

# Bayesian Hierarchical Model for cross calibration

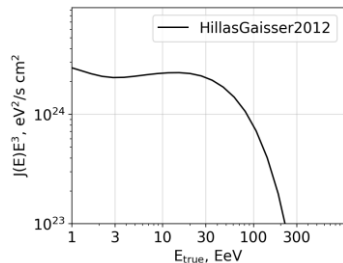
Anton Prosekin

TADML,  
October 17, 2025

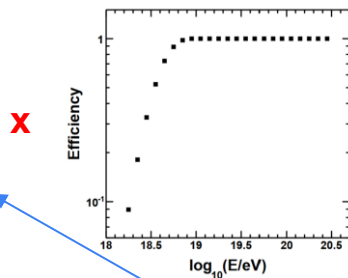


# Forward folding

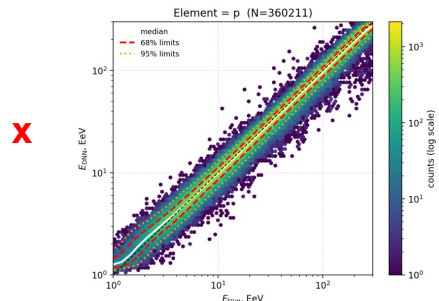
CR spectrum



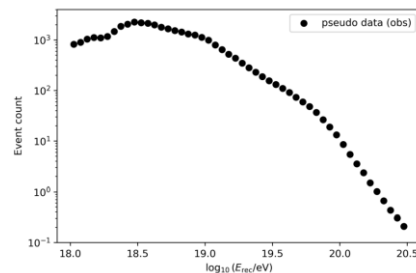
Efficiency



Response matrix/function



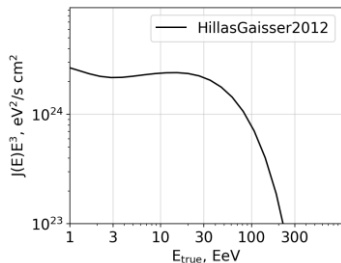
Event counts distribution



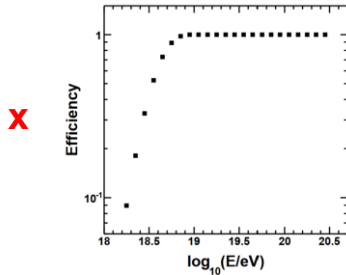
$$J^{\text{raw}}(E_{\text{SD}}; \mathbf{s}) = \frac{\int d\Omega \cos \theta \int dE \epsilon(E, \theta) J(E; \mathbf{s}) \kappa(E_{\text{SD}} | E; \theta)}{\int d\Omega \cos \theta}$$

# Forward folding: log-likelihood minimization

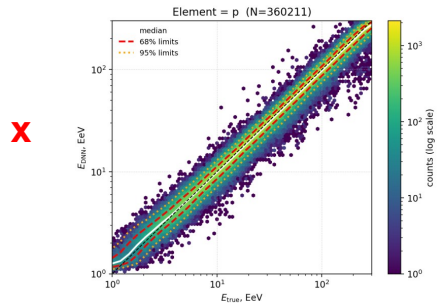
CR spectrum



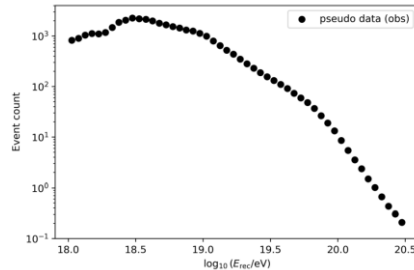
Efficiency



Response matrix/function



Event counts distribution



$$J^{\text{raw}}(E_{\text{SD}}; \mathbf{s}) = \frac{\int d\Omega \cos \theta \int dE \epsilon(E, \theta) J(E; \mathbf{s}) \kappa(E_{\text{SD}} | E; \theta)}{\int d\Omega \cos \theta}$$

binning

$$R_{ij} = \frac{\int_{\Delta E_i} dE_{\text{SD}} \int_{\Delta E_j} dE \int d\Omega \cos \theta \kappa(E_{\text{SD}} | E, \theta) \epsilon(E, \theta) J(E; \mathbf{s})}{\int_{\Delta E_j} dE \int d\Omega \cos \theta J(E; \mathbf{s})}$$

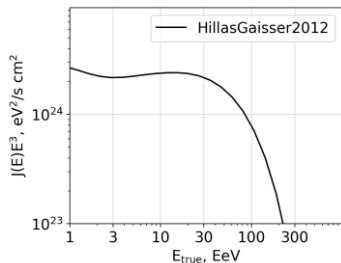
$$\nu_i = \sum_j R_{ij} \mu_j$$

Log Likelihood minimization:

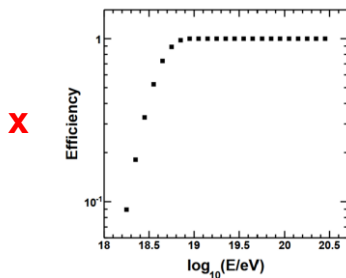
$$-\ln \mathcal{L}(\mathbf{s}) = \sum_i (\nu_i(\mathbf{s}) - N_i \ln \nu_i(\mathbf{s}))$$

# Forward folding: MCMC

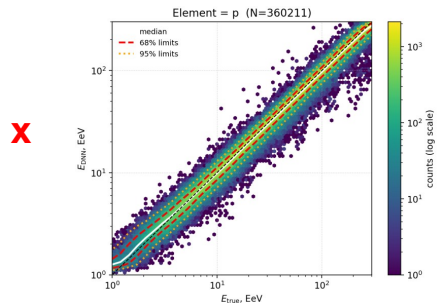
CR spectrum



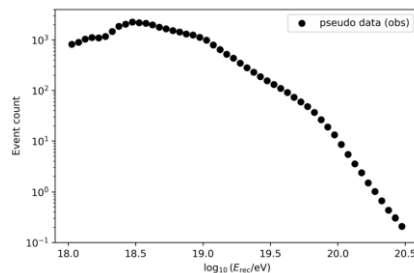
Efficiency



Response matrix/function



Event counts distribution



$$J^{\text{raw}}(E_{\text{SD}}; \mathbf{s}) = \frac{\int d\Omega \cos \theta \int dE \epsilon(E, \theta) J(E; \mathbf{s}) \kappa(E_{\text{SD}} | E; \theta)}{\int d\Omega \cos \theta} \quad \xrightarrow{\text{binning}} \quad R_{ij} = \frac{\int_{\Delta E_i} dE_{\text{SD}} \int_{\Delta E_j} dE \int d\Omega \cos \theta \kappa(E_{\text{SD}} | E, \theta) \epsilon(E, \theta) J(E; \mathbf{s})}{\int_{\Delta E_j} dE \int d\Omega \cos \theta J(E; \mathbf{s})}$$

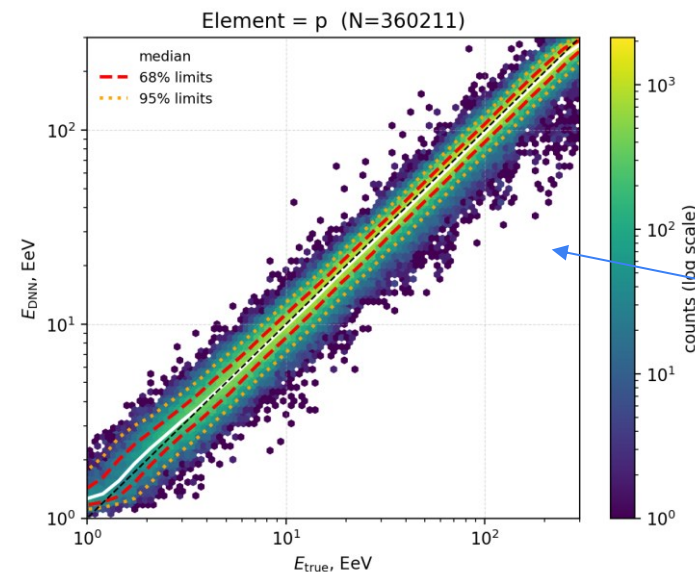
Log posterior sampling using MCMC:

$$p(\theta | \mathcal{D}) = \frac{p(\mathcal{D} | \theta) p(\theta)}{p(\mathcal{D})}$$

$$\nu_i = \sum_j R_{ij} \mu_j$$

$$\log p(\theta | D) \propto \sum_{i=1}^{N_{E, \text{rec}}} [N_i \log \nu_i(\theta) - \nu_i(\theta) - \log(N_i!)] + \log p(\theta)$$

# Response (resolution) matrix



$$\nu_i = \int dE f(E) \int du \varepsilon(E, u) \int_{\hat{E} \in \text{bin } i} \left[ \int ds p_{\text{det}}(\hat{E} | s, E, u) p_s(s | E, u) \right] d\hat{E}.$$

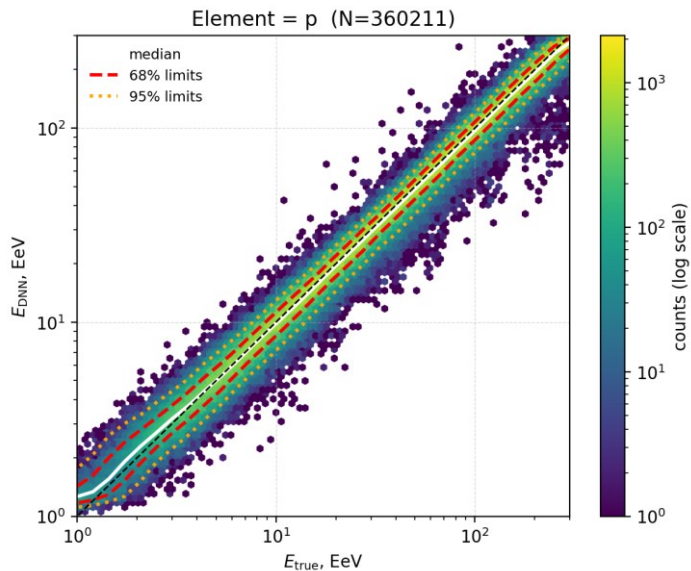
Resolution/smearing kernel:

$$\kappa(\hat{E} | E, u) = \int ds p_{\text{det}}(\hat{E} | s, E, u) p_s(s | E, u)$$

$$\nu_i = \int dE f(E) \int du \varepsilon(E, u) \int_{\hat{E} \in \text{bin } i} \kappa(\hat{E} | E, u) d\hat{E}$$

# Response (resolution) matrix

## Simulated resolution kernel



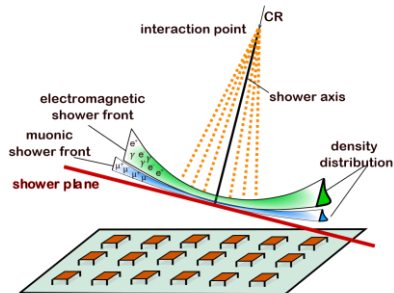
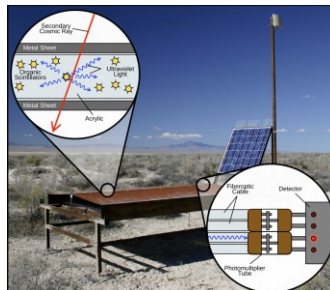
## Real resolution/smearing kernel:

$$\kappa(\hat{E} | E, u) = \int ds p_{\text{det}}(\hat{E} | s, E, u) p_s(s | E, u)$$

Detector fluctuations/smearing:

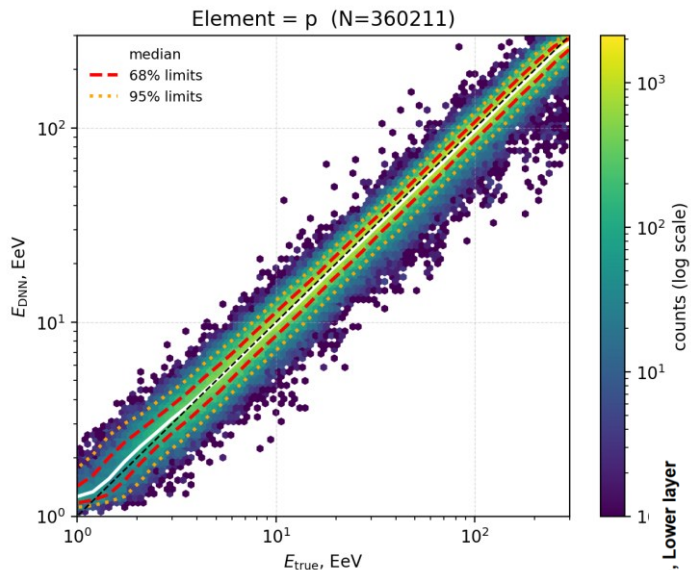
- sampling/LDF
- Scintillator/PMT

Shower size (s) fluctuations



# Response (resolution) matrix

## Simulated resolution kernel



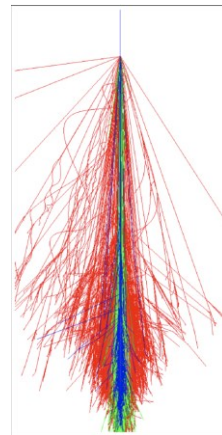
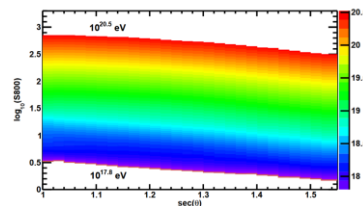
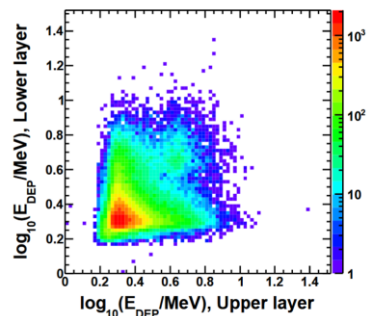
## Simulated resolution/smearing kernel:

$$\kappa(\hat{E} | E, u) = \int ds p_{\text{det}}(\hat{E} | s, E, u) p_s(s | E, u)$$

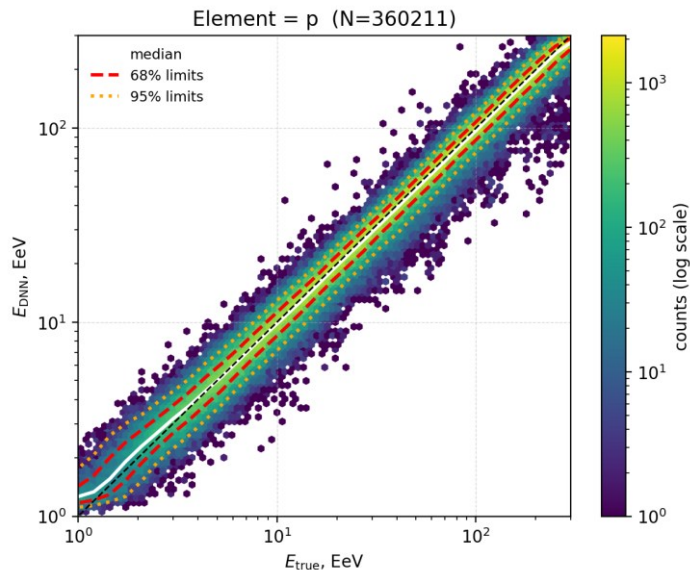
Detector fluctuations/smearing:

- Geant4
- Reconstruction (Std or DNN)

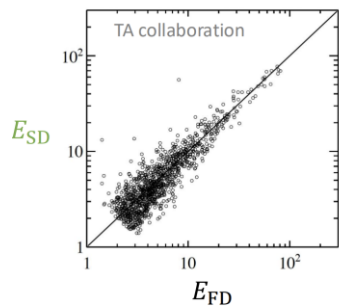
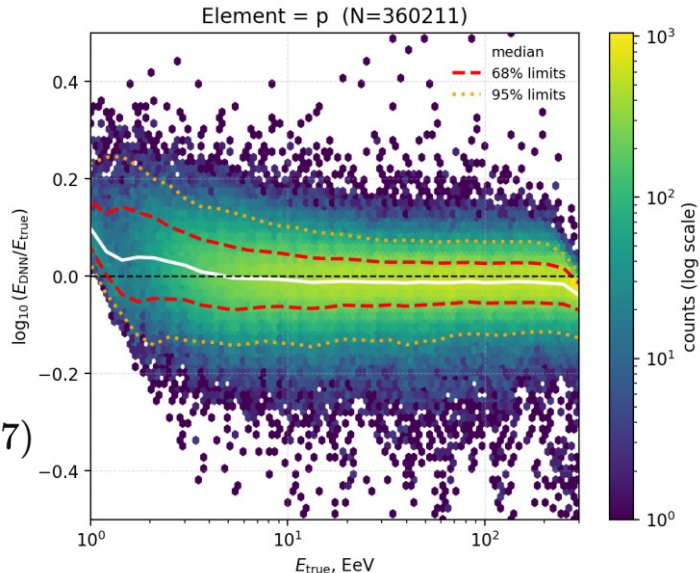
Shower size (s) fluctuations  
(CORSKA MC)



# Standard calibration



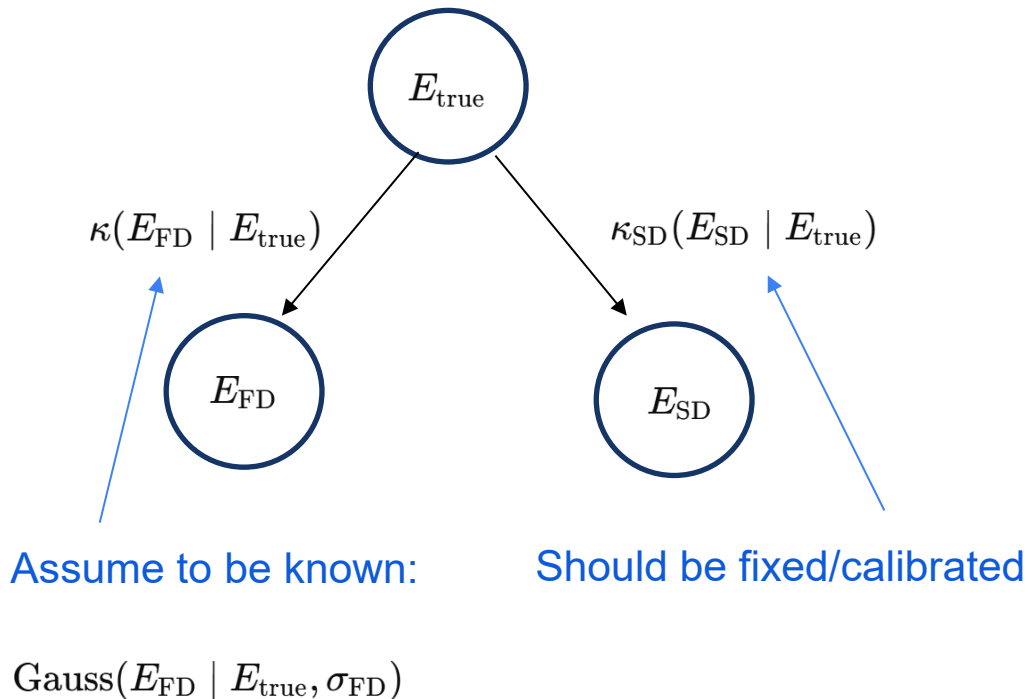
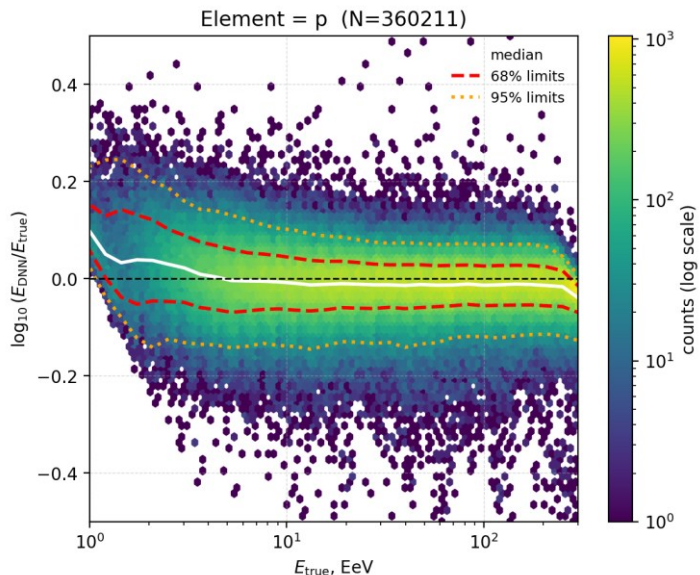
$\log_{10}(1.27)$



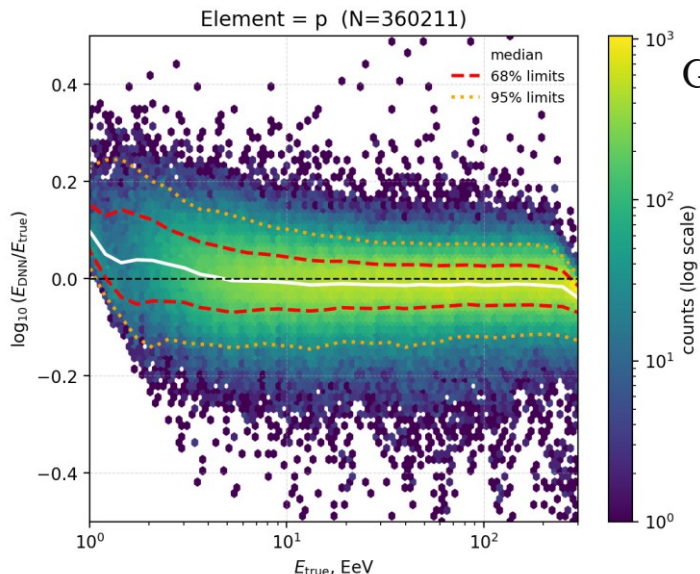
$$E = \frac{E_{SD}}{1.27}$$



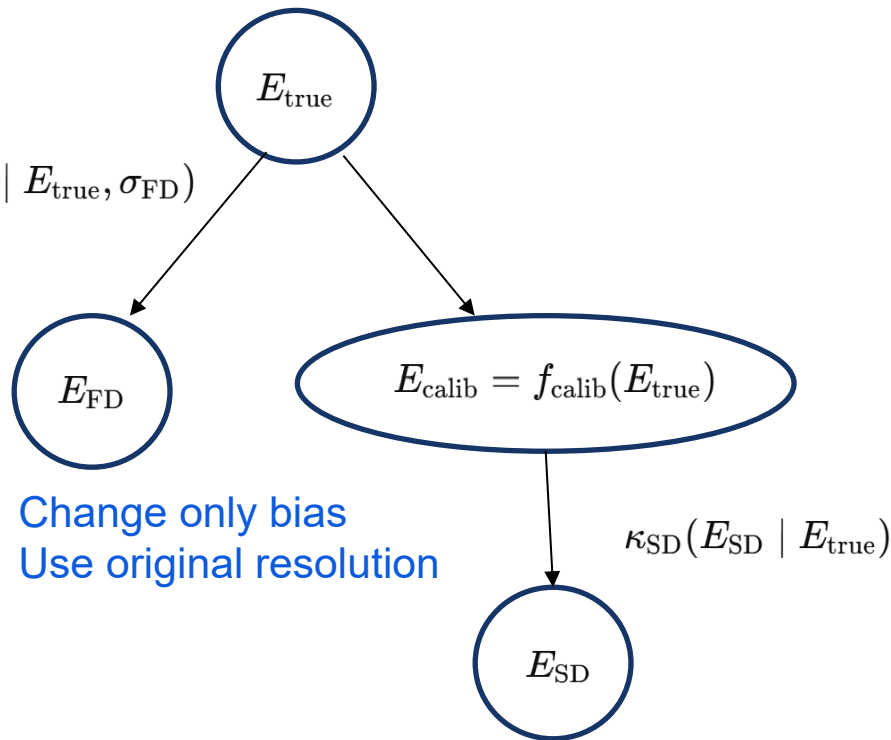
# Calibration with Hierarchical MCMC



# Calibration with Hierarchical MCMC



$$\text{Gauss}(E_{\text{FD}} | E_{\text{true}}, \sigma_{\text{FD}})$$



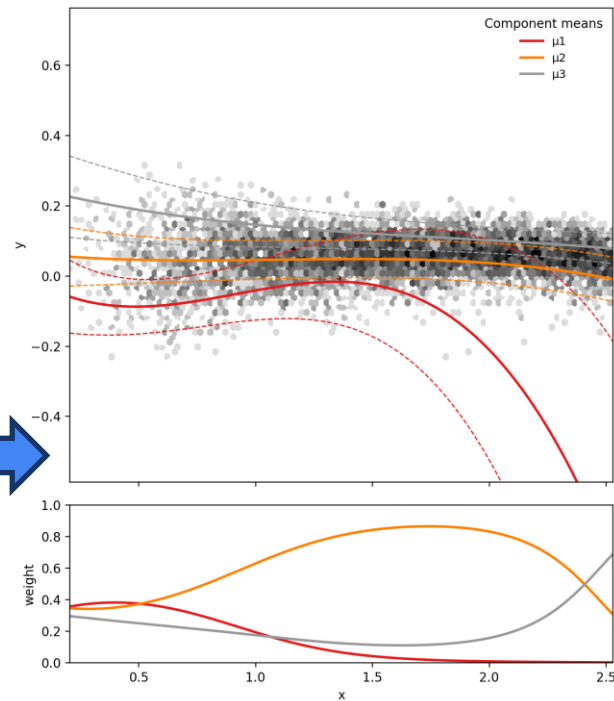
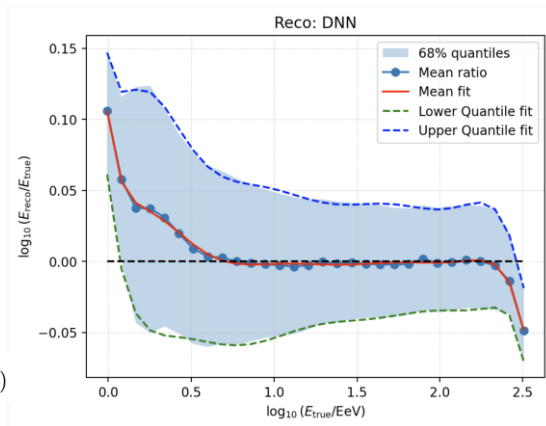
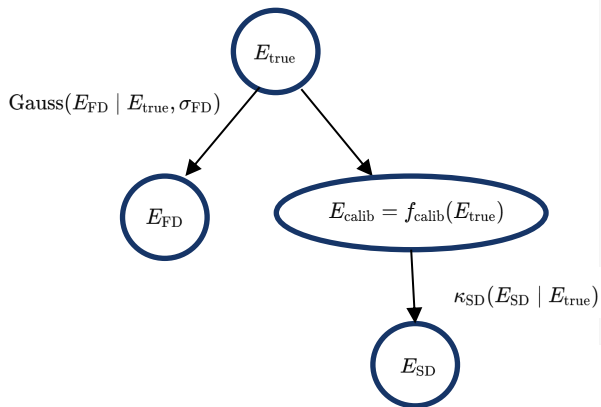
# Parametrization

Gauss (or Student-t) mixture model

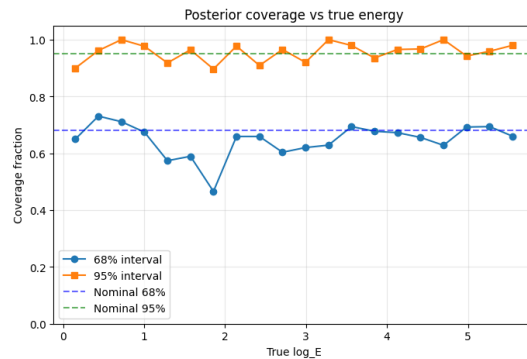
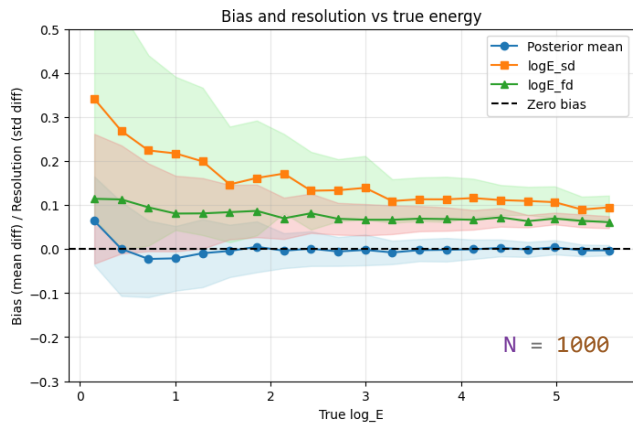
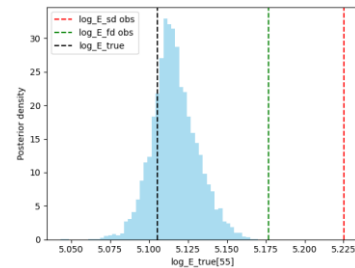
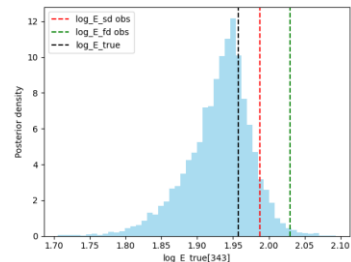
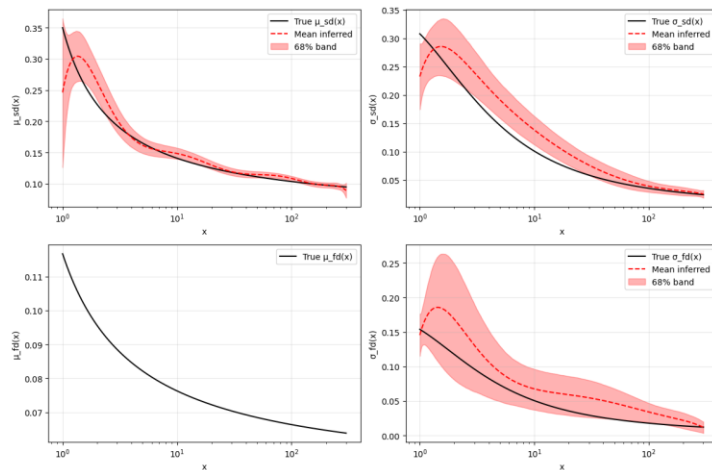
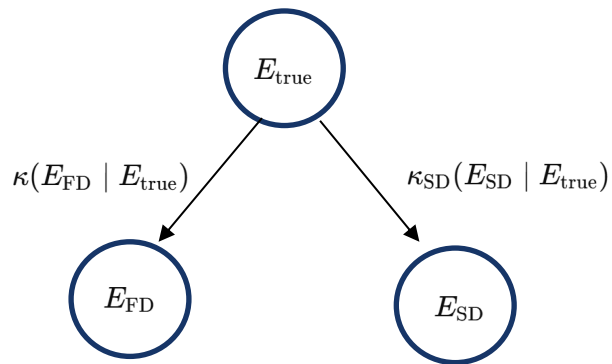
$$\kappa_{\text{SD}}(E_{\text{SD}} \mid E_{\text{true}}) = \sum_{n=1}^N w_n(x) G(r \mid \mu_n(x), \sigma_n(x))$$

$$r = \log_{10}(E_{\text{SD}}/E_{\text{true}}) \quad x = \log_{10} E_{\text{true}}$$

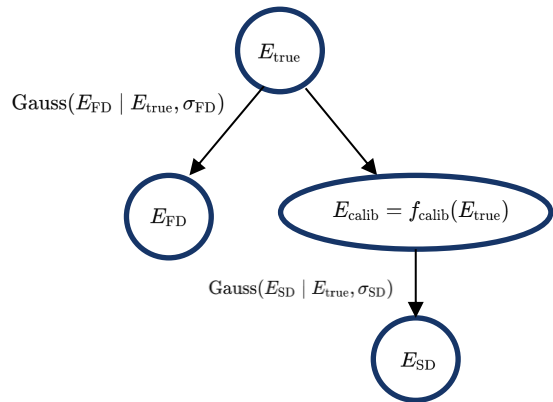
$$f_{\text{calib}} = a x^2 + b x + c$$



# Test: full resolution reconstruction



# Test: Gaussian, no energy dependence



Nsample = 1100

$a_{\text{true}} = 0$

$b_{\text{true}} = 0.1$

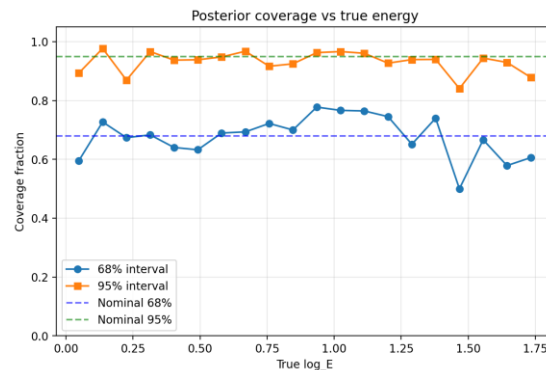
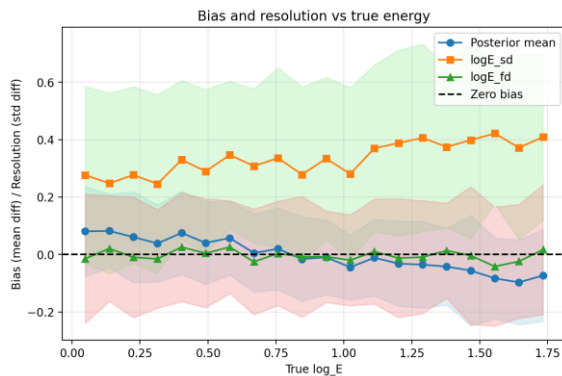
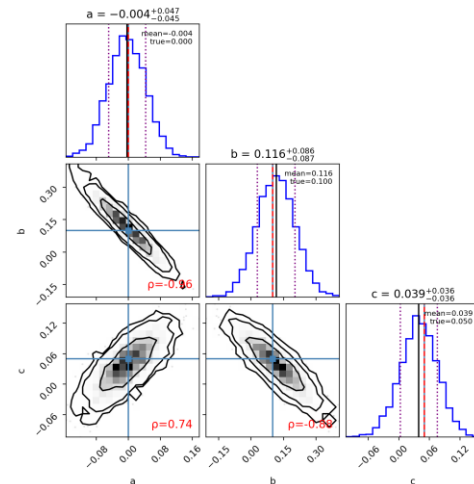
$c_{\text{true}} = 0.05$

$\mu_{\text{resid}} = 0.2$

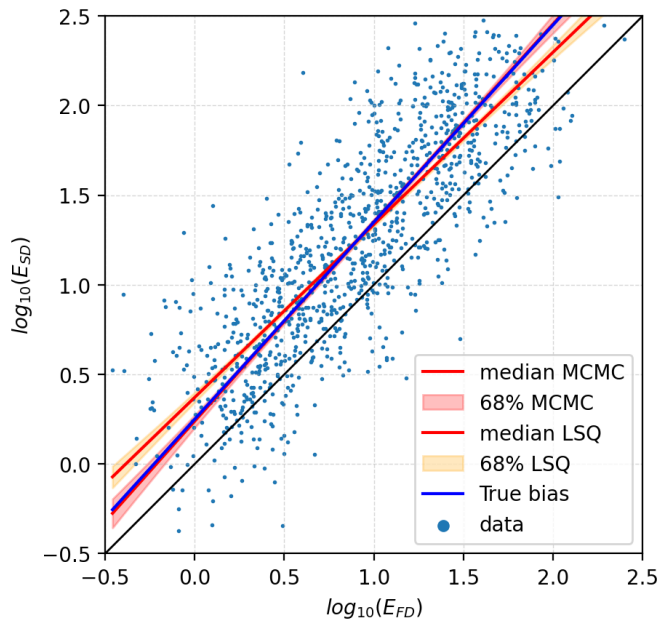
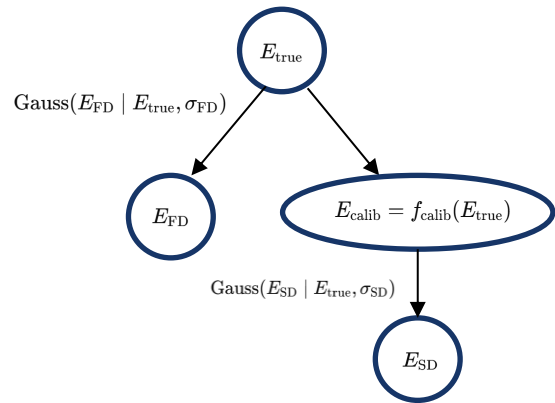
$\sigma_{\text{resid}} = 0.3$

$\sigma_{\text{fd\_frac}} = 0.2$

$$f_{\text{calib}} = ax^2 + bx + c$$



# MCMC vs least square minimization (LSQ)



MCMC:

$$\begin{aligned} a &= -0.004 \pm 0.045 \\ b &= 0.117 \pm 0.085 \\ c &= 0.038 \pm 0.035 \end{aligned}$$

LSQ:

$$\begin{aligned} a &= -0.001 \pm 0.033 \\ b &= -0.034 \pm 0.064 \\ c &= 0.372 \pm 0.028 \end{aligned}$$

Nsample = 1100

a\_true = 0

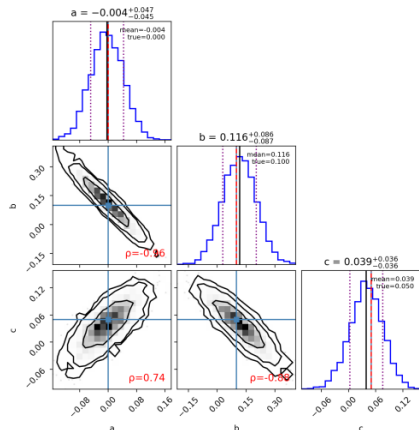
b\_true = 0.1

c\_true = 0.05

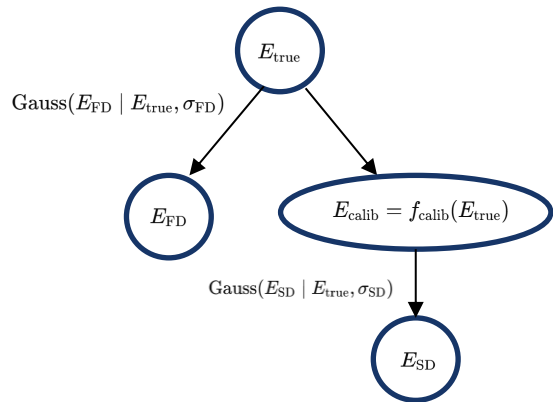
mu\_resid = 0.2

sigma\_resid = 0.3

sigma\_fd\_frac = 0.2



# MCMC vs least square minimization (LSQ)



Nsample = 1100

a\_true = 0

b\_true = 0.1

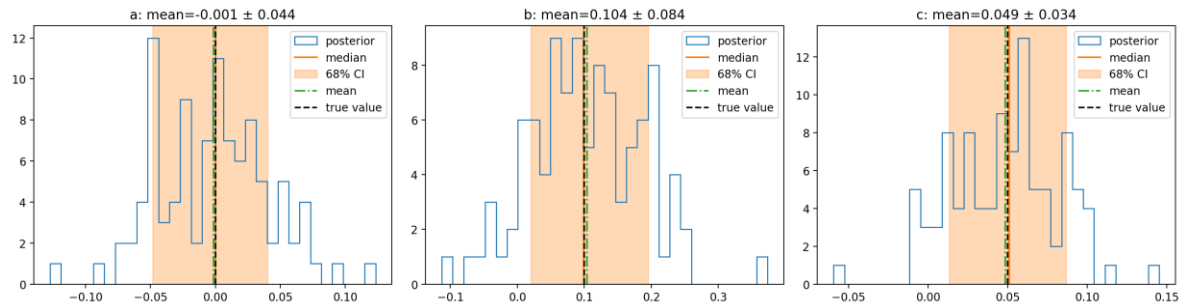
c\_true = 0.05

mu\_resid = 0.2

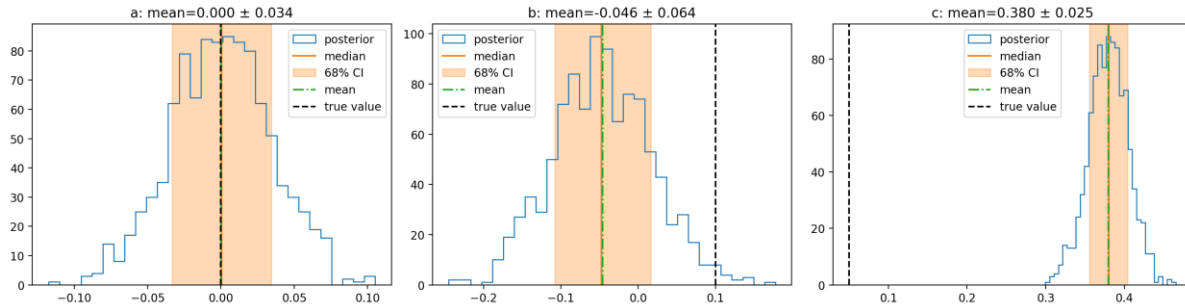
sigma\_resid = 0.3

sigma\_fd\_frac = 0.2

MCMC



LSQ



# Conclusions

- Calibration of the resolution function is required to correctly perform forward folding
- Auger used log-likelihood calibration on hybrid data; early work by Hans Dembinski showed that least-squares fits are biased
- MCMC is a powerful analog of the log-likelihood approach, as it does not require explicit multidimensional integration
- Bayesian MCMC calibration allows a natural extension of the MCMC forward folding method