

Analysis techniques of neutrino cross-section measurements in MiniBooNE

1. Introduction
2. Overview of MiniBooNE analysis
3. Neutrino xsec measurements
4. Conclusion

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1. Introduction
2. Overview of MiniBooNE analysis
3. Neutrino cross section measurements
4. Conclusion

1. Introduction

Purpose of this talk

- This talk is prepared for students who want to measure neutrino cross-sections for their PhD theses.
- It is especially focusing on **absolute flux-integrated topological differential cross section measurement [1]**. The most important **model-independent** result to provide to the community.

absolute: normalization is specified

flux-integrated: neutrino flux shape is not unfolded

topological: interaction is defined from final state particles

differential: cross section is function of measured kinematic variables

- MiniBooNE developed number of techniques necessary to measure these, and this talk covers technical aspects from the **CCQE**, **NCEL**, **NC1 π^0** , **CC1 π^+** , **CC1 π^0** , **antiCCQE**, **antiNCEL**. These are good reference but not the best, T2K should perform even better analyses!

[1] Can anybody invent a better name for this?

1. Introduction

Goal of neutrino cross section measurement

- Goal is to measure **model-independent cross section** as much as possible. This is what theorists want to study their models.

Model-independent cross-section is

- absolute (flux is not tuned from own measurement)
- the dependence of **signal channel MC** is minimum
- detector efficiency must be corrected (so it is detector model-dependent)
- no assumption on kinematics (cf neutrino energy reconstruction with lepton kinematics assume CCQE interaction and neutron at rest), which often means cross-section is function of measured variables (differential cross-section)

Formula of flux-integrated differential cross-section

$$\left(\frac{d\sigma}{dx}\right)_i = \frac{\sum_j U_{ij}(d_j - b_j)}{\varepsilon_i(\Phi T)\Delta x_i}$$

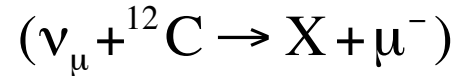
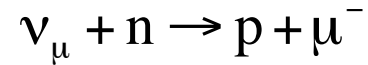
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2. Overview of MiniBooNE ν s measurements

CCQE (CC0 π)

PRD81(2010)092005

FERIMILAB-THESIS-2008-64



Signal definition: 1 μ + 0 π + N protons

Why we measure

- Test CCQE models

Why MiniBooNE measure

- Largest sample (~40%) to test detector efficiency, veto efficiency, event uniformity, timing, etc
- Best sample to study ν_e CCQE kinematics (=oscillation signal)
- ν_{μ} CCQE to constraint ν_e from μ -decay in oscillation sample

It is important to measure CCQE!

CCQE

2. Overview of MiniBooNE xs measurements

NCEL
PRD82(2011)092005
FERIMILAB-THESIS-2009-47

$$\nu_{\mu} + p \rightarrow \nu_{\mu} + p$$

$$\nu_{\mu} + n \rightarrow \nu_{\mu} + n$$

Signal definition: $0 \mu + 0 \pi + N$ protons

Why we measure

- Additional test of CCQE models
- Measurement of Δs
 - value is still controversial
 - connection of form factor (elastic) and PDF (inelastic)

$$\int_0^1 dx \Delta s(x) \equiv \Delta s \equiv G_A^s(Q^2 = 0)$$

Why MiniBooNE measure

- NCEL to constrain oil optical property
oil optical property is the largest detector systematics. NCEL was used to assign variation.

It is important to measure NCEL!

CCQE

NCEL



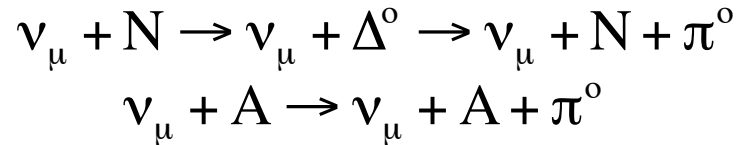
MiniBooNE collaboration,
PRD82(2011)092005

2. Overview of MiniBooNE ν s measurements

NC1 π^0

PRD81(2010)013005

FERIMILAB-THESIS-2010-49



Signal definition: 0 μ + 1 π^0 + N protons

Why we measure

- The biggest misID background for ν_e appearance experiments
- all oscillation experiments perform internal measurement to constrain

Why MiniBooNE measure

- π^0 mass peak for energy calibration
- measured π^0 kinematics is used to correct simulation

It is important to measure NC1 π^0 !



MiniBooNE collaboration,
PRD81(2010)013005

CCQE

NCEL

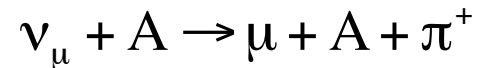
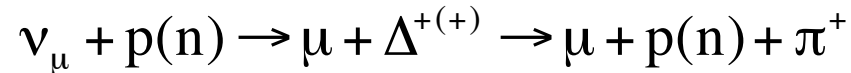
NC1 π^0

2. Overview of MiniBooNE ν s measurements

CC1 π^+

PRD83(2011)052007

FERIMILAB-THESIS-2009-27



Signal definition: 1 μ + 1 π^+ + N protons

Why we measure

- The biggest misID background for ν_{μ} disappearance experiments
- kinematic distortion by this background must be understood

Why MiniBooNE measure

- highest purity channel (~90%)
Michel electron tagging achieve extremely pure sample
- Constrain wrong sign background in anti-neutrino mode
CC1 π^+ in anti- ν mode is definitely from ν -contamination

It is important to measure CC1 π^+ !

CCQE

NCEL

NC1 π^0

CC1 π^+



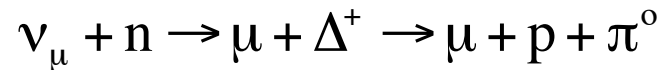
MiniBooNE collaboration,
PRD83(2011)052007

2. Overview of MiniBooNE ν s measurements

CC1 π^0

PRD83(2011)052009

FERIMILAB-THESIS-2010-9



Signal definition: 1 μ + 1 π^0 + N protons

Why we measure

- There is no coherent channel, so it is useful to understand coherent/resonance Δ production

Why MiniBooNE measure

- The last possibly measurable channels, by this, MiniBooNE measure 90% of interaction cross sections in ν -mode

It is important to measure CC1 π^0 !



MiniBooNE collaboration,
PRD83(2011)052009

CCQE

NCEL

NC1 π^0

CC1 π^+

CC1 π^0

2. Overview of MiniBooNE xs measurements

anti-CCQE

PRD88(2013)032001

FERIMILAB-THESIS-2013-14

Signal definition: $1 \mu + 0 \pi + N$ protons

$$\bar{\nu}_\mu + p \rightarrow n + \mu^+$$

$$(\bar{\nu}_\mu + {}^{12}\text{C} \rightarrow X + \mu^+)$$

$$(\bar{\nu}_\mu + {}^1\text{H} \rightarrow n + \mu^+)$$



MiniBooNE collaboration,
PRD88(2013)032001

Why we measure

- Additional test of CCQE models
- necessary measurement for CPV measurement

Why MiniBooNE measure

- To understand anti- ν_e CCQE interaction kinematics (oscillation signal)
- anti- ν_μ CCQE constraint for anti- ν_e from μ -decay
- Tune wrong sign components in anti- ν mode beam

It is important to measure antiCCQE!

CCQE

NCEL

NC1 π^0

CC1 π^+

CC1 π^0

anti
CCQE

2. Overview of MiniBooNE xs measurements

anti-NCEL

arXiv:1309.7257

FERIMILAB-THESIS-2012-29

$$\bar{\nu}_{\mu} + p \rightarrow \bar{\nu}_{\mu} + p$$

$$\bar{\nu}_{\mu} + n \rightarrow \bar{\nu}_{\mu} + n$$

Signal definition: 0 μ + 0 π + N protons



MiniBooNE collaboration,
arXiv:1309.7257

Why we measure

- Complete all 4 QE measurements (CCQE, NCEL, antiCCQE, antiNCEL)

Why MiniBooNE measure

- test beam-dump mode run (dark matter search)

anti- ν mode beam was used to test MiniBooNE dark matter sensitivity. antiNCEL was the biggest background.

It is important to measure antiNCEL!

CCQE

NCEL

NC1 π^0

CC1 π^+

CC1 π^0

anti
CCQE

anti
NCEL

2. Overview of MiniBooNE xs measurements

To be published

- CC inclusive cross section
- $1 \mu + 0 \pi + 1$ proton (2 track CCQE)

Possibly measured in T2K, but not in MiniBooNE

NC $1\pi^+$ ($0 \mu + 1 \pi^+ + N$ protons)

- another resonance only channel

$1 \mu + 1 \pi^+ + 1$ proton

- 3 track CC $1\pi^+$, high pure Δ^{++} measurement

$1 \mu + 0 \pi + 2$ protons

- high pure MEC?

CCQE

NCEL

NC $1\pi^0$

CC $1\pi^+$

CC $1\pi^0$

anti
CCQE

anti
NCEL

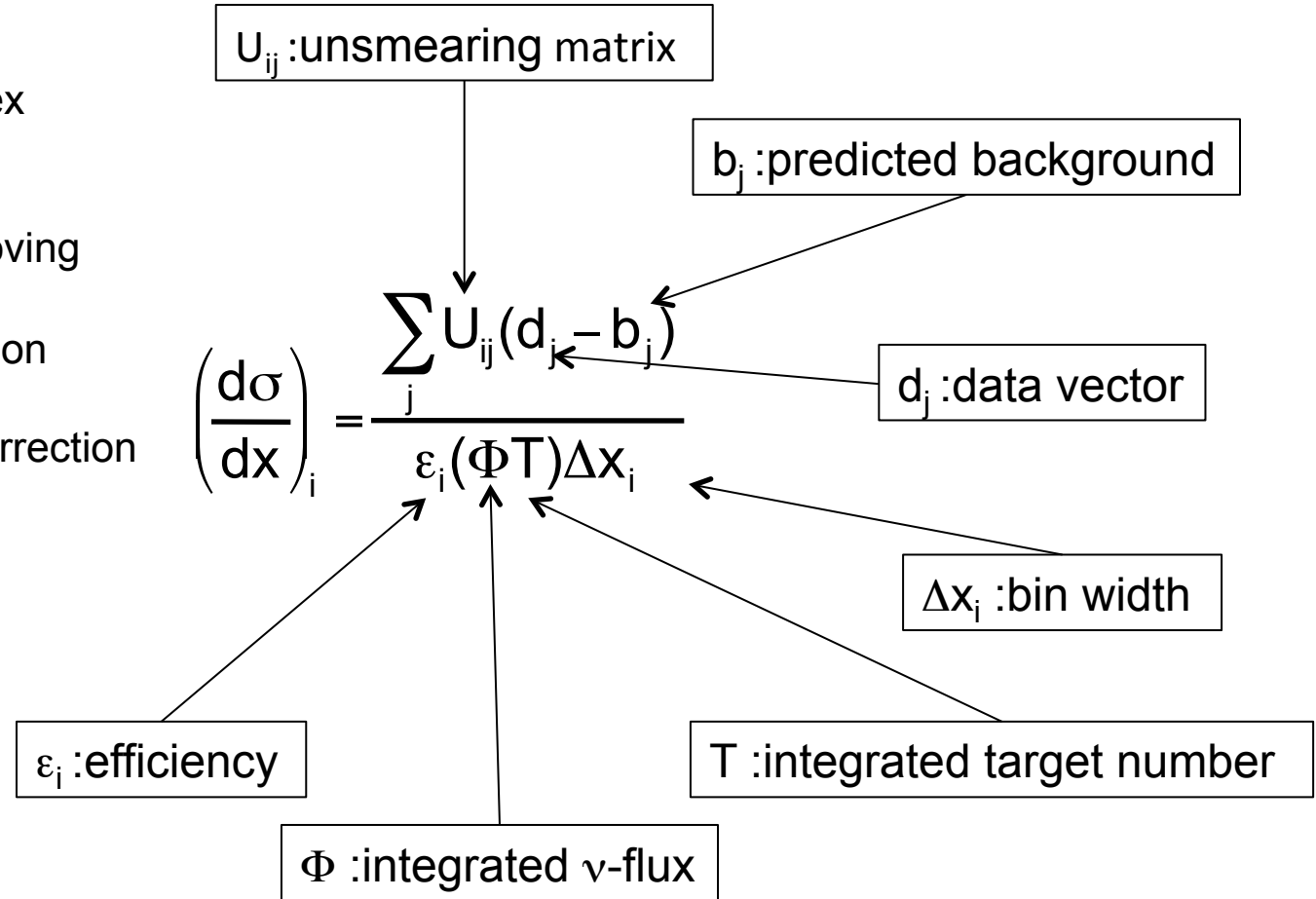
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3. Neutrino cross-section measurements

Absolute flux-integrated topological differential cross section formula

i : true index
 j : reconstructed index

- 3.1 Signal definition
- 3.2 Background removing
- 3.3 Unsmearing
- 3.4 Efficiency correction
- 3.5 Flux correction
- 3.6 Target number correction
- 3.7 Binning
- 3.8 Systematic errors
- 3.9 Data format



3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

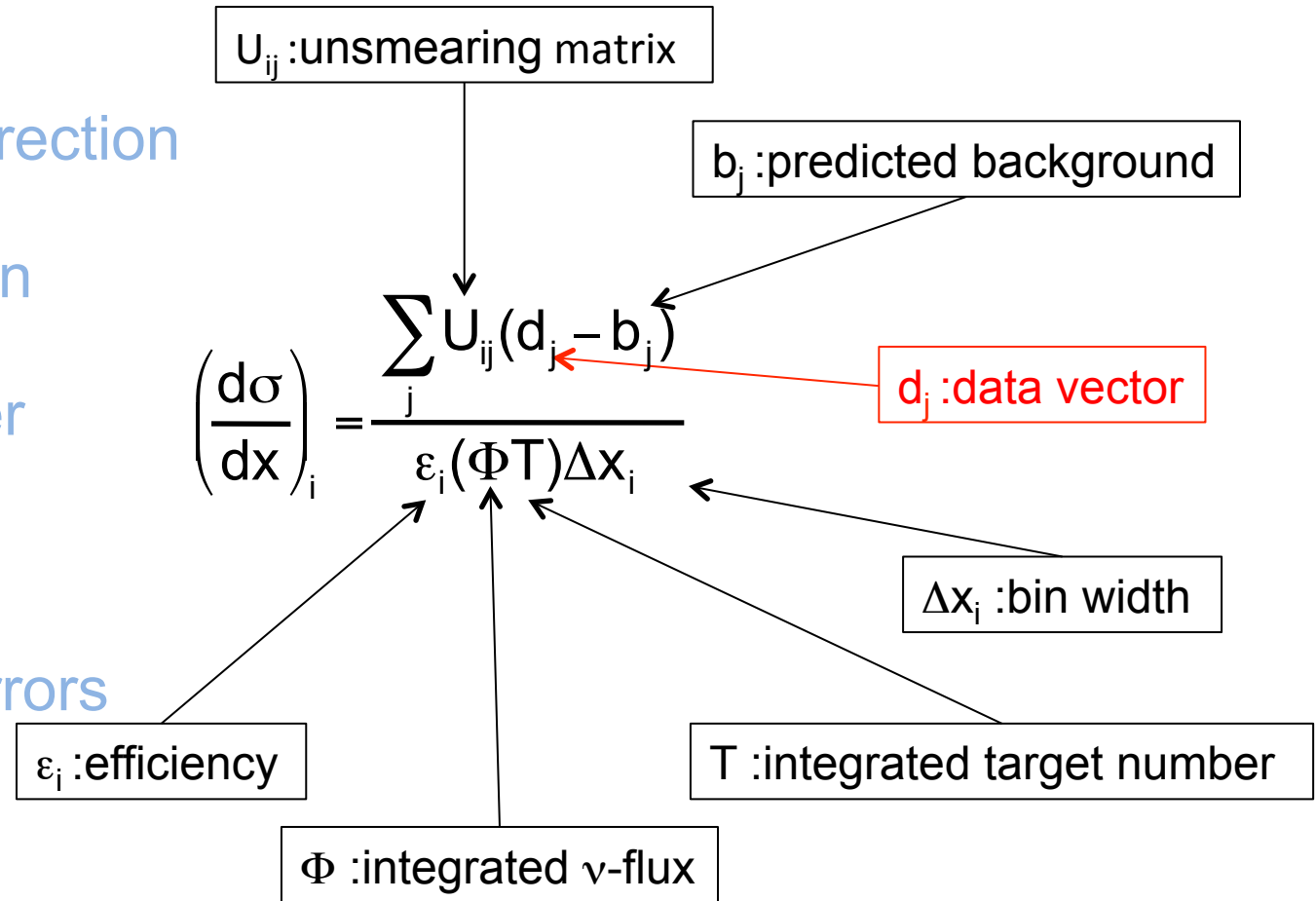
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3.0 Before you start cross section analysis...

Reconstructions and cuts

- You have good reconstruction and all cuts to select your data sample, congratulations, you are ready to measure cross sections!

(you can spend another year in grad school from there!)

3.0 Event selection

Reconstructions and cuts

- You have good reconstruction and all cuts to select your data sample, congratulations, you are ready to measure cross sections!
(you can spend another year in grad school from there!)

Good sample

- depends
- reasonable statistics to make few bins
- measure distributions as many as possible
- Purity $\sim 50\%$ or more (improving purity \rightarrow improving systematics)

Before you start systematics analysis, you should know rough total final error.

- loose/tight cuts
- change bin size
- sideband constraint
- which differential cross section to measure

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- change bin size
- sideband constraint
- which differential cross section to measure

We don't expect more statistics in near future, and most of analyses will be $\sim 10\text{-}20\%$ error. Any cross section data from T2K are unique and precious. **WE SHOULD PUBLISH ASAP!**

3.1 Signal definition

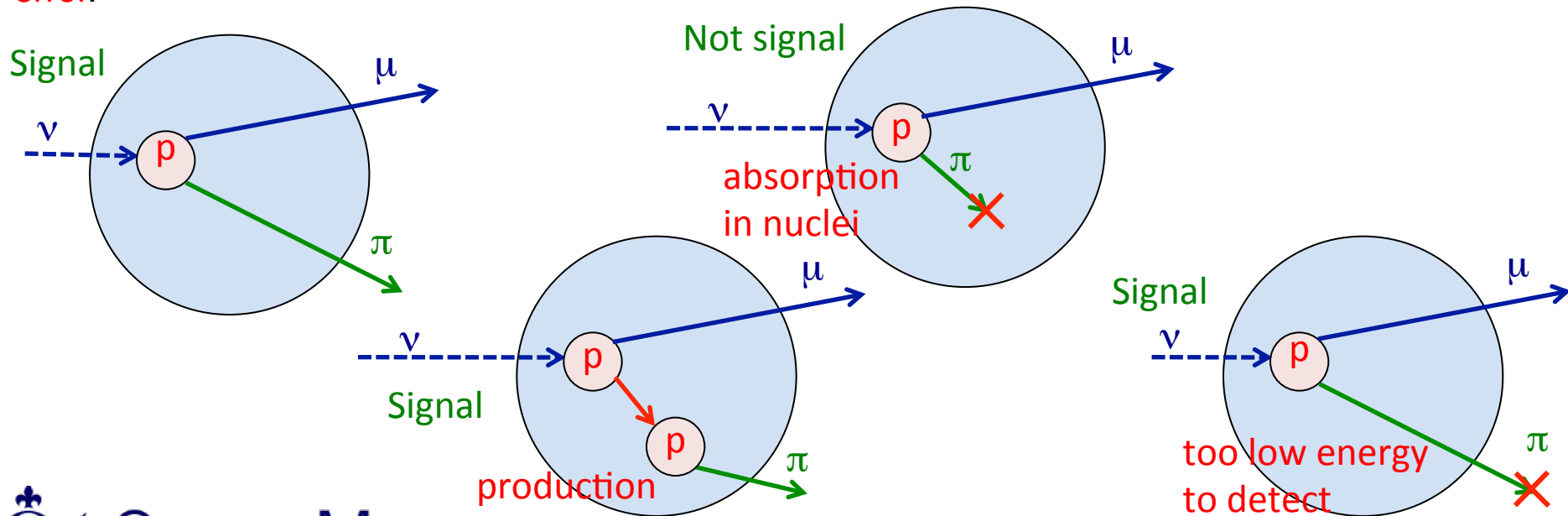
Topological cross section

ex) $CC1\pi^0 = "1 \mu + 1 \pi^0 + N \text{ protons}"$

CC1 π^0

- i. This definition includes π^0 production by final state interactions (FSIs).
- ii. This definition excludes CC1 π^0 interaction when π^0 is lost by FSIs.

This is the necessary definition for the theorists to understand final state interactions (FSIs) without biases. **Don't rely on the definition given by your interaction generator.** "Signal" needs to be added to signal MC, and "Not signal" needs to be removed from signal MC. By this definition, FSI error of pion shouldn't be big, **but detector pion absorption is part of final error.**

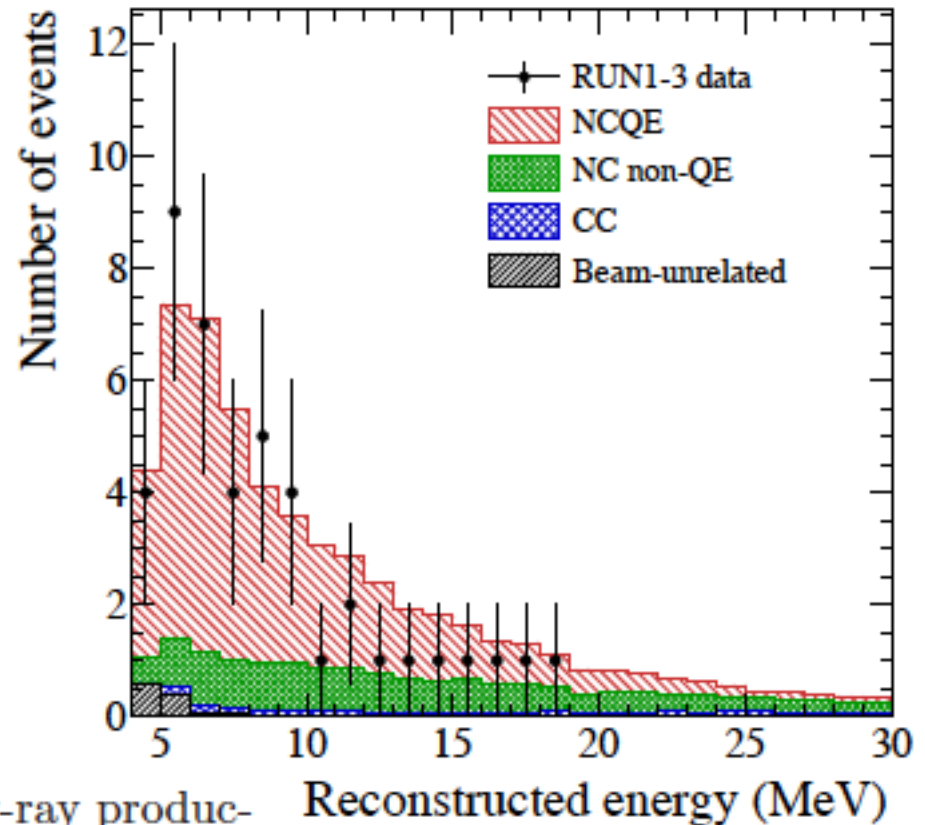


3.1 Signal definition

NCQE gamma measurement

- signal is defined to be gamma from NCQE interactions.
- gamma ray from FSI is not signal.

Alternatively, any gammas from any NC interaction can be defined “signal” (topological cross section), in this way, most of FSI error is gone (smaller systematics), signal statistics is higher, and data is less biased. Drawback is now theorist need to calculate FSI by themselves.



The systematic uncertainty on primary γ -ray production in signal (and the QE component of the CC background) comes from several sources. The largest contribution is from final-state nuclear interactions: NEUT assumes that the de-excitation γ -ray production is the same whether the final state contains a single nucleon or multiple nucleons. We estimate the systematic uncer-

3.1 Signal definition

Topological signal definition

1. Statistics is higher

You have more signal, so statistics is higher

2. Systematic error is lower

Nuclear effect is now signal, not error. Likewise, signal channel model error should be large.

3. Less biased

There is no cross-section model dependent selection nor correction. Data is less biased and preferred by theorists.

Theorists want to find how much MEC from our data using their state-of-the-art nuclear models. We are responsible to provide unbiased data containing these information.

If we use MEC model in our simulation for the selection of events, this mean we try to find how much MEC in our data based on our knowledge. This has 2 bad consequences,

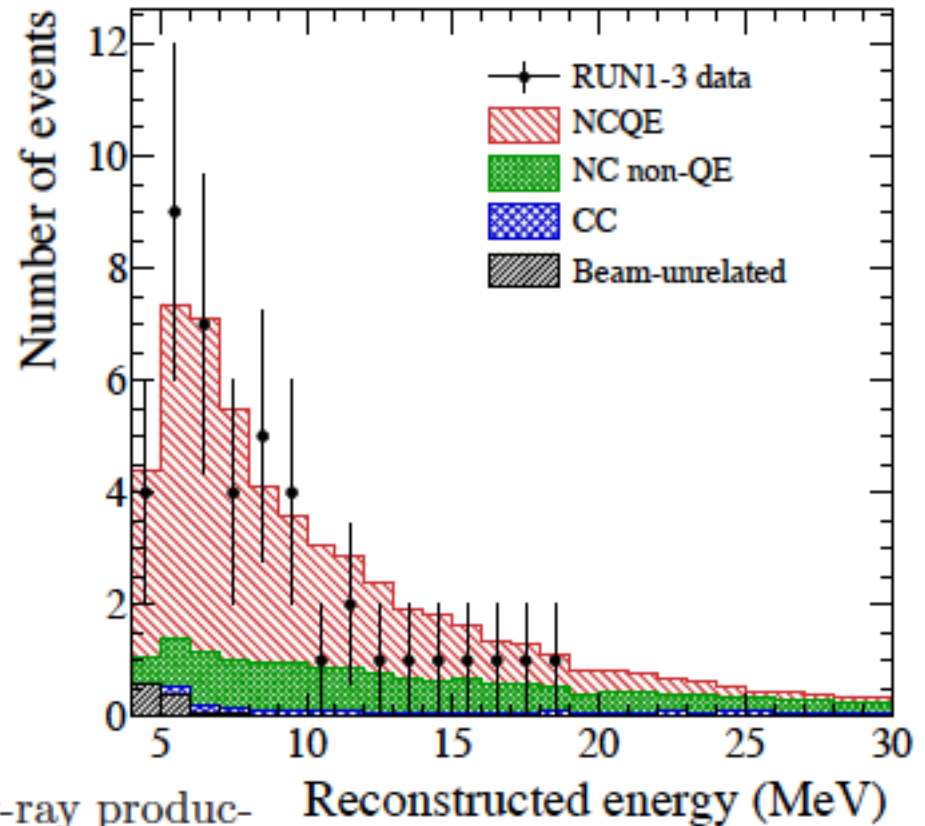
1. theorists lose jobs
2. result is wrong

3.1 Signal definition

NCQE gamma measurement

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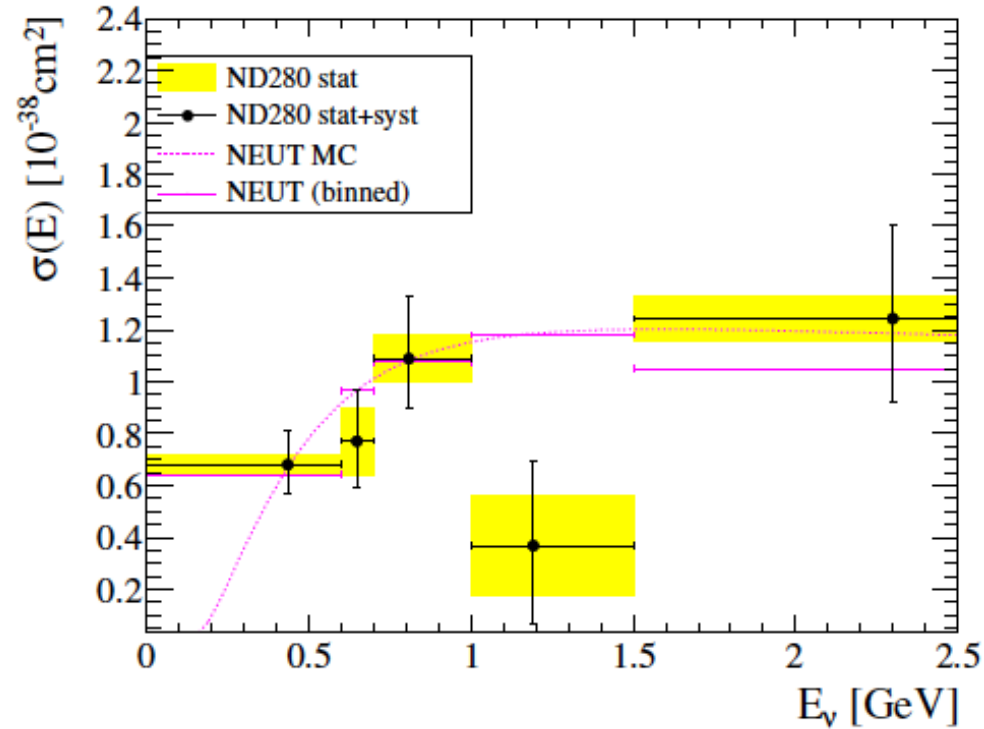
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3.1 Signal definition

Tracker CCQE analysis

- signal is defined from NEUT channel number.

Alternatively, signal can be defined “1 muon-like track”. You can measure total cross section by this way, too. Error should be smaller because any processes making “1 muon-like track” becomes signal. Total cross section has reconstruction bias so it’s still harder for theorists to get info of MEC from this shape of this.



parameter	nominal value	fitted value
CC1 π norm E1	1.63 ± 0.43	$1.34^{+0.34}_{-0.32}$
CC1 π norm E2	1.00 ± 0.40	$1.06^{+0.40}_{-0.40}$
M_A^{QE}	1.21 ± 0.20	$1.38^{+0.16}_{-0.16}$
M_A^{RES}	1.11 ± 0.11	$1.16^{+0.10}_{-0.10}$
CC Other Shape	0.00 ± 0.40	$0.02^{+0.40}_{-0.40}$
NC1 π^0 norm	1.19 ± 0.43	$1.03^{+0.40}_{-0.40}$
p_F	217.00 ± 30.38	$234.90^{+14.87}_{-15.05}$

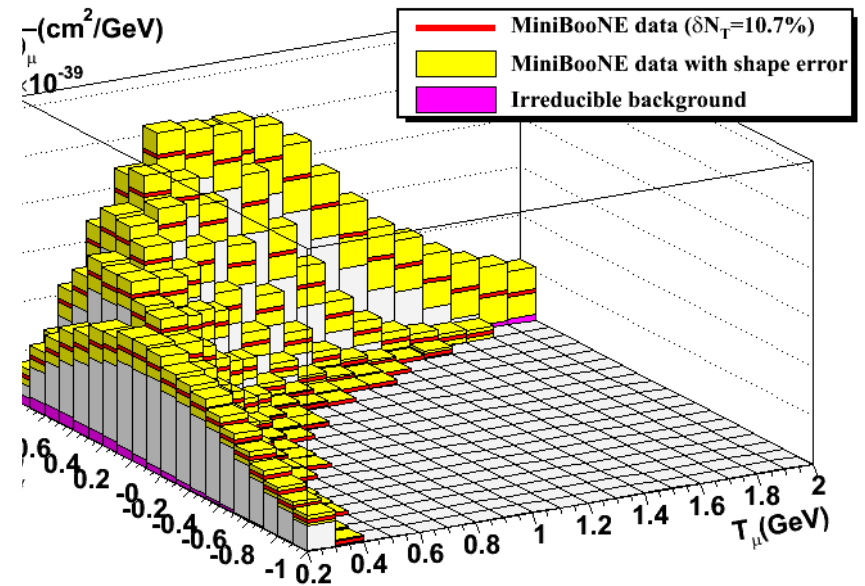
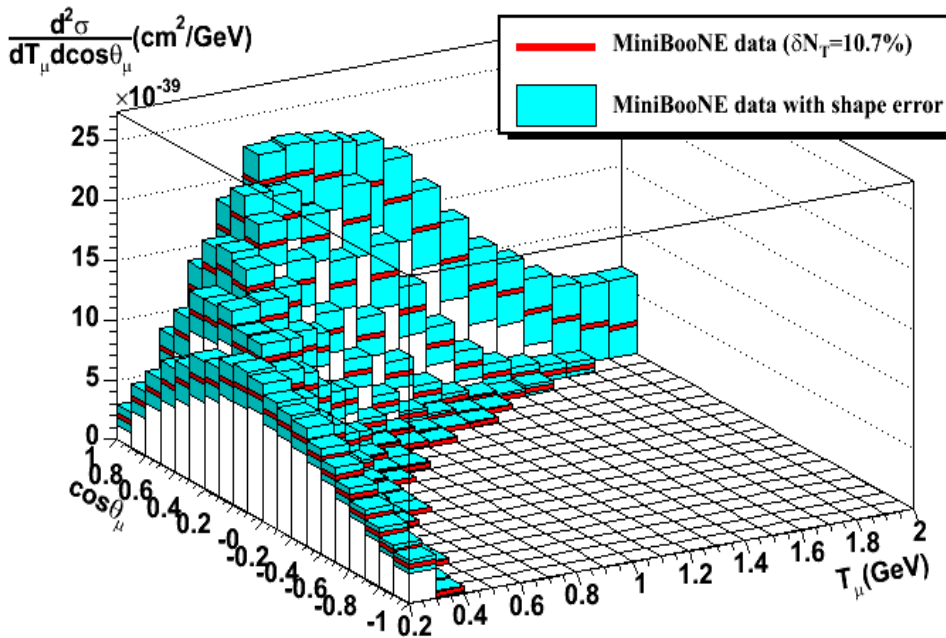
$$\begin{aligned}
 -2\ln\lambda(\theta) = & 2 \sum_{i=1}^{p_\mu - \cos(\theta_\mu) \text{ bins}} \left[N_i^{\text{predicted}}(\theta) - N_i^{\text{observed}} \right. \\
 & \left. + N_i^{\text{observed}} \ln \frac{N_i^{\text{observed}}}{N_i^{\text{predicted}}(\theta)} \right] \\
 & + \ln \frac{\pi_d(d)}{\pi_d(d_{\text{nominal}})} \\
 & + \ln \frac{\pi_f(f)}{\pi_f(f_{\text{nominal}})} \\
 & + \ln \frac{\pi_x(x)}{\pi_x(x_{\text{nominal}})}
 \end{aligned}$$

3.1 Signal definition

CCQE

Primary channel cross section

By subtracting CC1 π^+ without pion background distribution (from MC with sideband correction), “pure” CCQE cross section is published (which is not, due to 2p-2h). Pure channel cross section is helpful for some theorists who cannot simulate whole set of cross sections.



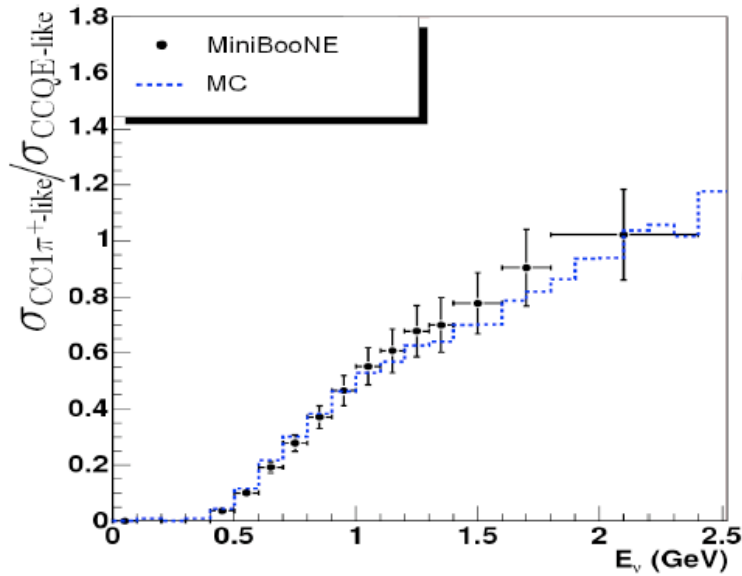
3.1 Signal definition

CC1 π^+ /CCQE

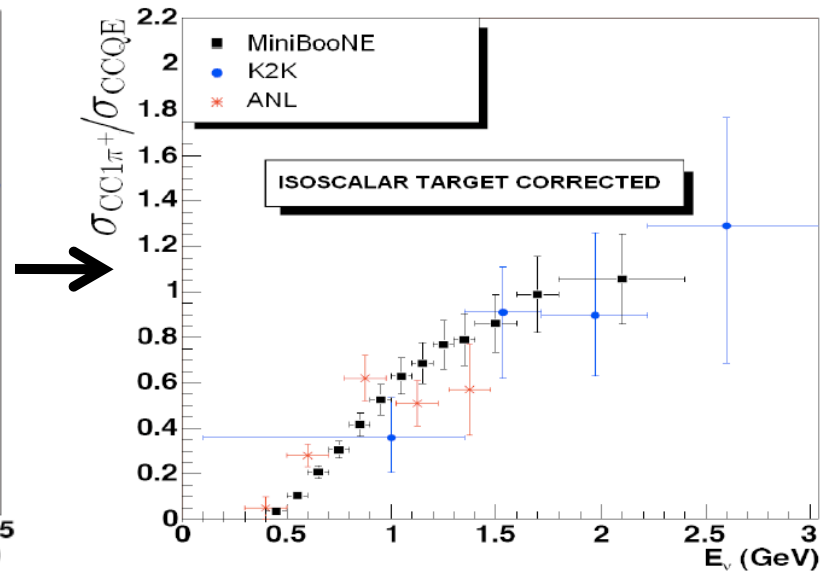
Primary channel cross section

Pure channel cross section is also useful to compare with other data (especially most of old data are published in pure channel cross section).

CC1 π^+ /CCQE topological cross section ratio



CC1 π^+ /CCQE primary channel cross section ratio



3.1 Signal definition

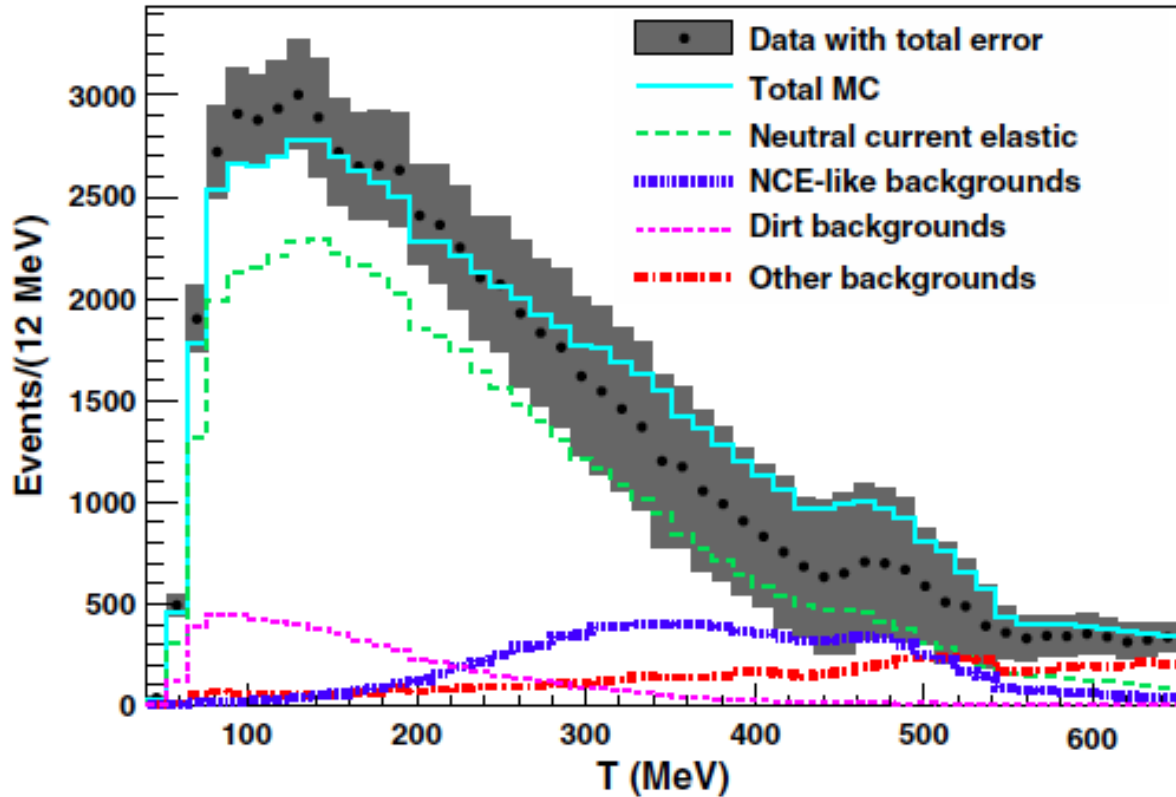
NCEL

Mixed target

Cross section result from mixed target is complicated. In NCEL, differential cross section is interpreted $\nu+N$, which means sum of $\nu+p$ (carbon), $\nu+p$ (hydrogen), and $\nu+n$ (carbon).

Forward folding

Migration matrices (T_p^{true} and T_p^{recon}) are provided for $\nu+p$ (carbon), $\nu+p$ (hydrogen), and $\nu+n$ (carbon), with and without FSI, so that theorists can fold their xsec models to compare with measured T_p .



3.1 Signal definition

Mixed target

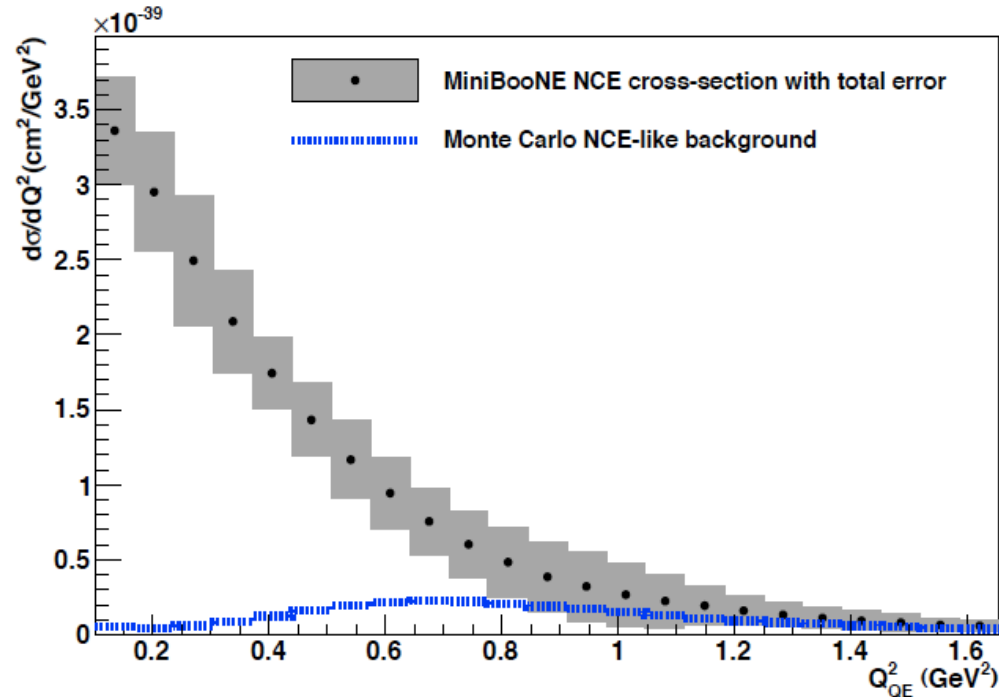
Cross section result from mixed target is complicated. In NCEL, differential cross section is interpreted $\nu+N$, which means sum of $\nu+p$ (carbon), $\nu+p$ (hydrogen), and $\nu+n$ (carbon).

NCEL

Efficiency difference

The efficiency difference of each interaction is provided to reproduce differential cross section from microscopic modes.

$$\begin{aligned} \frac{d\sigma_{\nu N \rightarrow \nu N}}{dQ^2} &= \frac{1}{7} C_{\nu p, H}(Q_{QE}^2) \frac{d\sigma_{\nu p \rightarrow \nu p, H}}{dQ^2} + \frac{3}{7} C_{\nu p, C}(Q_{QE}^2) \\ &\times \frac{d\sigma_{\nu p \rightarrow \nu p, C}}{dQ^2} + \frac{3}{7} C_{\nu n, C}(Q_{QE}^2) \frac{d\sigma_{\nu n \rightarrow \nu n, C}}{dQ^2}, \end{aligned} \tag{B6}$$



3.1 Measured variables

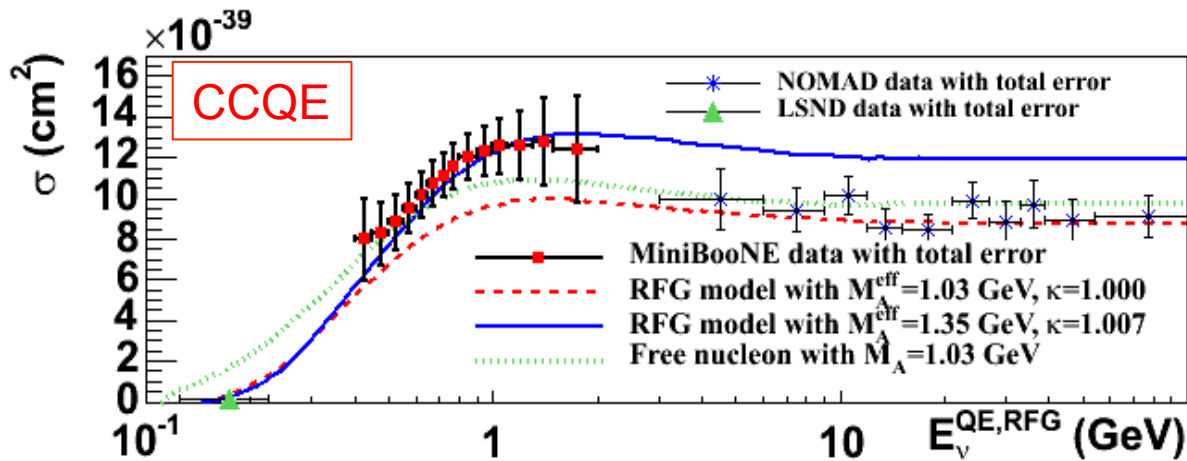
CCQE NCEL NC1 π^0 CC1 π^+ CC1 π^0 anti CCQE anti NCEL

Function of measured variables

Differential cross section results with function of measured variables (momentum, direction, etc) has **no reconstruction bias** and preferred.

Function of reconstructed variables

Cross section results of reconstructed variables (especially Q^2 and E_ν) are **model-dependent**. However, flux-unfolded total cross section function (function of true neutrino energy) is the only way to compare results with other experiments.



3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

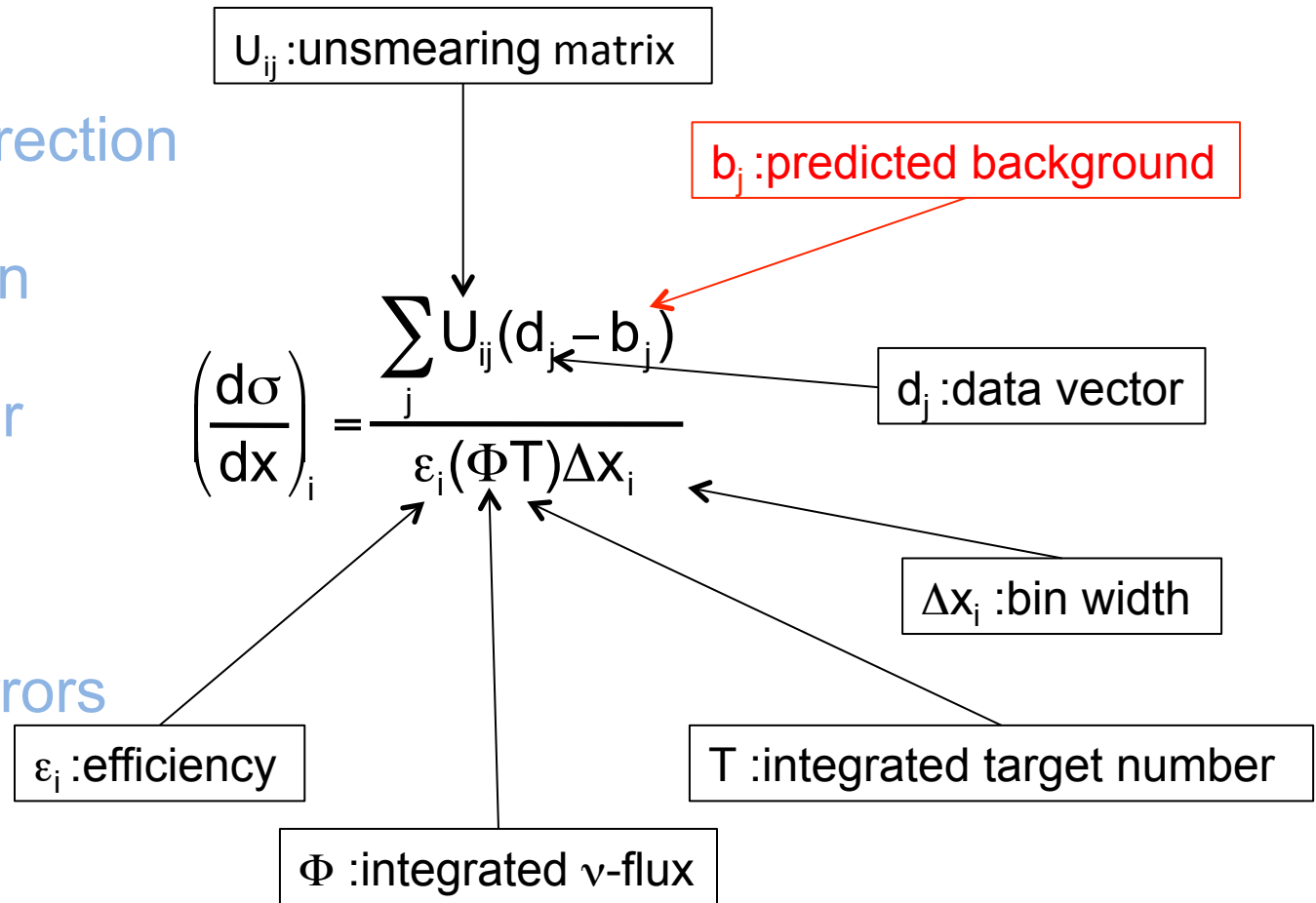
3.5 Flux correction

3.6 Target number

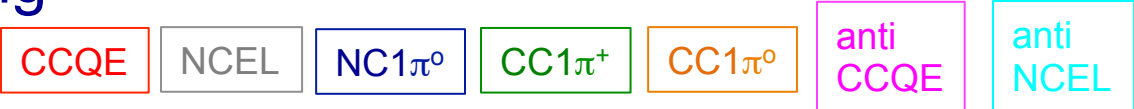
3.7 Binning

3.8 Systematic errors

3.9 Data format



3.2 Background removing



Background subtraction method

- It is preferred because **signal doesn't depend on signal MC explicitly**.
- Normalization of background must be known.

Background subtraction

$$d_i - b_i$$

Purity correction

$$d_i \times \frac{s_i}{s_i + b_i}$$

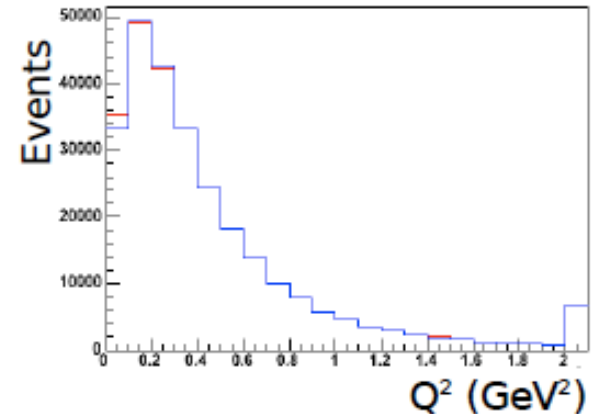
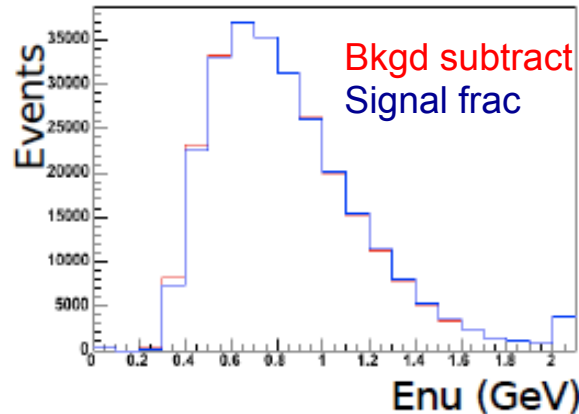
Purity correction method

- **Signal explicitly depends on signal MC** (=bad), potentially shape is distorted by signal to be measured (depending on size of error).
- However, if you don't know the normalization of background, this may be justified?

Background subtraction vs Purity correction

CCQE

For high purity CCQE sample (77%), difference is only large at low Q^2 , where background is ~30%



3.2 Background removing

CCQE
NCEL
NC1π⁰
CC1π⁺
CC1π⁰
anti CCQE
anti NCEL

Background subtraction method

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- Normalization of background must be known.

Background subtraction

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Purity correction

$$d_i \times \frac{s_i}{s_i + b_i}$$

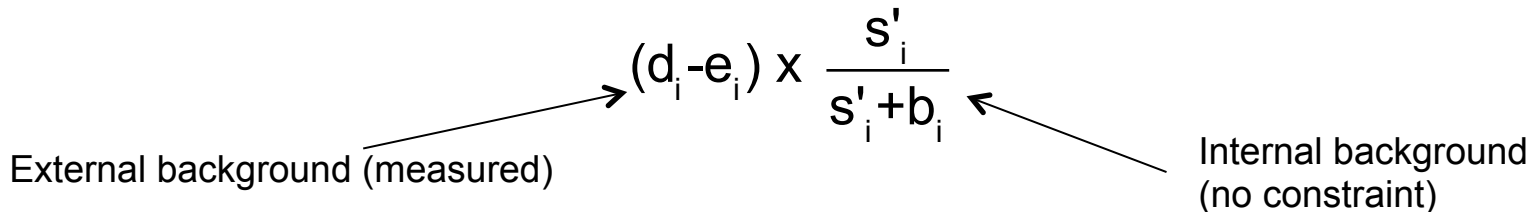
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NCEL
anti NCEL

Hybrid

- External background (=cosmic rays) are measured from sideband (=known normalization), and subtracted.
- Internal background (=NC1π⁺ where pion is below detection threshold) is removed by purity correction.



3.2 Background removing

CCQE NCEL CC1 π^+ CC1 π^0 anti CCQE anti NCEL

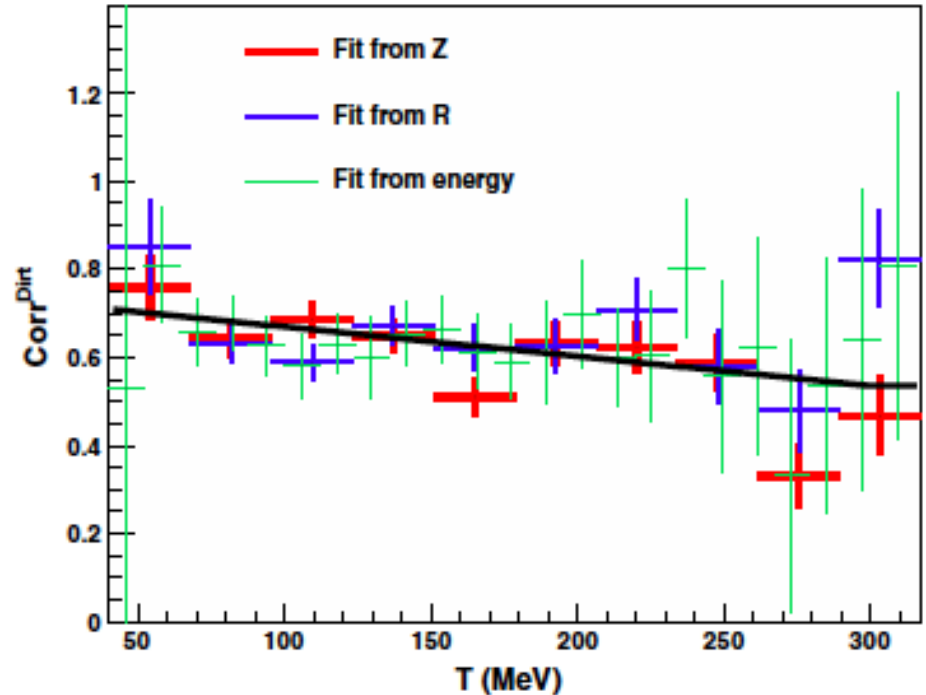
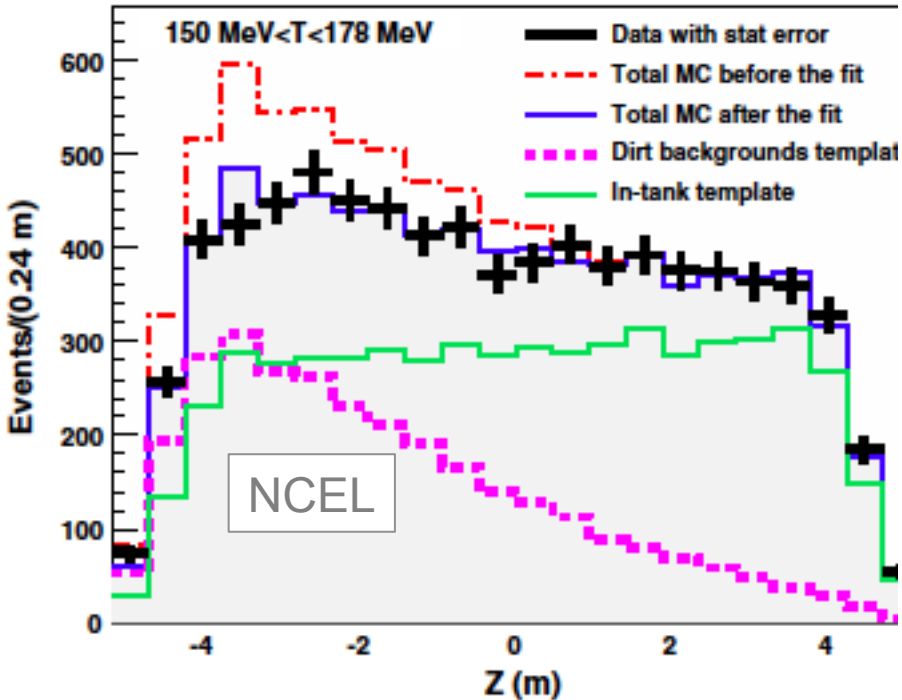
Data driven correction

- Almost all analyses use sideband data to correct background distribution in signal boxes

External background measurement NCEL

- NCEL analysis measure amount of background coming from outside, with function of R and Z, to extrapolate background in fiducial volume

External background enhanced sample



3.2 Background removing

CCQE

NCEL

CC1 π^+

CC1 π^0

anti
CCQE

anti
NCEL

Data driven correction

- Almost all analyses use sideband data to correct background distribution in signal boxes

Flux error double counting?

- For background dominant sample, background subtraction makes flux error larger.
- Way to avoid is to define background excursion after removing normalization correlated with signal (shape-only background subtraction)

$$\left(\frac{d\sigma}{dx}\right)_i = \frac{\sum_j U_{ij}(d_j - b_j)}{\varepsilon_i(\Phi T)\Delta x_i}$$

3.2 Background removing

CCQE

NCEL

CC1 π^+

CC1 π^0

anti
CCQE

anti
NCEL

Data driven correction

- Almost all analyses use sideband data to correct background distribution in signal boxes

Cross section error

- Main xsec error is the error assigned on background models
- Xsec models, such as FSIs?, change true-recon relationship for hadrons (smearing)
- Xsec models kinematics, i.e., cuts (=efficiency)

$$\left(\frac{d\sigma}{dx}\right)_i = \frac{\sum_j U_{ij}(d_j - b_j)}{\varepsilon_i(\Phi T)\Delta x_i}$$

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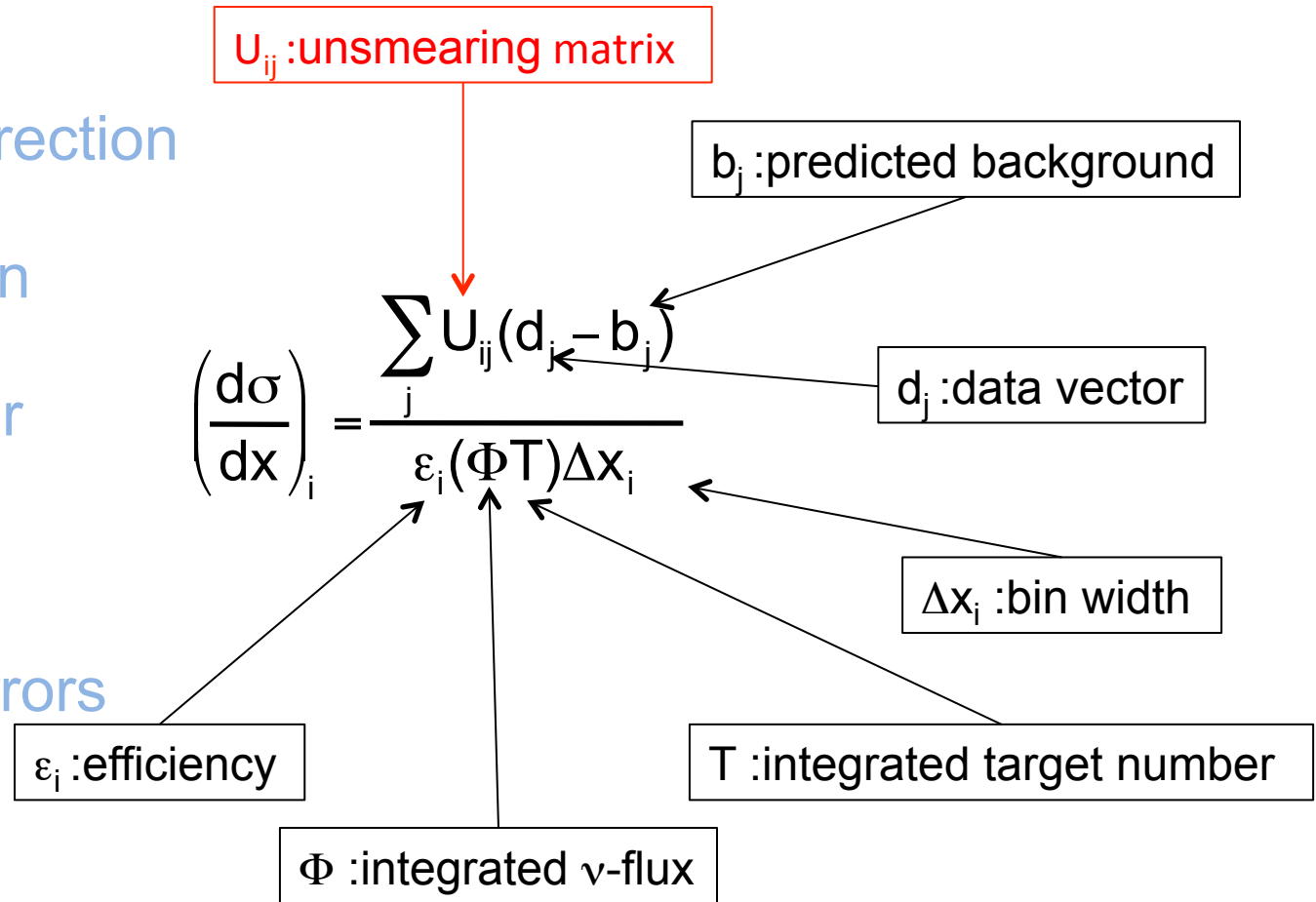
3.5 Flux correction

3.6 Target number

3.7 Binning

3.8 Systematic errors

3.9 Data format



3.3 Unsmearing

Unfolding

- The process removing the detector effects, mainly smearing and detector cut, is called **unfolding**. It is often easier to think by separating unfolding process to 2 parts, **unsmearing** and **efficiency correction**. We focus on unsmearing here.

Detector error

- Detector model affect smearing.

$$\left(\frac{d\sigma}{dx}\right)_i = \frac{\sum_j U_{ij}(d_j - b_j)}{\varepsilon_i(\Phi T)\Delta x_i}$$

3.3 Unsmearing

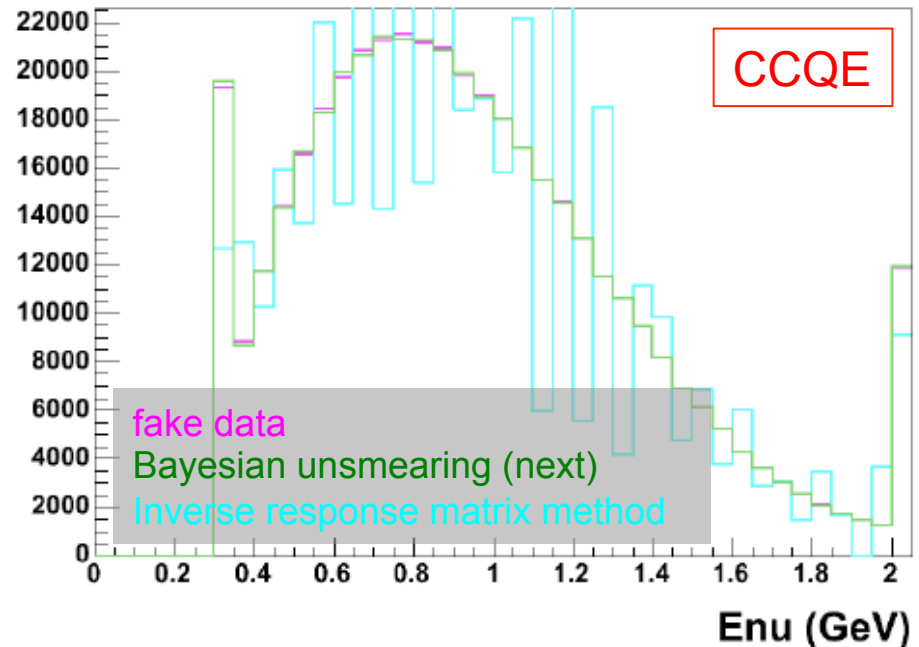
Inverse response matrix method

None

- Inverse response matrix method is the bias-free unfolding method, but this method doesn't work for anybody. Typically, it makes rapid oscillated solution (Gibb's phenomenon). Say, response matrix R gives the smearing and detector cut of true distribution a to measured distribution b in MC, it's inverse can be used to unfold data b to true distribution a

$$\beta_i = R_{ij}\alpha_j \rightarrow a_j = (R)_{ji}^{-1} b_i$$

Inverse response matrix method is very sensitive with MC statistics. It doesn't work for sparse matrix, it cannot handle large number of bins, it cannot deal histogram with zero-event bins. But all these are features for differential cross section!



3.3 Unsmearing

Tikhonov regularization method NC1 π^0

- The regularization term from the prior knowledge of distribution (e.g., how smooth is) can stabilize inverse response matrix. **The bias is introduced through the linear operator L and τ .**

$$\beta_i = R_{ij} \alpha_j \rightarrow a_j = (R)_{ji}^{-1} b_i \rightarrow (Ra - b)^T V(b)^{-1} (Ra - b) + \tau(La)^T (La) \sim 0$$

Regularization parameter τ should be chosen with care.

- too small τ doesn't regulate matrix inversion
- too large τ too much smooth out response matrix R

Solution is,

$$a = U' \cdot b + \left[\sum_i [(I - U') \cdot b]_i \right] \cdot s$$

$$U' = [R + \tau V \cdot R^{T-1} (L^T L)]$$

$$s_i = \frac{\sum_j [U' V R^{-1}]_{ij}}{\sum_{jk} [U' V R^{-1}]_{jk}}$$

3.3 Unsmearing

CCQE

NCEL

NC1 π^0 CC1 π^+ CC1 π^0 anti
CCQEanti
NCEL

Iterative Bayesian method

- Unsmearing is based on the Bayesian statistics, so **bias is introduced from MC knowledge**

- Efficiency ε is defined by true distribution after cut μ to true distribution before cut α .

M-matrix gives transformation from measured distribution to true distribution after cut. It give the true distribution after cut μ on projection on one axis.

$$\varepsilon_i = \frac{\mu_i}{\alpha_i} \quad \sum_{j=1}^n M_{ij} = \mu_i = \varepsilon_i \alpha_i$$

Now, define U-matrix by normalizing M-matrix with other axis,

$$U_{ij}^{0th} = \frac{M_{ij}}{\sum_{k=1}^n M_{kj}}$$

So, background subtracted data d^{0th} can be unsmearred and efficiency corrected to obtain unfolded cross section d^{1th}

$$d_i^{1st} = \frac{1}{\varepsilon_i} \sum_{j=1}^n U_{ij}^{0th} d_j^{0th}$$

3.3 Unsmearing

CCQE

NCEL

NC1 π^0 CC1 π^+ CC1 π^0 anti
CCQEanti
NCEL

Iterative Bayesian method

- Unsmearing is based on the Bayesian statistics, so **bias is introduced from MC knowledge**
- It is based on Bayes' theorem

$$M_{ij} = P(\text{recon}_j | \text{true}_i)P(\text{true}_i)$$

$$P(\text{true}_i | \text{recon}_j) = \frac{P(\text{recon}_j | \text{true}_i)P(\text{true}_i)}{\sum_k P(\text{recon}_j | \text{true}_k)P(\text{true}_k)} = \frac{M_{ij}}{\sum_k M_{kj}} = U_{ij}^{\text{0th}}$$

$$d_i^{\text{1st}} = \frac{1}{\varepsilon_i} \sum_{j=1}^n U_{ij}^{\text{0th}} d_j^{\text{0th}}$$

3.3 Unsmearing

CCQE

NCEL

NC1 π^0 CC1 π^+ CC1 π^0 anti
CCQEanti
NCEL

Iterative Bayesian method

- Unsmearing is based on the Bayesian statistics, so **bias is introduced from MC knowledge**

If initial guess μ (=prior probability of Bayesian statistics) is not so close to nature, we can improve U-matrix **by assuming $d^{1\text{th}}$ is close to nature**

$$\omega_i^{1\text{st}} = \frac{d_i^{1\text{st}}}{\alpha_i} \quad U_{ij}^{1\text{st}} = \frac{\omega_i^{1\text{st}} M_{ij}}{\sum_{k=1}^n (\omega_k^{1\text{st}} M_{kj})} \quad d_i^{2\text{nd}} = \frac{1}{\varepsilon_i} \sum_{j=1}^n U_{ij}^{1\text{st}} d_j^{0\text{th}}$$

- This iteration process usually converge <5 times. 0th iteration is not bad at all.
- Signal model dependence will become systematic error, this is done by varying M-matrix by changing systematics. So **signal cross error is part of final error**, but shape only.
- This method also fails if M-matrix is highly non-diagonal.
- 0th and 1st iteration difference of data is also included as systematic error (?).

Iterative Bayesian method works for, any number of bins, including zeros, sparse matrix, MxN matrix, background non-subtracted sample etc...

3.3 Unsmearing

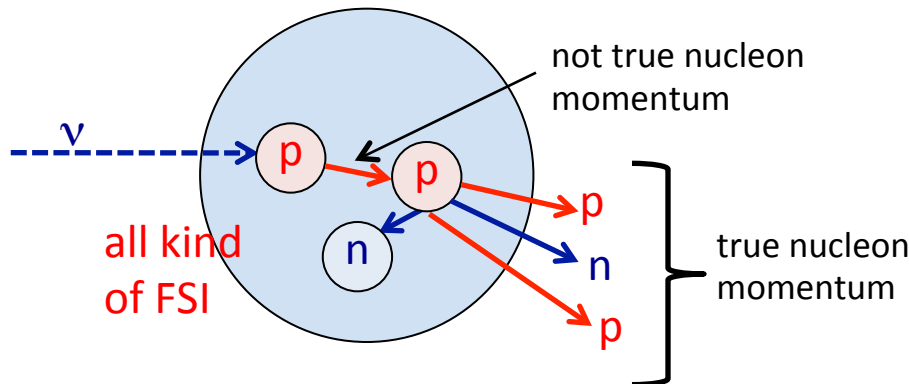
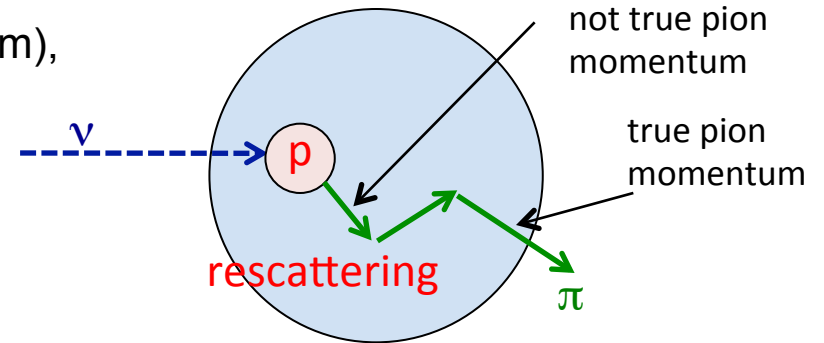
How to construct M-matrix? Measured variables

- It is desired to present differential cross section with function of measured quantities, such as muon energy, pion angle, etc, because **they are not biased by reconstruction.**

i. It is straightforward if you measure lepton kinematics.
True lepton kinematics are the true information for M-matrix.



ii. If you measure hadronic events (e.g., π^0 momentum), your “true” kinematics is after FSI, i.e., particle exiting the nuclei.



iii. “true” momentum is defined by sum of all outgoing nucleons, because that is the observables.

3.3 Unsmearing

How to construct M-matrix? Reconstructed variables

- The definition of true kinematics is tricky, because you have choices.

i. True Q^2 is defined by reconstructed Q^2 from true kinematics

CCQE
NCEL
anti CCQE
anti NCEL

For example, CCQE, true Q^2 is defined “reconstructed Q^2 from true muon energy and angle”, and we call it “ Q^2_{QE} ” to remind people this is reconstructed under QE assumption.

ii True Q^2 is defined by true Q^2 in MC

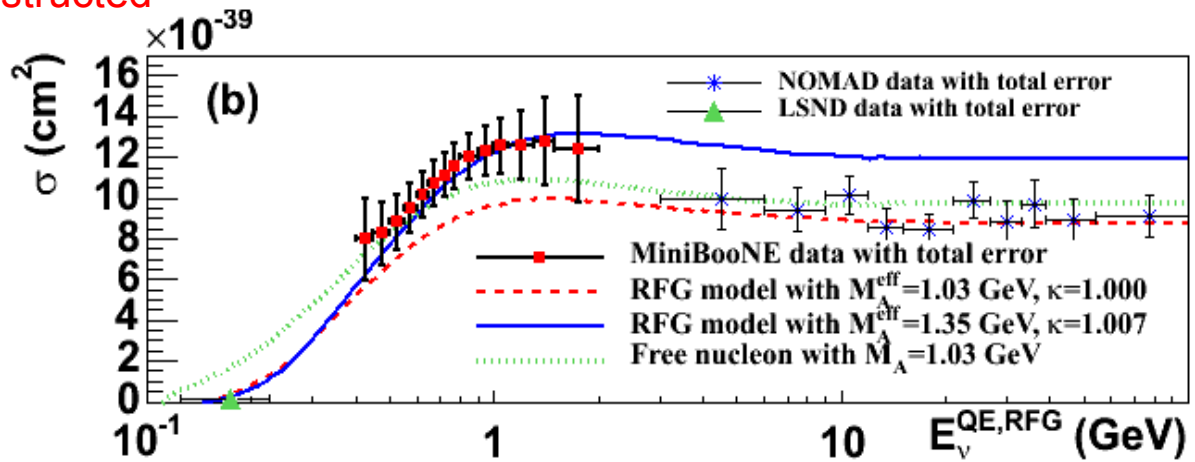
CC1 π^+
CC1 π^0

This may be useful to compare with old data, only presented by this way

iii True E_ν is defined by true E_ν in MC

CCQE
NC1 π^0
CC1 π^+
CC1 π^0
anti CCQE

For example, CCQE, E_ν is called “ $E_{\nu, QE, RFG}$ ” to remind people **this is reconstructed under QE assumption then unfolded by assuming RFG model.**



3.3 Unsmearing

No unsmearing $\text{NC1}\pi^0$

- If you know smearing is weak and statistics is low, no unsmearing may be the best option.

So what is the criteria?

3.3 Unsmearing

No unsmearing NC1 π^0

- If you know smearing is weak and statistics is low, no unsmearing may be the best option.

So what is the criteria?

Bias of unsmearing

- There is no perfect unfolding, unfolding method can be different depending on your distribution. **Biases** may be one of criteria.

i. Inverse response matrix method:

No bias, but it only works for very few bin histogram

$$\mathbf{B}^{\text{Inverse}} = 0$$

ii. Tikhonov regularization

Bias is introduced from linear function and regularization parameter. It also requires a fair amount of events.

$$\mathbf{B}^{\text{Tikhonov}} = \sum_i [\mathbf{U}' \cdot (\mathbf{R}\mathbf{a} - \mathbf{b})]_i$$

iii. Iterative Bayesian method

Bias is introduced from prior knowledge of MC. It works for any distribution.

$$\mathbf{B}^{\text{Bayesian}} = \sum_i [\mathbf{M} \cdot (\mathbf{R}\mathbf{a} - \mathbf{b})]_i$$

iv. No unsmearing

$$\mathbf{B}^{\text{NoUnsmear}} = \sum_i [(\mathbf{U} - \mathbf{I}) \cdot \mathbf{b}]_i$$

3.3 Unsmearing

π^0 kinematics

NC1 π^0

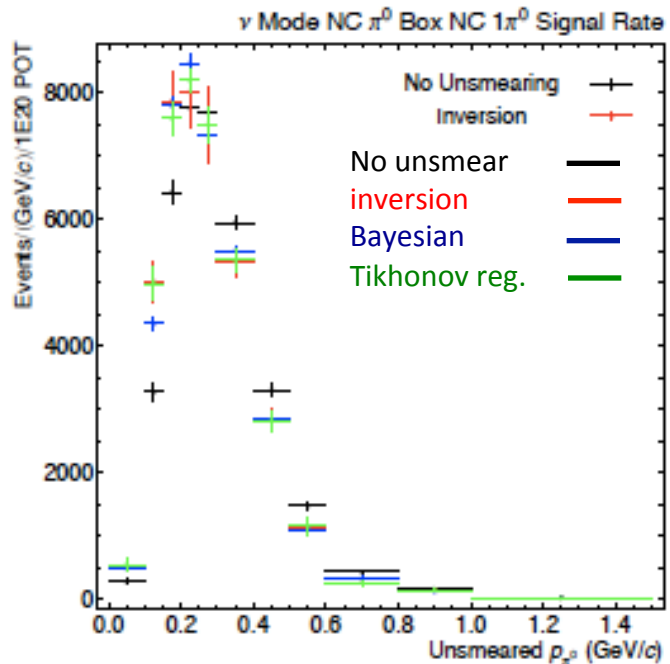
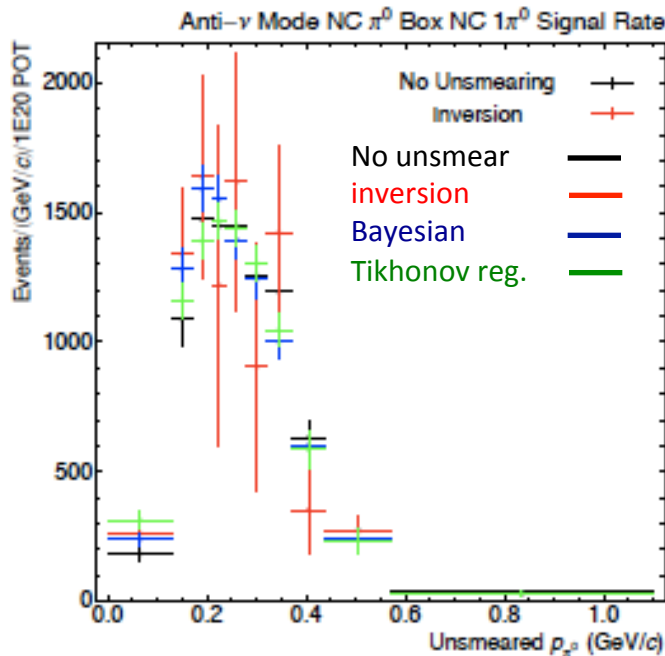
- Comparing biases and histograms by eyes, π^0 -kinematics are unfolded by 3 different methods.

$P_\pi(\nu)$: Tikhonov regularization

$\cos\theta_\pi(\nu)$: Iterative Bayesian

$P_\rho(\text{anti-}\nu)$: Iterative Bayesian

$\cos\theta_\pi(\text{anti-}\nu)$: No unsmearing



3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

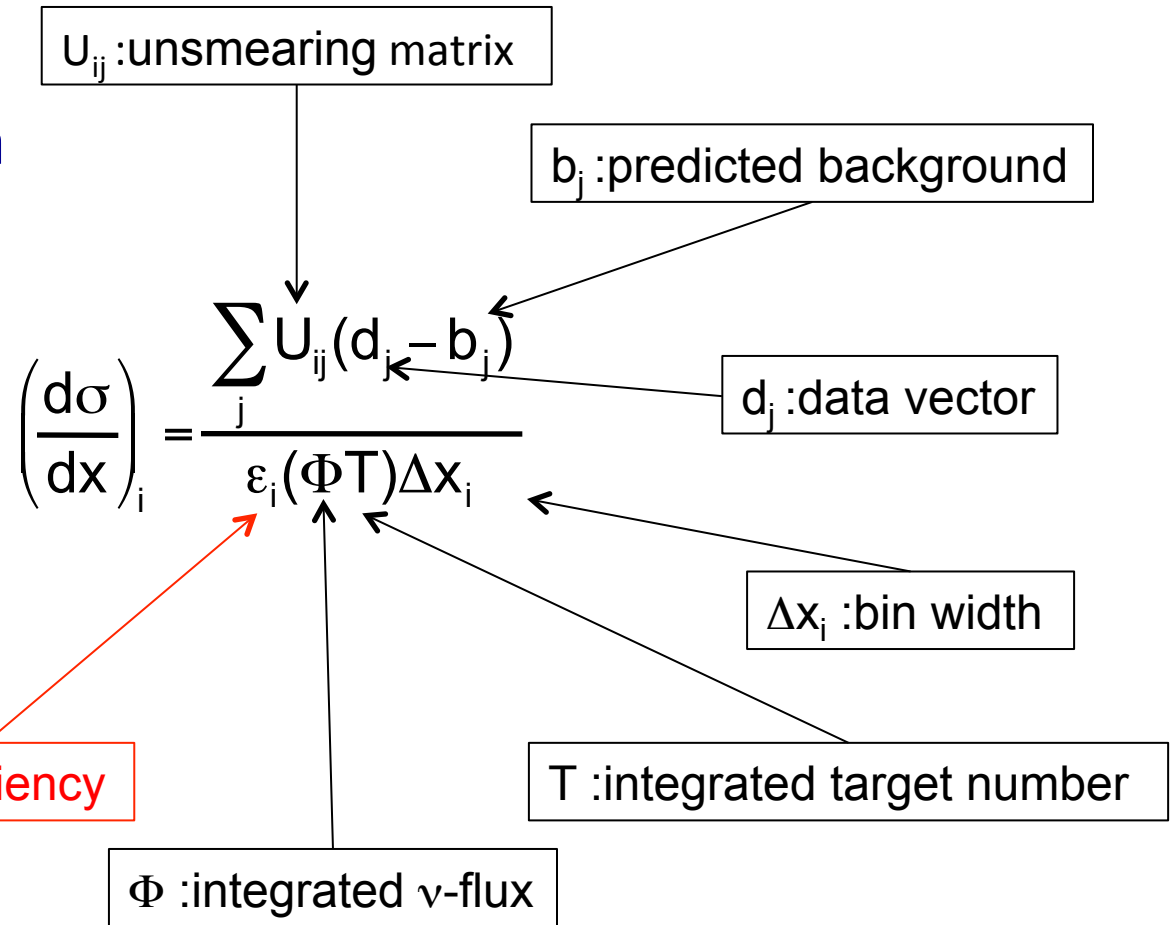
3.5 Flux correction

3.6 Target number

3.7 Binning

3.8 Systematic errors

3.9 Data format



3.4 Efficiency correction

CCQE	NCEL	NC1 π^0	CC1 π^+	CC1 π^0	anti CCQE	anti NCEL
------	------	-------------	-------------	-------------	-----------	-----------

Looks straightforward, no?

- The efficiency is defined as true distribution after cut μ divided by that before cut α .
Because of the nature of ratio, ϵ is insensitive with many systematic variations common for numerator and denominator such as flux error and cross section error. The **detector error** is important.

$$\left(\frac{d\sigma}{dx}\right)_i = \frac{\sum_j U_{ij}(d_j - b_j)}{\epsilon_i (\Phi T) \Delta x_i}$$

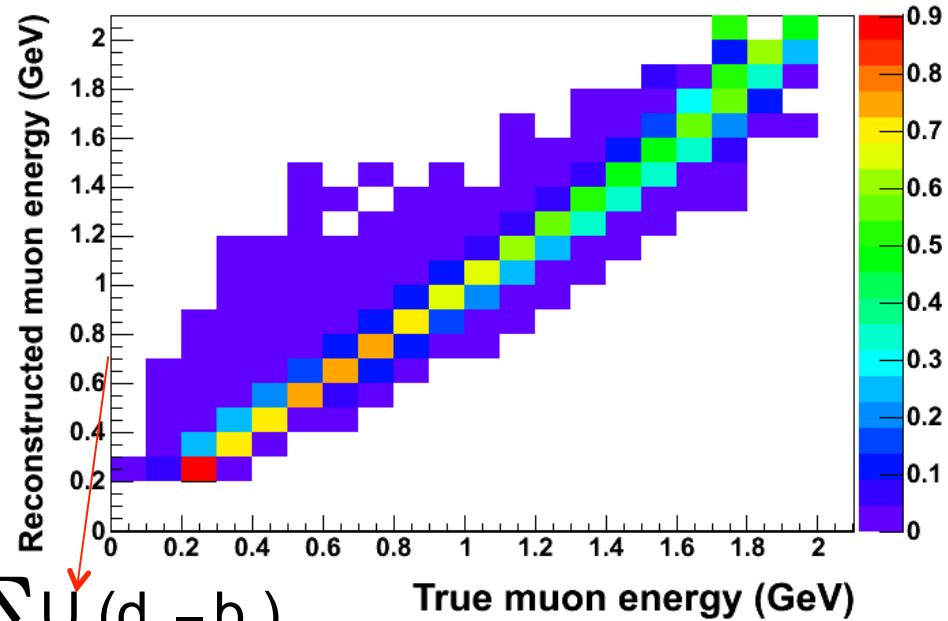
$$\epsilon_i = \frac{\mu_i}{\alpha_i} = \frac{N_i(\text{AfterCut})}{N_i(\text{BeforeCut})}$$

3.4 Efficiency correction

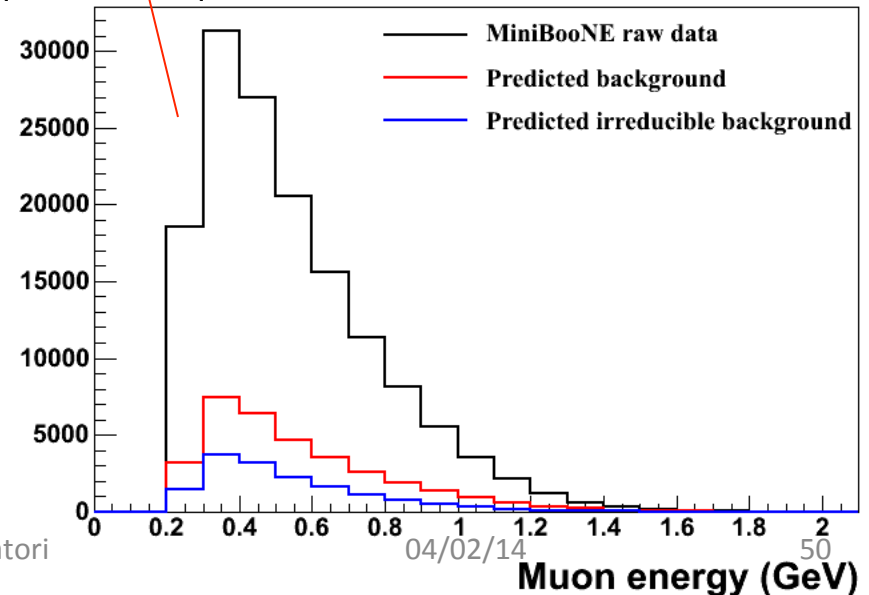
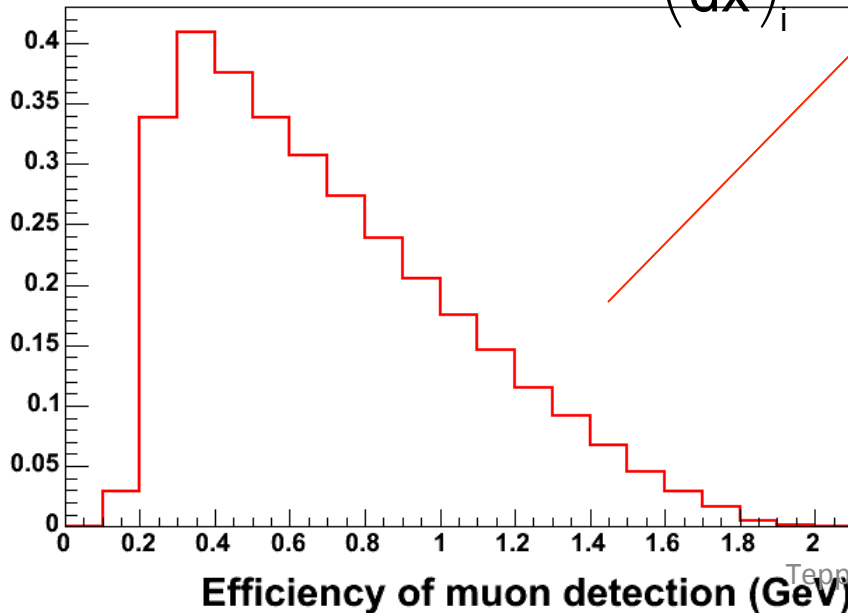
Muon energy unfolding

CCQE

- Because of the resolution of muon energy, it is possible to recover events outside of kinematic cut by unfolding process (detector model dependent)



$$\left(\frac{d\sigma}{dx}\right)_i = \frac{\sum_j U_{ij}(d_j - b_j)}{\epsilon_i(\Phi T)\Delta x_i}$$

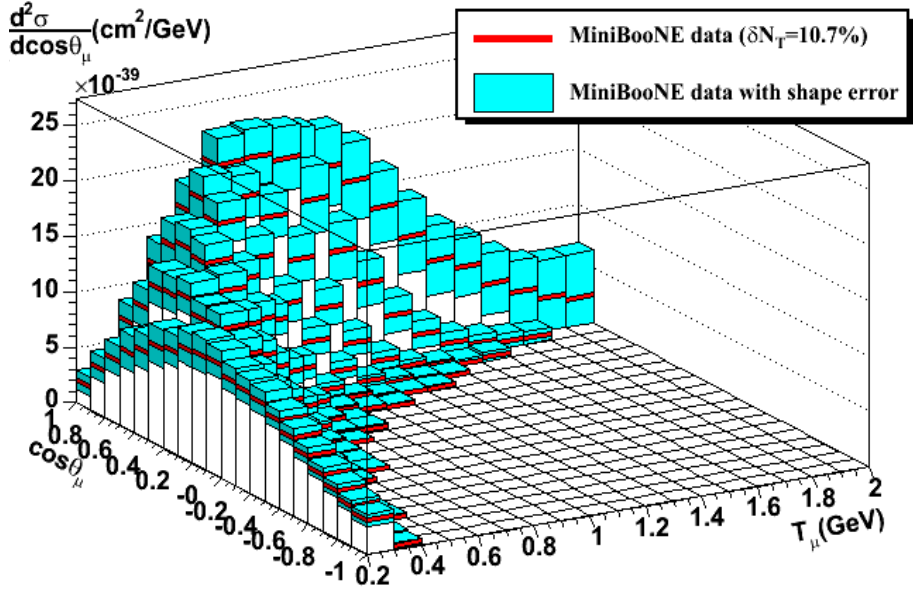
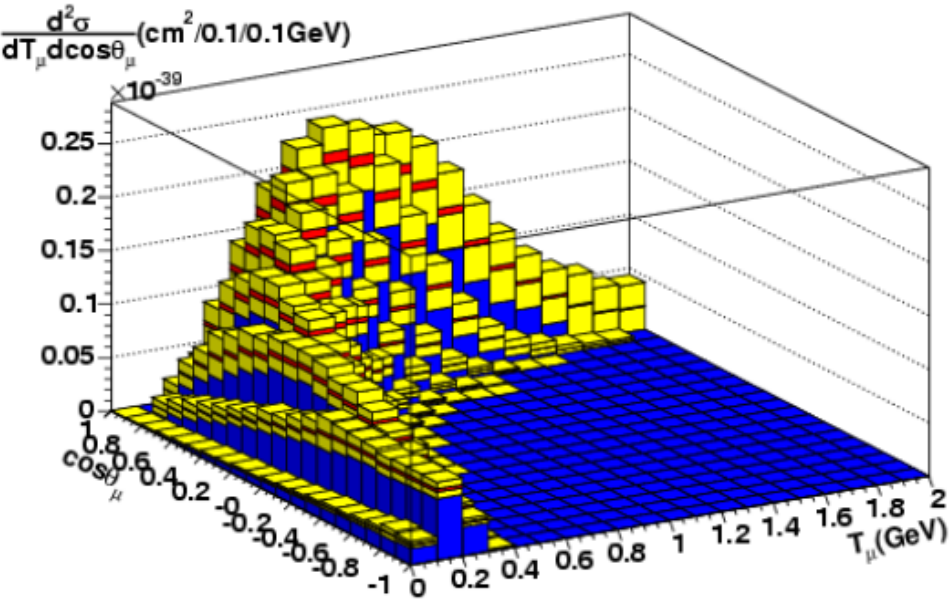


3.4 Efficiency correction

CCQE

Muon energy unfolding

- Because of the resolution of muon energy, it is possible to recover events outside of kinematic cut by unfolding process (detector model dependent)



3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

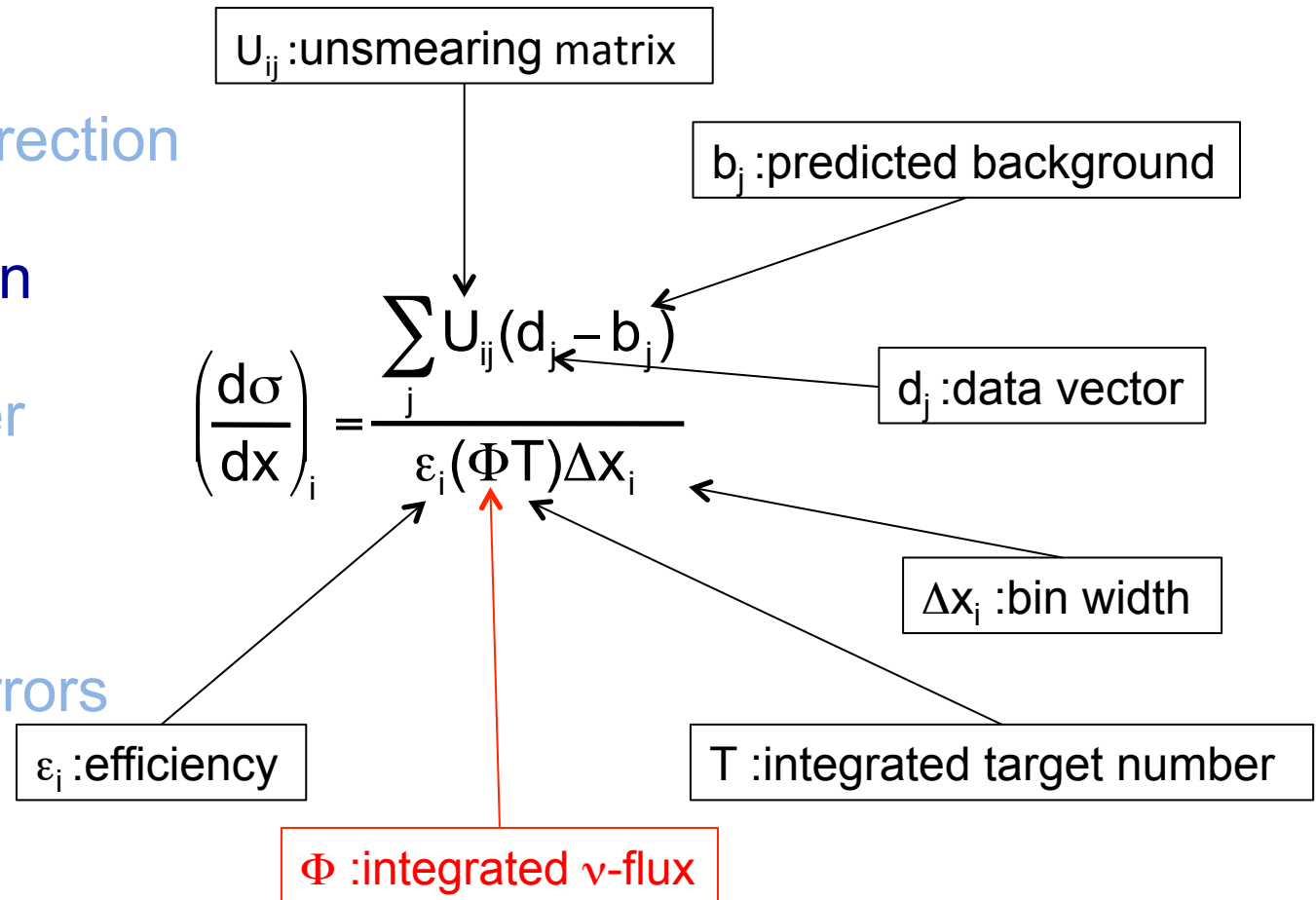
3.5 Flux correction

3.6 Target number

3.7 Binning

3.8 Systematic errors

3.9 Data format



3.5 Flux correction

Integral region of flux

- Flux is integrated and removed. There are many ways how to introduce flux error.

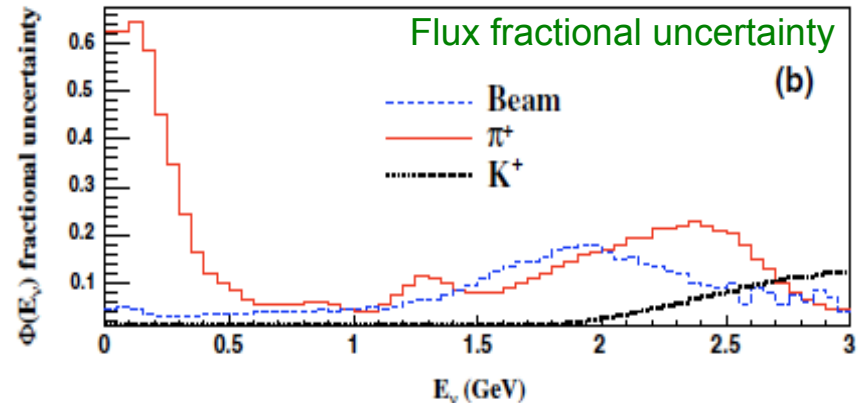
i. Flux is integrated in all spectrum region and it's variation is the flux error. This choice gives rather large flux error (e.g., ~12% for NC1 π^0).

NC1 π^0

ii. Cutoff for flux integration

CC1 π^0

Flux is integrated in [0.5-2.0] GeV, and error is variation of that. In this way, you can avoid flux variation at low energy which don't contribute to the channel. Error is smaller, ~7%.



CCQE

NCEL

anti
CCQE

anti
NCEL

iii. Flux is integrated all region, but flux error is calculated separately

Flux variation is calculated by variation of numerator of efficiency term (rate).

In this way, flux variation is automatically limited within the region relevant to cross section measurement. Both normalization and shape flux error are taken into account.

Error is smaller, ~8%.

3.5 Flux correction

Integral region of flux

- Flux is integrated and removed. There are many ways how to introduce flux error.

$$\left(\frac{d\sigma}{dx}\right)_i^F = \frac{\sum_j U_{ij}^F (d_j - b_j^F)}{\left(\frac{N_{\text{after}}}{N_{\text{before}}}\right)_i^F (\Phi^F T) \Delta x_i} \longrightarrow \left(\frac{d\sigma}{dx}\right)_i^F = \frac{\sum_j U_{ij}^F (d_j - b_j^F)}{\left(\frac{N_{\text{after}}}{N_{\text{before}}}\right)_i (\Phi T) \Delta x_i}$$



iii. Flux is integrated all region, but flux error is calculated separately

Flux variation is calculated by variation of numerator of efficiency term (rate).

In this way, flux variation is automatically limited within the region relevant to cross section measurement. Both normalization and shape flux error are taken into account.

Error is smaller, ~8%.

3.5 Flux correction

Integral region of flux

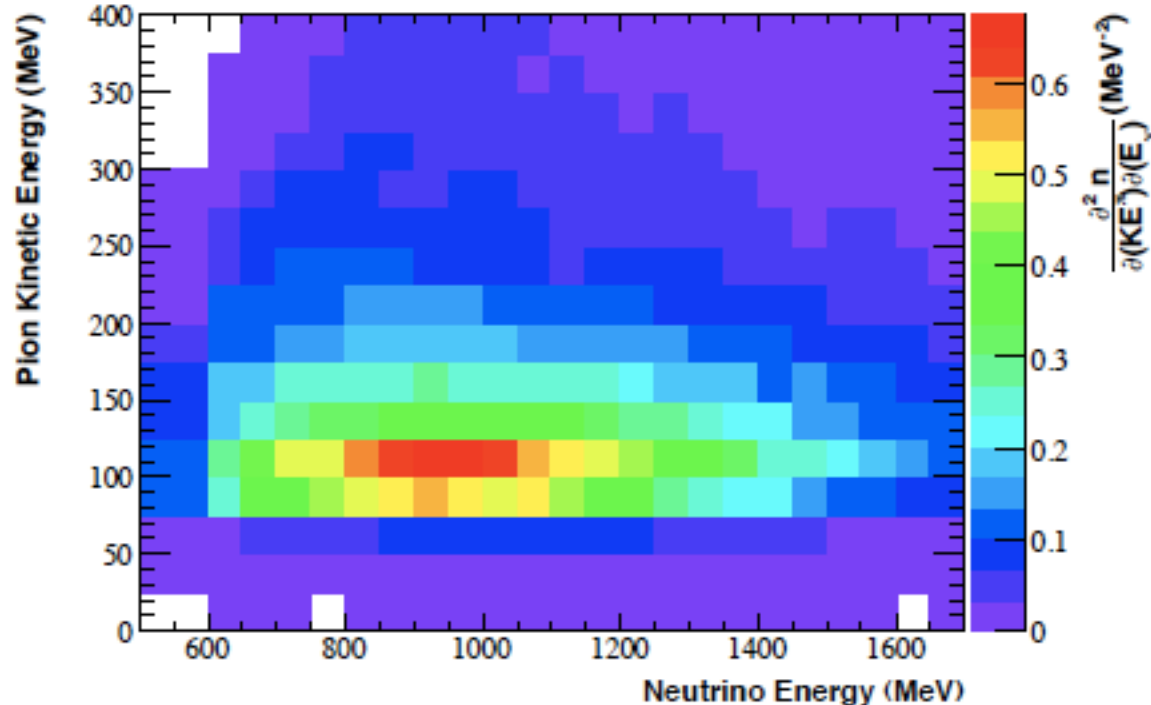
- Flux is integrated and removed. There are many ways how to introduce flux error.

iv. cross sections are function of neutrino energy

CC1 π^+

In this way, integrated flux in E_ν bin is unfolded in each bin of measured variables (e.g., pion kinetic energy), then flux error only relevant E_ν region apply to measured variables. This minimizes flux error at many region.

Pion kinetic energy- neutrino energy 2-dimensional cross section



3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

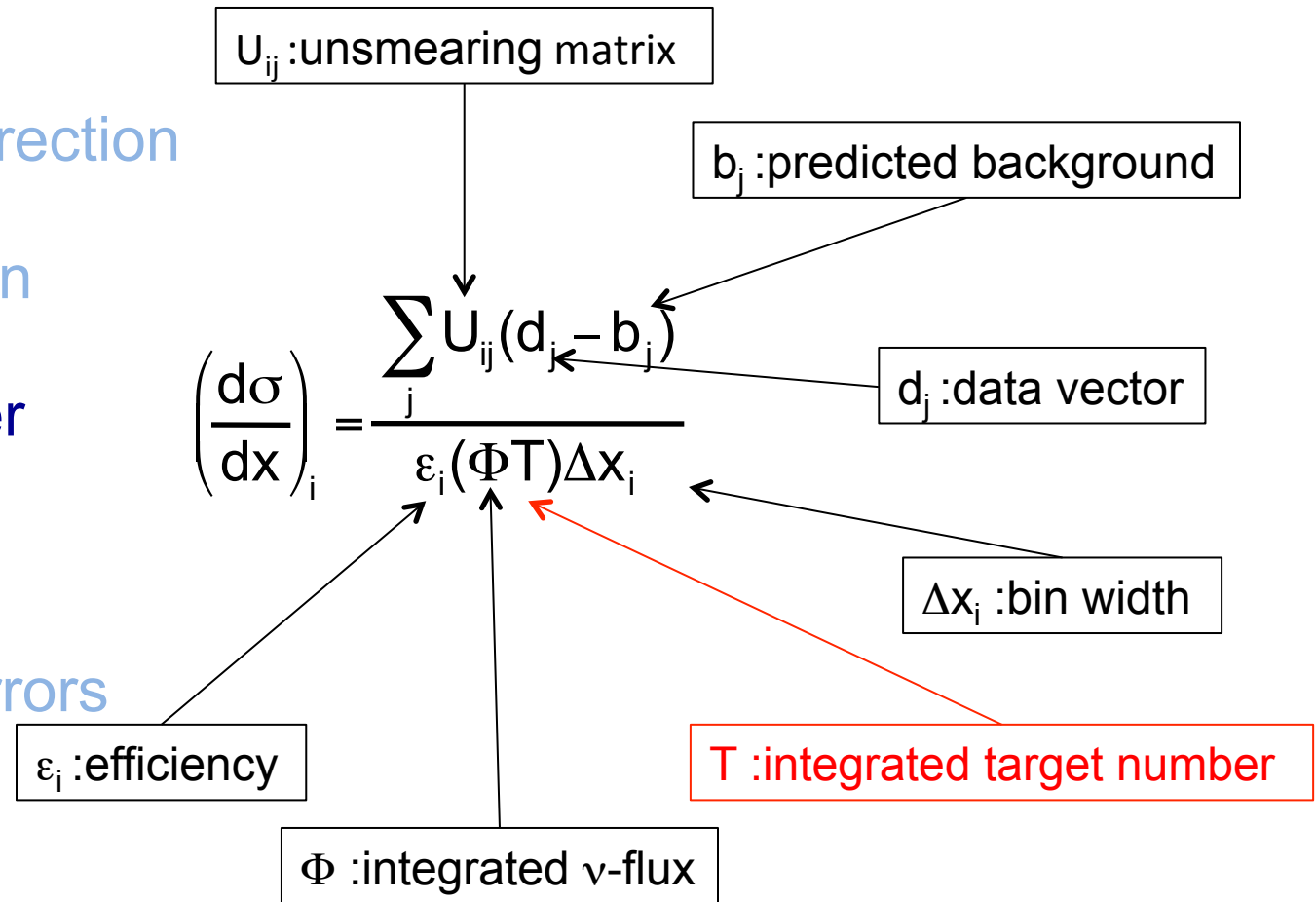
3.5 Flux correction

3.6 Target number

3.7 Binning

3.8 Systematic errors

3.9 Data format



3.6 Target number correction

What is the real fiducial volume?

- Fiducial cut is made based on reconstructed vertices.
- Fiducial volume is based on true dimension.

In MiniBooNE, fiducial cut is smaller than fiducial volume, to take account possible vertex mis-reconstruction.

- MiniBooNE is ~600cm radius sphere.
- MC is generated within 550cm sphere.
- The fiducial cut is 500cm sphere.

In this way, we can guarantee cross section is calculated in the region where we believe uniform.

- In general, **data-MC agreement is not enough for absolute cross section measurement.** Even data and MC perfectly agree in reconstructed spectrum, you need to worry the absolute calibration of vertex, target volume, and density.

3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

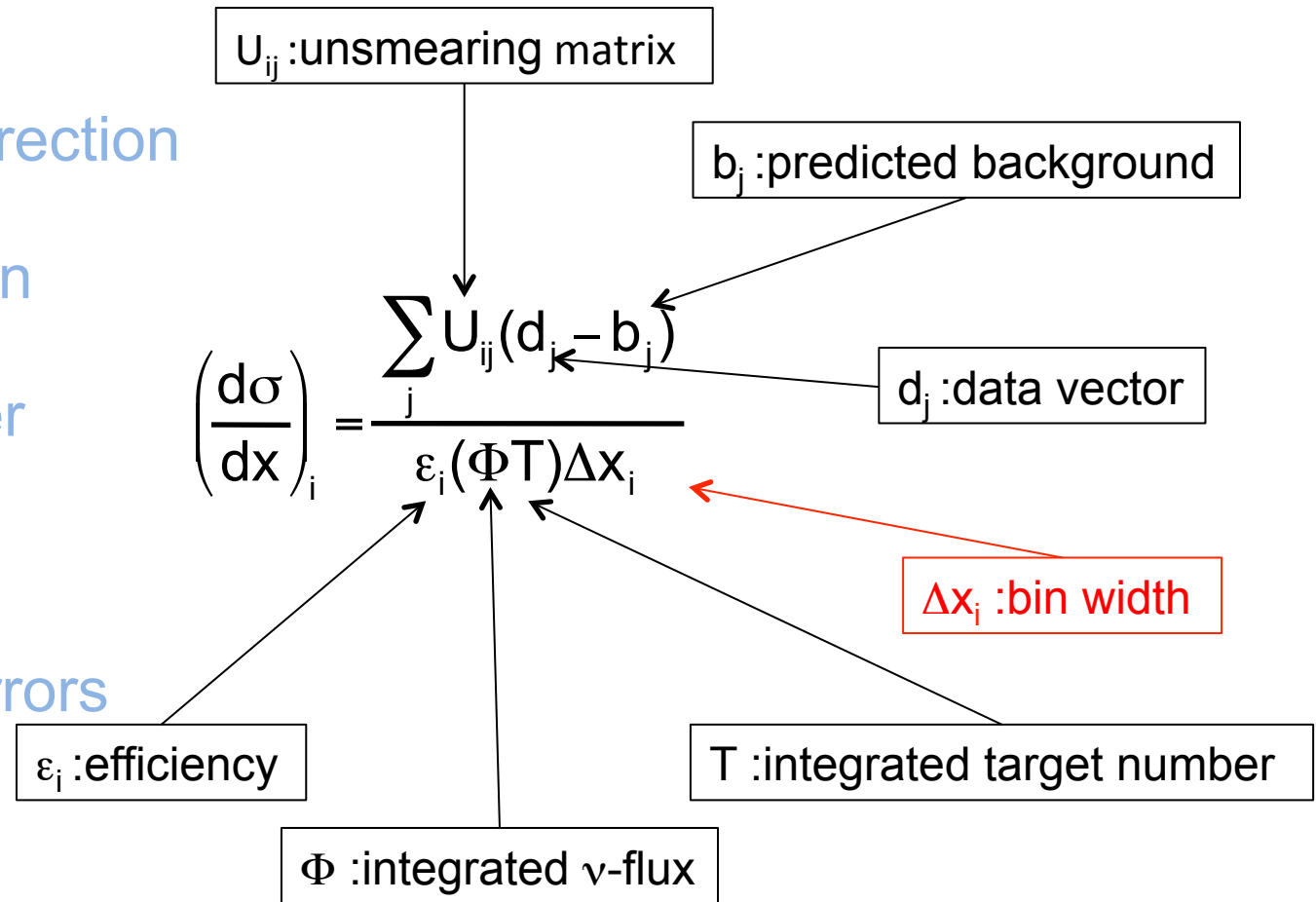
3.5 Flux correction

3.6 Target number

3.7 Binning

3.8 Systematic errors

3.9 Data format



3.7 Bin width

CCQE
NCEL
NC1 π^0
CC1 π^+
CC1 π^0
anti CCQE
anti NCEL

Statistics

Bin width is finer at high statistics region, and coarser at low statistics region.

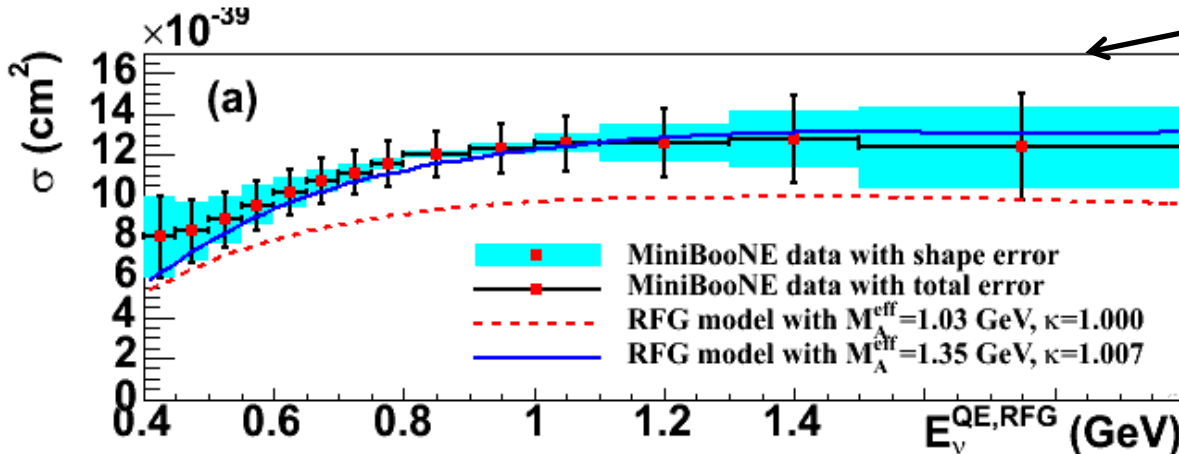
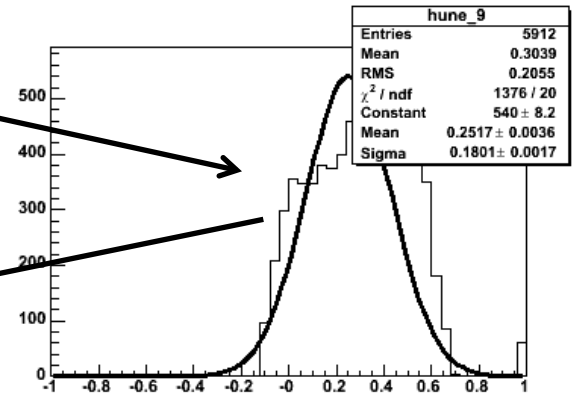
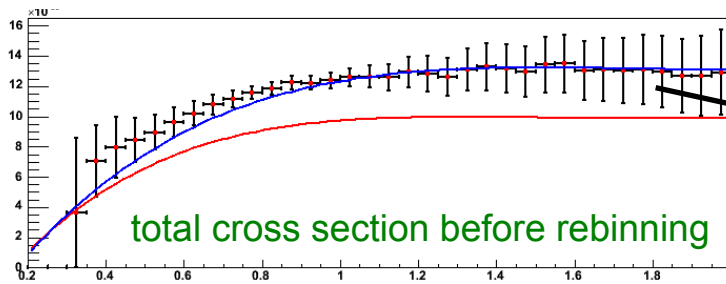
Systematics

Too fine bins with large shape systematic make no sense?

Reconstruction bias

CCQE

For CCQE analysis, reconstruction bias was added to bin resolution.



3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

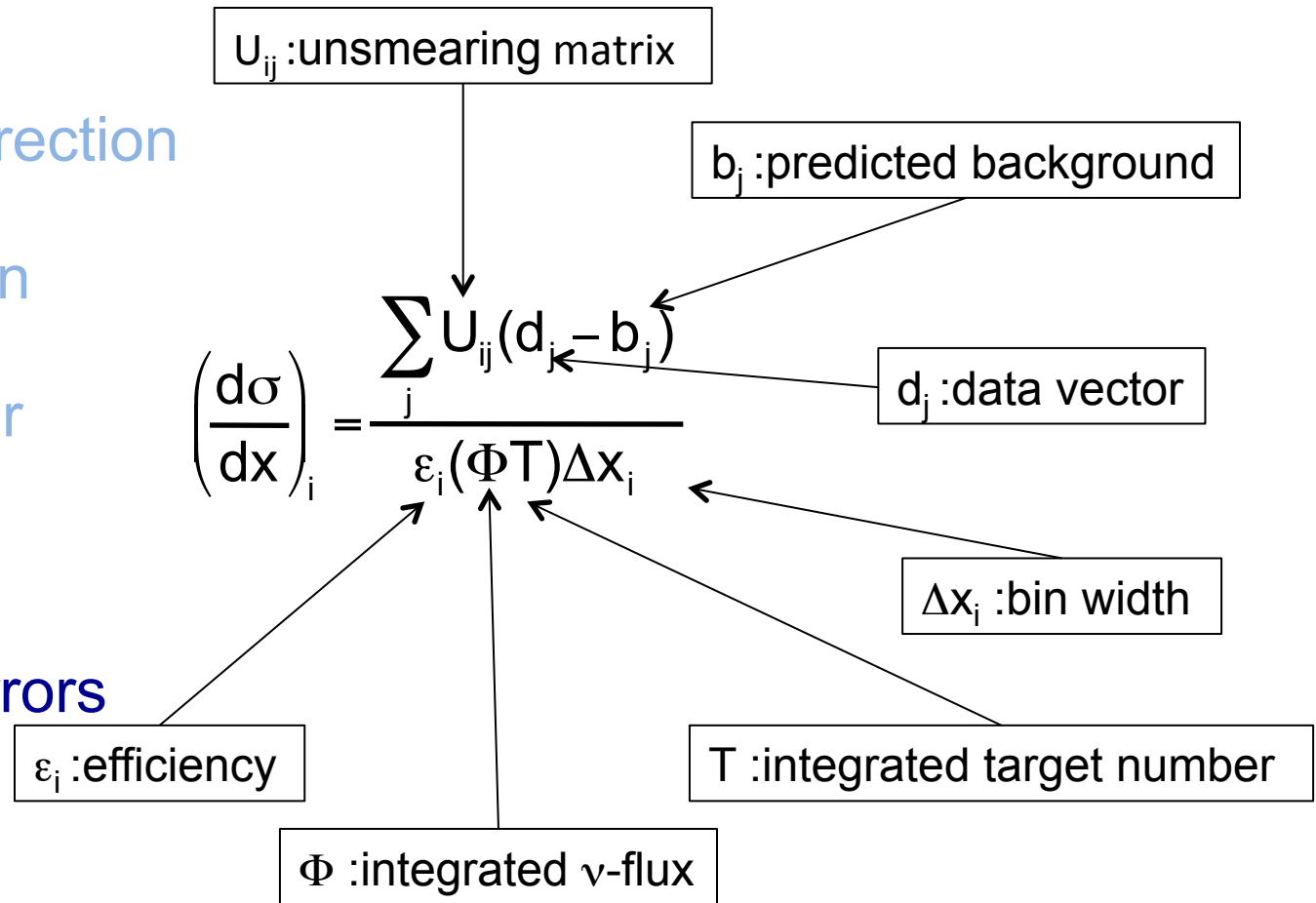
3.5 Flux correction

3.6 Target number

3.7 Binning

3.8 Systematic errors

3.9 Data format



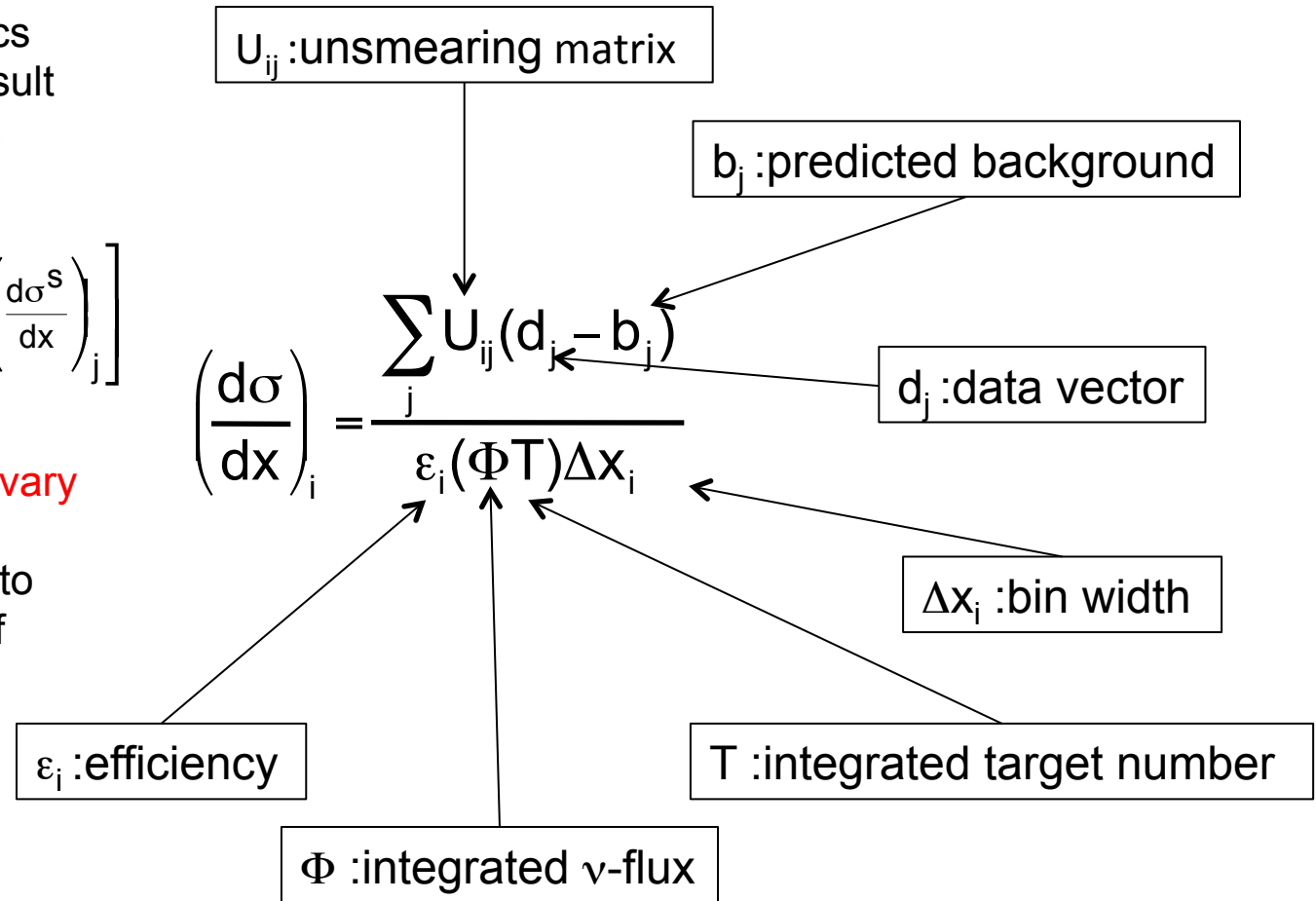
3.8 Systematic errors

Systematic error is calculated from the difference of systematics varied cross section result and central value cross section result.

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

4 parts are related with systematic error. **Don't vary all of them with all systematics!** You need to think about the effect of each term

- 3.8.1 background
- 3.8.2 U-matrix
- 3.8.3 Efficiency
- 3.8.4 Flux term
- 3.8.5 Target number



3.8 Systematic errors

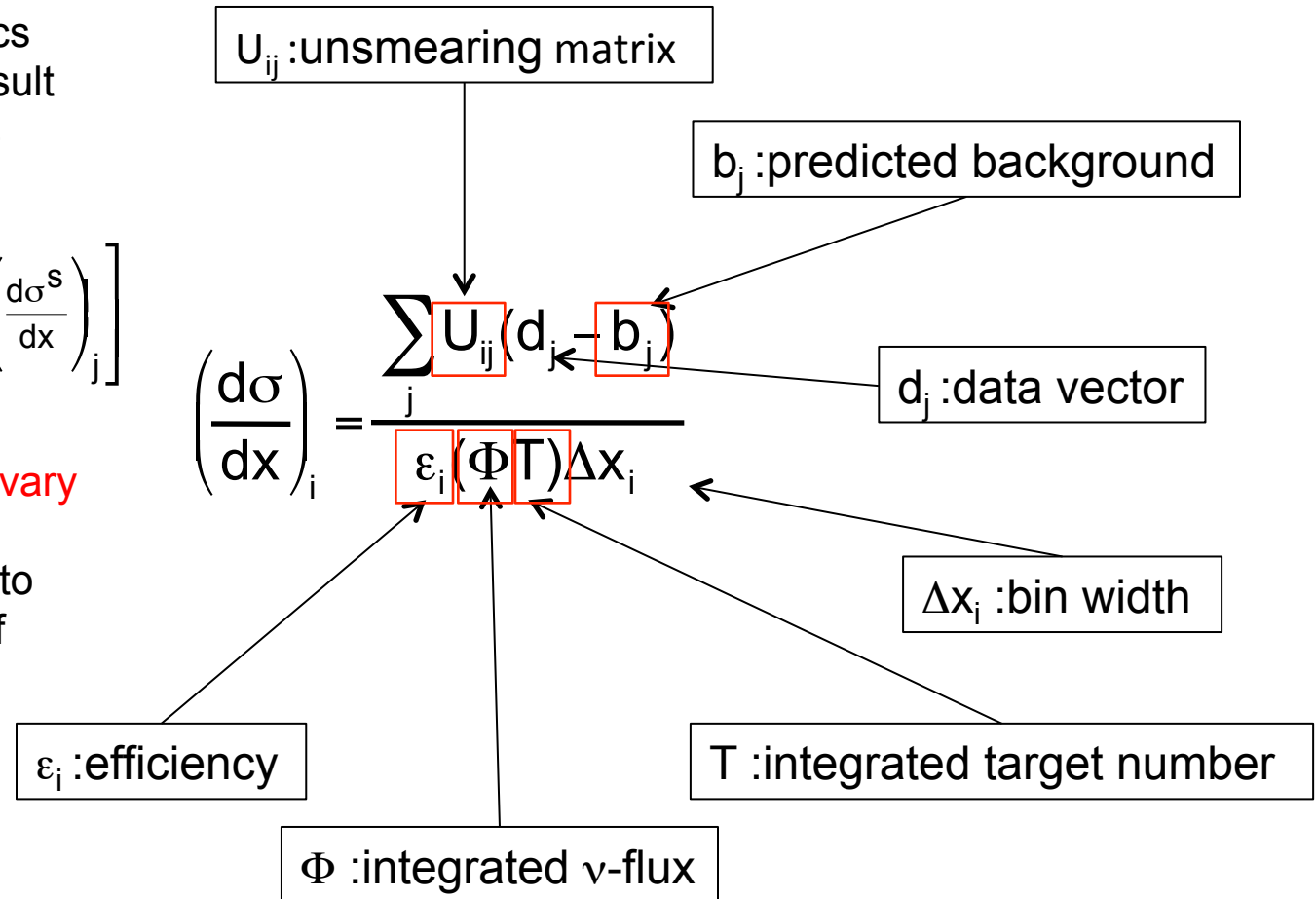
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- 3.8.1 background
- 3.8.2 U-matrix
- 3.8.3 Efficiency
- 3.8.4 Flux term
- 3.8.5 Target number

$$\left(\frac{d\sigma}{dx} \right)_i^S \neq \frac{\sum_j U_{ij}^S (d_j - b_j^S)}{\epsilon_i^S (\Phi^S T^S) \Delta x_i}$$



3.8 Systematic errors

Systematic error is calculated from the difference of systematics varied cross section result and central value cross section result.

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

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3.8.1 background

3.8.2 U-matrix

3.8.3 Efficiency

3.8.4 Flux term

3.8.5 Target number

$$\left(\frac{d\sigma}{dx} \right)_i = \frac{\sum_j U_{ij} (d_j - b_j)}{\varepsilon_i (\Phi T) \Delta x_i}$$

b_j : predicted background

To reduce cross section error

- higher purity
- sideband (then error is measurement)

3.8 Systematic errors

Systematic error is calculated from the difference of systematics varied cross section result and central value cross section result.

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

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3.8.1 background

3.8.2 U-matrix

3.8.3 Efficiency

3.8.4 Flux term

3.8.5 Target number

$$\left(\frac{d\sigma}{dx} \right)_i = \frac{\sum_j U_{ij} (d_j - b_j)}{\varepsilon_i (\Phi \Gamma) \Delta x_i}$$

b_j : predicted background

To reduce flux error

- use cancellation with signal MC
- purity correction method
- shape-only background subtraction method

3.8 Systematic errors

Systematic error is calculated from the difference of systematics varied cross section result and central value cross section result.

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

4 parts are related with systematic error. **Don't vary all of them with all systematics!** You need to think about the effect of each term

3.8.1 background

3.8.2 U-matrix

3.8.3 Efficiency

3.8.4 Flux term

3.8.5 Target number

U_{ij} : unsmearing matrix

$$\left(\frac{d\sigma}{dx} \right)_i = \frac{\sum_j U_{ij} (d_j - b_j)}{\varepsilon_i (\Phi T) \Delta x_i}$$

The error is dominated by detector error (flux error cancels). For Bayesian unfolding, **signal MC gives error here, too.**

3.8 Systematic errors

Systematic error is calculated from the difference of systematics varied cross section result and central value cross section result.

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

4 parts are related with systematic error. **Don't vary all of them with all systematics!** You need to think about the effect of each term

3.8.1 background

3.8.2 U-matrix

3.8.3 Efficiency

3.8.4 Flux term

3.8.5 Target number

Detector error goes here (flux and xsec error cancel).

$$\left(\frac{d\sigma}{dx} \right)_i = \frac{\sum_j U_{ij} (d_j - b_j)}{\epsilon_i (\Phi T) \Delta x_i}$$

Efficiency variation

3.8 Systematic errors

Systematic error is calculated from the difference of systematics varied cross section result and central value cross section result.

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

4 parts are related with systematic error. **Don't vary all of them with all systematics!** You need to think about the effect of each term

3.8.1 background

3.8.2 U-matrix

3.8.3 Efficiency

3.8.4 Flux term

3.8.5 Target number

Flux normalization error is here. You may need to apply cutoff to remove flux variation irrelevant for cross section measurement, to avoid overestimation of error.

$$\left(\frac{d\sigma}{dx} \right)_i = \frac{\sum_j U_{ij} (d_j - b_j)}{\epsilon_i (\Phi \Gamma) \Delta x_i}$$

Integrated flux variation

3.8 Systematic errors

Systematic error is calculated from the difference of systematics varied cross section result and central value cross section result.

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

4 parts are related with systematic error. **Don't vary all of them with all systematics!** You need to think about the effect of each term

3.8.1 background

3.8.2 U-matrix

3.8.3 Efficiency

3.8.4 Flux term

3.8.5 Target number

Precise definition of active volume may remove the bias of target number.
(**this is not simulated effect**, i.e., incorrect fiducial volume just give wrong answer)

$$\left(\frac{d\sigma}{dx} \right)_i = \frac{\sum_j U_{ij} (d_j - b_j)}{\varepsilon_i (\Phi T) \Delta x_i}$$

Total target number

3.8.6 Systematics error matrix production

Unisim

The error matrix can be made by changing one of systematics and calculate differential cross section ($d\sigma^S/dx$), then take a difference with differential cross section calculated with central value MC ($d\sigma/dx$).

$$E_{ij} = \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

Multisim

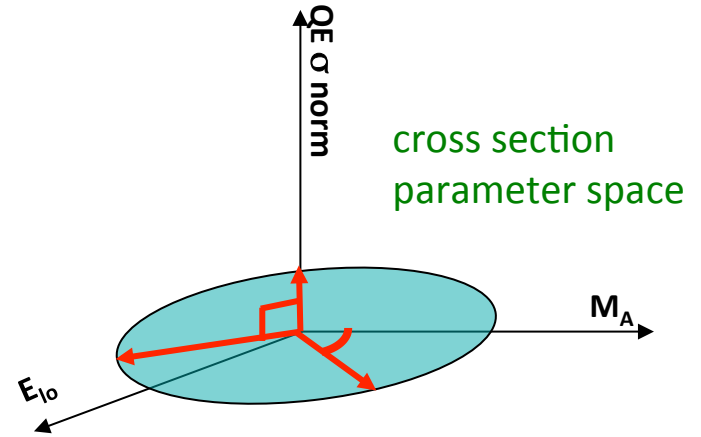
If there is a correlation between systematics (input error matrix), it should propagate correctly. In this case, number of $d\sigma^S/dx$ with different set of systematics drawn from input error matrix make many error matrices. Then, we take average of them to construct output error matrix.

$$E_{ij} = \frac{1}{M} \sum_s \left[\left(\frac{d\sigma}{dx} \right)_i - \left(\frac{d\sigma^S}{dx} \right)_i \right] \left[\left(\frac{d\sigma}{dx} \right)_j - \left(\frac{d\sigma^S}{dx} \right)_j \right]$$

3.8.6 Multisim

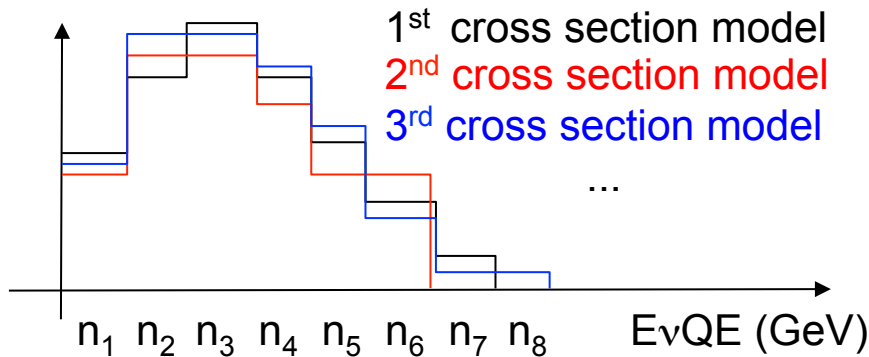
ex) cross section uncertainties

M_A QE	6%	↑ correlated
E_{lo} sf	2%	
QE σ norm	10%	uncorrelated



$$M_{input}(XS) = \begin{pmatrix} \text{var}(M_A) & \text{cov}(M_A, E_{lo}) & 0 \\ \text{cov}(M_A, E_{lo}) & \text{var}(E_{lo}) & 0 \\ 0 & 0 & \text{var}(\sigma - \text{norm}) \end{pmatrix}$$

cross section error for EvQE



repeat this exercise many times to create smooth error matrix for EvQE

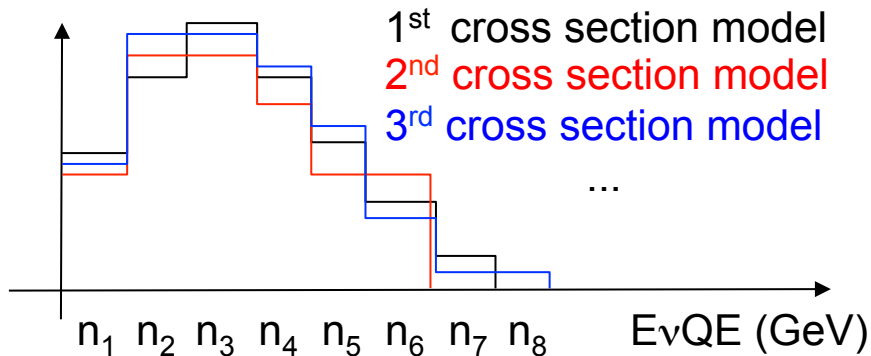
3.8.6 Multisim

Output cross section error matrix for EvQE

$$[\mathbf{M}_{\text{output}}(\mathbf{XS})]_{ij} \approx \frac{1}{S} \sum_k^S (\mathbf{N}_i^k(\mathbf{XS}) - \mathbf{N}_i^{\text{MC}})(\mathbf{N}_j^k(\mathbf{XS}) - \mathbf{N}_j^{\text{MC}})$$

$$\mathbf{M}_{\text{output}}(\mathbf{XS}) = \begin{pmatrix} \text{var}(n_1) & \text{cov}(n_1, n_2) & \text{cov}(n_1, n_3) & \cdots \\ \text{cov}(n_1, n_2) & \text{var}(n_2) & \text{cov}(n_2, n_3) & \cdots \\ \text{cov}(n_1, n_3) & \text{cov}(n_2, n_3) & \text{var}(n_3) & \cdots \\ \vdots & \vdots & \vdots & \ddots \end{pmatrix}$$

cross section error for EvQE



repeat this exercise many times to create smooth error matrix for EvQE

3.8.7 Statistics error

NC1 π^0

CC1 π^0

Statistical error propagation

Due to unfolding, there is data statistical error on off-diagonal term of error matrix. The diagonal statistical error can be propagated through Jacobian. It is weakened, and smoothly migrate to off diagonal. MC statistics can be transferred by similar way if it is large.

$$V_{ij} \left[\frac{d\sigma}{dx} \right] = V_{ij} \left[\frac{\sum U(d-b)}{\varepsilon(\Phi T) \Delta x} \right] = \left(\frac{\partial \left[\frac{\sum U(d-b)}{\varepsilon(\Phi T) \Delta x} \right]}{\partial d} \right)_{ki} V_{km} [d] \left(\frac{\partial \left[\frac{\sum U(d-b)}{\varepsilon(\Phi T) \Delta x} \right]}{\partial d} \right)_{mj}$$

Statistical error through Multisim

NCEL

Fake data set is made by applying fluctuation on data within data statistics. Then statistics multisim output error matrix is made from fake data set.

CCQE

CC1 π^+

anti
CCQE

anti
NCEL

Statistical error through detector error matrix

Detector error multisim MC set is made with data statistics (MiniBooNE historic reason), so the multisim output error matrix has ~data statistical error, too.

3.8.8 Correlated systematic errors between samples

Correlated errors between T2K cross section results

T2K used many MiniBooNE cross section results for the global fit, however, MiniBooNE data errors should be correlated because all of them are measured by the same beamline and the detector.

To avoid same mistake, we should provide correlated errors? For example

1. flux normalization
2. π/K ratio (Mark H.)

Providing these 2 numbers could reduce over all errors dramatically for the global fit.

3.1 Signal definition

3.2 Background removing

3.3 Unsmearing

3.4 Efficiency correction

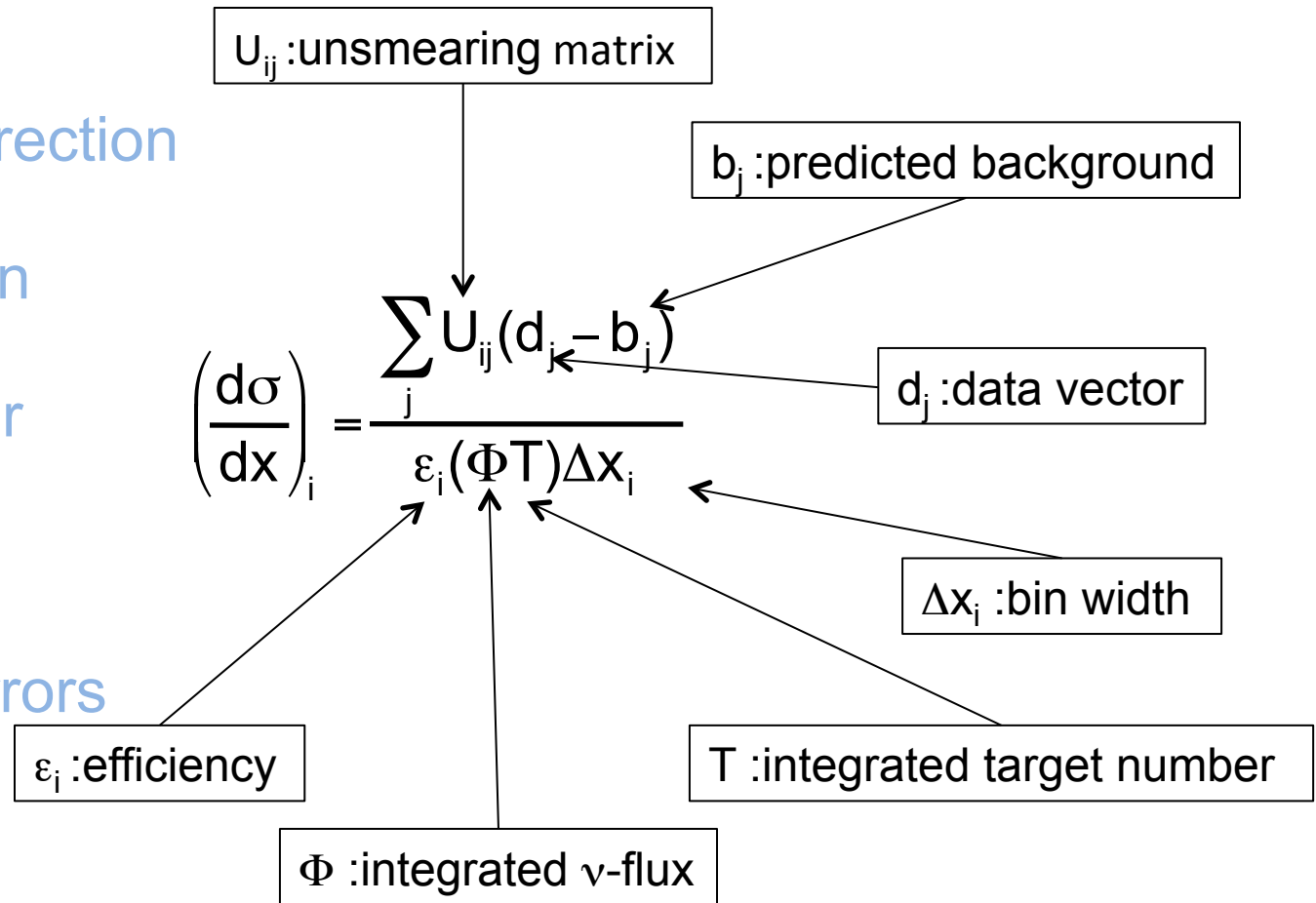
3.5 Flux correction

3.6 Target number

3.7 Binning

3.8 Systematic errors

3.9 Data format



3.9 Data format

CCQE

NCEL

NC1 π^0

CC1 π^+

CC1 π^0

anti
CCQE

anti
NCEL

Tables on MiniBooNE data release website

- In MiniBooNE, all cross section tables, as well as flux table, are released in website http://www-boone.fnal.gov/for_physicists/data_release/

Cross section

- Flux-integ

- Flux integ

- Flux-unfo

Additional t

- signal-like

cross section

- response

observed e

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Data Releases

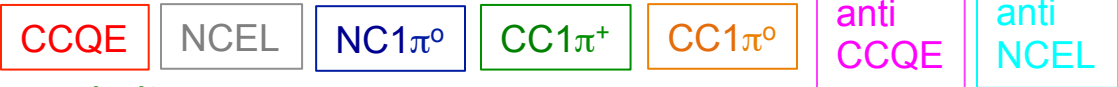
This page provides MiniBooNE data (histograms, error matrices, ntuples, etc) released in association with particular publications. Only the subset of MiniBooNE papers with released data are listed here. Refer to the [Publications](#) page for a complete list of MiniBooNE publications.

- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[Measurement of Muon Neutrino Induced Charged Current Neutral Pion Production Cross Sections on Mineral Oil at \$E_{\nu} = 0.5\text{-}2.0\$ GeV](#)", arXiv:1010.3264 [hep-ex], submitted to Phys. Rev. D
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[Measurement of the Neutrino Neutral Current Elastic Differential Cross Section](#)", arXiv:1007.4730 [hep-ex], submitted to Phys. Rev. D
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[First Measurement of the Muon Neutrino Charged Current Quasielastic Double Differential Cross section](#)", arXiv:1002.2680 [hep-ex], Phys. Rev. D81, 092005 (2010)
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[Measurement of \$\nu_{\mu}\$ and \$\bar{\nu}_{\mu}\$ induced neutral current single \$\pi^0\$ production cross sections on mineral oil at \$E_{\nu} \sim O\(1\$ GeV\)](#)", arXiv:0911.2063 [hep-ex], Phys. Rev. D81, 013005 (2010)
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[A Search for Electron Anti-Neutrino Appearance at the \$\Delta m^2 \sim 1\$ eV² Scale](#)", arXiv:0904.1958 [hep-ex], Phys. Rev. Lett. 103, 111801 (2009),
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[A Search for Muon Neutrino and Anti-Neutrino Disappearance in MiniBooNE](#)", arXiv:0903.2465 [hep-ex], Phys. Rev. Lett. 103, 061802 (2009)
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[Unexplained Excess of Electron-Like Events From a 1 GeV Neutrino Beam](#)", arXiv:0812.2243 [hep-ex], Phys. Rev. Lett. 102, 101802 (2009)
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[The Neutrino Flux Prediction at MiniBooNE](#)", arXiv:0806.1449 [hep-ex], Phys. Rev. D. 79, 072002 (2009)
- [Data Released](#) with A.A. Aguilar-Arevalo et al., "[A Search for Electron Neutrino Appearance at the \$\Delta m^2 \sim 1\$ eV² Scale](#)", arXiv:0704.1500 [hep-ex], Phys. Rev. Lett. 98, 231801 (2007)

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clusive

3.9 Data format



Tables on MiniBooNE data release website

- In MiniBooNE, all cross section tables, as well as flux table, are released in website http://www-boone.fnal.gov/for_physicists/data_release/

Data Release for A.A. Aguilar-Arevalo et al., "First Measurement of the Muon Neutrino Charged Current Quasielastic Double Differential Cross section", arXiv:1002:2680 [hep-ex], Phys. Rev. D81, 092005 (2010)

The following MiniBooNE information from the 2010 CCQE cross section paper is made available to the public:

• ν_{μ} CCQE cross sections:

- **MiniBooNE flux**
 - [table](#) of predicted MiniBooNE muon neutrino flux (Table V)
- **flux-integrated double differential cross section (Figure 13)**
 - [1D array](#) of bin boundaries partitioning the muon kinetic energy (top) and the cosine of the muon scattering angle (bottom)
 - [2D array](#) of the value of the double differential cross section in each bin in units of $10^{-41} \text{ cm}^2/\text{GeV}/\text{nucleon}$. The muon kinetic energy increases from left to right, and the cosine of the muon scattering angle decreases from top to bottom (Table VI)
 - [2D array](#) of the shape uncertainty of the double differential cross section in each bin in units of $10^{-42} \text{ cm}^2/\text{GeV}/\text{nucleon}$. The total normalization error is 10.7% (Table VII)
 - [2D array](#) of the predicted CCQE-like background double differential cross section in each bin in units of $10^{-41} \text{ cm}^2/\text{GeV}/\text{nucleon}$ (Table VIII)
- **flux-integrated single differential cross section in bins of Q^2 (Figure 14)**
 - [1D array](#) of bin boundaries partitioning the reconstructed four momentum transfer, Q^2
 - [1D array](#) of the value of the single differential cross section in each bin in units of $\text{cm}^2/\text{GeV}^2/\text{nucleon}$ (Table IX)
 - [1D array](#) of the shape uncertainty of the single differential cross section in each bin in units of $\text{cm}^2/\text{GeV}^2/\text{nucleon}$. The total normalization error is 10.7% (Table IX)
 - [1D array](#) of the predicted CCQE-like background single differential cross section in each bin in units of $\text{cm}^2/\text{GeV}^2/\text{nucleon}$ (Table IX)
- **flux-unfolded cross section as a function of neutrino energy (Figure 15)**
 - [1D array](#) of bin boundaries partitioning the neutrino energy
 - [1D array](#) of the value of the cross section in each bin in units of $\text{cm}^2/\text{nucleon}$ (Table X)
 - [1D array](#) of the shape uncertainty of the cross section in each bin in units of $\text{cm}^2/\text{nucleon}$. The total normalization error is 10.7% (Table X)
 - [1D array](#) of the total uncertainty of the cross section in each bin in units of $\text{cm}^2/\text{nucleon}$ (Table X)
 - [1D array](#) of the predicted CCQE-like background cross section in each bin in units of $\text{cm}^2/\text{nucleon}$ (Table X)

3.9 Data format

CCQE

NCEL

NC1 π^0

CC1 π^+

CC1 π^0

anti
CCQE

anti
NCEL

Tables on MiniBooNE data release website

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http://www-boone.fnal.gov/for_physicists/data_release/

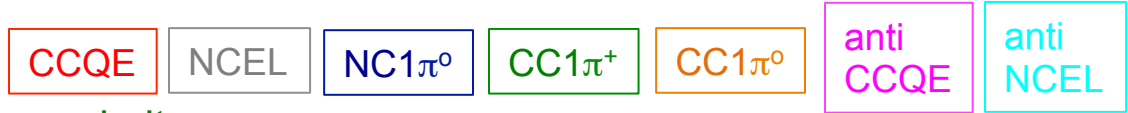
Cross section format

- Flux-integrated double differential cross section

- Flux integrated single differential cross section

190.0	326.5	539.2	901.8	1288	1633	1857	1874	1803	1636	1354	1047	794.0	687.9	494.3	372.5	278.3	227.4
401.9	780.6	1258	1714	2084	2100	2035	1620	1118	783.6	451.9	239.4	116.4	73.07	41.67	36.55	0	0
553.6	981.1	1501	1884	1847	1629	1203	723.8	359.8	156.2	66.90	26.87	1.527	19.50	0	0	0	0
681.9	1222	1546	1738	1365	909.6	526.7	222.8	81.65	35.61	11.36	0.131	0	0	0	0	0	0
765.6	1233	1495	1289	872.2	392.3	157.5	49.23	9.241	1.229	4.162	0	0	0	0	0	0	0
871.9	1279	1301	989.9	469.1	147.4	45.02	12.44	1.012	0	0	0	0	0	0	0	0	0
910.2	1157	1054	628.8	231.0	57.95	10.69	0	0	0	0	0	0	0	0	0	0	0
992.3	1148	850.0	394.4	105.0	16.96	10.93	0	0	0	0	0	0	0	0	0	0	0
1007	970.2	547.9	201.5	36.51	0.844	0	0	0	0	0	0	0	0	0	0	0	0
1003	813.1	404.9	92.93	11.63	0	0	0	0	0	0	0	0	0	0	0	0	0
919.3	686.6	272.3	40.63	2.176	0	0	0	0	0	0	0	0	0	0	0	0	0
891.8	503.3	134.7	10.92	0.071	0	0	0	0	0	0	0	0	0	0	0	0	0
857.5	401.6	79.10	1.947	0	0	0	0	0	0	0	0	0	0	0	0	0	0
778.1	292.1	33.69	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
692.3	202.2	17.42	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
600.2	135.2	3.624	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
497.6	85.80	0.164	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
418.3	44.84	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
348.7	25.82	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
289.2	15.18	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0

3.9 Data format

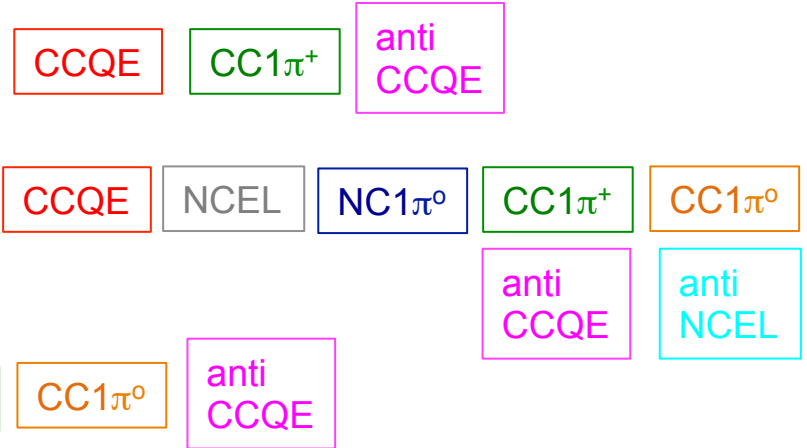


Tables on MiniBooNE data release website

- In MiniBooNE, all cross section tables, as well as flux table, are released in website http://www-boone.fnal.gov/for_physicists/data_release/

Cross section format

- Flux-integrated double differential cross section
- Flux integrated single differential cross section
- Flux-unfolded total cross section



Additional tables

- signal-like background tables are presented, so that people can use either exclusive cross section or effective cross section.



- response matrix R is presented so that people can calculate MiniBooNE observed energy spectrum



3.9 Data format

MINERvA style table?

- In MINERvA, data tables are all in “supplemental material” and the papers only have cross section plots, to fit in PRL page limit. I think this is clever.

3.9 Cross section errors

Cross section error format

NC1 π^0 CC1 π^+ CC1 π^0

- Complete error matrix of differential cross sections

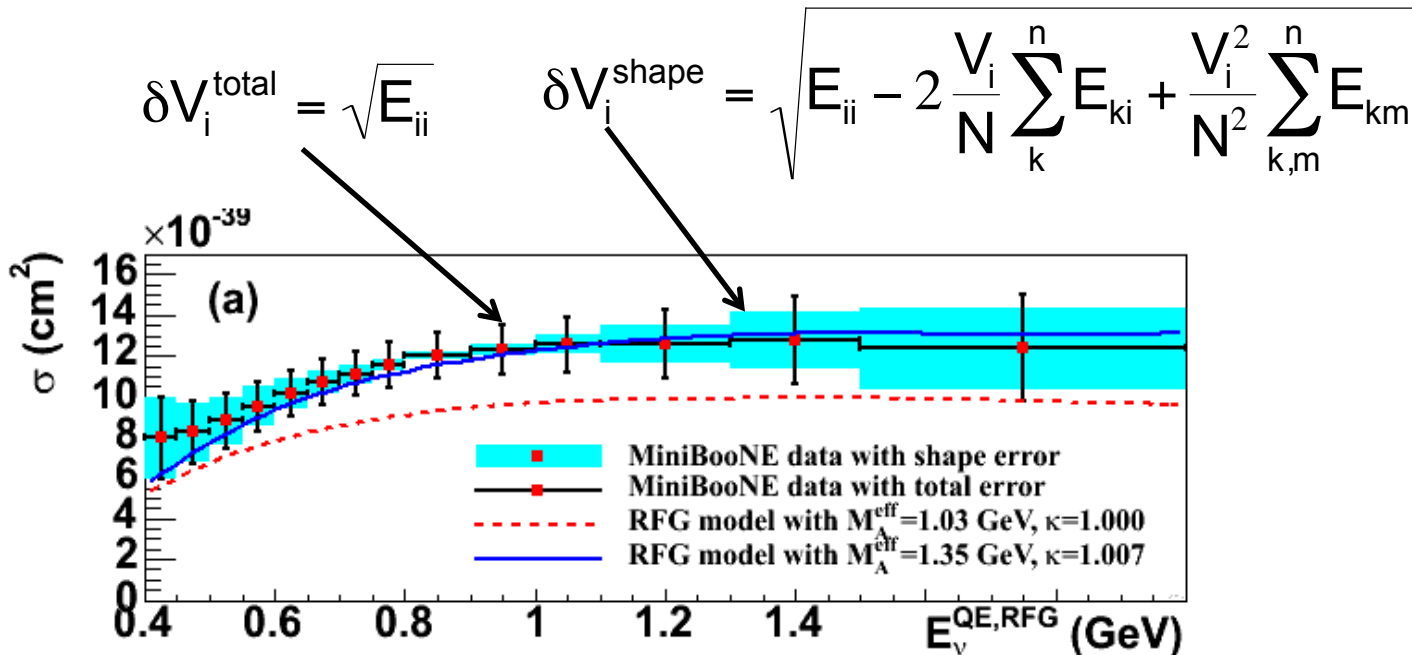
- Complete error matrix for reconstructed energy spectrum

NCEL

- Diagonal term of shape only error matrix and total normalization error.

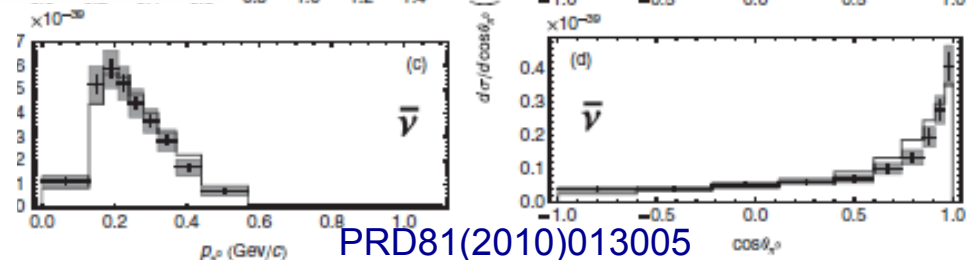
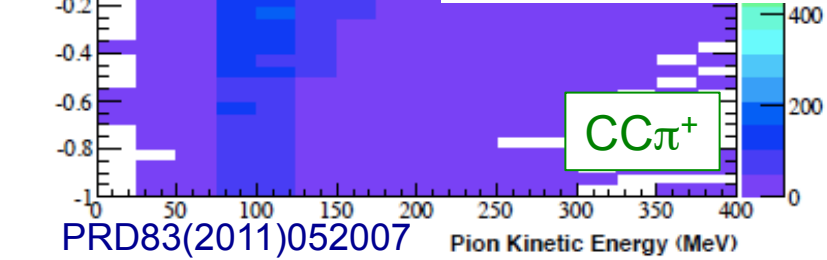
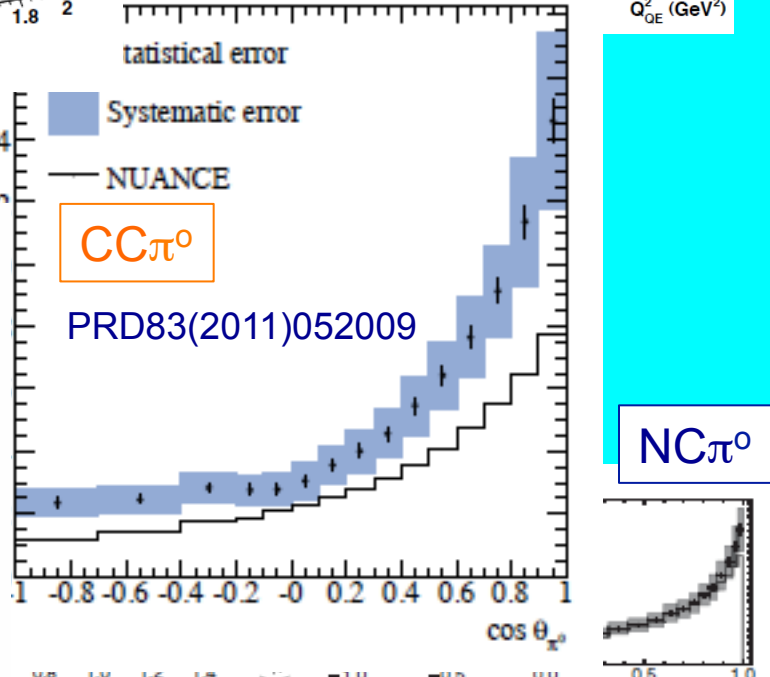
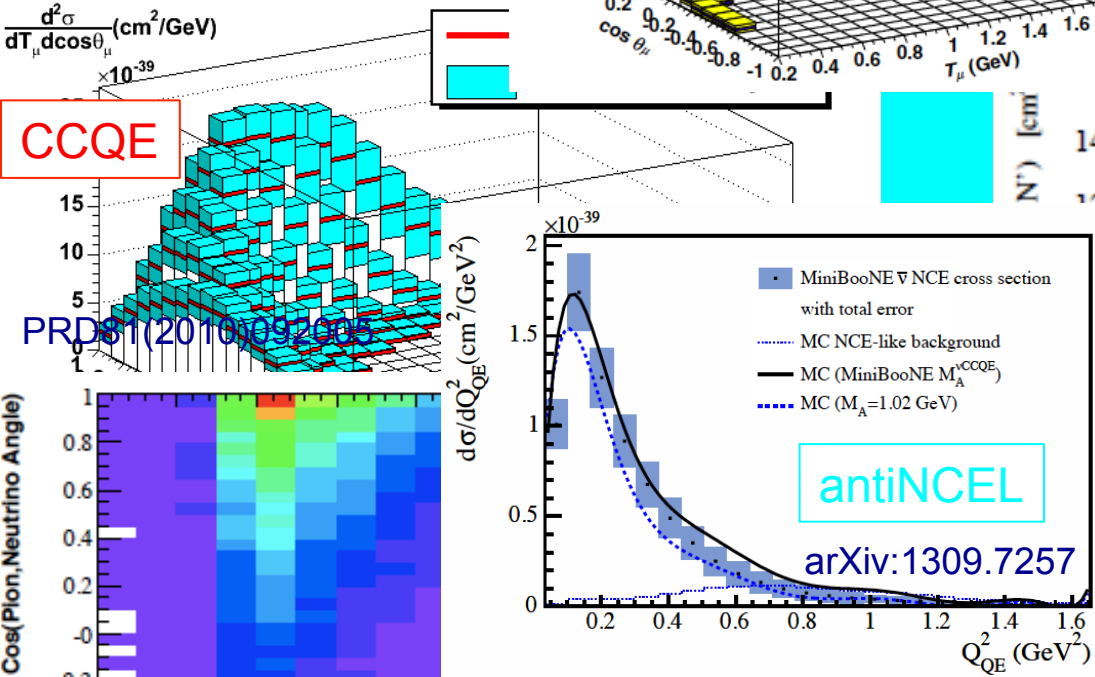
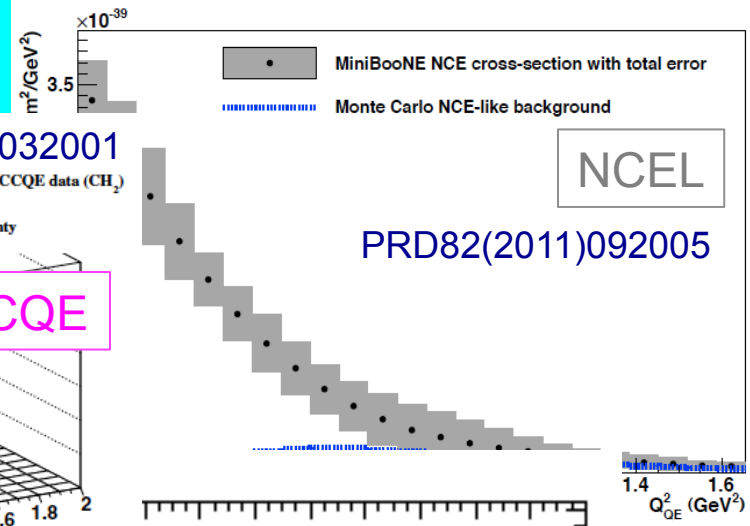
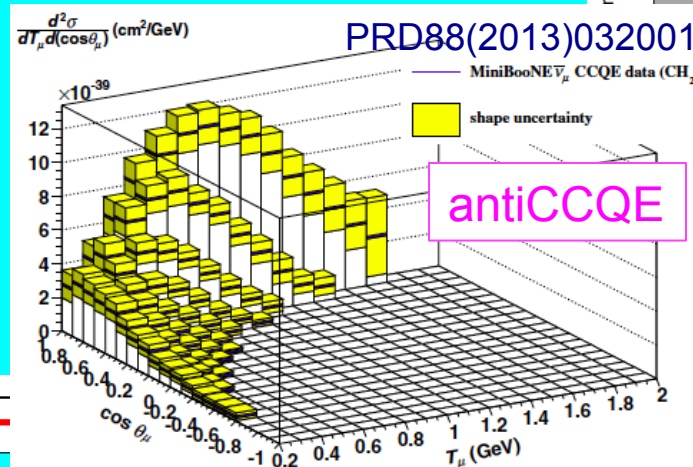
CCQE anti CCQE

Diagonal term of shape only error matrix has information of covariance of total error matrix, so this is a convenient way to show bin-bin correlation in 1-dimension



3.9 Data format

MiniBooNE flux-integrated differential cross section result gallery (over 800 citations)



Conclusion

Cross section analysis requires all set of different techniques and ideas.

Cross section analysis takes time.

MiniBooNE developed lots of techniques and ideas, those are useful start points for T2K analysis

Thank you for your attention!

3.2 Background removing

CCQE

NCEL

CC1 π^+

CC1 π^0

anti
CCQE

anti
NCEL

Data driven correction

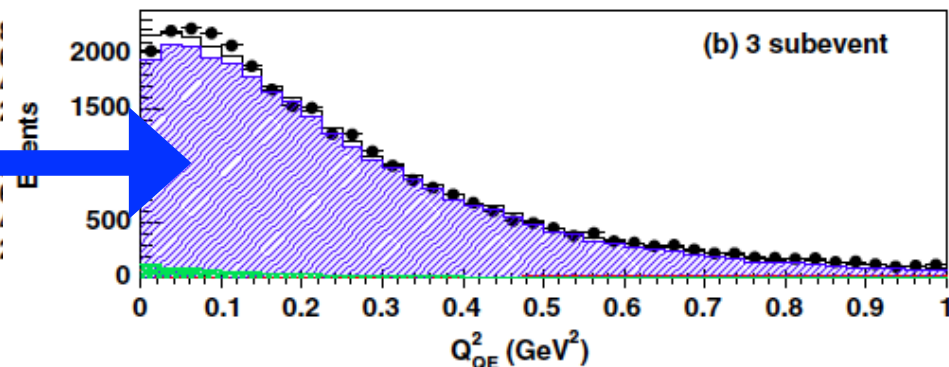
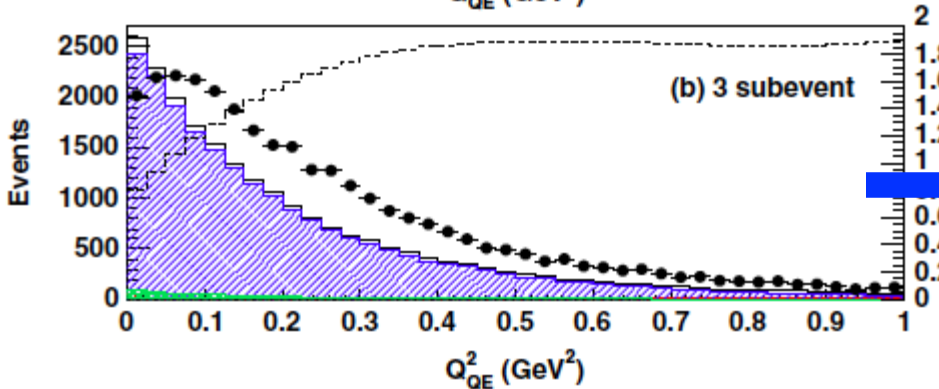
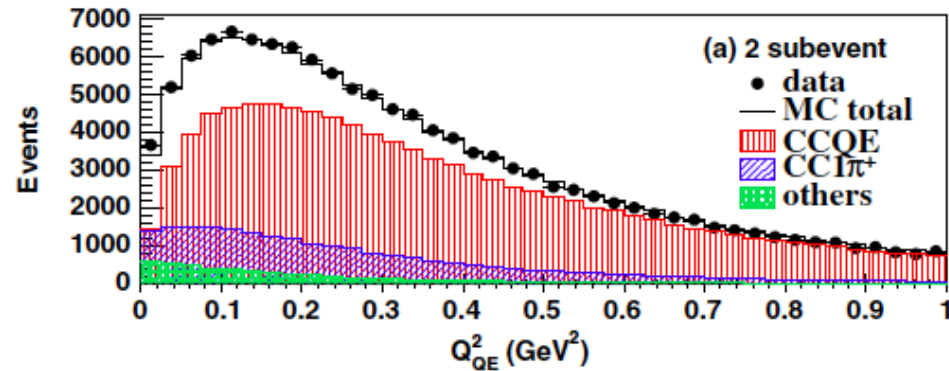
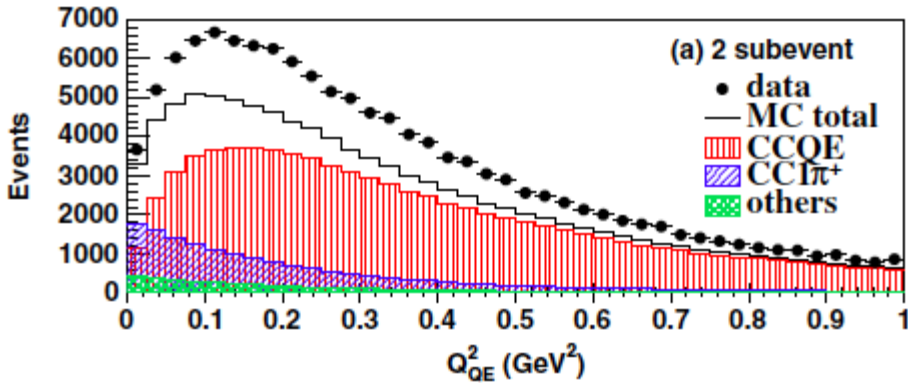
- Almost all analyses use sideband data to correct background distribution in signal boxes

CCQE-CC1 π^+ box simultaneous fit

CCQE

- CC1 π^+ box is >90% purity. Using this sample, background distribution in CCQE box is modified.

External background enhanced sample



3.2 Background removing

Cross section error of pion kinematics

CC1 π^+

- Purity is extremely high (~90%), so error from background is negligible.
- Signal is defined from pion in final state, so pion absorption is not systematic error.
 - In general, FSI is small error for all analyses
- The pion absorption in the oil is error.
 - Secondary scattering will be the largest error for hadron measurement.

I think you revise all errors to think which is the important error etc. If you run analysis machinery with all errors on, you will over-estimate systematic errors.

3.3 Unsmearing

How to construct M-matrix? Reconstructed variables

- The definition of true kinematics is tricky, because you have choice.

i. True Q^2 is defined by reconstructed Q^2 from true kinematics



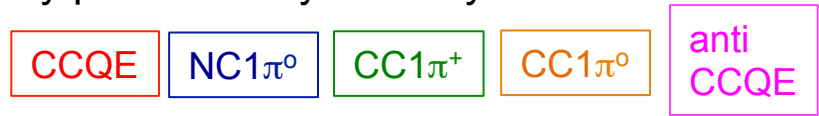
For example, CCQE, true Q^2 is defined “reconstructed Q^2 from true muon energy and angle”, and we call it “ Q^2_{QE} ” to remind people this is reconstructed under QE assumption.

ii True Q^2 is defined by true Q^2 in MC



This may be useful to compare with old data, only presented by this way

iii True E_ν is defined by true E_ν in MC



For example, CCQE, E_ν is called “ $E_\nu^{QE,RFG}$ ” to remind people **this is reconstructed under QE assumption then unfolded by assuming RFG model.**

Flux-unfolded total cross section

- it is important to unfold under RFG, otherwise reconstruction bias deviate cross section at the tail significantly.

