Summary of studies on limit setting with CLsGenerator and BayesianMCMC

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Validation of CLsGenerator without systematics and other checks

Validation of CLsGenerator without systematics

- When there's no systematics, an analytical solution for the limit can be found
 - ▶ This solution is used to validate CLsGenerator without systematics
- The analytical solution will also be derived and compared to CLsGenerator in the gaussian case.

Analytical solution (1)

- We have:
 - $N \sim \mathcal{P}(\mu s + b)$ under the s + b hypothesis
 - ▶ $N \sim \mathcal{P}(b)$ under the *b* hypothesis
- Consider first a $\mathit{CL}_{\mathsf{s+b}}$ limit. Upper limit μ_{up} at $1-\alpha$ confidence level given by

$$CL_{s+b} = \sum_{n=0}^{Nobs} P(n; \mu_{up}s + b) = \alpha$$

Poisson c.d.f. given by

$$\sum_{n=0}^{N_{\rm obs}} P(n; v) = 1 - F_{\chi^2}(2v; 2(N_{\rm obs} + 1))$$

Thus

$$\mu_{\sf up} = \frac{0.5 \times F_{\chi^2}^{-1} (1 - \alpha; 2(N_{\sf obs} + 1)) - b}{s}$$

Analytical solution (2)

• CLsGenerator computes CL_s limits, not CL_{s+b} . Previous formula can be generalized to CL_s by replacing α by $\alpha \times CL_b$. Indeed:

$$\begin{split} \frac{\mathit{CL}_{\mathsf{s}+\mathsf{b}}}{\mathit{CL}_{\mathsf{b}}} &= \frac{\displaystyle\sum_{n=0}^{N} \mathit{P}(n; \mu_{\mathsf{up}} s + b)}{\mathit{Nobs}} = \alpha \\ &\sum_{n=0}^{N} \mathit{P}(n; b) \\ \Rightarrow 1 - \mathit{F}_{\chi^2}\left(2(\mu_{\mathsf{up}} s + b); 2(\mathit{N}_{\mathsf{obs}} + 1)\right) = \alpha \times \left[1 - \mathit{F}_{\chi^2}\left(2b; 2(\mathit{N}_{\mathsf{obs}} + 1)\right)\right] \end{split}$$

• CL_b is independent of the signal, so we just need to replace α by $\alpha \times \mathit{CL}_b$, yielding

$$\mu_{\mathsf{up}} = \frac{0.5 \times F_{\chi^2}^{-1} \left(1 - \alpha \left(1 - F_{\chi^2} \left(2 b; 2 \left(N_{\mathsf{obs}} + 1\right)\right)\right); 2 \left(N_{\mathsf{obs}} + 1\right)\right) - b}{s}$$

Analytical solution in gaussian case (1)

- Gaussian approximation:
 - $N \sim \mathcal{N}(\mu s + b, \sqrt{\mu s + b})$ under the s + b hypothesis
 - ▶ $N \sim \mathcal{N}(b, \sqrt{b})$ under the *b* hypothesis
- Upper CL_{s+b} limit μ_{up} at $1-\alpha$ confidence level given by

$$\begin{split} \mathit{CL}_{\mathsf{s}+\mathsf{b}} &= \int_{0}^{\mathit{N}_{\mathsf{obs}}} \mathit{G}\left(\mathit{N}; \mu_{\mathsf{up}} \mathit{s} + \mathit{b}, \sqrt{\mu_{\mathsf{up}} \mathit{s} + \mathit{b}}\right) \mathsf{d}\mathit{N} = \alpha \\ &\Rightarrow \int_{0}^{\frac{\mathit{N}_{\mathsf{obs}} - (\mu_{\mathsf{up}} \mathit{s} + \mathit{b})}{\sqrt{\mu_{\mathsf{up}} \mathit{s} + \mathit{b}}}} \mathit{G}(\mathit{N}; 0, 1) \mathsf{d}\mathit{N} = \alpha \\ &\Rightarrow \Phi\left(\frac{\mathit{N}_{\mathsf{obs}} - (\mu_{\mathsf{up}} \mathit{s} + \mathit{b})}{\sqrt{\mu_{\mathsf{up}} \mathit{s} + \mathit{b}}}\right) = \alpha \end{split}$$

where Φ is the c.d.f. of the standard normal distribution

$$\Rightarrow N_{ ext{obs}} - (\mu_{ ext{up}} s + b) = -\sqrt{\mu_{ ext{up}} s + b} imes \Phi^{-1}(1 - lpha)$$

Analytical solution in gaussian case (2)

• Previous expression can be solved for μ_{up} :

$$\boxed{\mu_{\mathsf{up}} = \frac{\mathsf{N}_{\mathsf{obs}} - b}{s} + \frac{\left[\Phi^{-1}(1-\alpha)\right]^2}{2s} \left[1 + \sqrt{1 + 4\frac{\mathsf{N}obs}{\left[\Phi^{-1}(1-\alpha)\right]^2}}\right]} (\star)$$

• Previous formula can be generalized to CL_s by replacing α by $\alpha \times CL_b$ (as in the Poisson case). Indeed:

$$\begin{split} \frac{\mathit{CL}_{\mathsf{s}+\mathsf{b}}}{\mathit{CL}_{\mathsf{b}}} &= \frac{\int_{0}^{N_{\mathsf{obs}}} G\left(N; \mu_{\mathsf{up}} s + b, \sqrt{\mu_{\mathsf{up}} s + b}\right) \mathsf{d}N}{\int_{0}^{N_{\mathsf{obs}}} G\left(N; b, \sqrt{b}\right) \mathsf{d}N} = \alpha \\ &\Rightarrow \Phi\left(\frac{N_{\mathsf{obs}} - (\mu_{\mathsf{up}} s + b)}{\sqrt{\mu_{\mathsf{up}} s + b}}\right) = \alpha \times \Phi\left(\frac{N_{\mathsf{obs}} - b}{\sqrt{b}}\right) \end{split}$$

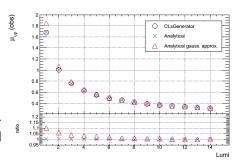
 CL_b is independent of the signal, so we just need to replace α by $\alpha \times \Phi\left(\frac{N_{\rm obs}-b}{\sqrt{b}}\right)$ in (\star) .

CLsGenerator vs analytical solutions

We take

- $N_{obs} = 1$
- b = 0.82
- s = 2.49
- No stat/syst uncertainty

and calculate limits for several luminosities (previous numbers correspond to $\mathscr{L}=1$)

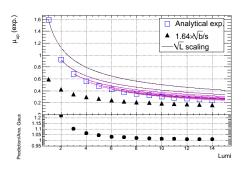


Conclusions:

- CLsGenerator and analytical agree at the 2 per mil level
- CLsGenerator and analytical gaussian approx. agree well when number of events not too small

Comparison to $1.64\frac{\sqrt{b}}{s}$ and scaling with luminosity

- $\mu_{\rm up} = 1.64 \frac{\sqrt{b}}{\rm s}$ is sometimes used to compute limits quickly.
 - From this formula, one often predicts that limit should scale as \sqrt{L}
- As before, we take $N_{obs}=1$, b=0.82, s=2.49 without systematics and compute limit for several luminosities



Conclusions:

- ► $1.64 \frac{\sqrt{s}}{b}$ quite different from analytical result.
- however, \sqrt{L} scaling is a good approximation when number of events not too small

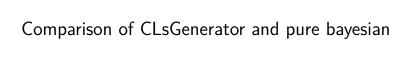
Where does $1.64\frac{\sqrt{b}}{5}$ come from ?

- Let's assume that
 - \mathbf{O} $N_{\text{obs}} = b$ (i.e. we consider an expected limit)
 - **2** $b \gg [\Phi^{-1}(1-\alpha)]^2$
 - ★ note that $\Phi^{-1}(1-\alpha) = 1.64$ for $\alpha = 0.05$, corresponding to $b \gg 2.7$
 - ★ this assumption corresponds more or less to the gaussian approx

Then, one sees directly from (\star) that

$$\mu_{\mathsf{up}} \simeq \Phi^{-1} (1-lpha) rac{\sqrt{b}}{s} = 1.64 rac{\sqrt{b}}{s}$$

• Remark: analytical solutions are more correct than the simple $1.64 \frac{\sqrt{b}}{s}$ and no more complicated to use in sensitivity/optimization studies \Rightarrow better to use them



Purpose of the study

- Purpose:
 - Always interesting to compare various methods
 - Comparing the two approaches tells us how sensitive the analysis is on the treatment of systematics
 - \star CL_s and bayesian with uniform prior are equivalent without systematics \Rightarrow differences arise only from the different treatment of systematics
- Bayesian implementation validated by comparing it to CLsGenerator/analytical without systematics

Bayesian implementation

- Statistical model built using RooFit/RooStat
 - ▶ It is exactly the same as the CLsGenerator (mclimit) one
 - ▶ It is build from the same input files as CLsGenerator
- Marginal likelihood is

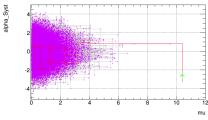
$$\mathscr{L}_{\mathsf{m}}(\mu) = \int \frac{(\mu s + \sum_{i} b_{i})^{N_{obs}}}{N_{obs}!} \, \mathrm{e}^{-(\mu s + \sum_{i} b_{i})} \prod_{j} \mathrm{e}^{-\frac{\eta_{j}^{2}}{2}} \prod_{i} \mathrm{e}^{-\frac{(\gamma_{i} - 1)^{2}}{2\sigma_{i}^{2}}} \, \mathrm{e}^{-\frac{(\gamma_{s} - 1)^{2}}{2\sigma_{s}^{2}}} \prod_{j} \mathrm{d}\eta_{j} \prod_{i} \mathrm{d}\gamma_{i} \mathrm{d}\gamma_{s}$$

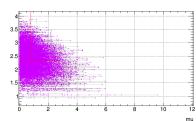
Marginalization done by Markov Chain Monte Carlo

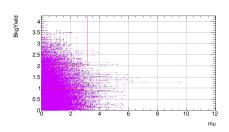
Visualization of Markov Chain

 \bullet Plots below shows markov chain for following configuration: $_{\text{+bg Bkg }0.82\ 1}$

```
+bg Bkg 0.82 1
.syst Syst 0.02 -0.1
+sig Sig 2.49 0.4
+data 1
```

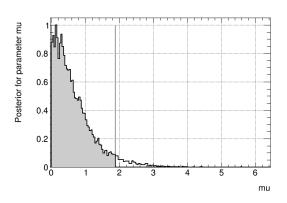






Posterior

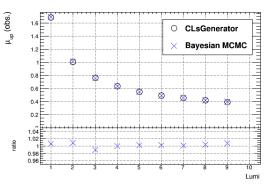
 Plot below shows posterior and 95% CL interval for same configuration as in previous slide



Validation of bayesian implementation

 CLsGenerator and bayesian (uniform prior) should be the same without systematics

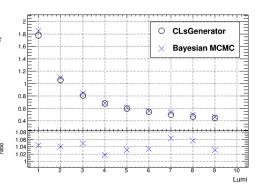
- Configuration
 - $N_{obs} = 1$
 - b = 0.82
 - s = 2.49
 - No stat/syst uncertainty

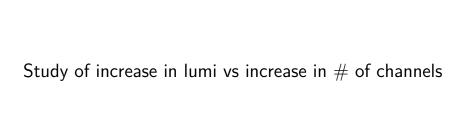


Conclusion: bayesian implementation validated

CLsGenerator/bayesian comparison with systematics

- Configuration
 - $N_{obs} = 1$
 - b = 0.82
 - s = 2.49
 - ► Stat uncert. on *b*: 0.5
 - Stat uncert. on s: 0.3
 - Syst uncert. on s:
 - ★ up=0.05
 - ★ down=-0.05
 - ► Syst uncert. on *b* (100% correlated with *s*):
 - ★ up=0.3
 - \star down=-0.3
- Effect of systematics treatment difference starts to be visible
- Need to study the effect of systematics further in both cases





Lumi vs # channels (1)

- We expect that increasing luminosity by factor N is equivalent to increasing number of channels (having the same sensitivity) by the same factor (demo. on next slides)
- Here we check that CLsGenerator and BayesianMCMC are in agreement with this expectation
 - ▶ Validates in both cases implementation of channel combination

Demonstration equivalence lumi/#channels (CLsGenerator)

- Single channel: $q_{\mu} = 2 \left[\mu s N_{\text{obs}} \ln \frac{\mu s + b}{b} \right]$
- *N* channels: $q'_{\mu} = 2 \left[\mu \sum_{c:channels} s_c \sum_{c:channels} N^c_{obs} \ln \frac{\mu s_c + b_c}{b_c} \right]$
 - Assume yields are the same in all channels $(s_c = s, b_c = b, N_{\text{obs}}^c = N_{\text{obs}})$: $q'_{\mu} = 2 \left[\mu Ns N_{\text{obs}} \ln \left(\prod_{c:channels} \frac{\mu s_c + b_c}{b_c} \right) \right] = N \times 2 \left[\mu s N_{\text{obs}} \ln \frac{\mu s + b}{b} \right]$ $\Rightarrow q'_{\mu} = Nq_{\mu}$
- Multiplying lumi. by N: $q''_{\mu}=2\left[\mu s''-N''_{\mathrm{obs}}\ln\frac{\mu s''+b''}{b''}\right]$
 - ightharpoonup s'' = Ns, b'' = Nb and $N''_{obs} = NN_{obs}$

$$\Rightarrow$$
 $q''_{\mu} = N \times 2 \left[\mu s - N_{\mathrm{obs}} \ln \frac{\mu s + b}{b} \right] = N q_{\mu}$

• Conclusion: $q''_{\mu} = q'_{\mu}$

Demonstration equivalence lumi/#channels (BayesianMCMC)

- Single channel: $\mathscr{L}(\mu) = \frac{(\mu s + b)^{N_{\text{obs}}}}{N_{\text{tot}}} e^{-(\mu s + b)}$
- N channels:

$$\mathscr{L}'(\mu) = \prod_{c:channels} \mathscr{L}_c(\mu) = \prod_{c:channels} \frac{(\mu s_c + b_c)^{N_{obs}^c}}{N_{obs}^c} e^{-(\mu s_c + b_c)}$$

Assume yields are the same in all channels ($s_c = s$, $b_c = b$, $N_{\rm obs}^c = N_{\rm obs}$):

$$\Rightarrow \mathcal{L}'(\mu) = \left[\frac{(\mu s + b)^{N_{\text{obs}}}}{N_{\text{obs}}!} e^{-(\mu s + b)}\right]^{N} = \frac{1}{(N_{\text{obs}}!)^{N}} (\mu s + b)^{NN_{\text{obs}}} e^{-N(\mu s + b)}$$

- Multiplying lumi. by N: $\mathscr{L}''(\mu) = \frac{(\mu s'' + b'')^{N''_{obs}}}{N''_{obs}!} e^{-(\mu s'' + b'')}$
 - s'' = Ns, b'' = Nb and $N''_{obs} = NN_{obs}$

$$\Rightarrow \mathcal{L}''(\mu) = \frac{(\mu Ns + Nb)^{NN_{\text{obs}}}}{(NN_{\text{obs}})!} e^{-(\mu Ns + Nb)} = \frac{N^{NN_{\text{obs}}}}{(NN_{\text{obs}})!} (\mu s + b)^{NN_{\text{obs}}} e^{-N(\mu s + b)}$$

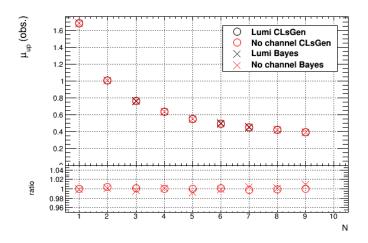
• Conclusion: $\mathscr{L}''(\mu) \propto \mathscr{L}'(\mu)$ so inference on μ is the same

Lumi vs # channels (2)

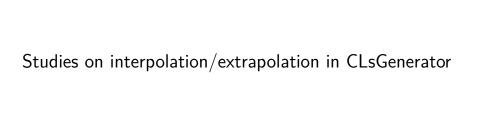
- We consider this initial situation (no stat./syst. uncertainties):
 - $N_{obs} = 1$
 - b = 0.82
 - s = 2.49
- Then we compare two limits:
 - first one computed from one channel with
 - ★ $N_{obs} = 1 \times N$
 - ★ $b = 0.82 \times N$
 - ★ $s = 2.49 \times N$

- second one computed from N identical channels with
 - ★ $N_{obs} = 1$
 - ★ b = 0.82
 - **★** *s* = 2.49

Lumi vs # channels (3)



 Conclusion: expectation verified in both CLsGenerator and BayesianMCMC



Interpolation/extrapolation of systematics (1)

- For one sample and one systematic uncertainty we know
 - ► N_{nom}: nominal yield
 - ▶ N_{\uparrow} : yield with systematic varied 1σ up
 - ▶ N_{\downarrow} : yield with systematic varied 1σ down
- However, for the purpose of setting limits we need a continuous parametrization of the yield: $N(\eta)$
 - ▶ η defined such that $N(\eta=0)=N_{nom},\ N(\eta=+1)=N_{\uparrow}$ and $N(\eta=-1)=N_{\downarrow}$
- How do we interpolate for $\eta = [-1,1]$ and extrapolate for $\eta < -1$ and $\eta > 1$?

Interpolation/extrapolation of systematics (1)

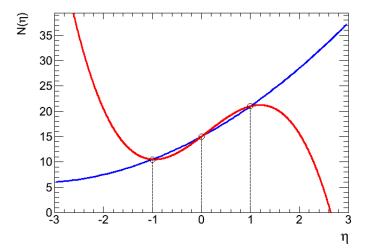
- For one sample and one systematic uncertainty we know
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- $oldsymbol{\bullet}$ How do we interpolate for $\eta=[-1,1]$ and extrapolate for $\eta<-1$ and $\eta>1$?

Interpolation/extrapolation of systematics (1)

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 - ► N_{nom}: nominal yield
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- How do we interpolate for $\eta = [-1,1]$ and extrapolate for $\eta < -1$ and $\eta > 1$?

Interpolation/extrapolation of systematics (2)

Example : $N_{nom}=15,~N_{\uparrow}=21,~N_{\downarrow}=10.5$



Interpolation/extrapolation of systematics (3)

• Rather than using $N_{nom,\uparrow,\downarrow}$, let's use

$$f^{\uparrow} = \frac{N_{\uparrow} - N_{nom}}{N_{nom}}$$

$$f^{\downarrow} = \frac{N_{\downarrow} - N_{nom}}{N_{nom}}$$

$$f^{syst}(\eta) = \frac{N(\eta)}{N_{nom}}$$

One has

$$f^{syst}(\eta = 0) = 1$$

$$f^{syst}(\eta = -1) = 1 + f^{\downarrow}$$

$$f^{syst}(\eta = 1) = 1 + f^{\uparrow}$$

 Goal: find an inter/extrapolation algorithm such that these relations are satisfied (at least approximately)

Interpolation/extrapolation of systematics (3)

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Interpolation/extrapolation of systematics (3)

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$$f^{\downarrow} = \frac{N_{\downarrow} - N_{nom}}{N_{nom}}$$

$$f^{syst}(\eta) = \frac{N(\eta)}{N_{nom}}$$

One has

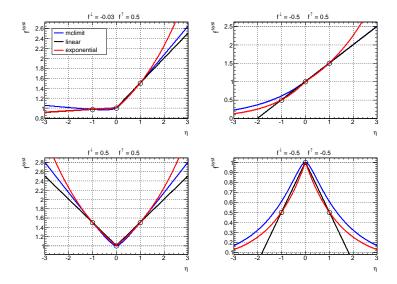
$$f^{syst}(\eta = 0) = 1$$

$$f^{syst}(\eta = -1) = 1 + f^{\downarrow}$$

$$f^{syst}(\eta = 1) = 1 + f^{\uparrow}$$

• Goal: find an inter/extrapolation algorithm such that these relations are satisfied (at least approximately)

First look at mclimit, linear and exponential algorithms



Closer look at mclimit algorithm (1)

- What does mclimit give for $\eta = -1, 0, +1$?
 - $f^{syst}(\eta = 0) = 1$
 - if $f^{\uparrow} > 0$: $f^{syst}(\eta = +1) = 1 + f^{\uparrow}$
 - if $f^{\downarrow} > 0$: $f^{syst}(\eta = -1) = 1 + f^{\downarrow}$
 - if $f^{\uparrow} < 0$: $f^{syst}(\eta = +1) = e^{f^{\uparrow}}$
 - if $f^{\downarrow} < 0$: $f^{syst}(\eta = -1) = e^{f^{\downarrow}}$

In the last two cases, $f^{syst} \simeq 1 + f^{\uparrow(\downarrow)}$ only if $f^{\uparrow(\downarrow)} \simeq 0$ (if $f^{\uparrow(\downarrow)} \simeq -1$, difference can be large).

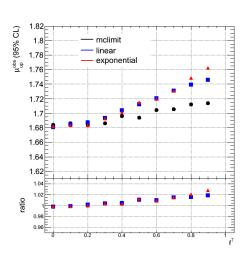
Closer look at mclimit algorithm (2)

- What does mclimit give when uncertainties are symetric: $f^{\uparrow} = -f^{\downarrow}$?
 - $\eta > 0$:
 - ★ $f^{\uparrow} \ge 0$: $f^{syst} = 1 + \eta f^{\uparrow}$
 - ★ f^{\uparrow} < 0: $f^{syst} = e^{\eta f^{\uparrow}}$
 - $\eta < 0$:
 - ★ $f^{\downarrow} \ge 0$: $f^{syst} = 1 \eta f^{\downarrow}$
 - $\star f^{\downarrow} < 0$: $f^{syst} = e^{-\eta \dot{f}^{\downarrow}}$

mclimit is equivalent to linear for $\eta>0$ if $f^\uparrow\geq 0$ or $\eta<0$ if $f^\downarrow\geq 0$

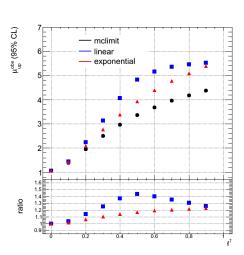
Comparison of algorithms: test 1

- $N_{obs} = 1$
- b = 0.82
- s = 2.49
- No stat uncertainty
- No syst uncertainty on s
- Syst uncertainty on b: $f^{\uparrow} = -f^{\downarrow} = 0, 0.1, 0.2, \dots, 0.9$



Comparison of algorithms: test 2

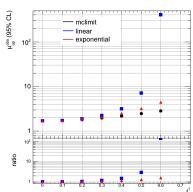
- $N_{obs} = 100$
- b = 100
- s = 20
- No stat uncertainty
- No syst uncertainty on s
- Syst uncertainty on b: $f^{\uparrow} = -f^{\downarrow} = 0, 0.1, 0.2, \dots, 0.9$



Comparison of algorithms: test 3

- $N_{obs} = 1$
- b = 0.82
- s = 2.49
- No stat uncertainty
- Syst uncertainty on s and b (100% correlated): $f^{\uparrow} = -f^{\downarrow} = 0.01, 0.2, 0.0$

$$f^{\uparrow} = -f^{\downarrow} = 0, 0.1, 0.2, \dots, 0.6$$

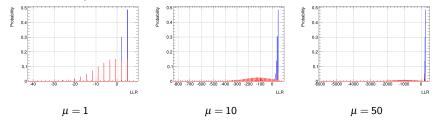


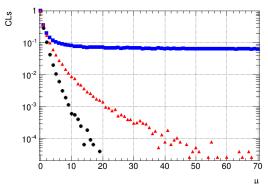
Comments:

- $q_{\mu} = 2\left(\mu s N_{obs} \ln \frac{\mu s + b}{b}\right)$
- When $f^{\downarrow} \ll 0$, $f^{syst} = 0$ quite frequently $\Rightarrow s = b = 0 \Rightarrow N_{obs} = 0 \Rightarrow q_{\mu} = 2\mu s$ (for both hypothesis) \Rightarrow CL_{s+b} and CL_s can't go to very low values as μ increases

Comparison of algorithms: test 3 (cont'd)

Consider the point $f^{\uparrow} = -f^{\downarrow} = 0.6$





Comparison of algorithms: test 3 (cont'd)

- After having studied a few situations by varying systematics on signal, backgrounds and correlations, it appears that one should be very careful with linear inter/extrapolation
 - when signal systematic uncertainty is large
 - when in addition background systematics are large effect is more pronounced
 - when in addition background systematics are large and correlated to signal systematics effect can be dramatic (see two previous slides)
- Remark: effect exists only if f^{\uparrow} and/or f^{\downarrow} (for the signal at least) is negative

Comparison of algorithms: LHCP mumu channel

Here we use 4tmm.dat from David (with +data 1)

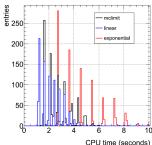
Limits

	mclimit	linear	expo
Expected median	1.70	1.70	1.69
Expected $\pm 1\sigma$	1.44-2.08	1.43-2.07	1.43-2.07
Expected $\pm 2\sigma$	1.26-3.52	1.27-3.53	1.27-3.54
Observed	1.67	1.68	1.67

Conclusion: the three algorithms give compatible results

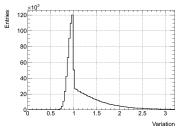
CPU time

ran
clgen.observedSigStrengthFor95excl(0,1e5,cls)
in identical conditions several
times to make this distrib



Understanding f^{syst} distribution: exponential case

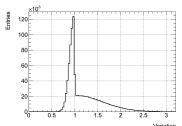
• f^{syst} sometimes looks very weird. Let's consider exponential algo with $f^{\downarrow} = -0.1$ and $f^{\uparrow} = 0.6$ (on the plot, Variation= f^{syst})



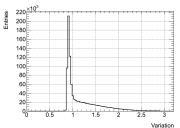
- Where does this funny looking shape comes from ?
 - We have $f^{syst}=(1+f^{\uparrow})^{\eta}$ for $\eta>0$ $((1+f^{\uparrow})^{-\eta}$ for $\eta<0$) with $\eta \sim \mathcal{N}(0,1)$
 - It's straightforward to show that $f^{syst} \sim \frac{1}{\sqrt{2\pi}|\ln(1+f^{\uparrow(\downarrow)})|f^{syst}|} e^{-\frac{1}{2}\left(\frac{\ln f^{syst}}{\ln(1+f^{\uparrow(\downarrow)})}\right)^2}$
 - i.e. $f^{syst} \sim \text{piecewise } \log \mathcal{N}$
- Note: understanding the shape of a distribution doesn't mean that such a distribution is desirable \rightarrow is it?

Understanding f^{syst} distribution: linear and mclimit cases

• linear case is obvious: $f^{syst} \sim$ piecewise gaussian truncated at 0. Using values of previous slide we have

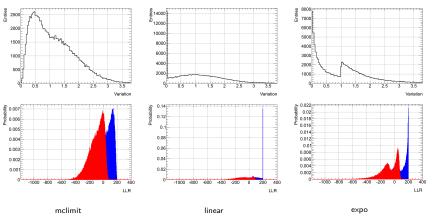


 mclimit case: doesn't seem to be possible to determine analytic solution. Using values of previous slide we have



Coming back to test 2

- In some extreme cases (very large systematics), linear and expo give really really worrying distributions
- Let's consider the point $f^{\uparrow} = -f^{\downarrow} = 0.9$ and $\mu = 5.5$ of test 2.



Using truncated piecewise gaussian (linear) and piecewise log N
 (expo) seems to be absurd →mclimit seems to be much more
 reasonable in such large systematics case

Preliminary conclusions (1)

- mclimit, linear and exponential algorithms give similar results for small systematics
 - Any choice seems to be rather safe in this case
- Differences can be significant for large systematics
 - linear shouldn't be used when signal systematics (and background ?) are large
 - expo shouldn't be used when systematics on background and/or signal are large and f^{\downarrow} or $f^{\uparrow} \ll 0$
 - mclimit seems to be the safest choice when systematics are large
- Argument of continuous derivative which at first sight seemed important only for MINUIT is actually also important when no fitting is done as it provides distributions of f^{syst} that are smooth (particularly important for large systematics).

Preliminary conclusions (2)

- mclimit:
 - pros: f^{syst} smooth, $f^{syst}(\eta) \neq 0 \ \forall \eta$
 - cons: $f^{syst}(\eta = \pm 1) \neq 1 + f^{\uparrow(\downarrow)}$ when $f^{\uparrow(\downarrow)} < 0$,
- linear:
 - pros: simple, fast
 - ▶ cons: $f^{syst}(\eta)$ =0 in some cases (\Rightarrow problem when signal syst large)
- expo:
 - ▶ pros: simple, $f^{syst}(\eta) \neq 0 \ \forall \eta$
 - cons: f^{syst} can be very discontinuous for large $f^{\uparrow(\downarrow)}$, slow

Preliminary conclusions (3)

• Systematics are such a pain! It's good that we're not affected too much by them in the same-sign analysis

Ideas for further studies (1)

- Try linear interpolation and expo extrapolation
- Alternative to TMath::Power() for exponential case ?
- Try $\eta \sim log \mathcal{N}$
 - Note that $\eta \sim log \mathcal{N} + linear$ is equivalent to $\eta \sim \mathcal{N} + expo$ (in both cases $f^{syst} \sim log \mathcal{N}$)
 - * As $\eta \sim \mathcal{N}+$ expo is quite slow, could be interesting to see if it's faster in terms of CPU to do $\eta \sim \log \mathcal{N}+$ linear
- Try $\eta \sim$ Gamma

Ideas for further studies (2)

Statistical model

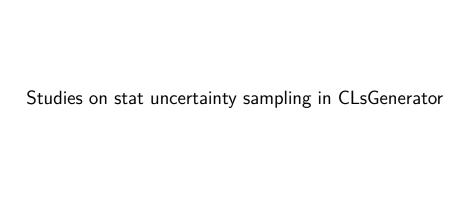
$$\mathscr{L}(\mu, \{\eta\}, \{\gamma\}) = \frac{(\mu s + \sum_i b_i)^{N_{obs}}}{N_{obs}!} e^{-(\mu s + \sum_i b_i)}$$
 where

- $b_i = b_i(\{\eta\}, \{\gamma\}) = b_i^{nom} \times \prod_j f_{ij}(\eta_j) \times \gamma_i$
- $s = s(\{\eta\}, \{\gamma\}) = s^{nom} \times \prod_j f_j^s(\eta_j) \times \gamma_s$

and

- $\eta_j \sim \mathcal{N}(0,1)$
 - $\gamma_i \sim \mathcal{N}(1, \sigma_i)$ and $\gamma_s \sim \mathcal{N}(1, \sigma_s)$
 - When generating pseudo exp, N_{obs} generated according to marginal likelihood:

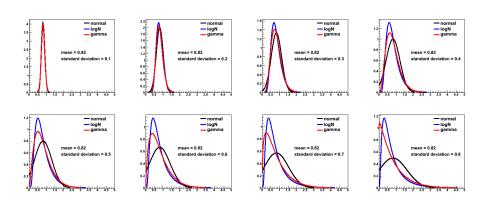
$$\begin{split} \mathscr{L}_{\text{m}}(\mu) &= \int \frac{(\mu s + \Sigma_{i} \, b_{i})^{N_{obs}}}{N_{obs}!} \, e^{-(\mu s + \Sigma_{i} \, b_{i})} \prod_{j} e^{-\frac{\eta_{j}^{2}}{2}} \prod_{i} e^{-\frac{(\gamma_{i} - 1)^{2}}{2\sigma_{i}^{2}}} \, e^{-\frac{(\gamma_{s} - 1)^{2}}{2\sigma_{s}^{2}}} \prod_{j} \mathrm{d}\eta_{j} \prod_{i} \mathrm{d}\gamma_{i} \mathrm{d}\gamma_{s} \\ & \text{However, } \mathscr{L}(\mu, \{\eta\} = 0, \{\gamma\} = 1) \text{ is used to calculate the test} \\ & \Rightarrow \text{try calculating the test with marginal likelihood} \end{split}$$



Purpose of the study

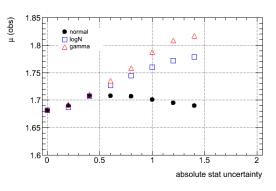
- Nuisance parameters for stat uncertainty sampled from normal distribution by default
- This choice is arbitrary and other distributions could be used
 - ▶ In this study normal is compared to logN and gamma
- Following results obtained with the mclimit inter/extrapolation

Comparison of distributions



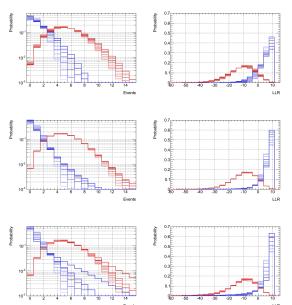
Comparison of samplings for particular case (1)

- N_{obs} = 1
- b = 0.82
- s = 2.49
- No syst uncertainty
- Stat uncertainty on b = 0.1, ..., 1.4



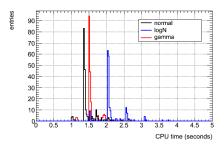
Comparison of samplings for particular case (2)

Normal (top), logN (middle) and Gamma (bottom)



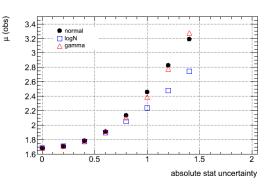
Comparison of samplings for particular case (3)

- Truncation at 0 in normal case doesn't smear distributions but shifts them
 - ▶ Happens for both b and s+b distribs $\Rightarrow \mu$ doesn't change much
- ullet logN smears less than gamma \Rightarrow better μ for logN than for gamma
- What is the best choice ? logN ?



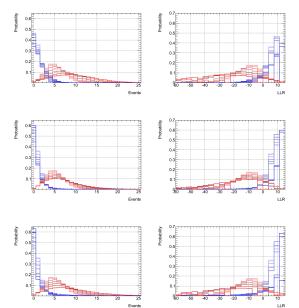
Comparison of samplings for another particular case (1)

- N_{obs} = 1
- b = 0.82
- s = 2.49
- No syst uncertainty
- Stat uncertainty on s and b = 0.1,...,1.4



Comparison of samplings for another particular case (2)

Normal (top), logN (middle) and Gamma (bottom)



Comparison of samplings for another particular case (3)

- logN smears less than gamma \Rightarrow better μ for logN than for gamma (as in previous case)
- Normal case now very different from previous case
 - ▶ s+b distribution is now "more smeared" than b alone distribution \Rightarrow have to go to larger μ to reach given confidence level