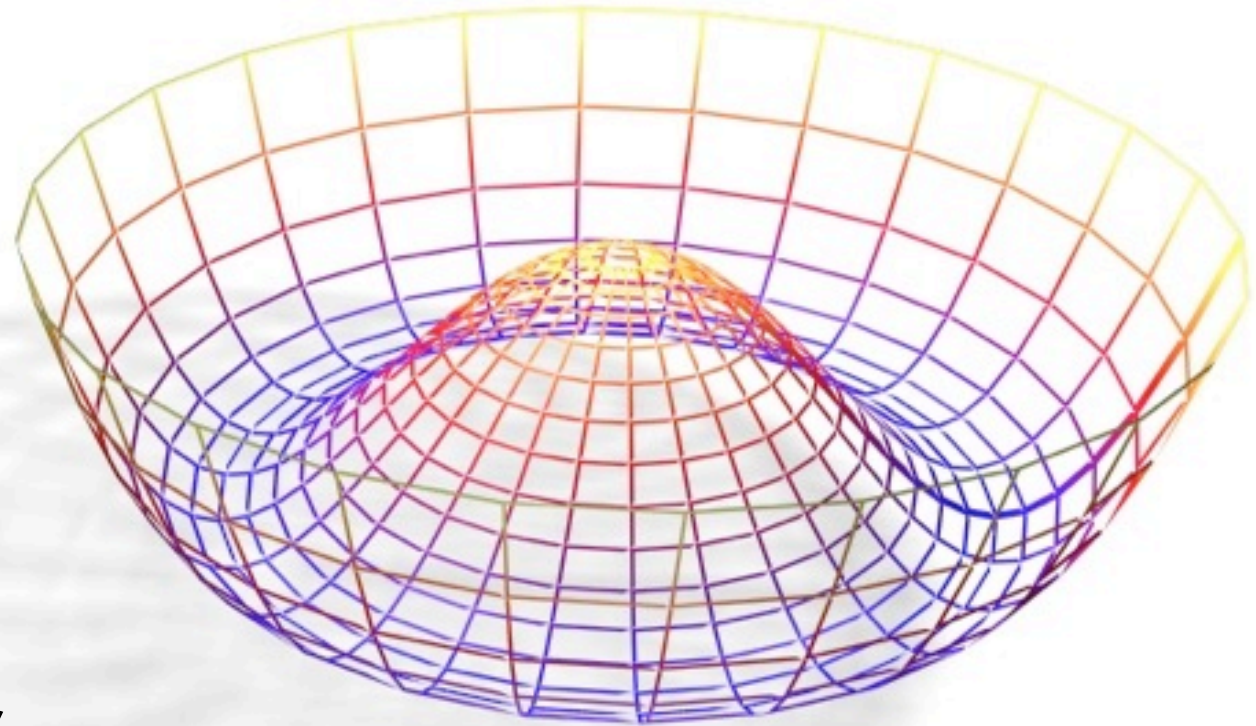
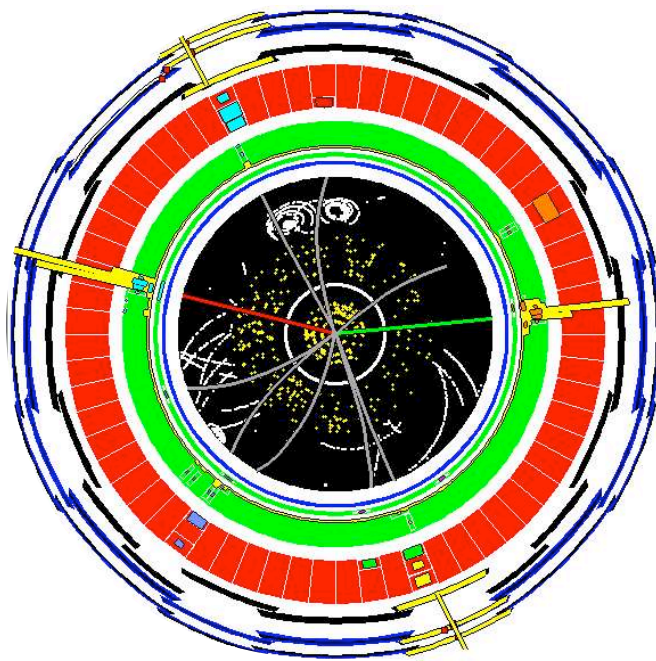
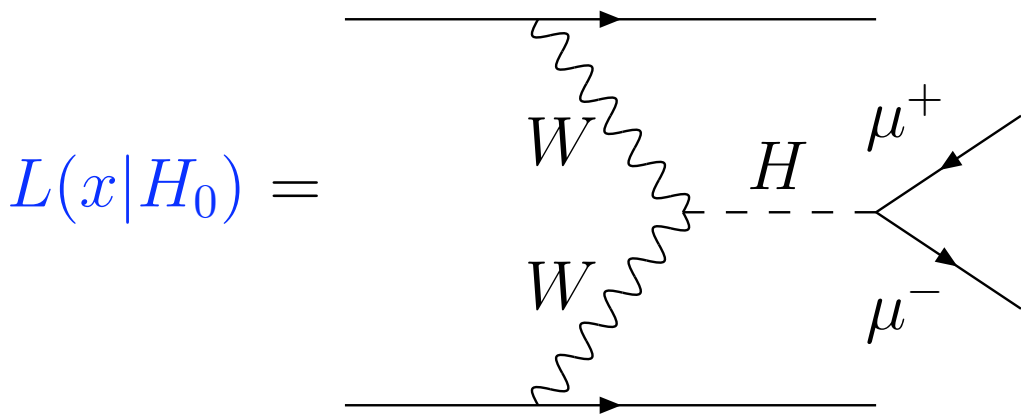




Thoughts on the Matrix Element Method



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In [hep-ph/0605268] Tilman and I used the Neyman-Pearson lemma to establish a formal maximum expected significance using MEM.

- region of the data that maximizes power of a **simple hypothesis test** is given by the likelihood ratio

$$\frac{P(x|H_1)}{P(x|H_0)} > k_\alpha$$

Expected significance: you don't need to match specific observations $\{x_i\}$.

- the MC integration is always “forward” [generate ϕ , smear via $W(x|\phi)$]

What we really care about computing is the distribution of this ratio, not the numerator or the denominator

- **theme**: instead of computing a cross-section, we compute a formal statistical quantity at some order in perturbation theory

Today: some generalizations of this idea

Channel: a subset of the data defined by some selection requirements.

- ▶ eg. all events with 4 electrons with energy > 10 GeV
- ▶ n : number of events observed in the channel
- ▶ ν : number of events expected in the channel

Discriminating variable: a property of those events that can be measured and which helps discriminate the signal from background

- ▶ for MEM, this is observed kinematics and particle ID information
- ▶ $f(x)$: the p.d.f. of the discriminating variable x , ie. $\int d\phi |M|^2 W(x|\phi)$

$$\mathcal{D} = \{x_1, \dots, x_n\}$$

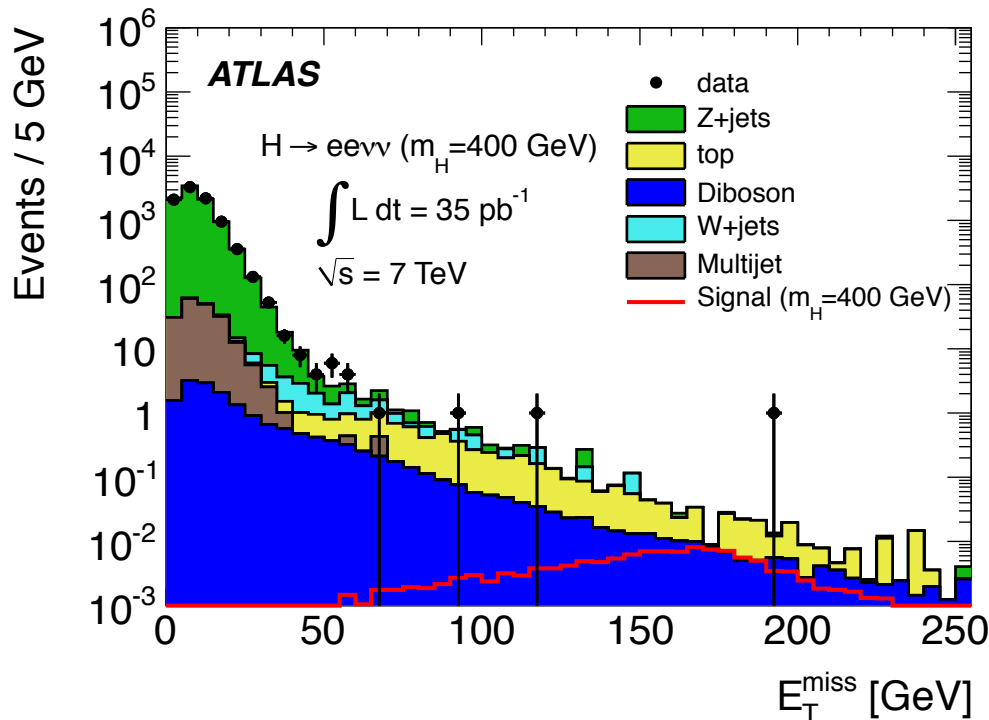
Marked Poisson Process:

$$\mathbf{f}(\mathcal{D}|\nu) = \text{Pois}(n|\nu) \prod_{e=1}^n f(x_e)$$

Sample: a sample of simulated events corresponding to particular type interaction that populates the channel.

▶ statisticians call this a mixture model

$$f(x) = \frac{1}{\nu_{\text{tot}}} \sum_{s \in \text{samples}} \nu_s f_s(x), \quad \nu_{\text{tot}} = \sum_{s \in \text{samples}} \nu_s$$



Note, $f(x)$ is a normalized pdf, so all rate information due to acceptance & tagging encoded in ν

$$\nu = L\sigma = L \int d\phi |\mathcal{M}(\phi)|^2 W(x|\phi)$$

$$f(x) = \frac{1}{\sigma} \int d\phi |\mathcal{M}(\phi)|^2 W(x|\phi)$$

What to do for reducible backgrounds, where M, W uncertain?

Parameters of interest (μ): parameters of the theory that modify the rates and shapes of the distributions, eg.

- ▶ the mass of a hypothesized particle
- ▶ the “signal strength” $\mu=0$ no signal, $\mu=1$ predicted signal rate

Nuisance parameters (θ or α_p): associated to uncertainty in:

- ▶ response of the detector (calibration)
 - typically ignored in MEM, need $W(x | \phi) \rightarrow W(x | \phi, \theta)$
- ▶ theoretical uncertainties

Lead to a parametrized model: $\nu \rightarrow \nu(\alpha), f(x) \rightarrow f(x|\alpha)$

$$\mathbf{f}(\mathcal{D}|\alpha) = \text{Pois}(n|\nu(\alpha)) \prod_{e=1}^n f(x_e|\alpha)$$

Control Regions: Some channels are not populated by signal processes, but are used to constrain the nuisance parameters

Constraint Terms: Often auxiliary measurements for certain nuisance parameters summarized / idealized as

$$f_p(a_p | \alpha_p) \quad \text{for } p \in \mathcal{S}$$

Simultaneous Multi-Channel Model: Several disjoint regions of the data are modeled simultaneously. Identification of common parameters across many channels requires coordination between groups such that meaning of the parameters are really the same.

$$\mathbf{f}_{\text{tot}}(\mathcal{D}_{\text{sim}}, \mathcal{G} | \boldsymbol{\alpha}) = \prod_{c \in \text{channels}} \left[\text{Pois}(n_c | \nu_c(\boldsymbol{\alpha})) \prod_{e=1}^{n_c} f_c(x_{ce} | \boldsymbol{\alpha}) \right] \cdot \prod_{p \in \mathcal{S}} f_p(a_p | \alpha_p)$$

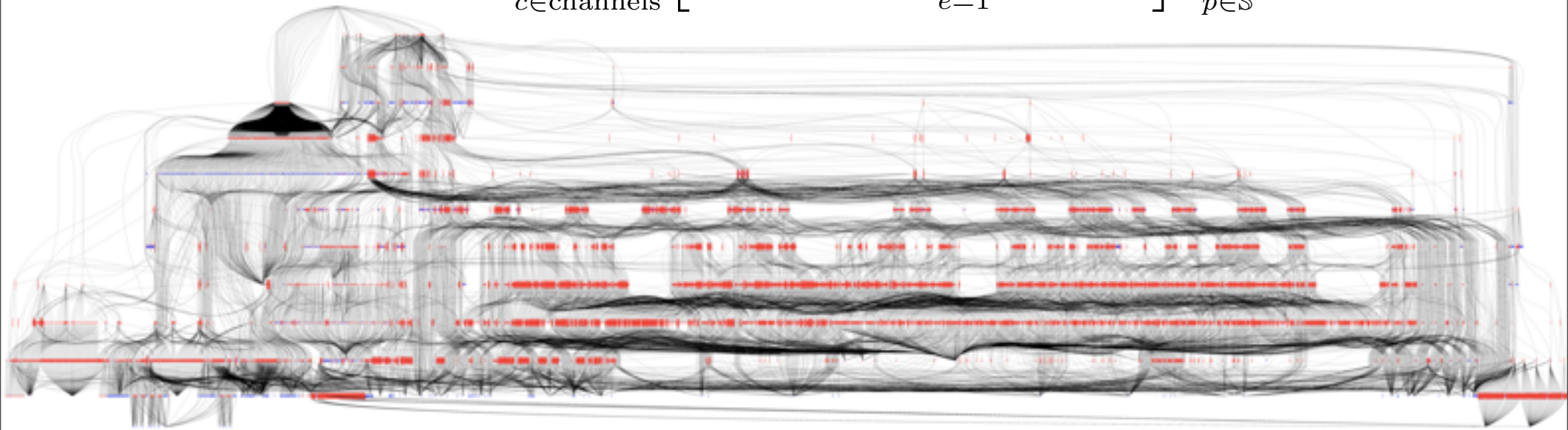
where

$$\mathcal{D}_{\text{sim}} = \{\mathcal{D}_1, \dots, \mathcal{D}_{c_{\text{max}}}\}, \quad \mathcal{G} = \{a_p\} \text{ for } p \in \mathcal{S}$$

Roofit / RooStats: is the modeling language (C++) which provides technologies for collaborative modeling

- ▶ provides technology to publish likelihood functions digitally
- ▶ and more, it's the full model so we can also generate pseudo-data

$$\mathbf{f}_{\text{tot}}(\mathcal{D}_{\text{sim}}, \mathcal{G} | \boldsymbol{\alpha}) = \prod_{c \in \text{channels}} \left[\text{Pois}(n_c | \nu_c(\boldsymbol{\alpha})) \prod_{e=1}^{n_c} f_c(x_{ce} | \boldsymbol{\alpha}) \right] \cdot \prod_{p \in \mathbb{S}} f_p(a_p | \alpha_p)$$



To incorporate MEM approaches directly into common statistical machinery (used for Higgs, SUSY) need interface to Roofit/RooStats

- ▶ specifically, need a class that inherits from RooAbsPdf

Matrix Element Method

The Matrix Element Method consist in minimizing a likelihood.

The likelihood for N events is defined as $L(\alpha) = e^{-N \int \bar{P}(x, \alpha) dx} \prod_{i=1}^N \bar{P}(x_i; \alpha)$

The best estimate of the parameter α is obtained through a maximisation of the likelihood. It is common practice to minimize $-\ln(L(\alpha))$ with respect to α ,

$$-\ln(L) = -\sum_{i=1}^N \ln(\bar{P}(x_i; \alpha)) + N \int \bar{P}(x, \alpha) dx$$

In general, the probability that an event is accepted depends on the characteristics of the measured event, and not on the process that produced it. The measured probability density $\bar{P}(x, \alpha)$ can be related to the produced probability density $P(x, \alpha)$:

I don't actually understand this... $\int P(x, \alpha) dx = 1$ if P is a pdf, and Poisson is not e^{-N} , it is
$$\text{Pois}(n|\nu) = \frac{\nu^n e^{-\nu}}{n!}$$

I would write:
$$\mathbf{f}(\mathcal{D}|\alpha) = \text{Pois}(n|\nu(\alpha)) \prod_{e=1}^n f(x_e|\alpha)$$

$$-\ln L(\alpha) = \underbrace{\nu(\alpha) - n \ln \nu(\alpha)}_{\text{extended term}} - \sum_{e=1}^n \ln f(x_e) + \underbrace{\ln n!}_{\text{constant}}$$

The simple hypothesis test case

Special case of the general probability model (no nuisance parameters)

$$Q = \frac{L(x|H_1)}{L(x|H_0)} = \frac{\prod_i^{N_{chan}} Pois(n_i | s_i + b_i) \prod_j^{n_i} \frac{s_i f_s(x_{ij}) + b_i f_b(x_{ij})}{s_i + b_i}}{\prod_i^{N_{chan}} Pois(n_i | b_i) \prod_j^{n_i} f_b(x_{ij})}$$

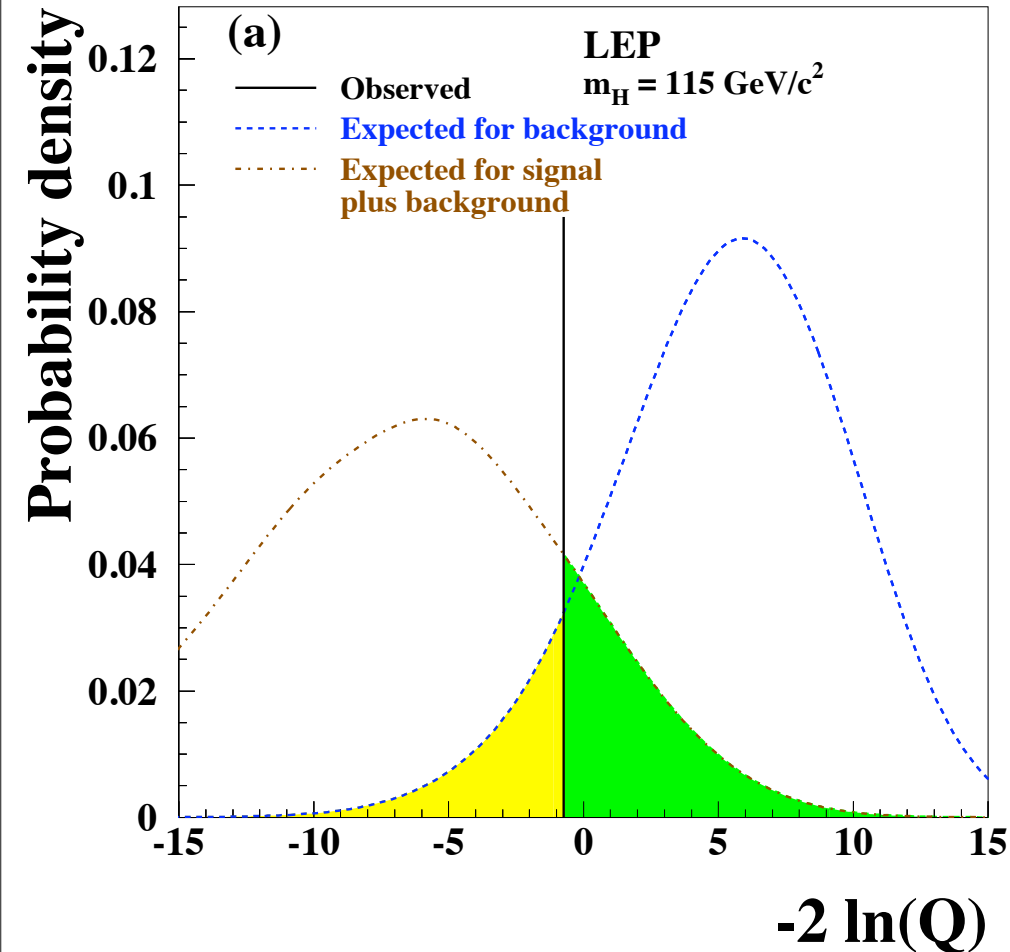
$$q = \ln Q = -s_{tot} + \sum_i^{N_{chan}} \sum_j^{n_i} \ln \left(1 + \frac{s_i f_s(x_{ij})}{b_i f_b(x_{ij})} \right)$$

Instead of simply counting events, the optimal test statistic is equivalent to adding events weighted by

$\ln(1 + \text{signal}/\text{background})$

The test statistic is a map $q:\text{data} \rightarrow \mathbb{R}$

By repeating the experiment many times, you obtain a distribution for q



There is a clever trick for bootstrapping from distribution of q for a single event to the distribution for an experiment with N events

$$Q = \frac{L(x|H_1)}{L(x|H_0)} = \frac{\prod_i^{N_{chan}} \text{Pois}(n_i | s_i + b_i) \prod_j^{n_i} \frac{s_i f_s(x_{ij}) + b_i f_b(x_{ij})}{s_i + b_i}}{\prod_i^{N_{chan}} \text{Pois}(n_i | b_i) \prod_j^{n_i} f_b(x_{ij})}$$
$$q = \ln Q = -s_{tot} + \sum_i^{N_{chan}} \sum_j^{n_i} \ln \left(1 + \frac{s_i f_s(x_{ij})}{b_i f_b(x_{ij})} \right)$$

For N events, use Fourier transform to perform N convolutions

$$\rho_{N,i}(q) = \underbrace{\rho_{N,i}(q) \oplus \cdots \oplus \rho_{N,i}(q)}_{N \text{ times}} = \mathcal{F}^{-1} \left\{ [\mathcal{F}(\rho_{1,i})]^N \right\}$$

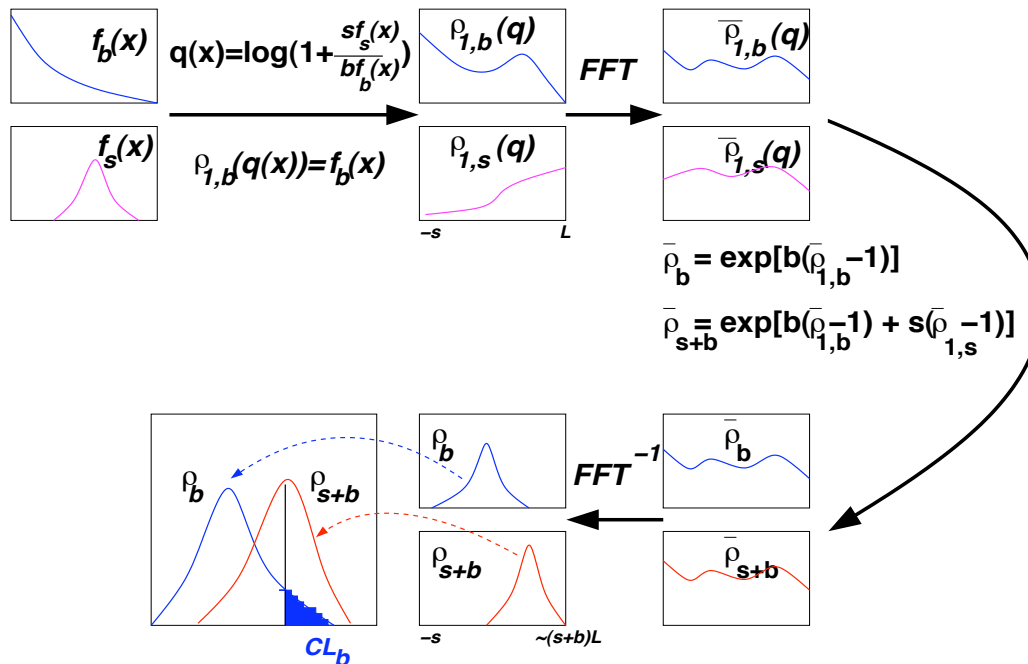
To include Poisson fluctuations on N for a given luminosity, one can exponentiate

$$\rho_i(q) = \sum_{N=0}^{\infty} P(N; L\sigma_i) \cdot \rho_{N,i}(q) = \mathcal{F}^{-1} \left\{ e^{L\sigma_i [\mathcal{F}(\rho_{1,i}(q)) - 1]} \right\}$$

There is a clever trick for bootstrapping from distribution of q for a single event to the distribution for an experiment with N events

$$Q = \frac{L(x|H_1)}{L(x|H_0)} = \frac{\prod_i^{N_{chan}} Pois(n_i | s_i + b_i) \prod_j^{n_i} \frac{s_i f_s(x_{ij}) + b_i f_b(x_{ij})}{s_i + b_i}}{\prod_i^{N_{chan}} Pois(n_i | b_i) \prod_j^{n_i} f_b(x_{ij})}$$

$$q = \ln Q = -s_{tot} + \sum_i^{N_{chan}} \sum_j^{n_i} \ln \left(1 + \frac{s_i f_s(x_{ij})}{b_i f_b(x_{ij})} \right)$$



Hu and Nielsen's CLFFT used Fourier Transform and exponentiation trick to transform the log-likelihood ratio distribution for one event to the distribution for an experiment

When we go beyond simple hypothesis tests to **parametrized families** of distributions, there is no **uniformly most powerful** test in general

- ▶ The most common generalization of the likelihood ratio test statistic is to keep null in numerator and best fit in denominator [Feldman-Cousins]
- ▶ In the presence of nuisance parameters, it is the profile likelihood ratio

$$\lambda(\mu) = \frac{L(\mu, \hat{\theta}(\mu))}{L(\hat{\mu}, \hat{\theta})} = \frac{f(\mathcal{D} | \mu, \hat{\theta}(\mu; \mathcal{D}))}{f(\mathcal{D} | \hat{\mu}, \hat{\theta})}$$

- ▶ The Fourier exponentiation trick doesn't work anymore, but the asymptotically the distributions are known

G. Cowan, K. C., E. Gross, O. Vitells.
Eur. Phys. J., C71 2011. [arXiv:1007.1727](https://arxiv.org/abs/1007.1727)

Specifically, I'd like to incorporate experimental uncertainty into the transfer functions: $W(x | \phi) \rightarrow W(x | \phi, \theta)$

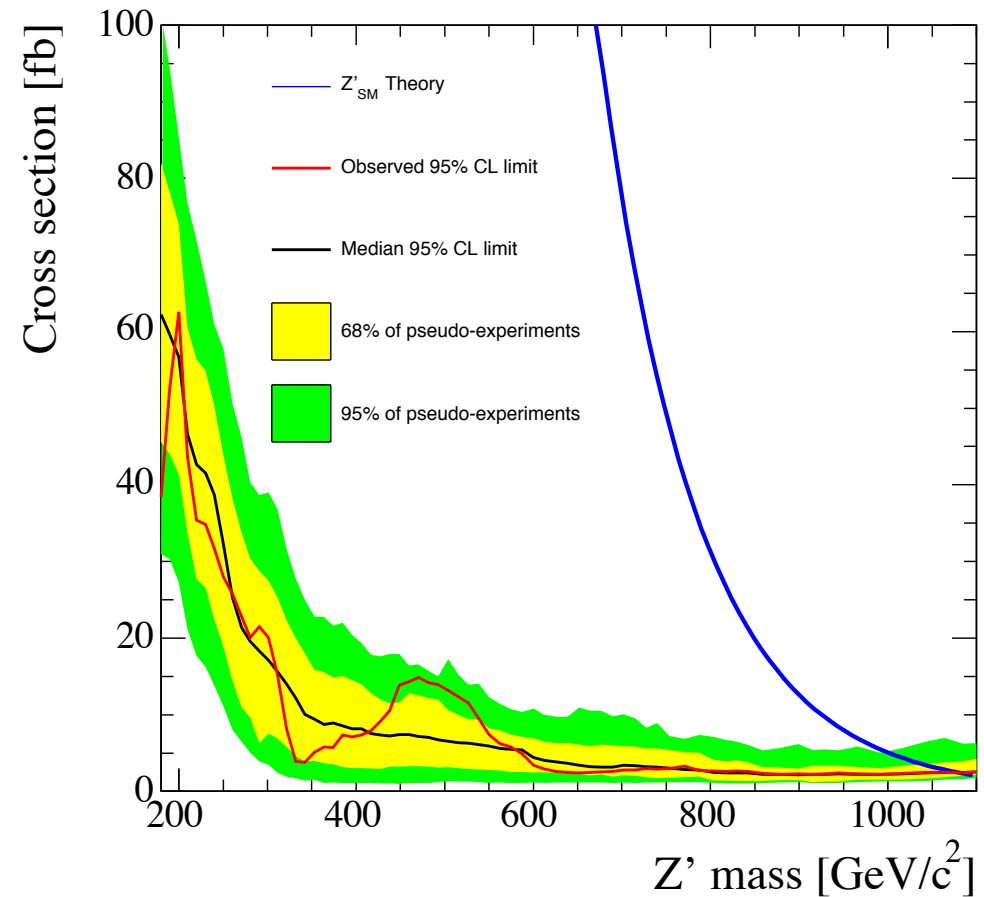
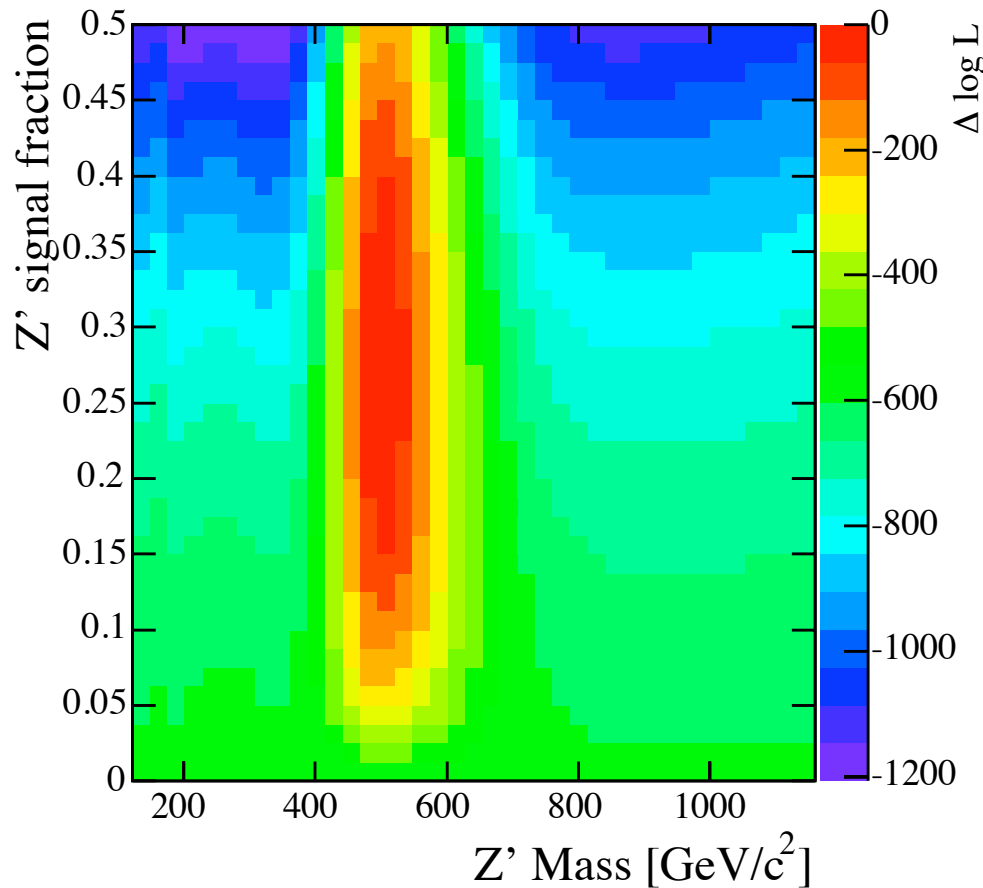
We directly integrated MEM Likelihood into limit-setting procedure

- ▶ Included interference of Z' and Z/γ
- ▶ 2-d Feldman-Cousins instead of “raster scan”

CDF Collaboration $Z' \rightarrow \mu^+ \mu^-$
Phys.Rev.Lett. 106 (2011)
arXiv:1101.4578

We present a search for a new narrow, spin-1, high mass resonance decaying to $\mu^+ \mu^- + X$, using a matrix element based likelihood and a simultaneous measurement of the resonance mass and production rate. In data with 4.6 fb^{-1} of integrated luminosity collected by the CDF detector in

CDF Run II Preliminary



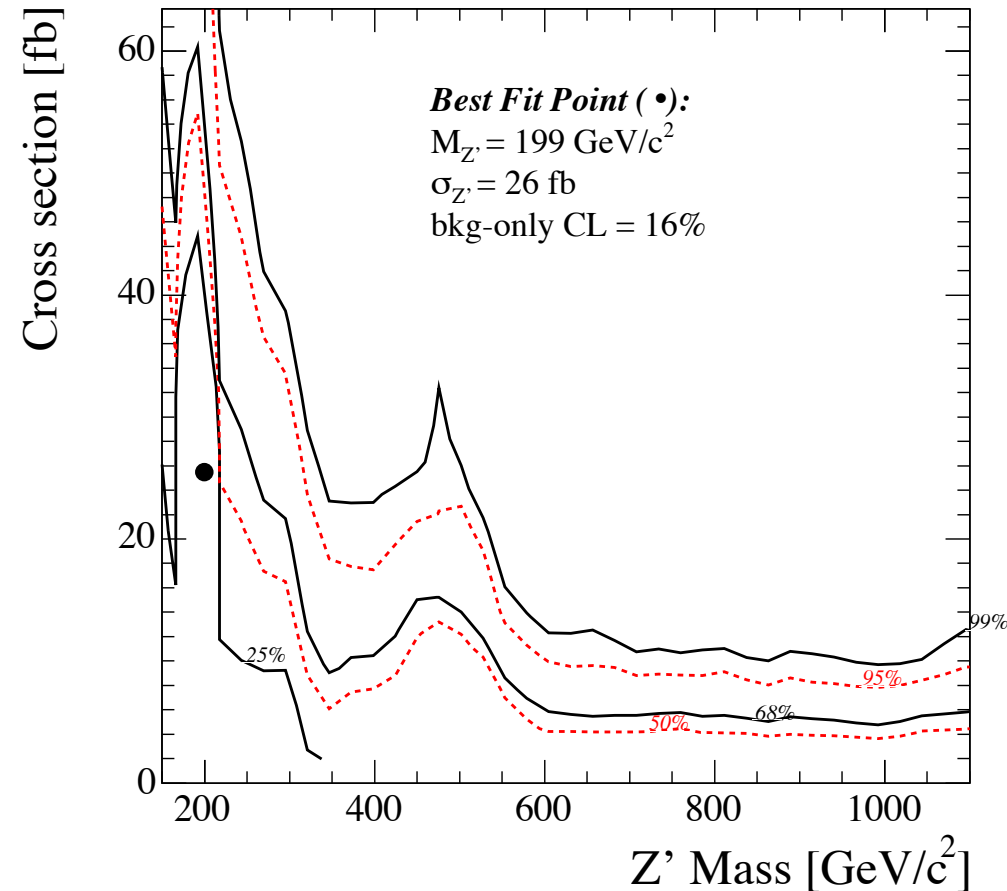
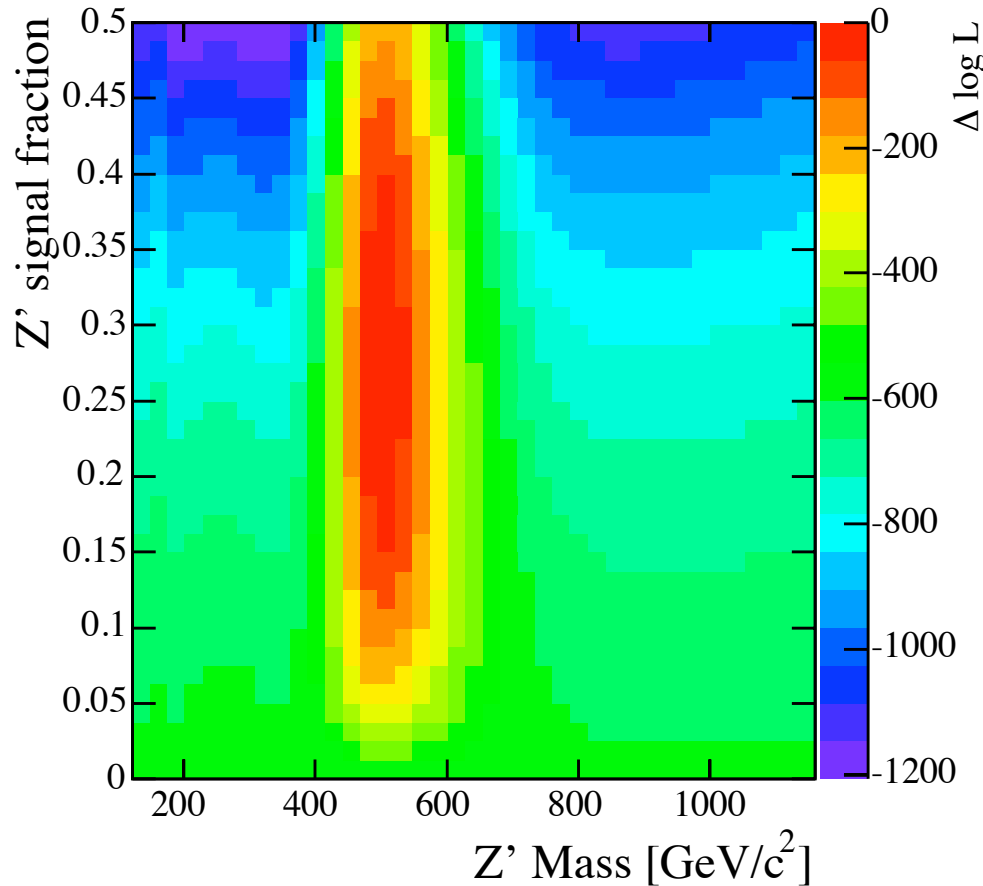
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Similar to the Neyman-Pearson lemma for simple hypothesis tests is the Cramér-Rao bound for the covariance of an (unbiased) estimator

$$\text{cov}[\hat{\boldsymbol{\alpha}}|\boldsymbol{\alpha}] \geq I_{\mu\nu}^{-1}(\boldsymbol{\alpha})$$

where $I_{\mu\nu}$ is the Fisher Information matrix

$$I_{\mu\nu}(\boldsymbol{\alpha}) = \int p(\mathbf{x}|\boldsymbol{\alpha}) \frac{\partial \ln p(\mathbf{x}|\boldsymbol{\alpha})}{\partial \alpha_{\mu}} \frac{\partial \ln p(\mathbf{x}|\boldsymbol{\alpha})}{\partial \alpha_{\nu}} d\mathbf{x} = E [\partial_{\mu} \ln L(\boldsymbol{\alpha}) \partial_{\nu} \ln L(\boldsymbol{\alpha}) | \boldsymbol{\alpha}]$$

In the case of our Marked Poisson model, this is given by

$$I_{\mu\nu}(\boldsymbol{\alpha}) \rightarrow \int dx \frac{\partial \nu(\boldsymbol{\alpha}) f(x|\boldsymbol{\alpha})}{\partial \alpha_{\mu}} \frac{\partial \nu(\boldsymbol{\alpha}) f(x|\boldsymbol{\alpha})}{\partial \alpha_{\nu}} \frac{1}{\nu(\boldsymbol{\alpha}) f(x|\boldsymbol{\alpha})}$$

B. Allanach, K.C.
[in prep.]

The integral through the transfer function is easy in the “forward” direction

- Evaluating derivative would be aided by importance sampling

Bayesian / Frequentist often comes down to integrate vs. maximize

- ▶ true momenta ϕ plays role of “nuisance parameters”
- ▶ Lorentz-invariant phase space $d\phi$ plays role of prior [w/ frequency interpretation]

Perhaps the “Profiled” MEM is even more powerful?

- ▶ note, similarity to constrained fit, but also use $|M(\phi)|^2$

		Likelihood	
		$ M(\phi) ^2 W(x/\phi)$	$W(x/\phi)$
$\int d\phi$	Typical Matrix Element Method		N/A
\sup_{ϕ}	“Profiled” MEM		Constrained fit (two-stage: $x \rightarrow \phi \rightarrow \alpha$)

Consider a simple case where some interaction characterized by M produces particles of energy e_i

- ▶ the matrix element is represented by Gaussian: $G(e|M, \sigma_m)$
- ▶ the transfer function is a simple Gaussian: $G(x|e, \sigma_e)$

$$P(\{x_i\}|M, \{e_i\}) = \prod_i G(e_i|M, \sigma_m) G(x_i|e_i, \sigma_e)$$

One can find the maximum likelihood estimators

$$\hat{e}_i = x_i \quad \hat{M} = \frac{1}{n} \sum_i \hat{e}_i = \bar{x}$$

and the estimators are consistent [as $n \rightarrow \infty$, expectation = true value]

$$E[\hat{M}] = M$$

... so far so good.

Consider a simple case where some interaction characterized by M produces particles of energy e_i

- ▶ the matrix element is represented by **falling exponential**
- ▶ the transfer function is a simple Gaussian: $G(x|e, \sigma_e)$

$$P(\{x_i\} | M, \{e_i\}) = \prod_i \frac{1}{M} e^{-e_i/M} G(x_i | e_i, \sigma_e)$$

One can find the maximum likelihood estimators

$$\hat{M} = \frac{\bar{x} + \sqrt{\bar{x}^2 - 4\sigma_e^2}}{2}$$

but the estimator is ***inconsistent!***

$$E[\hat{M}] \neq M$$

This is a general problem if you add more parameters as you add more data, the estimator can be biased even in limit of infinite data!

Jet-levels: Parton → Hadron → Reconstructed

- ▶ it may be beneficial to factorize these stages for transfer function
 - $W(\text{Reco}|\text{Parton}) \rightarrow W(\text{Reco}|\text{Hadron}) W(\text{Hadron} | \text{Parton})$

To deal with extra jet radiation, will need to deal with ME-PS matching

- ▶ “Poor-man’s MEM”:
 - store large sample at hadron level, only apply $W(\text{reco}|\text{hadron})$
 - implementation is trivial, but phase space integration is inefficient
- ▶ MLM Matching
 - basically requires $N_{\text{jet}} @ \text{hadron-level} = N_{\text{parton}}$ defined at some scale
 - alignment of reco jet algorithm with matching procedure would mean $N_{\text{jet}}=N_{\text{parton}}$
 - if $W(\text{reco}|\text{parton})$ encodes jet reconstruction inefficiencies, then $\sum d\phi |M_n|^2$ for $n \geq n_{\text{jet}}$

A phase space integration idea

The biggest practical issue with the matrix element method is that it is very computationally intensive.

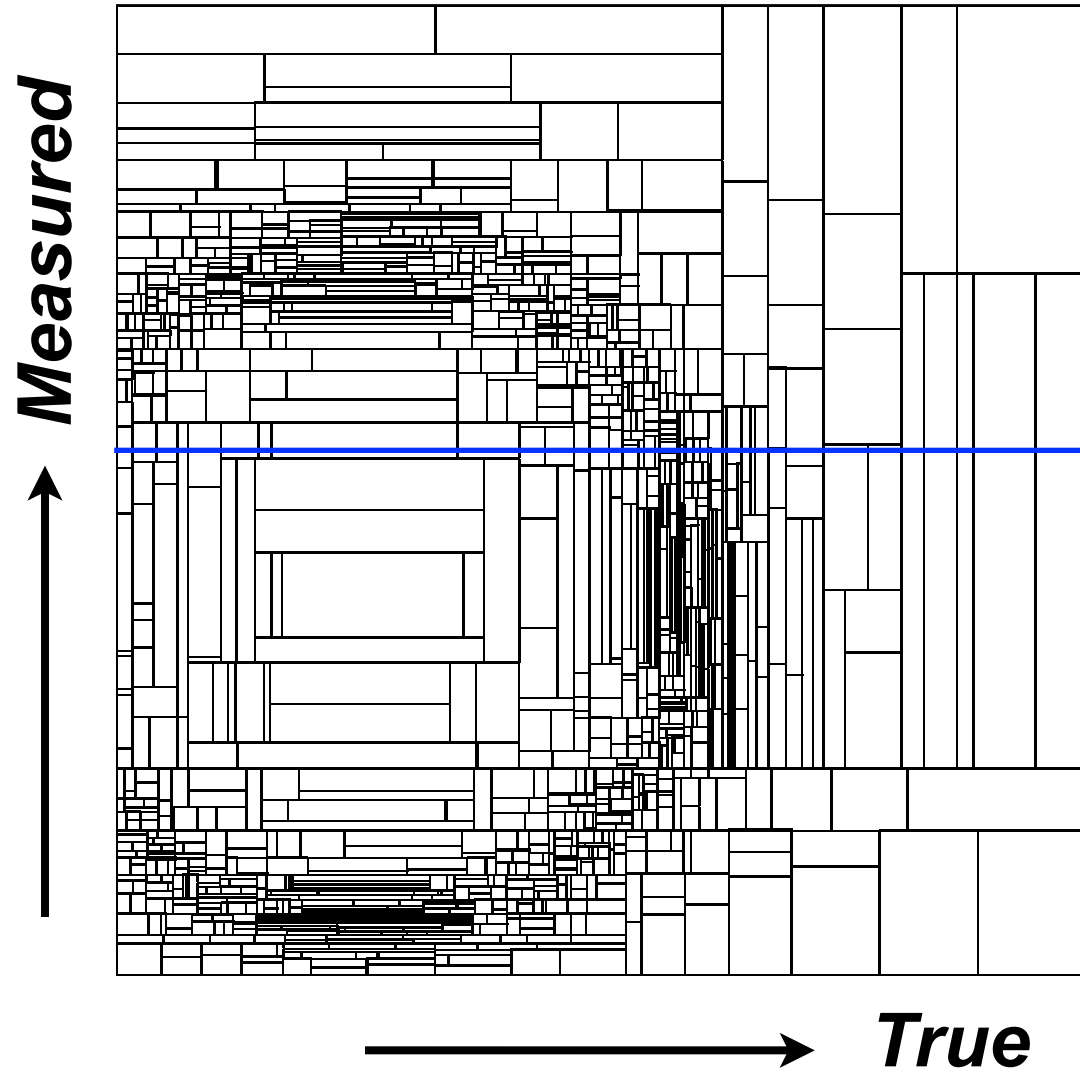
Normally, integration over degrees of freedom in matrix element requires a new Vegas grid for each measurement!

Instead, integrate the joint distribution

- ▶ save joint grid

Then for each measurement

- ▶ take a slice through the grid
- ▶ induced importance sampling



In Bayes's theorem, $P(A|B) = \frac{P(B|A)P(A)}{P(B)}$, the normalization $P(B)$ often called "evidence". Similar to MEM with $B \rightarrow x, A \rightarrow \phi$

$$P(x) = \frac{1}{\sigma} \int d\phi |\mathcal{M}(\phi)|^2 W(x|\phi)$$

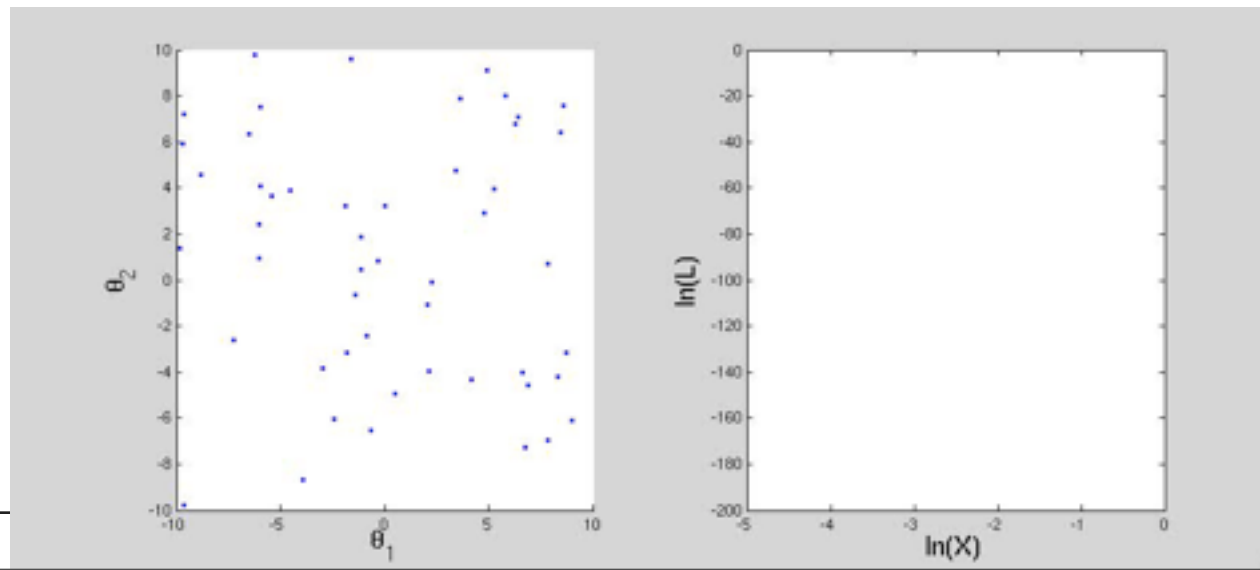
Nested sampling:

An algorithm originally aimed primarily at the Bayesian evidence computation (Skilling, 2006):

$$X(\lambda) = \int_{\mathcal{L}(\theta) > \lambda} P(\theta) d\theta$$

$$P(d) = \int d\theta \mathcal{L}(\theta) P(\theta) = \int_0^1 X(\lambda) d\lambda$$

Feroz et al (2008), [arxiv: 0807.4512](https://arxiv.org/abs/0807.4512), Trotta et al (2008), [arxiv: 0809.3792](https://arxiv.org/abs/0809.3792)



(animation
courtesy of
David Parkinson)

MEM natural procedure that provides most powerful test in case of simple hypothesis tests

- In that case, what we want to integrate is a ratio
- noted difficulty when there are irreducible backgrounds

For parametrized model, Cramér-Rao bound is similar to Neyman-Pearson

- Showed explicit form of what we need to calculate in that case
- Showed CDF Z' example for MEM embedded in Feldman-Cousins including interference effects

To include experimental uncertainties, parametrize transfer functions!

- MEM codes should provide interfaces to RooFit/RooStats

Considered “Profiled” MEM as alternative to traditional MEM

- leads to inconsistent estimators and Neyman-Scott phenomena

Some thoughts on “MEPSM” for matching partons

Two thoughts on PS integration: “induced grid” & nested sampling

The Gaussian case:

```
f2[x1_, x2_, M_, e1_, e2_] :=
  Exp[-(e1 - M)^2 / (2 sm^2)] / (Sqrt[2 Pi] sm) * Exp[-(e2 - M)^2 / (2 sm^2)] / (Sqrt[2 Pi] sm) *
  Exp[-(x1 - e1)^2 / (2 se^2)] / (Sqrt[2 Pi] se) * Exp[-(x2 - e2)^2 / (2 se^2)] / (Sqrt[2 Pi] se)
```

```
Solve[D[Log[f2[x, y, M, e1, e2]], e1] == 0 && D[Log[f2[x, y, M, e1, e2]], e2] == 0 &&
  D[Log[f2[x, y, M, e1, e2]], M] == 0, {M, e1, e2}]
```

$$\left\{ \left\{ M \rightarrow \frac{x+y}{2}, e1 \rightarrow -\frac{-se^2 x - 2 sm^2 x - se^2 y}{2 (se^2 + sm^2)}, e2 \rightarrow -\frac{-se^2 x - se^2 y - 2 sm^2 y}{2 (se^2 + sm^2)} \right\} \right\}$$

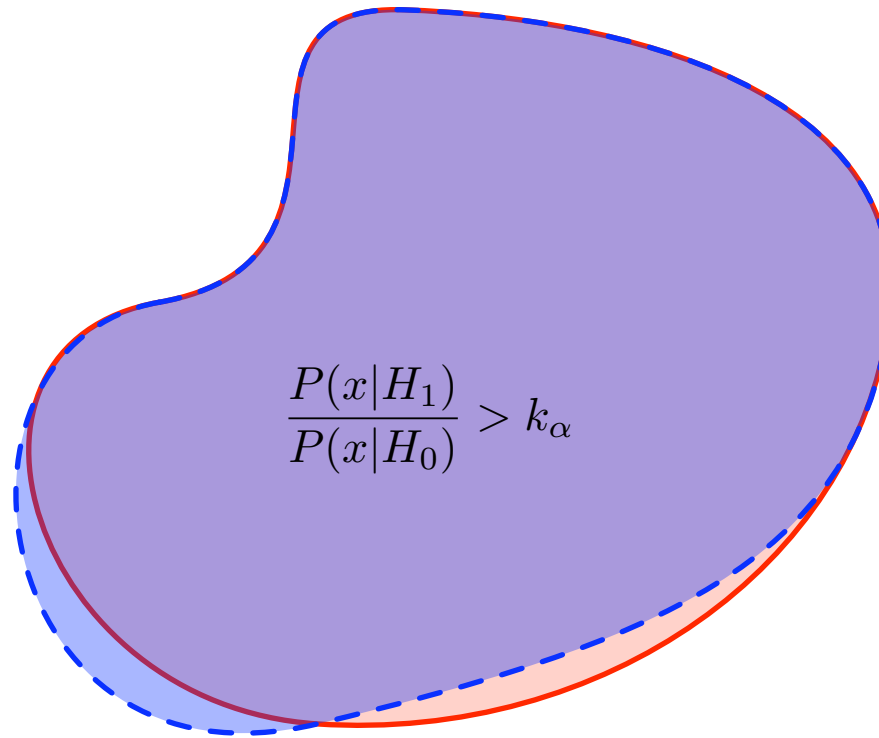
The exponential case:

```
In[23]:= g[x1_, x2_, M_, e1_, e2_] :=
  Exp[-e1/M] / M * Exp[-e2/M] / M * Exp[-(x1 - e1)^2 / (2 se^2)] / (Sqrt[2 Pi] se) *
  Exp[-(x2 - e2)^2 / (2 se^2)] / (Sqrt[2 Pi] se)
```

```
In[34]:= Solve[D[Log[g[x, y, M, e1, e2]], e1] == 0 && D[Log[g[x, y, M, e1, e2]], e2] == 0 &&
  D[Log[g[x, y, M, e1, e2]], M] == 0, {M, e1, e2}]
```

$$\text{Out[34]= } \left\{ \left\{ M \rightarrow \frac{1}{4} \left(x + y - \sqrt{-16 se^2 + x^2 + 2 xy + y^2} \right), e1 \rightarrow \frac{1}{2} \left(\frac{3x}{2} - \frac{y}{2} - \frac{1}{2} \sqrt{-16 se^2 + x^2 + 2 xy + y^2} \right), \right. \right. \\ \left. e2 \rightarrow \frac{1}{2} \left(-\frac{x}{2} + \frac{3y}{2} - \frac{1}{2} \sqrt{-16 se^2 + x^2 + 2 xy + y^2} \right) \right\}, \left\{ M \rightarrow \frac{1}{4} \left(x + y + \sqrt{-16 se^2 + x^2 + 2 xy + y^2} \right), \right. \\ \left. e1 \rightarrow \frac{1}{2} \left(\frac{3x}{2} - \frac{y}{2} + \frac{1}{2} \sqrt{-16 se^2 + x^2 + 2 xy + y^2} \right), e2 \rightarrow \frac{1}{2} \left(-\frac{x}{2} + \frac{3y}{2} + \frac{1}{2} \sqrt{-16 se^2 + x^2 + 2 xy + y^2} \right) \right\} \right\}$$

A short proof of Neyman-Pearson



$$\frac{P(x|H_1)}{P(x|H_0)} > k_\alpha$$

$$P(\text{blue crescent} | H_0) = P(\text{red crescent} | H_0)$$

$$\frac{P(x|H_1)}{P(x|H_0)} < k_\alpha$$

$$\frac{P(x|H_1)}{P(x|H_0)} > k_\alpha$$

$$P(\text{blue crescent} | H_1) < P(\text{blue crescent} | H_0)k_\alpha$$

$$P(\text{red crescent} | H_1) > P(\text{red crescent} | H_0)k_\alpha$$

$$P(\text{blue crescent} | H_1) < P(\text{red crescent} | H_1)$$

The new region region has less power.