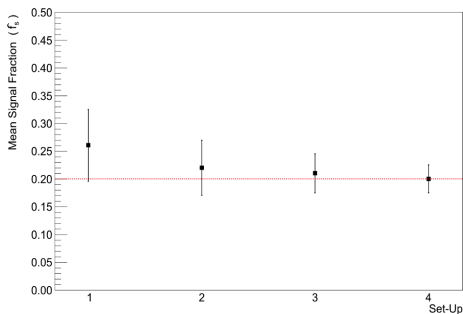


MEM - simultaneous fit - signal fraction

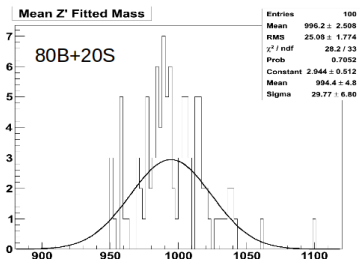
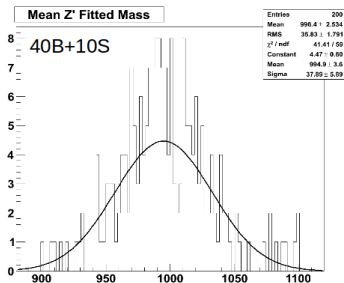
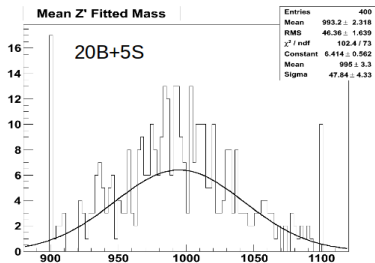


MEM - simultaneous fit - signal fraction

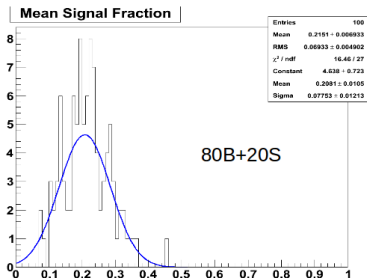
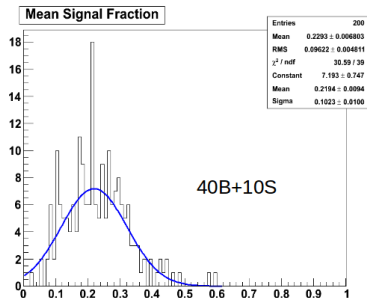
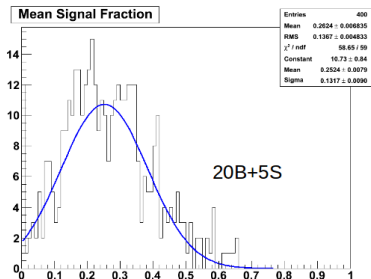
Estimated signal fraction with different pseudo experiments tests with different statistics (S+B ensemble) for a fixed signal fraction



Classical analysis - simultaneous fit - mass



Classical analysis - simultaneous fit - signal fraction



The same bias observed !

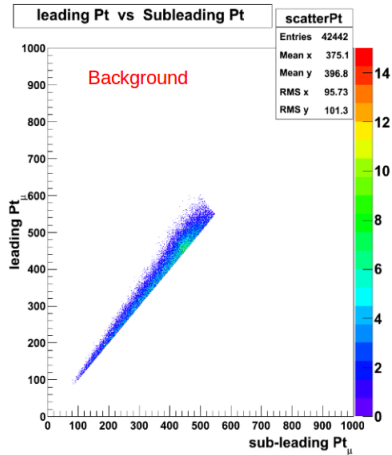
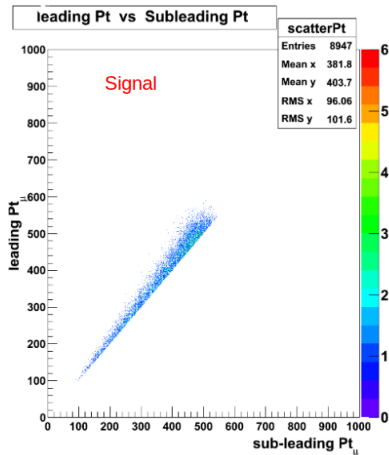
Conclusion

- A generic likelihood-based analysis approach with Matrix element method has been developed.
- The method can be used for parameter estimation (mass, cross-section) or hypotheses testing (significance, exclusion limits, spin-parity etc.).
- It is adapted and validated on MC for a search analysis $Z' \rightarrow \mu^- \mu^+$
 - Mass estimator is ok! estimator is biased for signal fraction in low statistics cases.
 - The same bias was observed with shape analysis.
 - The bias disappears when we increase the size of S+B numbers in the samples.
 - Sensitivity gain $\approx 10 - 20\%$ comparison to shape analysis
- Application to another process like $H \rightarrow ZZ \rightarrow 4l$ can follow a very similar way.

Back-Up

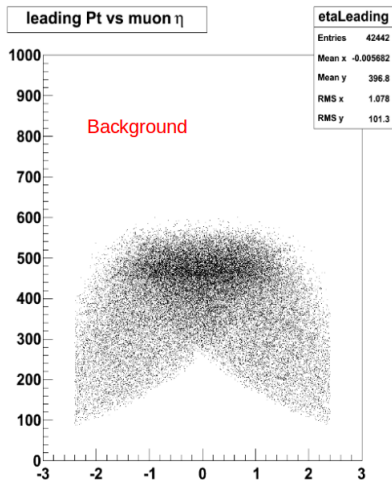
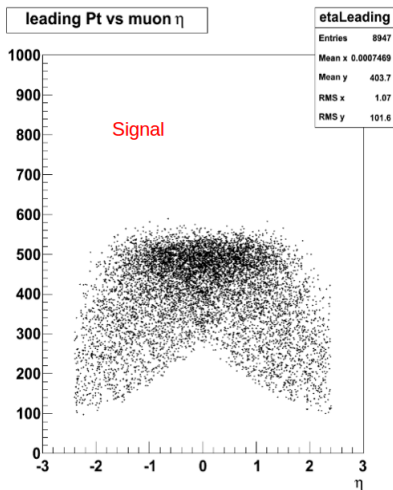
Kinematical Distributions

Leading P_t^μ vs. subleading P_t^μ for selected events.



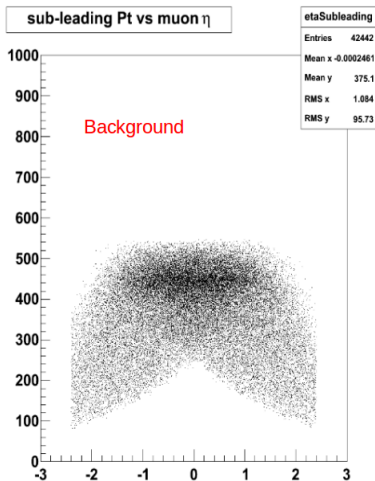
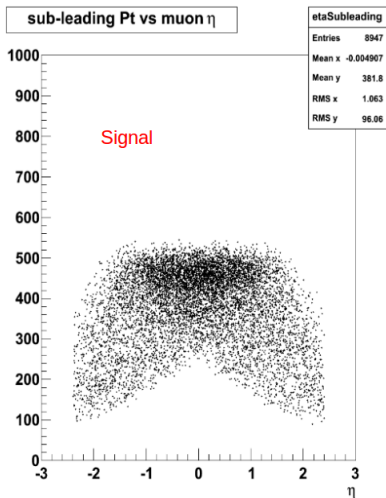
Kinematical Distributions

Leading P_t^μ vs. η^μ for selected events.



Kinematical Distributions

Sub-leading P_t^μ vs. η^μ for selected events.



MEM : Fitted Fraction vs. Fitted Mass

