

A Joint ND280-SK 1R_μ-SK 1R_e Fit using MCMC

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February 17, 2014

Abstract

An analysis of the Run 1–4 T2K data is performed with a Markov Chain Monte Carlo. The data included in the analysis are the ND280 ν_μ , SK 1R_μ, and SK 1R_e samples. When fitting with only T2K data, the best fit point for the oscillation parameter is $\Delta m_{32}^2 = 2.491 \times 10^{-3} \text{ eV}^2$, $\sin^2 \theta_{23} = 0.520$, and $\sin^2 \theta_{13} = 0.0377$ for normal hierarchy and $\delta_{cp} = 0$, with 90% credible intervals of $2.34\text{--}2.69 \times 10^{-3} \text{ eV}^2$, $0.445\text{--}0.595$, and $0.0230\text{--}0.0600$, respectively. When fitting with the reactor constraint, the best fit point for normal hierarchy is $\Delta m_{32}^2 = 2.510 \times 10^{-3} \text{ eV}^2$, $\sin^2 \theta_{23} = 0.527$, and $\delta_{cp} = -1.551$. The 90% credible interval for δ_{cp} excludes $0.45\text{--}2.66$ for the normal hierarchy and $0.15\text{--}3.04$ for inverted hierarchy. Other interpretations of the data are also discussed.

T2K-TN-171

Contents

19	1 Introduction	4
20	2 Event Selection	5
21	2.1 ND280 Tracker ν_μ	5
22	2.2 SK 1R _e and 1R _{μ}	7
23	3 Systematic Errors	9
24	3.1 Flux	9
25	3.2 Cross Section	10
26	3.3 ND280 Detector	10
27	3.4 Final State Interactions (ND280 only)	12
28	3.5 SK Detector	13
29	4 Monte Carlo Predictions and Pre-fit Data/MC comparison	14
30	4.1 ND280	14
31	4.2 SK 1R _{μ}	18
32	4.3 SK 1R _e	18
33	5 Adaptive Kernel Density Method	24
34	6 Fitter Validation	24
35	6.1 Nominal Data Set	25
36	6.2 Toy Experiments	25
37	6.3 Marginalization Induced Biases	28
38	6.4 Fake Data Set Fits	34
39	6.4.1 T2K Only Fits	34
40	6.4.2 T2K with Reactor Constraint	38
41	6.5 Comparison with BANFF Matrix Fit	38
42	7 Fit Results	40
43	7.1 T2K Run 1–4 Data Fit	40
44	7.2 T2K Run 1–4 Data Fit With Reactor Constraint	47

45	A Additional Fake Data Set Plots	57
46	A.1 T2K Only	57
47	A.2 T2K with Reactor Constraint	57
48	A.3 Comparison with VALOR	62

1 Introduction

This technical note describes a fit to the ND280 tracker ν_μ , SK 1R $_\mu$, and SK 1R $_e$ Run 1–4 data using the Markov Chain Monte Carlo method. A description of the Markov Chain method can be found in [1].

This analysis uses two new features compared to the method described the referenced note; instead of reweighting the predicted Monte Carlo (MC) spectra using binned pdf templates, the individual MC events are weighted event-by-event, according to the relevant variable(s) for the tweak being applied. Then, when all weights have been calculated, the MC events are binned to create the predicted spectra.

Additionally, the method to find the best fit point has been changed, due to the increased number of interesting oscillation variables. The fitter now uses an adaptive kernel density method to smooth the posterior and find the maximal point. This method is described in Section 5.

The Bayesian probability function used to fit the data depends on the data sample and flux, cross section, detector, and final state interactions (FSI) systematics, which will be described in subsequent sections. This function has the form:

$$\begin{aligned}
-\ln(P) = & \sum_i^{ND280bins} N_i^p(\vec{b}, \vec{x}, \vec{f}, \vec{d}) - N_i^d + N_i^d \ln[N_i^d/N_i^p(\vec{b}, \vec{x}, \vec{f}, \vec{d})] \\
& + \sum_i^{SK1R_\mu bins} N_i^p(\vec{b}, \vec{x}, \vec{skd}) - N_i^d + N_i^d \ln[N_i^d/N_i^p(\vec{b}, \vec{x}, \vec{skd})] \\
& + \sum_i^{SK1R_e bins} N_i^p(\vec{b}, \vec{x}, \vec{skd}) - N_i^d + N_i^d \ln[N_i^d/N_i^p(\vec{b}, \vec{x}, \vec{skd})] \\
& + \frac{1}{2} \sum_i^{E_\nu bins} \sum_j^{E_\nu bins} \Delta b_i (V_b^{-1})_{i,j} \Delta b_j \\
& + \frac{1}{2} \sum_i^{xsecpars} \sum_j^{xsecpars} \Delta x_i (V_x^{-1})_{i,j} \Delta x_j \\
& + \frac{1}{2} \sum_i^{fsipars} \sum_j^{fsipars} \Delta f_i (V_f^{-1})_{i,j} \Delta f_j \\
& + \frac{1}{2} \sum_i^{nd280det} \sum_j^{nd280det} \Delta d_i (V_d^{-1})_{i,j} \Delta d_j \\
& + \frac{1}{2} \sum_i^{skdet} \sum_j^{skdet} \Delta skd_i (V_{skd}^{-1})_{i,j} \Delta skd_j
\end{aligned} \tag{1}$$

66 where V_{ij} represents covariance matrices constraining systematic parameters la-
67 beled by b for flux, x for cross section, f for FSI, d for ND280 detector, and skd for
68 SK detector. N_i^p is the number of predicted events in a particular bin, given the
69 values of the systematic parameters, and N_i^d is the number of data events.

70 2 Event Selection

71 2.1 ND280 Tracker ν_μ

72 The 2013 tracker ν_μ selection is described in T2K-TN-152 [2]. The charged-current
73 inclusive (CCInc) is divided into three subsamples: charged-current $0-\pi$ (CC 0π),
74 charged-current single π^+ (CC 1π), and charged-current other (CCoth). The sample
75 is subdivided in order to isolate topologies of interest for constraining cross section
76 systematics.

77 The inclusive sample is defined by the following cuts:

- 78 1. Good Data Quality: the global ND280 data quality flag must be good

- 79 2. Bunching: Tracks considered part of the same event must be in the same beam
80 bunch
- 81 3. TPC Quality and Fiducial Volume: There must be at least one track beginning
82 in FGD1's fiducial volume, and entering a TPC with at least 18 vertical TPC
83 clusters
- 84 4. Backwards-going and TPC1 veto: if there is activity in TPC1, or if the end
85 position of the highest momentum track is more upstream than the start po-
86 sition, the track is vetoed
- 87 5. Broken Tracks FGD1: Events are rejected when the muon candidate's z start
88 position is more than 425 mm away from the FGD1 upstream edge and in the
89 same event where at least one "FGD-only" track with its start position out the
90 FGD1 fiducial volume exists.
- 91 6. Muon PID: The highest momentum negative track in the event must be muon-
92 like, according to TPC PID

93 The $CC0\pi$ sample is further defined by rejecting events with any pion recon-
94 structed in the TPC, any electrons or positrons in the TPC, or any Michel electrons
95 or pions reconstructed in the FGDs.

96 The $CC1\pi$ sample is further defined by rejecting events with negative pions or
97 electrons or positrons in the TPC and selecting events where there is one recon-
98 structed positive pion or one Michel electron reconstructed in the TPCs and FGDs.

99 The $CC0th$ sample contains all other $CCInc$ events not in the $CC0\pi$ or $CC1\pi$
100 samples.

101 The binning for the samples chosen for fitting is finer than the binning from the
102 2012 analysis, and is chosen to be as fine as possible while still requiring at least 25
103 MC events in each bin. The binning procedure is described in [2]. The bins are:

- 104 • $CC0\pi$ and $CC0th$
- 105 – p_μ (MeV): 0, 300, 400, 500, 600, 700, 800, 900, 1000, 1250, 1500, 2000,
106 3000, 5000, 30000
- 107 – $\cos\theta$: -1.0, 0.6, 0.7, 0.8, 0.85, 0.9, 0.92, 0.94, 0.96, 0.98, 0.99, 1.0
- 108 • $CC1\pi$

109 – p_μ (MeV): 0, 300, 400, 500, 600, 700, 800, 900, 1000, 1250, 1500, 2000,
110 5000, 30000
111 – $\cos\theta$: -1.0, 0.6, 0.7, 0.8, 0.85, 0.9, 0.92, 0.94, 0.96, 0.98, 0.99, 1.0

112 The data samples are shown in Figure 1. Table 1 gives the number of events in
113 the 0–30 GeV muon momentum region for the three samples and the CC inclusive
114 total sample.

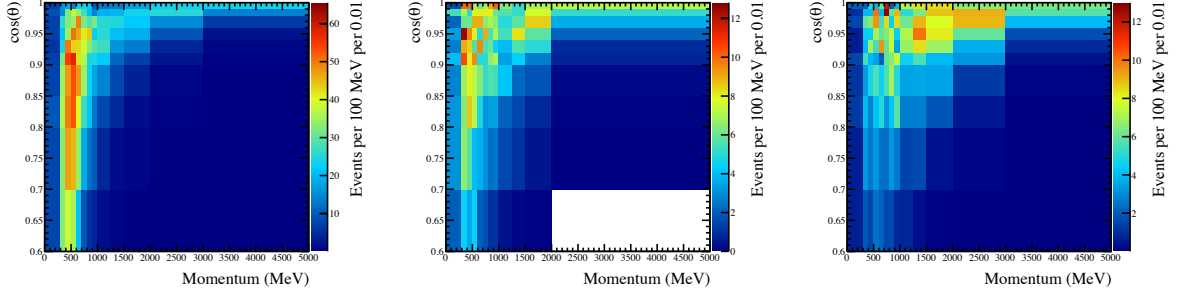


Figure 1: The data samples for this analysis. Shown in (a) is the CC0 π sample, in (b) the CC1 π sample, and in (c) the CCoth sample.

Table 1: Number of data events in the three subsamples and the inclusive sample.

CC0 π	CC1 π	CCo θ	CCInc
17369	4047	4173	25589

115 2.2 SK 1R $_e$ and 1R $_\mu$

116 The selection for the SK data samples in 2013 is described in TN-148 [3]. For the
117 1R $_e$ events, the selection is as follows:

- 118 1. Fully-contained fiducial volume
- 119 2. One ring found by the ring counting algorithm
- 120 3. The ring is identified as electron-like by the PID algorithm
- 121 4. Visible energy (E_{vis}) is greater than 100 MeV

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5. Zero decay electrons
6. Reconstructed neutrino energy (E_{rec}) is less than 1250 MeV
7. fitQun π^0 cut of $\ln(L_{\pi^0}/L_e) < 175 - 0.875 \times m_{\pi^0}$

There are 28 total events in this sample.
For the $1R_\mu$ events, the selection is as follows:

1. Fully-contained fiducial volume event
2. One ring found by the ring counting algorithm
3. The ring is identified by the PID as muon-like
4. Reconstructed momentum is greater than 200 MeV/c
5. Number of decay electrons is equal or less than one

There are 120 total events in this sample.

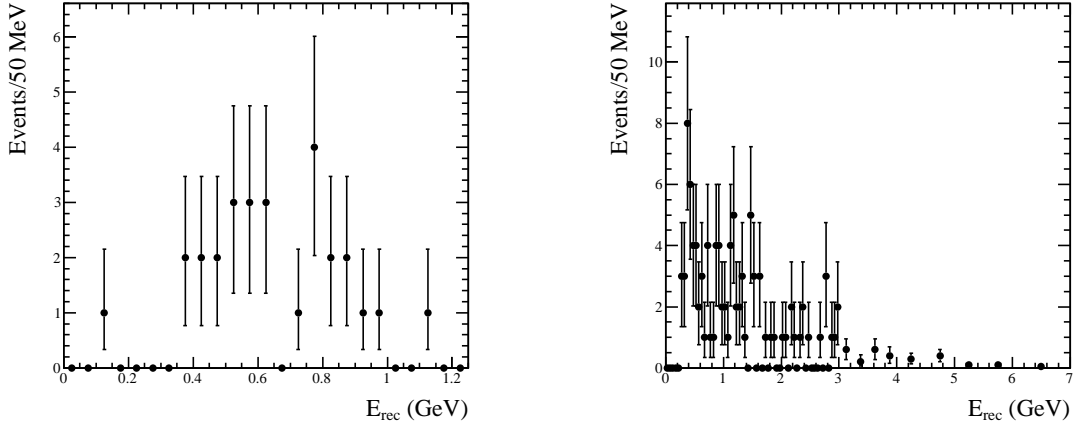


Figure 2: SK data samples for Runs 1–4. Left plot shows $1R_e$ and right plot $1R_\mu$. The fit window for the $1R_\mu$ events extends to 30 GeV, but no events are found above 7 GeV, so the data is only shown up to this limit for clarity.

3 Systematic Errors

3.1 Flux

The flux systematic errors are from TN-099 [4]. The covariance matrix is binned in 11 bins for ν_μ , 5 bins for $\bar{\nu}_\mu$, 7 bins for ν_e , and 2 bins for $\bar{\nu}_e$ for both ND280 and SK as follows, in true neutrino energy (GeV):

- ν_μ : 0.0, 0.4, 0.5, 0.6, 0.7, 1.0, 1.5, 2.5, 3.5, 5.0, 7.0, 30.0
- $\bar{\nu}_\mu$: 0.0, 0.7, 1.0, 1.5, 2.5, 30.0
- ν_e : 0.0, 0.5, 0.7, 0.8, 1.5, 2.5, 4.0, 30.0
- $\bar{\nu}_e$: 0.0, 2.5, 30.0

Figure 3 shows the flux covariance matrix.

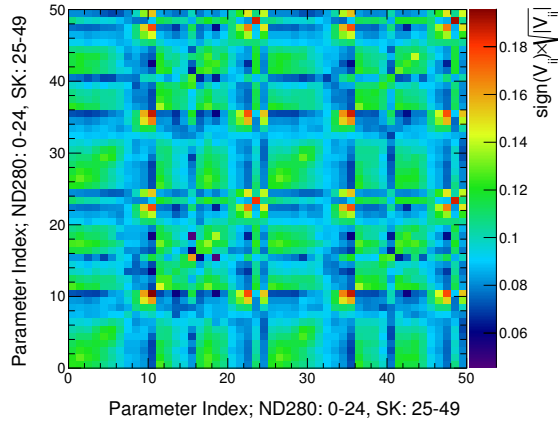


Figure 3: The flux covariance matrix used in the analysis. The bin indices are as follows: ND280 ν_μ (0-10), ND280 $\bar{\nu}_\mu$ (11-15), ND280 ν_e (16-22), ND280 $\bar{\nu}_e$ (23-24), SK ν_μ (25-35), SK $\bar{\nu}_\mu$ (36-40), SK ν_e (41-47), and SK $\bar{\nu}_e$ (48-49), with the energy divisions for the neutrino types given in the text.

Flux weights are applied on an event-by-event basis to the MC events depending on the true neutrino energy of the event.

3.2 Cross Section

The cross section parameterization is largely unchanged from the 2012 analysis. The relevant parameters are given in Table 2. All parameters are independent from one another, excepting M_A^{RES} , CC1 π E1, and NC1 π^0 , which have correlations between them as detailed in [5].

The two types of systematic, shape and normalization, are treated differently. For the shape parameters, the treatment is different between ND280 and SK. At ND280 a spline is created using T2KReWeight for each MC event. This spline is then evaluated for the desired reweighting value of the parameter, and that weight is applied to the event. At SK, splines are created in binned E_{rec} and E_{true} . Each MC event is weighted according to the evaluated spline for the kinematic bin of that event. For the normalization parameters, the event is simply weighted by the value of the parameter.

3.3 ND280 Detector

The detector systematics for this analysis are described in the tracker selection technical note [2]. For an MCMC analysis, the method of reanalyzing every event for every step was computationally prohibitive, taking approximately 3s to reweight each step. Therefore, a covariance matrix approach was used, similar to the 2012 method. The covariance matrix was produced by 2000 throws of the inputs for the detector systematics, and the full detector systematic analysis was used for each throw. The covariance for each bin of the matrix was calculated as

$$V_{ij} = \frac{1}{2000} \sum_{n=1}^{2000} \frac{(N_n^{\text{reweighted},i} - N^{\text{average},i})(N_n^{\text{reweighted},j} - N^{\text{average},j})}{N^{\text{average},j} N^{\text{average},i}} \quad (2)$$

where $N^{\text{average},i}$ is the average of the 2000 throws.

The binning for the detector systematic covariance matrix was chosen to be coarser than the binning used for fitting the data, in order to reduce the number of parameters used in the fit, especially as the size of the detector systematic errors is typically smaller than the size of the flux and cross section errors. The binning chosen for all samples has seven bins in momentum and five bins in $\cos\theta$ and is as follows:

Table 2: NIWG 2012a cross section parameters for the fit, showing the applicable range of neutrino energy, nominal value and prior error. The type of systematic (shape or normalization) is also shown.

Parameter	E_ν Range	Nominal	Error	Class
M_A^{QE}	all	1.21 GeV/ c^2	0.45	shape
M_A^{RES}	all	1.41 GeV/ c^2	0.11	shape
p_F ^{12}C	all	217 MeV/ c	30	shape
E_B ^{12}C	all	25 MeV	9	shape
SF ^{12}C	all	0 (off)	1 (on)	shape
CC Oth shape ND280	all	0.0	0.40	shape
p_F ^{16}O	all	225 MeV/ c	30	shape
E_B ^{16}O	all	27 MeV	9	shape
SF ^{16}O	all	0 (off)	1 (on)	shape
CC Oth shape SK	all	0.0	0.40	shape
W-Shape	all	0.0	0.20	shape
Pionless Delta Decay	all	0.0	0.2	shape
CCQE E1	$0 < E_\nu < 1.5$	1.0	0.11	norm
CCQE E2	$1.5 < E_\nu < 3.5$	1.0	0.30	norm
CCQE E3	$E_\nu > 3.5$	1.0	0.30	norm
CC1 π E1	$0 < E_\nu < 2.5$	1.15	0.43	norm
CC1 π E2	$E_\nu > 2.5$	1.0	0.40	norm
CC Coh	all	1.0	1.0	norm
NC1 π^0	all	0.96	0.43	norm
NC 1 π^\pm	all	1.0	0.3	norm
NC Coh	all	1.0	0.3	norm
NC other	all	1.0	0.30	norm
ν_μ/ν_e	all	1.0	0.03	norm
$\nu/\bar{\nu}$	all	1.0	0.40	norm

- 173 • p_μ (MeV): 0, 300, 500, 600, 700, 1000, 2000, 30000
- 174 • $\cos\theta$: -1, 0.85, 0.9, 0.94, 0.98, 1.0

175 The covariance matrix is shown in Figure 4. In this matrix, bins 0-34 cover the
 176 CC0 π sample, 35-69 the CC1 π sample, and 70-104 the CCoth sample. Within each
 177 sample, the bins iterate over $\cos\theta$ from low to high for the lowest momentum bin,
 178 then from low to high for the second lowest momentum bin, etc.

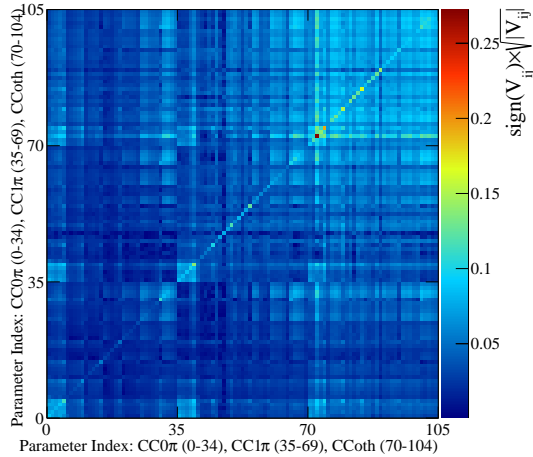


Figure 4: The detector covariance matrix used in the analysis. In this matrix, bins 0-34 cover the CC0 π sample, 35-69 the CC1 π sample, and 70-104 the CCoth sample. Within each sample, the bins iterate over $\cos\theta$ from low to high for the lowest momentum bin, then from low to high for the second lowest momentum bin, etc.

179 To apply this systematic, each event is weighted by the value according to the
 180 bin corresponding to the event's reconstructed momentum and angle.

181 3.4 Final State Interactions (ND280 only)

182 In previous ND280 analyses, the final state interaction systematics were combined
 183 with the detector systematics. However, due to the new treatment of the detector
 184 systematics, the FSI is no longer included. For this analysis, the six FSI parameters
 185 described in [5] (Pion production, 'PION_PROD'; pion absorption 'PION_ABS';
 186 low and high energy charge exchange, 'CEX_LO' and 'CEX_HI'; and low and high

187 energy inelastic interactions, ‘INEL_LO’ and ‘INEL_HI’) are treated as indepen-
 188 dent. That is

$$\begin{aligned}
 W_{FSI}(\sigma_{INEL_LO}, \sigma_{INEL_HI}, \sigma_{PION_PROD}, \sigma_{PION_ABS}, \sigma_{CEX_LO}, \sigma_{CEX_HI}) = \\
 W(\sigma_{INEL_LO}) \times W(\sigma_{INEL_HI}) \times W(\sigma_{PION_PROD}) \times \\
 W(\sigma_{PION_ABS}) \times W(\sigma_{CEX_LO}) \times W(\sigma_{CEX_HI})
 \end{aligned}$$

189 A covariance matrix was created from the variations in Table 1 of [5], and is
 190 shown in Figure 5.

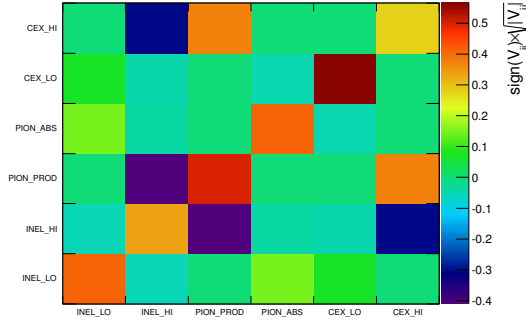


Figure 5: The FSI covariance matrix used in the analysis. The parameters are defined in the NIWG 2012a technical note. [5]

191 For each parameter, a spline is created using T2KReWeight for each MC event.
 192 This spline is then evaluated for the desired reweighting value of the parameter, and
 193 that weight is applied to the event.

194 3.5 SK Detector

195 The SK detector systematics are correlated between the $1R_e$ and $1R_\mu$ samples, as
 196 described in TN-186 [6]. The first 12 parameters are for the $1R_e$ sample, in four
 197 sets of three energy bins (0–0.35; 0.35–0.8; 0.8–1.25 GeV) for the signal ν_e , beam
 198 ν_μ CC, beam ν_e CC, and NC events. The next 6 parameters are for $1R_\mu$: three
 199 energy bins (0–0.4; 0.4–1.1; 1.1–30 GeV) for ν_μ CCQE, one bin for ν_μ CCnQE, one
 200 bin for ν_e CC, and one bin for NC events. The final bin is the energy scale error.

201 The covariance matrix is shown in Figure 6. The matrix contains the FSI+SI errors
 202 for SK.

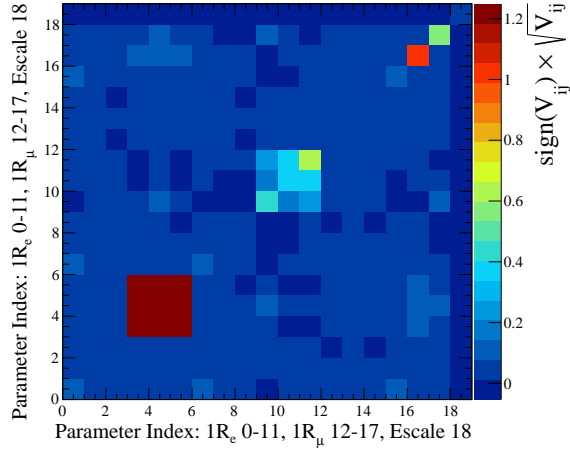


Figure 6: The SK detector covariance matrix used in the analysis. The errors for $1R_e$ are in bins 0-11, $1R_\mu$ in bins 12-17, and the energy scale error in bin 18.

203 4 Monte Carlo Predictions and Pre-fit Data/MC 204 comparison

205 4.1 ND280

206 This analysis uses Production 5E/F MC to generate the predicted spectra for the
 207 samples. The raw MC undergoes two tunings to generate the initial predicted
 208 distributions. First, the events are tuned according to the 11bv3.2 tuning including
 209 Run 4 data. Secondly, the events are tuned for the non-nominal values of the
 210 cross section parameters M_A^{RES} , CC1 π E1, and NC1 π 0 according to a fit to the
 211 MiniBoone CC1 π data as described in [5]. Table 3 gives the number of events in
 212 the 0–30 GeV/c muon momentum region for the data and the MC.

213 The nominal MC prediction for the is shown in Figure 7. The ratio of data
 214 to nominal MC is shown in Figure 8. Projections of the data and nominal MC in
 215 momentum and angle are shown in Figures 9 and 10. Generally, the MC predicts

Table 3: Number of data events in the three subsamples and the inclusive sample.

	CC0 π	CC1 π	CCoth	CCInc
Data	17369	4047	4173	25589
MC	19978.2	4953.2	4544.26	29475.6
Data/MC Ratio	0.869	0.817	0.918	0.868

216 a larger number of events than the data, with the effect more pronounced in the
 217 CC0 π and CC1 π samples than in the CCoth sample.

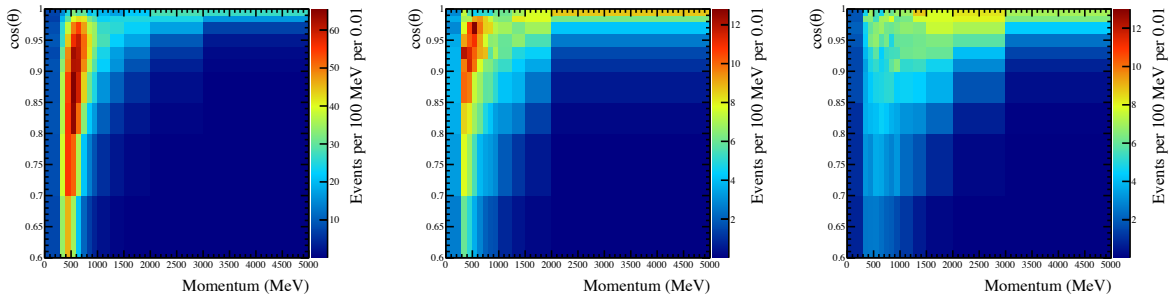


Figure 7: The nominal number of MC predicted events in the p - $\cos\theta$ binning used for the fit. The highest momentum and backwards angle bins are not shown for clarity. Shown in (a) is the CC0 π sample, in (b) the CC1 π sample, and in (c) the CCoth sample.

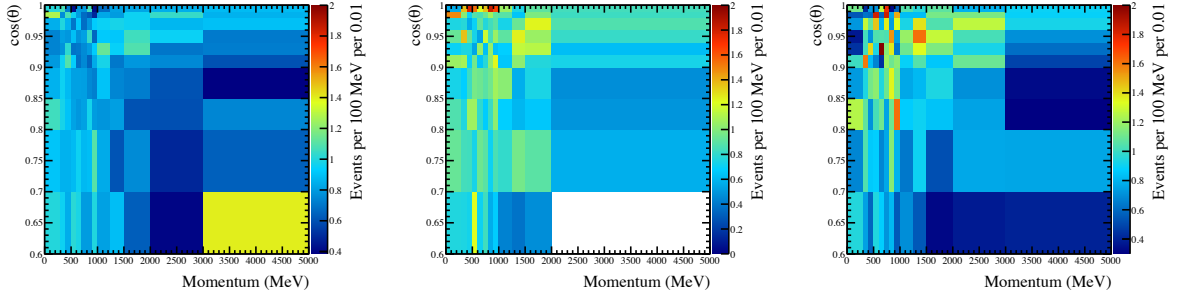


Figure 8: The ratio between the data events and the nominal number of MC events in the p - $\cos\theta$ binning used for the fit. Shown in (a) is the $CC0\pi$ sample, in (b) the $CC1\pi$ sample, and in (c) the $CCo\theta$ sample.

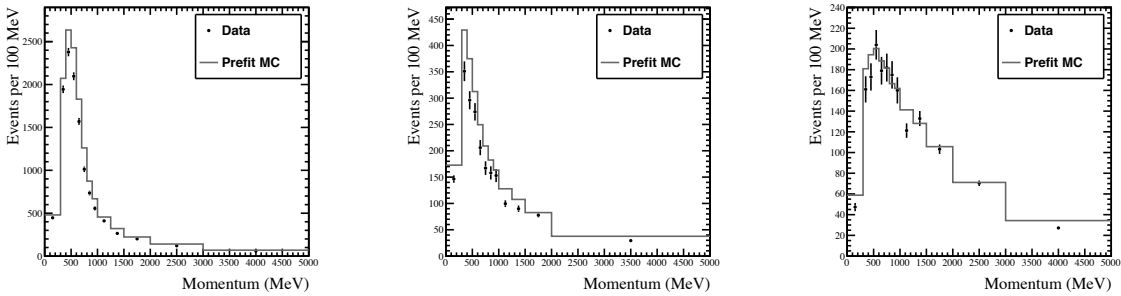


Figure 9: The data and predicted number of MC events projected onto the momentum axis. Shown in (a) is the $CC0\pi$ sample, in (b) the $CC1\pi$ sample, and in (c) the $CCo\theta$ sample.

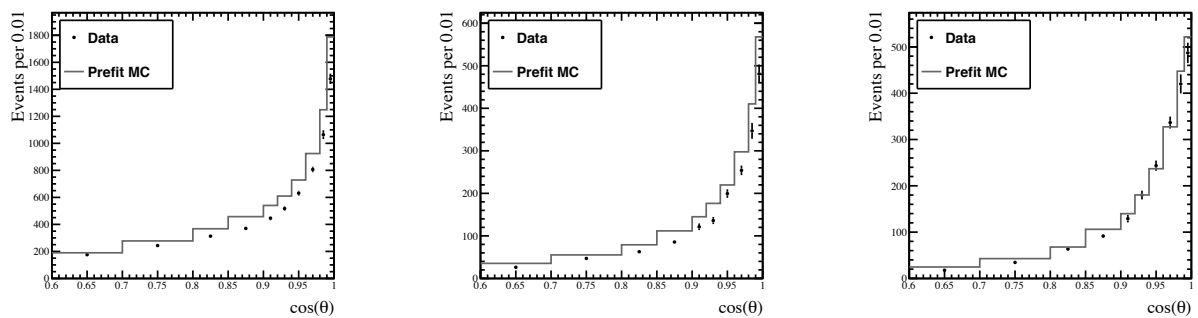


Figure 10: The data and predicted number of MC events projected onto the $\cos \theta$ axis. Shown in (a) is the CC0 π sample, in (b) the CC1 π sample, and in (c) the CCoth sample.

218 4.2 SK 1R_μ

219 This analysis uses SKMC v13a to generate the predicted spectra for the samples.
220 The raw MC undergoes two tunings to generate the initial predicted distributions.
221 First, the events are tuned according to the 11bv3.2 tuning including Run 4 data.
222 Secondly, the events are tuned for the non-nominal values of the cross section pa-
223 rameters M_A^{RES} , CC1π E1, and NC1π0 according to a fit to the MiniBoone CC1π
224 data as described in [5]. Table 4 gives the number of events in the 0–30 GeV
225 reconstructed energy range, broken down by sample type and interaction mode.
226 Additionally, Table 5 shows the number of predicted events by sample type, after
227 tuning by the BANFF v5 ND280 fit.

228 4.3 SK 1R_e

229 This analysis uses SKMC v13a to generate the predicted spectra for the samples.
230 The raw MC undergoes two tunings to generate the initial predicted distributions.
231 First, the events are tuned according to the 11bv3.2 tuning including Run 4 data.
232 Secondly, the events are tuned for the non-nominal values of the cross section pa-
233 rameters M_A^{RES} , CC1π E1, and NC1π0 according to a fit to the MiniBoone CC1π
234 data as described in [5]. Table 6 gives the number of events in the 0–1250 MeV
235 reconstructed energy range, broken down by sample type and interaction mode.
236 Additionally, Table 7 shows the number of predicted events by sample type, after
237 tuning by the BANFF v5 ND280 fit.

238 Table 8 shows the number of data events and predicted MC events and their
239 ratios for the two samples, using PDG2012 values for the oscillation parameters;
240 Figure 11 shows the same graphically as a function of E_{rec} , along with the unoscil-
241 lated spectra. Figure 12 shows a scan of the total rates as a function of oscillation
242 parameters.

Table 4: Top: Oscillated rates for $1R_\mu$, tuned by NIWG2012 for 6.57×10^{20} POT. Oscillation parameters used: $\sin^2 \theta_{23} = 0.5$, $\sin^2 \theta_{13} = 0.0251$, $\sin^2 \theta_{12} = 0.311$, $\Delta m_{12}^2 = 7.5 \times 10^{-5}$ eV, $\Delta m_{32}^2 = 2.4 \times 10^{-3}$ eV, $\delta_{cp} = 0$. Bottom: Unoscillated rates for $1R_\mu$. All mixing angles set to zero.

	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
CCQE	73.583	0.035	4.782	0.002	0.198
CC 1π	41.398	0.029	2.949	0.002	0.081
CC coherent	0.897	0.001	0.247	0.000	0.005
CCn π	6.558	0.004	0.404	0.000	0.001
CC other	2.175	0.003	0.100	0.000	0.001
NC π^0	0.945	0.032	0.054	0.004	0.000
NC $\pi^{+/-}$	4.638	0.131	0.262	0.016	0.000
NC coherent	0.018	0.000	0.001	0.000	0.000
NC other	2.764	0.112	0.158	0.012	0.000
Sample Totals	132.977	0.348	8.956	0.036	0.285
Total Rate	142.603				
	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
CCQE	367.066	0.038	9.710	0.002	0.000
CC 1π	81.343	0.031	4.143	0.002	0.000
CC coherent	2.138	0.001	0.462	0.000	0.000
CCn π	7.465	0.004	0.461	0.000	0.000
CC other	2.304	0.003	0.107	0.000	0.000
NC π^0	0.945	0.032	0.054	0.004	0.000
NC $\pi^{+/-}$	4.638	0.131	0.262	0.016	0.000
NC coherent	0.018	0.000	0.001	0.000	0.000
NC other	2.764	0.112	0.158	0.012	0.000
Sample Totals	468.681	0.353	15.358	0.036	0.000
Total Rate	484.428				

Table 5: Top: Oscillated rates for $1R_\mu$, tuned by BANFF2013 v5 for 6.57×10^{20} POT. Oscillation parameters used: $\sin^2 \theta_{23} = 0.5$, $\sin^2 \theta_{13} = 0.0251$, $\sin^2 \theta_{12} = 0.311$, $\Delta m_{12}^2 = 7.5 \times 10^{-5}$ eV, $\Delta m_{32}^2 = 2.4 \times 10^{-3}$ eV, $\delta_{cp} = 0$. Bottom: Unoscillated rates for $1R_\mu$. All mixing angles set to zero.

	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
Sample Totals	116.642	0.259	7.866	0.024	0.275
Total Rate	125.067				
	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
Sample Totals	431.753	0.263	13.992	0.024	0.000
Total Rate	446.032				

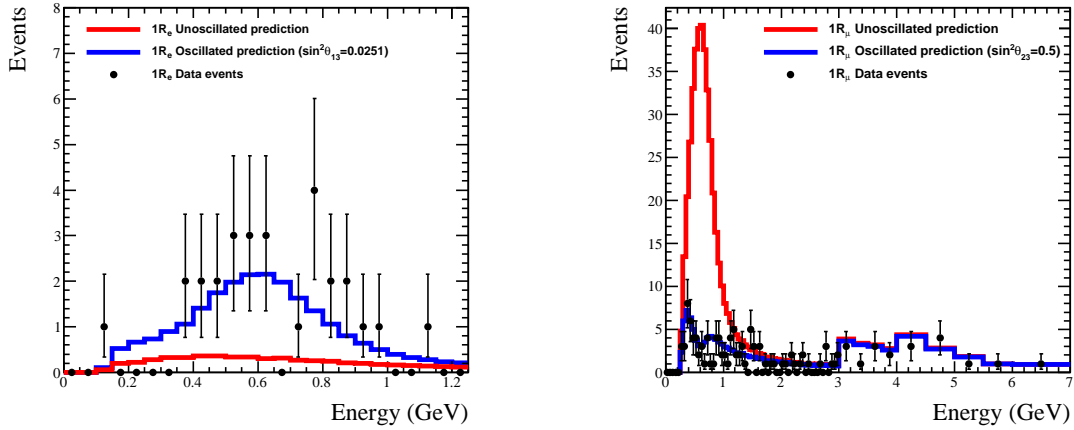


Figure 11: Oscillated (blue) and unoscillated (red) spectra for $1R_\mu$ (left) and $1R_e$ (right) samples. Rates are tuned by NIWG2012 for 6.57×10^{20} POT. Oscillation parameters used: $\sin^2 \theta_{23} = 0.5$, $\sin^2 \theta_{13} = 0.0251$, $\sin^2 \theta_{12} = 0.311$, $\Delta m_{12}^2 = 7.5 \times 10^{-5}$ eV, $\Delta m_{32}^2 = 2.4 \times 10^{-3}$ eV, $\delta_{cp} = 0$.

Table 6: Top: Oscillated rates for $1R_e$, tuned by NIWG2012 for 6.57×10^{20} POT, using the fitQun π^0 cut. Oscillation parameters used: $\sin^2 \theta_{23} = 0.5$, $\sin^2 \theta_{13} = 0.0251$, $\sin^2 \theta_{12} = 0.311$, $\Delta m_{12}^2 = 7.5 \times 10^{-5}$ eV, $\Delta m_{32}^2 = 2.4 \times 10^{-3}$ eV, $\delta_{cp} = 0$. Bottom: Unoscillated rates for $1R_e$. All mixing angles set to zero.

	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
CCQE	0.050	2.276	0.001	0.098	14.989
CC1 π	0.021	0.952	0.000	0.053	2.970
CC coherent	0.000	0.009	0.000	0.007	0.044
CCn π	0.001	0.050	0.000	0.003	0.030
CC other	0.000	0.008	0.000	0.000	0.002
NC π^0	0.475	0.015	0.024	0.002	0.000
NC $\pi^{+/-}$	0.149	0.004	0.008	0.000	0.000
NC coherent	0.181	0.005	0.016	0.001	0.000
NC other	0.329	0.010	0.013	0.001	0.000
Sample Totals	1.207	3.329	0.062	0.165	18.036
Total Rate	22.798				
	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
CCQE	0.050	2.471	0.001	0.104	0.365
CC1 π	0.021	1.010	0.000	0.056	0.040
CC coherent	0.000	0.010	0.000	0.007	0.001
CCn π	0.001	0.052	0.000	0.003	0.000
CC other	0.000	0.008	0.000	0.000	0.000
NC π^0	0.475	0.015	0.024	0.002	0.000
NC $\pi^{+/-}$	0.149	0.004	0.008	0.000	0.000
NC coherent	0.181	0.005	0.016	0.001	0.000
NC other	0.329	0.010	0.013	0.001	0.000
Sample Totals	1.206	3.585	0.062	0.175	0.406
Total Rate	5.434				

Table 7: Top: Rates for oscillated $1R_e$ using the fitQun π^0 cut and tuned by BANFF2013 v5 for 6.57×10^{20} POT. Oscillation parameters used: $\sin^2 \theta_{23} = 0.5$, $\sin^2 2\theta_{13} = 0.1$, $\sin^2 2\theta_{12} = 0.8704$, $\Delta m_{12}^2 = 7.6 \times 10^{-5}$ eV, $\Delta m_{32}^2 = 2.4 \times 10^{-3}$ eV, $\delta_{cp} = 0$. Bottom: Rates for unoscillated $1R_e$ using the fitqun π^0 cut and tuned by BANFF2013 v5. Only $\sin^2 2\theta_{13} = 0.0$; other oscillation parameters remain the same.

	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
Sample Totals	0.946	3.114	0.067	0.152	17.331
Total Rate	21.610				
	ν_μ	ν_e	$\bar{\nu}_\mu$	$\bar{\nu}_e$	ν_e signal
Sample Totals	0.946	3.364	0.067	0.161	0.410
Total Rate	4.947				

Table 8: Number of data events in the SK samples, with MC tuned by NIWG2012 for 6.57×10^{20} POT, using the fitQun π^0 cut for $1R_e$. Oscillation parameters used: $\sin^2 \theta_{23} = 0.5$, $\sin^2 \theta_{13} = 0.0251$, $\sin^2 \theta_{12} = 0.311$, $\Delta m_{12}^2 = 7.5 \times 10^{-5}$ eV, $\Delta m_{32}^2 = 2.4 \times 10^{-3}$ eV, $\delta_{cp} = 0$.

	$1R_e$	$1R_\mu$
Data	28	120
MC	22.798	142.603
Data/MC Ratio	1.228	0.841

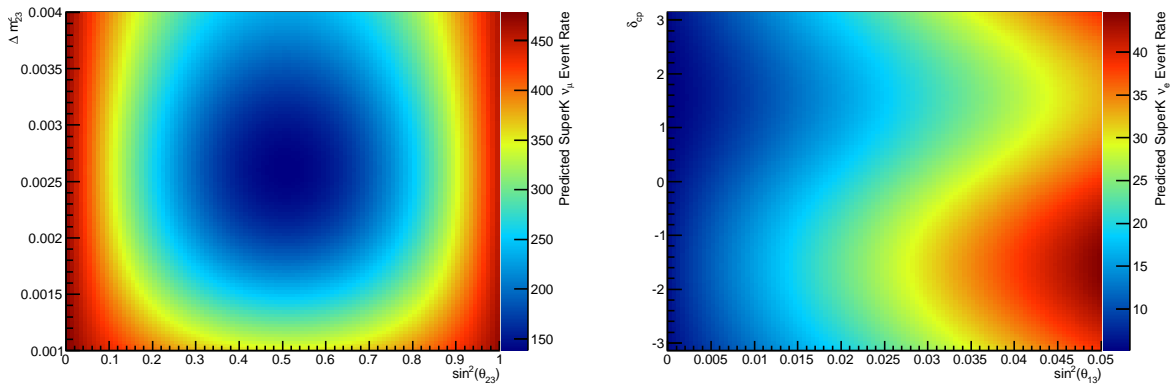


Figure 12: A scan of event rates for Run 1–Run 4 data of 6.57×10^{20} POT. Left shows $1R_\mu$ scanning over $\sin^2 \theta_{23}$ and Δm_{32}^2 ; right shows $1R_e$ scanning over $\sin^2 \theta_{13}$ and δ_{cp} . Other oscillation parameter are fixed at $\sin^2 \theta_{23} = 0.5$ (for $1R_e$), $\sin^2 \theta_{13} = 0.0251$ (for $1R_\mu$), $\sin^2 \theta_{12} = 0.311$, $\Delta m_{12}^2 = 7.5 \times 10^{-5}$ eV, $\Delta m_{32}^2 = 2.4 \times 10^{-3}$ eV (for $1R_e$), $\delta_{cp} = 0$ (for $1R_\mu$)

5 Adaptive Kernel Density Method

The primary result of a Bayesian analysis such as this one is the whole posterior; however, it is desirable to summarize the result with a best fit point. Here, it is defined as the point of maximum probability density in oscillation parameter space. In the previous MCMC analysis, there were only two oscillation parameters of interest, and the best fit point was determined by the maximum bin of the binned 2D posterior in those parameters. This analysis, however, has four oscillation parameters of interest, and as a result, binning the posterior and finding the maximum bin quickly runs into a problem of bin statistics. Therefore, this analysis uses a kernel density estimation (KDE) technique to turn a set of discrete points into a smooth continuous density surface. Minuit [7] is then used to find the point of maximum density.

The kernel density estimator at a point x is defined as:

$$\hat{f}(x) = \frac{1}{nh} \sum_{i=1}^n K\left(\frac{x - x_i}{h}\right) \quad (3)$$

where $x_1, x_2 \dots x_n$ are discrete points and K is the kernel function. This analysis uses a gaussian kernel function, with bandwidth h becoming the σ of the gaussian:

$$\hat{f}(x) = \frac{1}{n} \sum_{i=1}^n \frac{1}{\sigma\sqrt{2\pi}} e^{-\left(\frac{x-x_i}{\sqrt{2}\sigma}\right)^2} \quad (4)$$

For optimum smoothing, we use an adaptive kernel density estimator that adjusts the bandwidth to the local density of points as detailed in [8]. In this method, the bandwidth is inversely proportional to the local density of points—producing a larger bandwidth in areas of low density and a smaller bandwidth in areas of high density—which means that low density areas are not undersmoothed and high density areas are not oversmoothed.

6 Fitter Validation

This analysis has been validated with three methods: using a nominal data set, an ensemble of toy experiments and a series of common fake data sets shared between joint oscillation analyzers.

6.1 Nominal Data Set

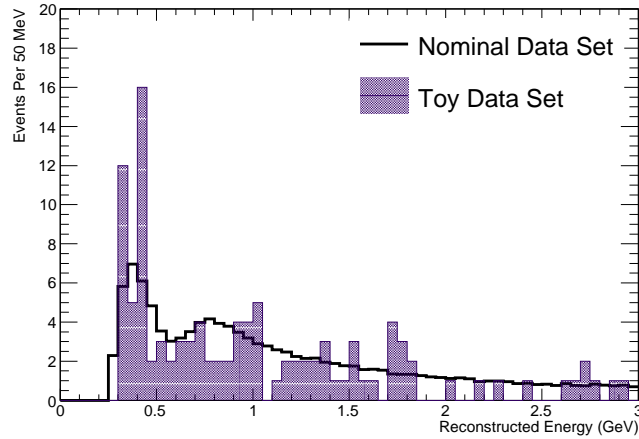


Figure 13: Nominal data set compared to a toy data set.

270 A nominal dataset is defined to be a toy experiment generated from the PDF in
 271 such a way that there are no statistical or systematic fluctuation as illustrated in
 272 Figure 13. This is achieved by reweighting the PDF to nominal values of systematic
 273 parameters, along with the chosen oscillation parameter values, and required protons
 274 on target, but instead of drawing randomly from the PDF, the PDF is considered
 275 as the dataset. This produces a dataset free from statistical fluctuations, which,
 276 when fit, should result in parameters free from bias. Figures 14 and 15 show the
 277 results of a fit to a nominal dataset using 20 million MCMC steps. Figure 14
 278 shows the best fit values of all systematic parameters and their posterior error, and
 279 Figure 15 shows the fractional residual of each systematic parameter. Both plots
 280 show minimal bias in the parameters, and are complimentary to the toy experiment
 281 results in section 6.2.

282 Figures 16 and 17 show credible intervals and best fit values constructed from
 283 the nominal posterior distributions. Also plotted are the true parameter values of
 284 the nominal data set.

285 6.2 Toy Experiments

286 Toy experiments are produced by throwing fake datasets from both SK and ND280
 287 PDFs. Data sets are generated from poisson fluctuations of a particular underlying

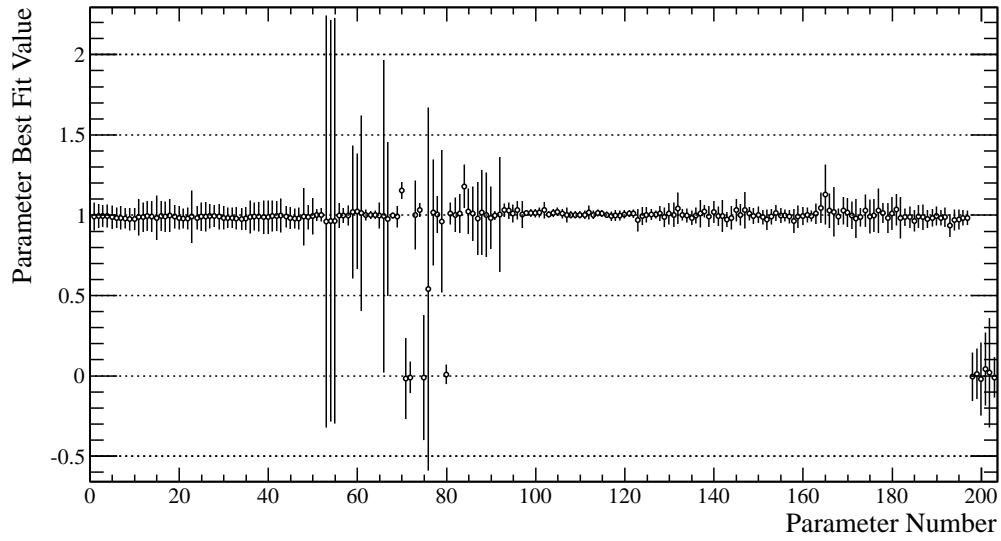


Figure 14: Nominal best fit values from systematic parameters. Error bars are the posterior error. Most parameters have either a true central value of 1 or 0.

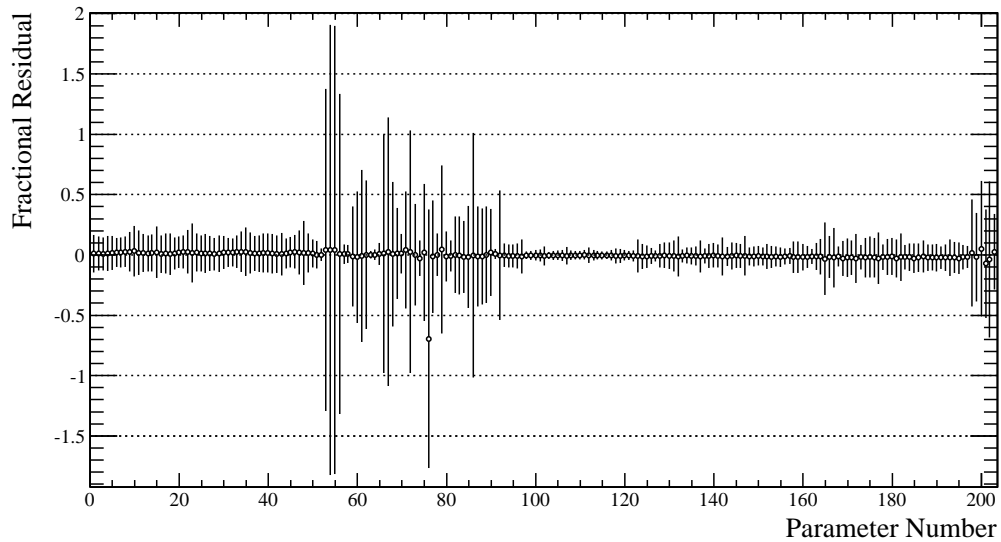


Figure 15: Nominal best fit value subtracted from the true central value, divided by the best fit value. Shows the fractional shift from the true value of each systematic parameter. All parameters stay within 10% of the true value.

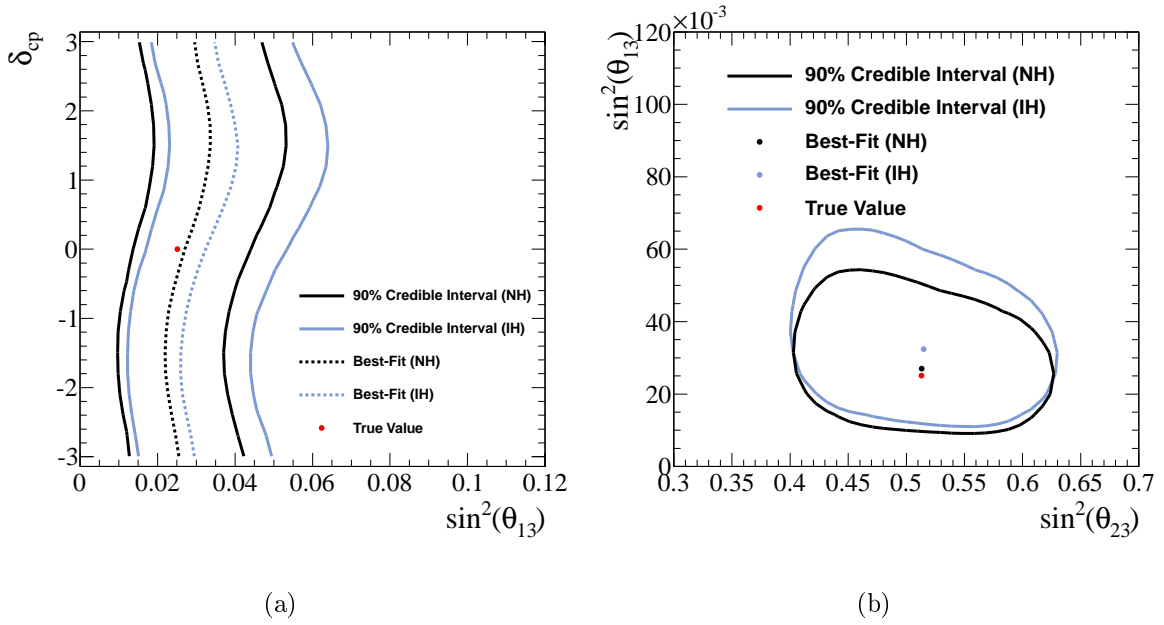


Figure 16: Nominal data fit contours. In (a), best fit lines are constructed in slices of δ_{cp} , and the value at $\delta_{cp} = 0$ is positively offset from the true value due to marginalization of the spectral function as shown in Figure 21.

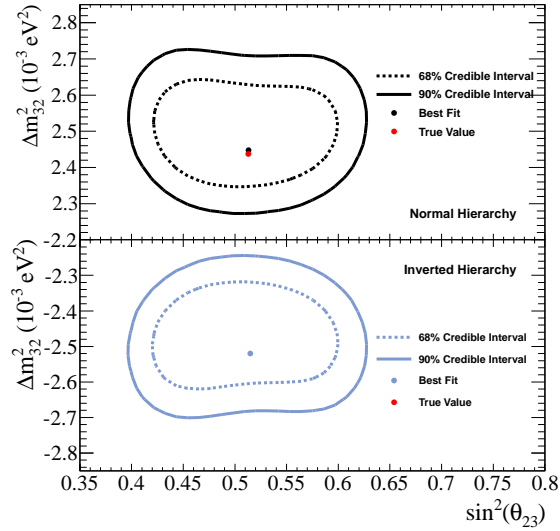


Figure 17: $\sin^2(\theta_{23})$ vs Δm_{32}^2 separated into both hierarchies for a nominal data fit.

288 PDF. To create this underlying PDF, all systematic parameters are kept at nomi-
 289 nal values; however, systematic fluctuation is introduced by randomly throwing the
 290 central values for the systematic penalty terms when performing the fit. In Equa-
 291 tion 5, p^{nom} is the central value which is thrown separately for each toy dataset,
 292 according to the prior PDF for that systematic, including the correlation between
 293 related systematics. Toy experiments are fit using a minimum of 10^6 steps to allow
 294 the production of many fits, whilst ensuring adequate convergence.

$$-\ln P = \sum_{i=0}^n \sum_{j=0}^n \frac{1}{2} (p_i^{prop} - p_i^{nom}) V_{ij}^{-1} (p_j^{prop} - p_j^{nom}) \quad (5)$$

295 To test the fitter for bias and correct error determination, the following definition
 296 is used to construct pull distributions for all parameters:

$$pull = \frac{\mu_{fit} - \mu_{true}}{\sigma_{fit}} \quad (6)$$

297 The best fit and post-fit error for nuisance parameters are extracted from the toy
 298 posterior distributions by constructing a 1D marginal distribution for each paramete-
 299 r and fitting a gaussian to a restricted range defined by $\mu \pm rms$ of the histogram.
 300 For oscillation parameter pulls, the best fit is found using the 3D posterior mode
 301 at $\delta_{cp} = 0$ described in Section 5. Because the 1D posterior distributions for the
 302 oscillation parameters are non-gaussian, the RMS is used as a better estimate of
 303 the error.

304 The post-fit error σ_{fit} of each parameter for every toy experiment was plotted
 305 against the prior error and, where available, the ND280 BANFFv2 post-fit error
 306 value [9] in Figure 18. Shown in Figure 19 is the mean of the pull distributions for
 307 the toys, constructed from Equation 6. The plots in Figure 20 show the oscillation
 308 parameter pull distributions.

309 6.3 Marginalization Induced Biases

310 To extract the best fit values and errors necessary for pull calculations, the 1D
 311 marginal posterior for each parameter is constructed as described in section 6.2.
 312 This method means that for each parameter, the best fit estimate and error is
 313 found marginalizing all other parameters. In doing so, any non-gaussian behavior
 314 and correlations with parameters with non-gaussian behavior can cause apparent

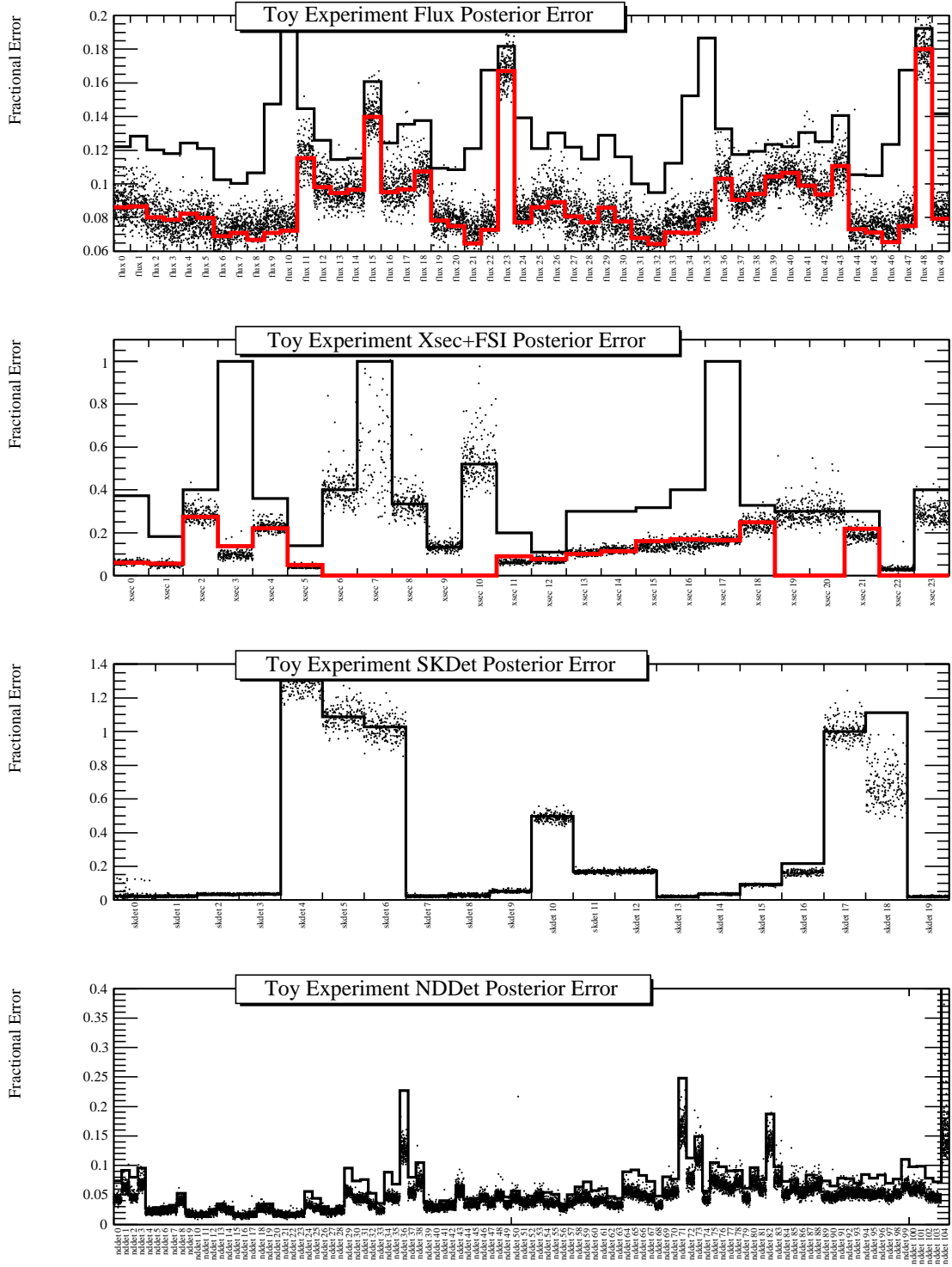


Figure 18: Black line: prior error on parameter. Red line: BANFFv2 post-fit error (where applicable). Black points: posterior error from toy experiments. These plots show how the power to constrain parameter errors is in good agreement with the BANFFv2 post-fit values.

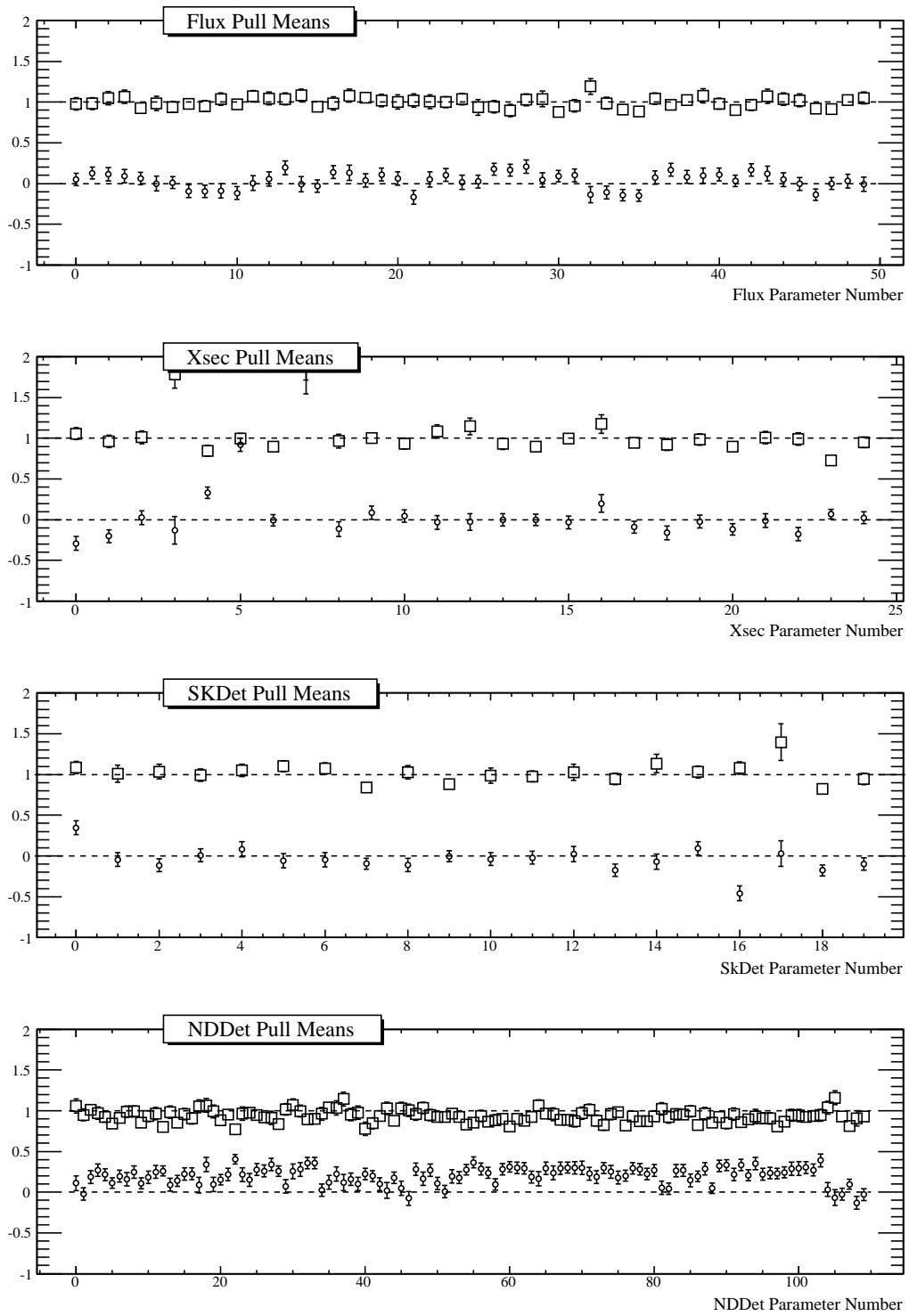


Figure 19: Pull means

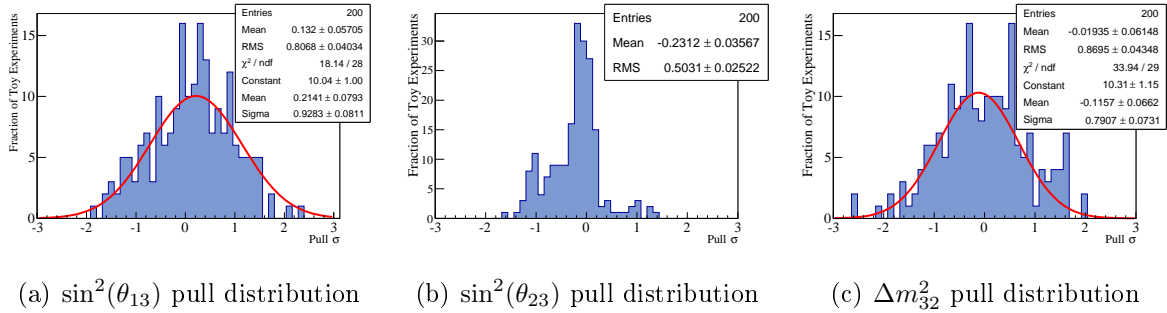


Figure 20: Oscillation parameter pulls for $\delta_{cp} = 0$. Since there is no sensitivity to fit for δ_{cp} , the pull distribution is omitted.

biases in the mean of the pull distributions. In this analysis, there are several parameter pulls which are not within 1σ of 0. These are:

- Quasi-Elastic Axial Mass (M_A^{QE}): this parameter is correlated with the ND280 spectral function parameter, which is both non-gaussian and one-sided (Figure 22).
- Fermi momentum: this parameter is highly correlated with spectral function, which likewise produces an apparent shift as with M_A^{QE} . (Figure 23).
- Spectral function for carbon and oxygen: these are parameters which are defined to be between 0 and 1, and have a distinctly non-gaussian shape in the posteriors.
- SK Energy Scale: the energy scale is a unique parameter in that it shifts the reconstructed energy of events from both SK samples. A high enough shift will cause an event to migrate to an adjacent bin. This behavior causes a non-gaussian posterior distribution for the energy scale parameter. Although the posterior mode shows there is negligible bias, fitting a gaussian to a non-gaussian distribution causes a bias in the resulting pull distribution (Figure 24).
- CCnQE ν_μ Normalization: this parameter is correlated with the oscillation parameters. Since these parameters have non-gaussian posterior distributions, marginalizing them affects the posteriors of correlated parameters. This manifests in the CCnQE ν_μ normalization parameter as a small negative shift in the central value.

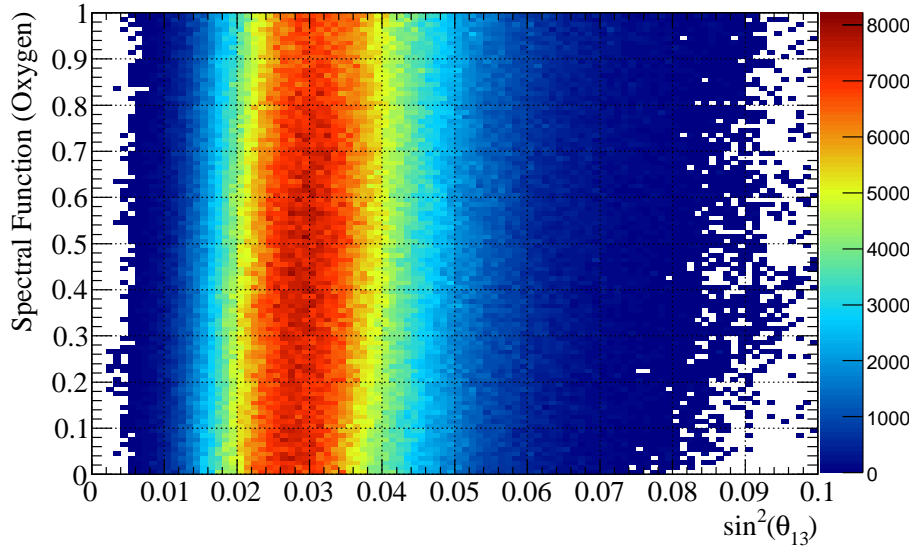


Figure 21: Joint posterior for $\sin^2(\theta_{13})$ and the oxygen spectral function. When marginalizing the spectral function, due to the correlations between both parameters and the boundary at 0, a shift in probability to positive values is caused in the 1D marginal posterior of $\sin^2(\theta_{13})$. Plot constructed from a nominal data set posterior.

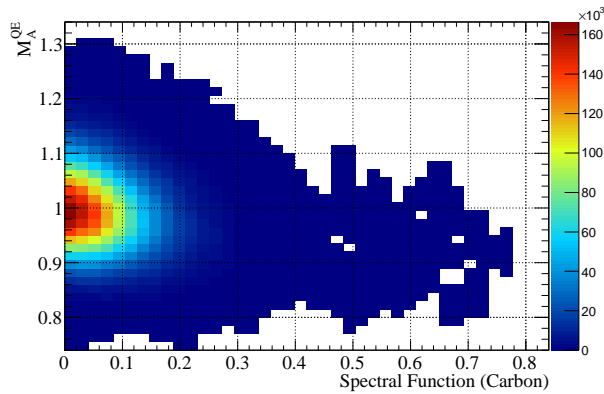


Figure 22: Correlation between quasi-elastic axial mass and spectral function parameters for carbon. Plot constructed from a nominal data set posterior.

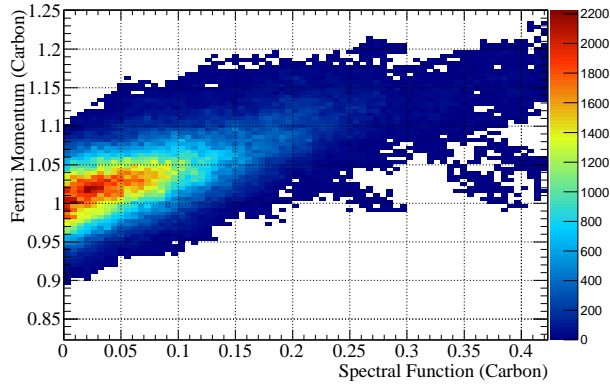


Figure 23: Correlation between fermi momentum and spectral function parameters for carbon. Plot constructed from a nominal data set posterior.

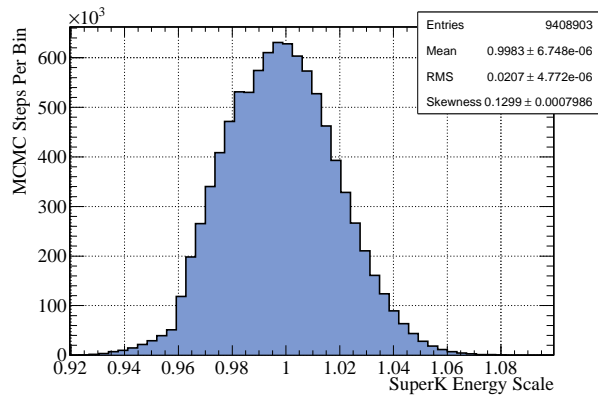


Figure 24: Non-gaussian posterior distribution of the SK energy scale parameter. Plot constructed from a nominal data set posterior.

6.4 Fake Data Set Fits

A series of 6 fake data sets (FDS) were produced by the VaLOR group and distributed to the joint oscillation analyzers. The parameter values used to generate these data sets are denoted in Table 9. Best-fit points for T2K only fits are found using the adaptive kernel density estimator method with δ_{cp} fixed at the VaLOR best fit value. When including the reactor constraint, the best fit is found in 4 dimensions. The $\sin^2(\theta_{13}) - \delta_{cp}$ best fit line is drawn for fits without reactor constraint. It is constructed by finding the maximum density in 3D for steps along the δ_{cp} posterior.

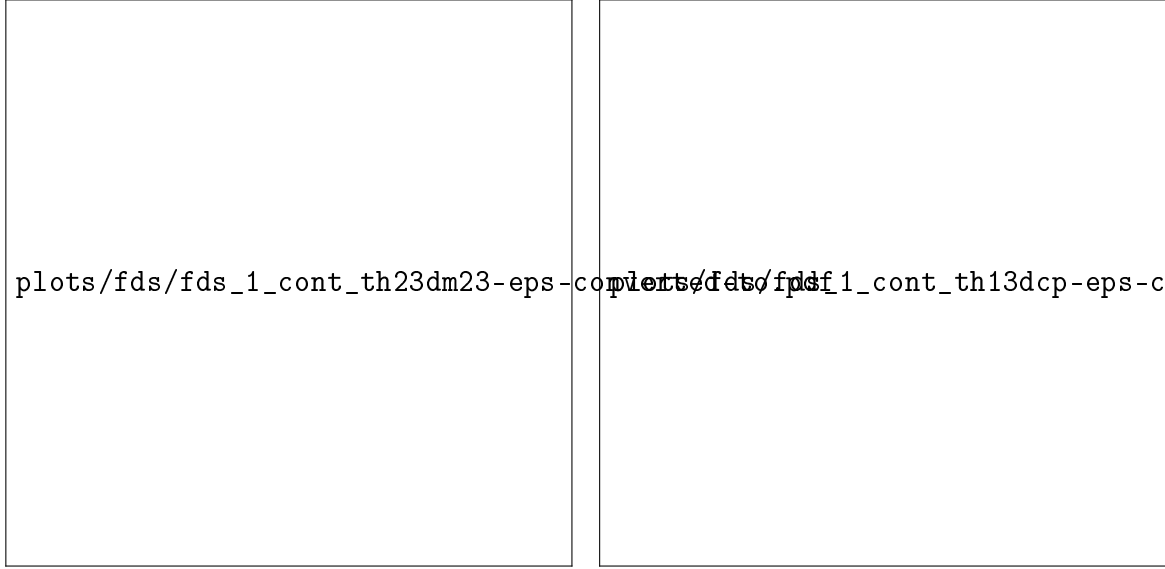
Table 9: Table showing the configuration of the fake data sets provided by the VaLOR group. **Bold** elements highlight the defining parameter value of that data set.

Fake Data Set	Mass Hierarchy	$\sin^2(\theta_{23})$	Δm_{32}^2	$\sin^2(\theta_{13})$	δ_{cp}	Systematic
0	NH	0.513	2.4375	0.0251	0	Nominal
1	NH	0.37	2.4375	0.0251	0	Random Throws
2	NH	0.513	2.75	0.0251	0	Random Throws
3	NH	0.513	2.4375	0.04	0	Random Throws
4	NH	0.513	2.4375	0.0251	$-\pi/2$	Nominal
5	IH	0.513	2.4375	0.0251	0	Nominal

6.4.1 T2K Only Fits

One example of the 2D contours in $\sin^2(\theta_{23}) - \Delta m_{32}^2$ and $\sin^2(\theta_{13}) - \delta_{cp}$ is shown in Figure 25. The contours for all other datasets are contained in Appendix A. There is generally good agreement between the two fitters, and between the fitters and the input values, as shown in Table 10. Generally, the MaCh3 fitter finds a higher value of $\sin^2(\theta_{13})$ than the VaLOR fit; this difference is consistent with the size of shifts coming from the marginalization over spectral function.

There is an interesting discrepancy between the two fitters in FDS1, where the input value was an off-maximal value of $\sin^2 \theta_{23} = 0.37$. MaCh3 finds the best fit value in the lower octant, where VaLOR finds the best fit value in the upper octant. This discrepancy is explained in Figure 26, which shows the full marginal posterior in $\sin^2 \theta_{23}$, and there is greater posterior density in the lower octant. However, if



(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 25: Fake Data Set 1

358 the posterior is restricted to a smaller window around the best fit points in $\sin^2 \theta_{13}$
 359 and Δm_{32}^2 —a technique similar to the profiling method of the minimizer fit—there
 360 is greater posterior density in the upper octant. Thus, the difference in the best fit
 361 points comes from the difference in the methods of the fitters.



Figure 26: The full marginal posterior of $\sin^2(\theta_{23})$ for FDS1 (cyan) compared with a “restricted posterior” constructed from MCMC steps taken only from a small region around the best fit point of $\sin^2(\theta_{13})$ and Δm_{32}^2 (darker blue). Restricting the posterior to points only around the most probable regions of the marginalized oscillation parameters is similar in approach to the frequentist profiling technique. Red arrows indicate the 1D posterior mode for each distribution. This exercise highlights the difference in best fit points between analyses.

Table 10: Normal hierarchy best-fit comparison table between MaCh3 and VaLOR for all fake data sets with no reactor constraint. MaCh3 values of Δm_{32}^2 have been rescaled to enable comparison with the Fogli convention used by VaLOR.

FDS	$\Delta m_{32}^2 \times 10^{-3}$	$\sin^2(\theta_{23})$	$\sin^2(\theta_{13})$	δ_{cp}
0 True	2.4375	0.513	0.0251	0
0 VaLOR	2.413	0.513	0.0364	-0.0825
0 MaCh3	2.419	0.522	0.0385	-0.0825
1 True	2.4375	0.37	0.0251	0
1 VaLOR	2.327	0.619	0.0152	1.585
1 MaCh3	2.268	0.409	0.0259	1.585
2 True	2.75	0.513	0.0251	0
2 VaLOR	2.578	0.508	0.0185	-0.0179
2 MaCh3	2.598	0.508	0.0200	-0.0179
3 True	2.4375	0.513	0.04	0
3 VaLOR	2.583	0.568	0.0572	1.087
3 MaCh3	2.578	0.535	0.0642	1.087
4 True	2.4375	0.513	0.0251	0
4 VaLOR	2.466	0.526	0.0464	-2.564
4 MaCh3	2.468	0.526	0.0494	-2.564
5 True	2.4375	0.513	0.0251	0
5 VaLOR	2.53	0.511	0.0246	2.367
5 MaCh3	2.56	0.511	0.0232	2.367

6.4.2 T2K with Reactor Constraint

The application of a prior constraint from reactor experiments can provide increased sensitivity to the oscillation parameters. For these toy datasets, the constraint was applied as the true input $\sin^2 2\theta_{13}$ constraint, ± 0.01 , the PDG 2013 error. Figure 27 shows one example of this process, for FDS 0. The plots for the other FDS are in Appendix A.



Figure 27: Fake Data Set 0

6.5 Comparison with BANFF Matrix Fit

The fitter can also be configured to constrain the SK flux and cross-section uncertainties using the BANFF matrix instead of using the ND280 data directly. A comparison of the contours and best fit points (Figure 28) produced with both methods when fitting fake data set 5 was made and the results show negligible difference between the two results.

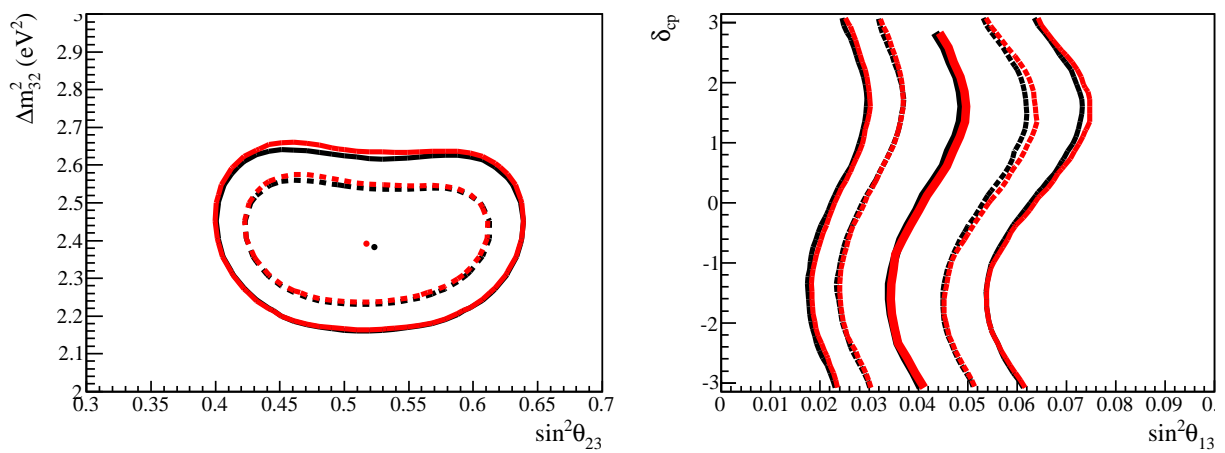


Figure 28: The comparison of contours and best fit points between the BANFF extrapolation (red) and the simultaneous fit with ND280 data methods. The contours suggest that the simultaneous fitting method yields a smaller uncertainty, however the difference is considered negligible.

7 Fit Results

This section details the results obtained from fitting simultaneously the SK and ND280 Run 1–4 data, totalling 6.57×10^{20} and 5.9×10^{20} protons on target respectively. For these fits, the solar sector oscillation parameters are $\sin^2 \theta_{12} = 0.311 \pm 0.017$ and $\Delta m_{21}^2 = 7.5 \pm 0.2 \times 10^{-5} \text{ eV}^2$.

7.1 T2K Run 1–4 Data Fit

The data samples were first fit using T2K data alone, with a Markov chain of 1.8×10^7 steps after burn-in. For this type of fit, since there is little constraint in δ_{cp} , the best fit point is found by fixing δ_{cp} at 21 steps in its range, and fixing the parameter in the 4D adaptive kernel estimation to find the best fit in 3D for the other oscillation parameters. Table 11 shows the best fit parameters in the $\delta_{cp} = 0$ slice. Credible regions are produced in 2D for several different sets of parameters; these contours are produced marginalized over all other parameters, but constructed separately for normal and inverted hierarchies. Figure 7.1 shows the contours in $\sin^2(\theta_{23})$ – Δm_{32}^2 space. Figure 30(a) shows the contours in $\sin^2(\theta_{13})$ – δ_{cp} space, where the best fit is shown as a line connecting the best fit values in the slices of δ_{cp} . Figure 30(b) shows the contours in $\sin^2(\theta_{23})$ – $\sin^2(\theta_{13})$.

Figure 31 shows the 1D credible intervals for $\sin^2(\theta_{13})$, $\sin^2(\theta_{23})$, and Δm_{32}^2 , where all other parameters are marginalized.

Figure 7.1 shows the best fit spectra of the Run 1–4 SK data constrained by the ND280 data, for $1R_\mu$ and $1R_e$ samples. The best fit spectra is determined via a marginalization method. The fit posterior is sampled randomly 2500 times, and with each sample the parameter values are used to calculate the expected event rate per bin of the energy spectra; this is essentially marginalizing over all parameters, oscillation included, to find the posterior distribution in each energy bin. The combination of all the samples creates a distribution of event rates for each bin. Finally, for each bin, a gaussian is fitted around the peak of the event rate distribution, and the mean of the fit is taken to be the predicted value for that bin. Most bins take on a gaussian shape, but in some bins, especially near the oscillation maxima in the $1R_\mu$ sample, the distribution is non-gaussian, due to the influence of the nearby physical boundary in $\sin^2 \theta_{23}$.

Table 11: Best-fit values for oscillation parameters extracted from the marginal posterior of the Run 1–4 data.

	$ \Delta m_{32}^2 $	$\sin^2(\theta_{23})$	$\sin^2(\theta_{13})$	δ_{cp}
Normal Hierarchy	2.491	0.520	0.0377	0 (fixed)
Inverted Hierarchy	2.571	0.520	0.0454	0 (fixed)

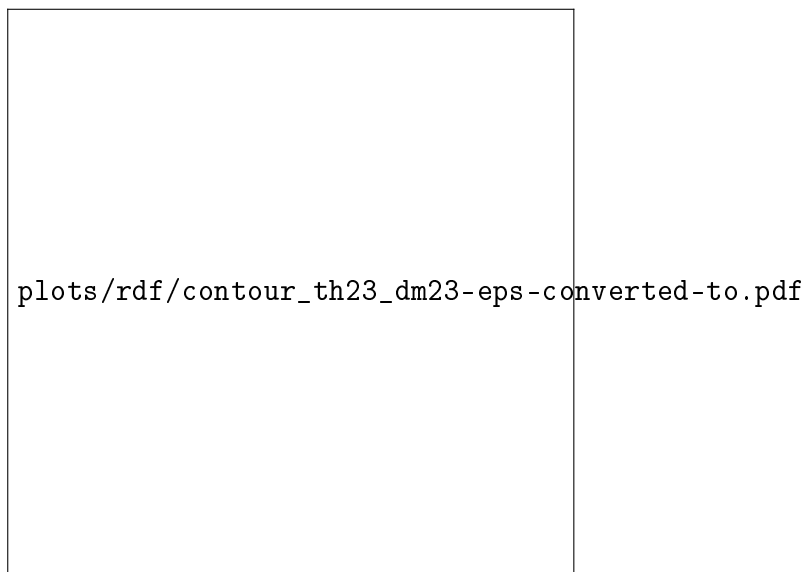


Figure 29: Run 1–4 data fit 2D contours in $\sin^2(\theta_{23})$ – Δm_{32}^2 space.

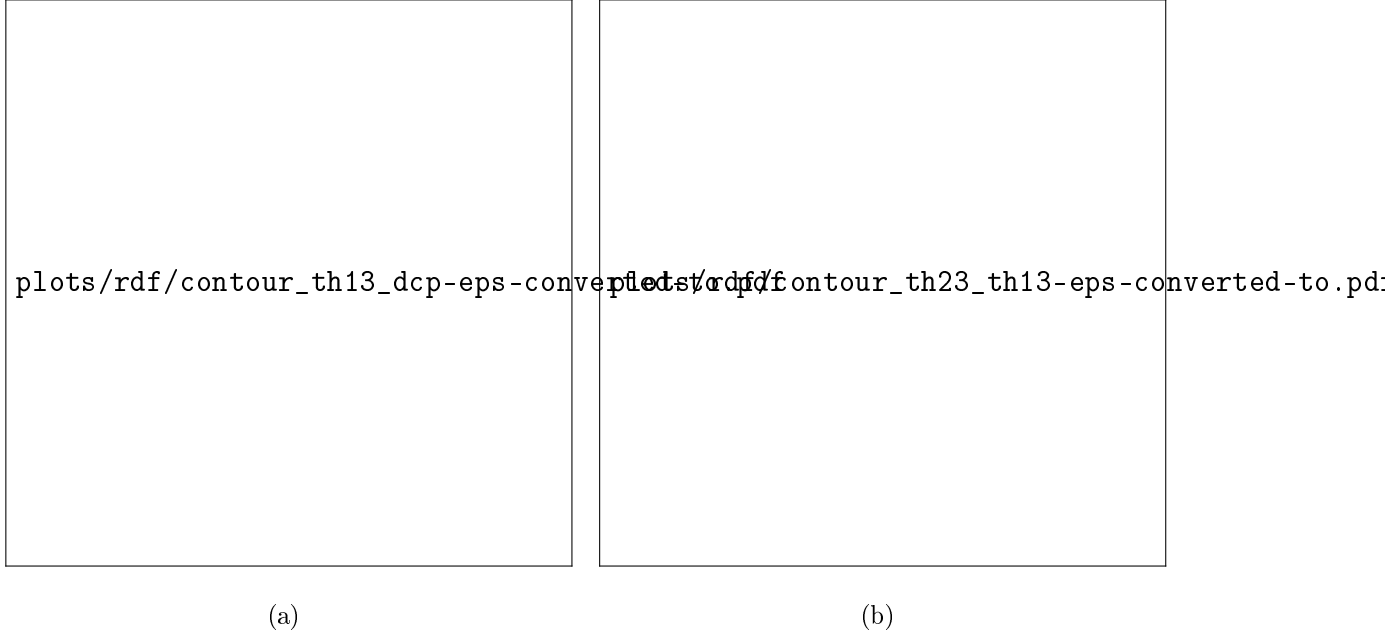


Figure 30: Run 1–4 data fit 2D contours in (a) $\sin^2(\theta_{13})-\delta_{cp}$ space and (b) $\sin^2(\theta_{23})-\sin^2(\theta_{13})$ space.

405 Figures 33 and 34 show the momentum and angle distributions for ND280 with
 406 the pre-fit MC prediction and post-fit spectra, calculated in the same way as for the
 407 SK spectra.

408 A goodness-of-fit is calculated as in [?], where at each chain sample used for
 409 the best fit spectra, a fake dataset is thrown from the MC prediction for that
 410 sample. The log likelihood ratio between the fake dataset and the MC prediction is
 411 calculated, as is the log likelihood ratio between the real data and the MC prediction.
 412 A p-value is calculated as the percentage of samples for which the data better fit
 413 the MC prediction than the fake data. In order to have $N > 10$ in each bin,
 414 a requirement for this method, the $1R_\mu$ sample is rebinned into five bins (0–0.4;
 415 0.4–0.7; 0.7–1.0; 1.0–2.0; and 2.0–30.0 GeV) and the $1R_e$ sample is considered as
 416 one bin only. The ND280 sample is considered in the bins used to fit the data.
 417 This means that the overall p-value is completely dominated by the ND280 sample.
 418 Figure 35 shows the ND280, $1R_\mu$, $1R_e$, and total distributions for the quantity
 419 $\ln L_{data} - \ln L_{throw}$; the p-value is the percentage of this distribution above zero. The
 420 p-values are: ND280-only, 0.044; SK $1R_e$, 0.32; SK $1R_\mu$, 0.35; and all samples, 0.036.
 421 These values indicate no disagreement with data for the SK samples. The value for

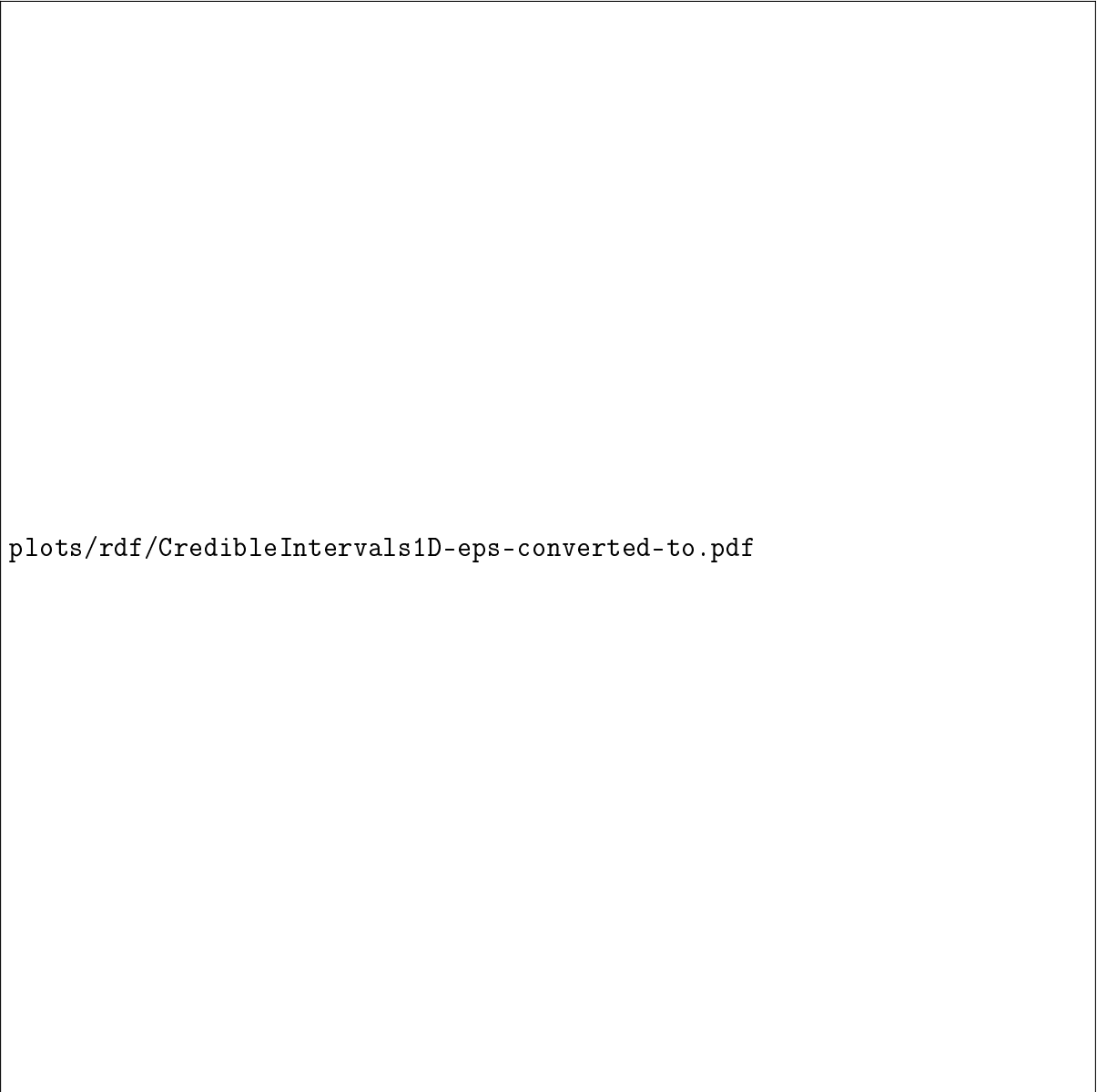


Figure 31: Credible intervals in 1D for $\sin^2(\theta_{13})$, $\sin^2(\theta_{23})$, and $|\Delta m_{32}^2|$. The PDFs for the angles are shown for normal hierarchy, inverted hierarchy, and marginalized over the hierarchies. The PDF for the mass splitting is shown only for normal and inverted hierarchies. The 90% credible intervals are shown by the dotted lines and given in the plot legends.

plots/rdf/bfs_rdf1_t2konly-eps-converted-to.pdf

Figure 32: Run 1–4 data best fit spectra for SuperK $1R_\mu$ and $1R_e$ samples.

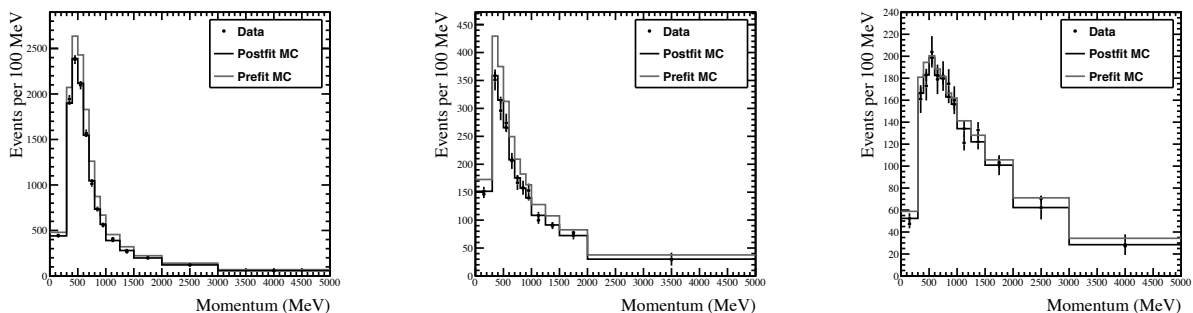


Figure 33: The data (black points) and pre-fit (grey) and post-fit (black) predicted number of MC events projected onto the momentum axis. Shown left-to-right are the $CC0\pi$ sample, the $CC1\pi$ sample, and the $CCo\theta$ sample.

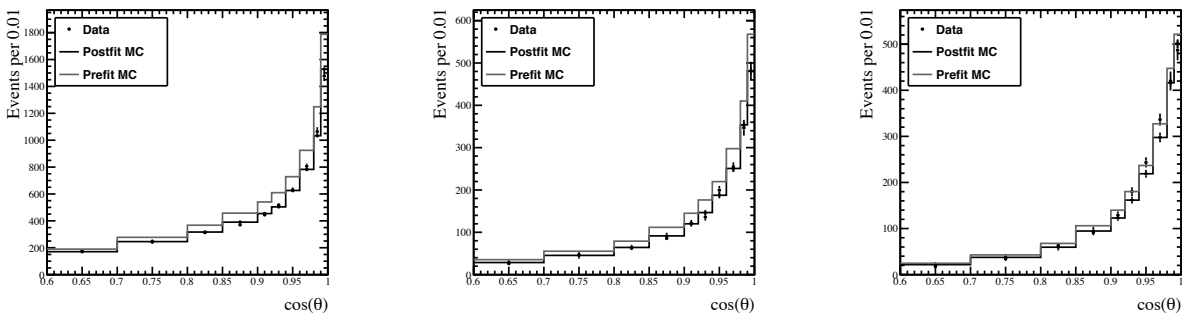


Figure 34: The data (black points) and pre-fit (grey) and post-fit (black) predicted number of MC events projected onto the $\cos\theta$ axis. Shown left-to-right are the $CC0\pi$ sample, the $CC1\pi$ sample, and the $CCo\theta$ sample.

422 the ND280 samples is somewhat low, indicating some disagreement; however, this is
423 a known effect (see [9]), and the agreement between the results of the ND280 fits for
424 both MaCh3 and the minimizer BANFFv2 fit and the data are nearly equivalent.



Figure 35: Goodness-of-fit distributions for the three different samples in the fit and the summed total. The p-value is the percentage of each distribution which is greater than zero.

7.2 T2K Run 1–4 Data Fit With Reactor Constraint

The data samples were then fit using T2K data in combination with the PDG 2013 reactor gaussian constraint of $\sin^2(2\theta_{13}) = 0.095 \pm 0.01$, with a Markov chain of 3.168×10^7 steps after burn-in. For this type of fit, the best fit point is found with a 4D adaptive kernel estimate of the oscillation parameters of interest. Table 12 shows the best fit parameters. Credible regions are produced in 2D for several different sets of parameters; these contours are produced marginalized over all other parameters, but constructed separately for normal and inverted hierarchies. Figure 36 shows the contours in $\sin^2(\theta_{23})-\Delta m_{32}^2$ space. Figure 37(a) shows the contours in $\sin^2(\theta_{13})-\delta_{cp}$ space. Figure 37(b) shows the contours in $\sin^2(\theta_{23})-\sin^2(\theta_{13})$.

Figure 38 shows the 1D credible intervals for $\sin^2(\theta_{13})$, $\sin^2(\theta_{23})$, and Δm_{32}^2 , where all other parameters are marginalized.

Table 12: Best-fit values for oscillation parameters extracted from the marginal posterior of the Run 1-4 data fit with reactor constraint.

	$ \Delta m_{32}^2 $	$\sin^2(\theta_{23})$	$\sin^2(\theta_{13})$	δ_{cp}
Normal Hierarchy	2.510	0.527	0.0247	-1.551
Inverted Hierarchy	2.553	0.531	0.0249	-1.596

plots/rdf/reactor_contour_th23_dm23-eps-converted-to

Figure 36: Run 1–4 data fit with reactor constraint 2D contours in $\sin^2(\theta_{23})-\Delta m_{32}^2$ space.

The goodness-of-fit was repeated for the reactor constrained data. Figure 41 shows the ND280, $1R_{\mu}$, $1R_e$, and total distributions for the quantity $\ln L_{data} - \ln L_{throw}$; the p-value is the percentage of this distribution above zero. The p-values are: ND280-only, 0.044; SK $1R_e$, 0.44; SK $1R_{\mu}$, 0.33; and all samples, 0.042. These values indicate no disagreement with data for the SK samples. It is interesting that the p-value for SK $1R_e$ increases slightly for this fit as compared to the T2K-only fit, despite the fact that the predicted number of events for the T2K-only fit is closer to the number of data events. This is due to the fact that the reactor constraint

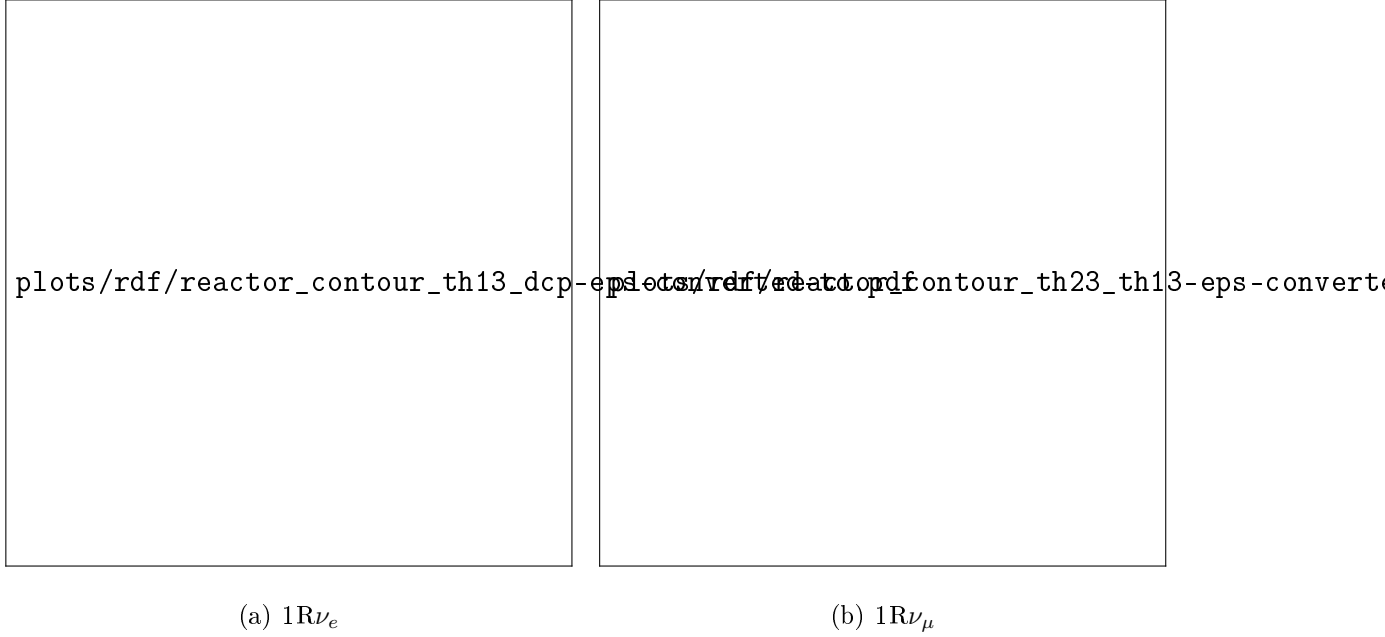


Figure 37: Run 1-4 data fit with reactor constraint 2D contours in (a) $\sin^2(\theta_{13})-\delta_{cp}$ space and (b) $\sin^2(\theta_{23})-\sin^2(\theta_{13})$ space.

445 narrows the distribution of allowed events significantly, and therefore the predicted
 446 spectra from the throws do not move as far from the data point as they do for the
 447 T2K-only fit.

448 The addition of the reactor constraint to the T2K data also produces some sensi-
 449 tivity in δ_{cp} . Figure 42 shows the δ_{cp} posterior for the normal hierarchy, considered
 450 alone; the inverted hierarchy, considered alone; and marginalizing over the hierar-
 451 chies. Figure 43 shows the δ_{cp} posterior when considering the normal and inverted
 452 hierarchies jointly. Each of these methods answers a slightly different question about
 453 the preferred region for the value of δ_{cp} , and caution should be used when using these
 454 plots to describe them correctly. Table 13 enumerates the 90% allowed regions for
 455 the different methods.

456 The constraint on δ_{cp} can also be considered separating the lower and upper oc-
 457 tant, as in Figure 4 of [?]. This is shown in Figure 44. Unlike the MINOS data, the
 458 best fit point remains constant at $\approx -\pi/2$ for all of the choices of octant and hier-
 459 archy. However, some are more preferred than others; the inverted hierarchy/lower
 460 octant choice is excluded completely at the 68% level and nearly completely at the
 461 90% level. By contrast, nearly all of the normal hierarchy/upper octant is allowed

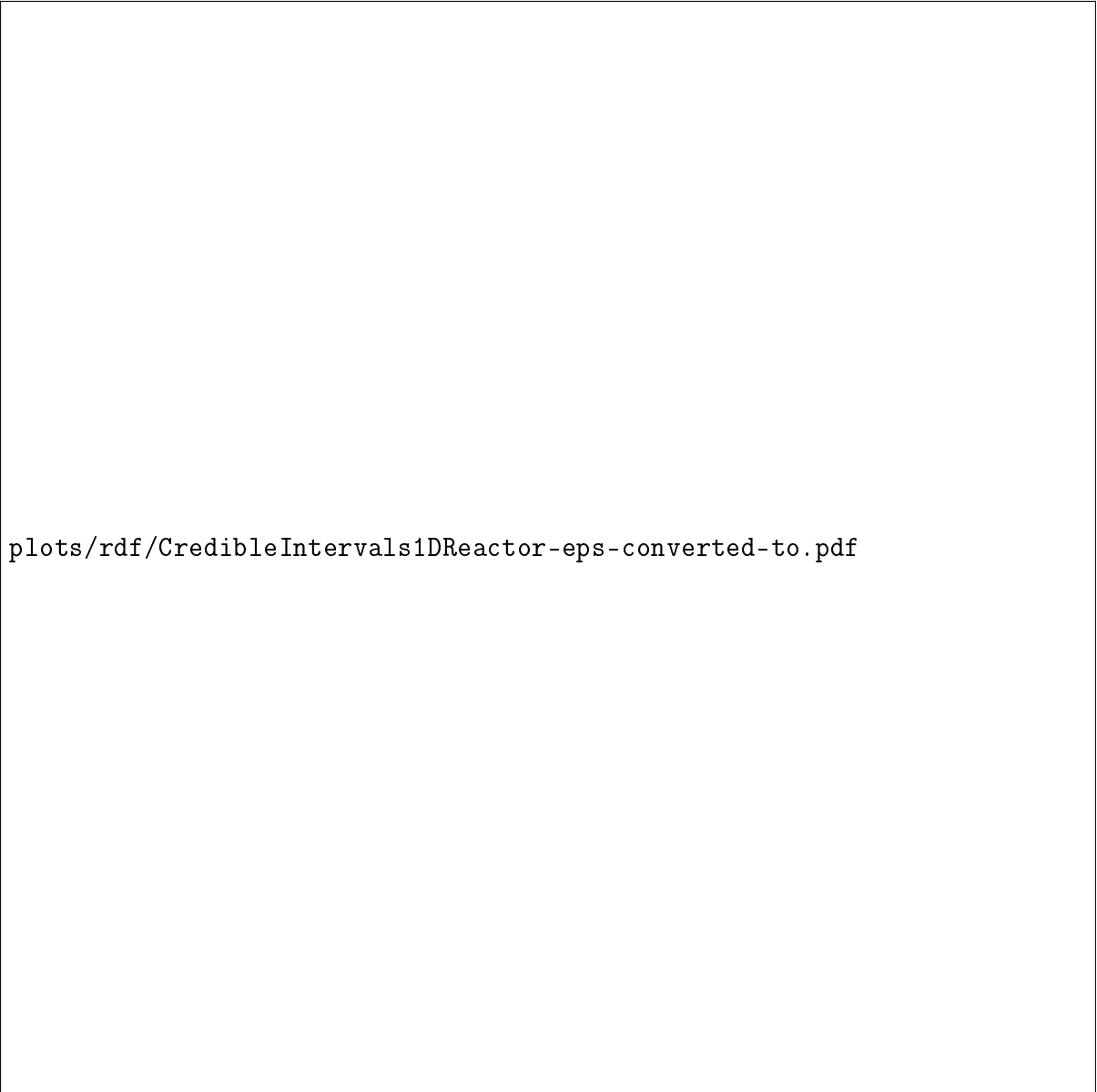

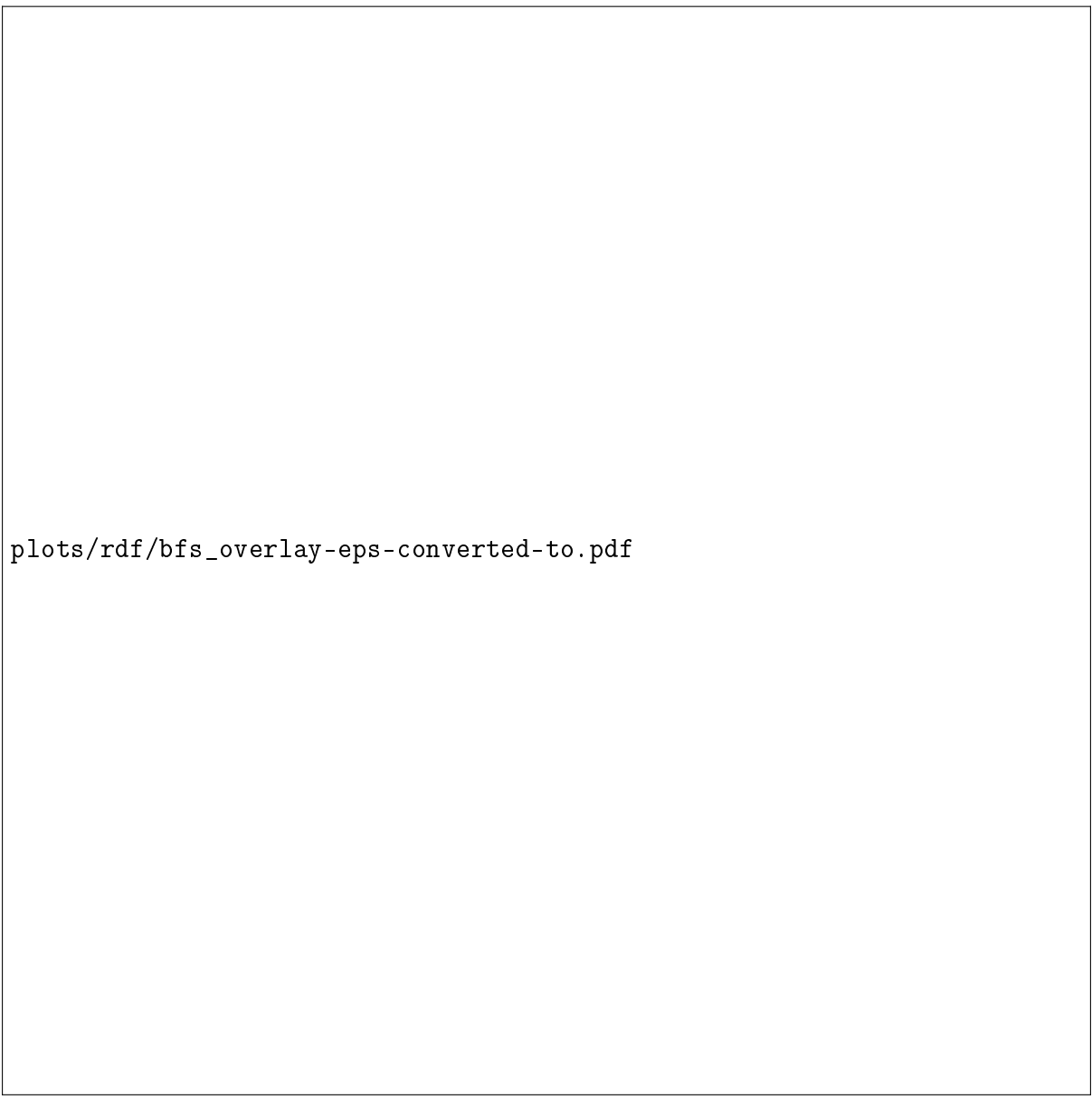


Figure 38: Credible intervals in 1D for $\sin^2(\theta_{13})$, $\sin^2(\theta_{23})$, and $|\Delta m_{32}^2|$, using the reactor constraint. The PDFs for the angles are shown for normal hierarchy, inverted hierarchy, and marginalized over the hierarchies. The PDF for the mass splitting is shown only for normal and inverted hierarchies. The 90% credible intervals are shown by the dotted lines and given in the plot legends.



plots/rdf/bfs_rdf1_reactor-eps-converted-to.pdf

Figure 39: Run 1–4 data best fit spectra for SuperK ν_μ and ν_e samples with reactor constraint.



plots/rdf/bfs_overlay-eps-converted-to.pdf

Figure 40: Comparison of best fit spectra of T2K data with and without reactor constraint applied.



Figure 41: Goodness-of-fit distributions for the three different samples in the fit and the summed total. The p-value is the percentage of each distribution which is greater than zero.

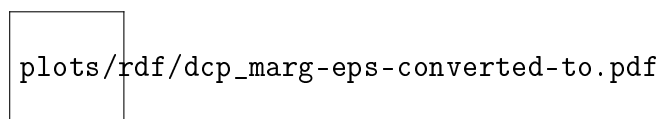


Figure 42: The posterior probability for δ_{cp} , marginalized over all other parameters. The red curve shows the posterior for the normal hierarchy only; the blue curve for the inverted hierarchy only; and the black curve marginalized over the hierarchies. The grey bands show the 68% and 90% credible intervals for the posterior marginalized over δ_{cp} .

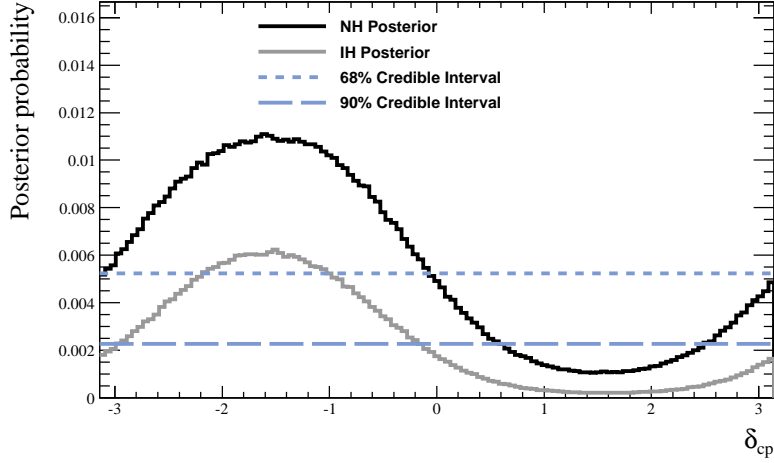


Figure 43: The posterior probability for δ_{cp} , for the normal and inverted hierarchies considered jointly. The dotted lines show the 68% and 90% credible intervals, where the allowed region is the region of the posterior above the line.

462 at the 90% level.

Table 13: The 90% allowed credible interval for different methods of constructing the δ_{cp} posterior.

Method	90% Allowed Credible Interval
Normal Hierarchy ONLY	$[-\pi, 0.45] \cup [2.66, \pi]$
Inverted Hierarchy ONLY	$[-\pi, 0.15] \cup [3.04, \pi]$
Marginalized Hierarchy	$[-\pi, 0.38] \cup [2.79, \pi]$
Joint Hierarchy	$[-\pi, 0.68]$ (NH) \cup $[2.49, \pi]$ (NH) \cup $[-2.99, -0.08]$ (IH)

463 The Markov chain also provides an interesting and natural way to compare the
464 mass hierarchies. Figure 45 shows the 1D posterior for Δm_{32}^2 . In this framework, the
465 integral of posterior where $\Delta m_{32}^2 > 0$ gives the probability that the true hierarchy
466 is normal; for this analysis, that probability is 69.1%, or about a 2.24:1 preference
467 of the data for the normal hierarchy. This is interesting, but not significant enough
468 to draw any firm conclusions. A similar number can be produced for the preference
469 of $\sin^2(\theta_{23}) > 0.5$ or < 0.5 ; the data prefers $\sin^2(\theta_{23}) > 0.5$ at 2.87:1. Again,

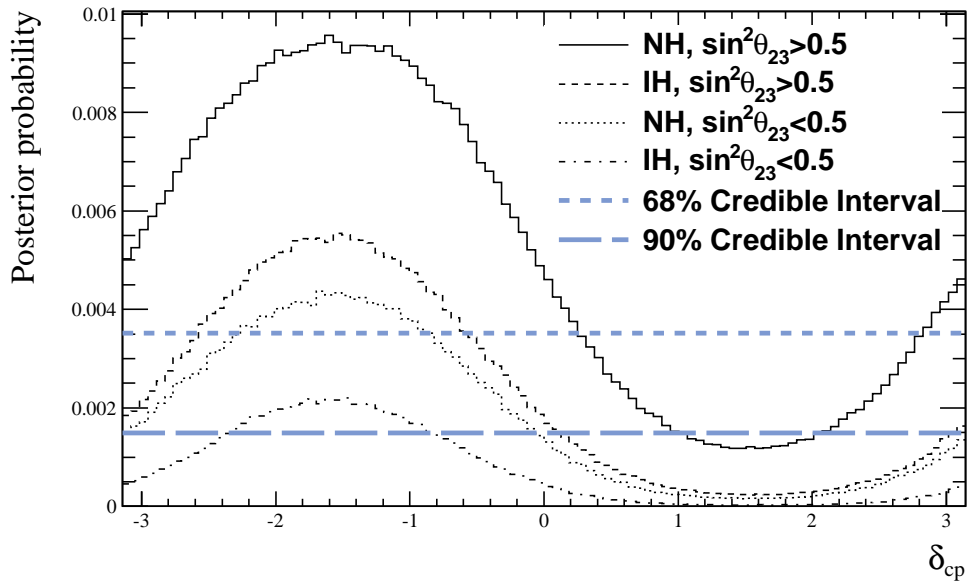


Figure 44: The marginalized δ_{cp} posteriors, for normal and inverted hierarchies, as well as $\sin^2 \theta_{23} > 0.5$ or < 0.5 . The four choices are considered jointly for setting the credible interval levels. The allowed region is the region of the posterior above the line.

interesting, but not significant.

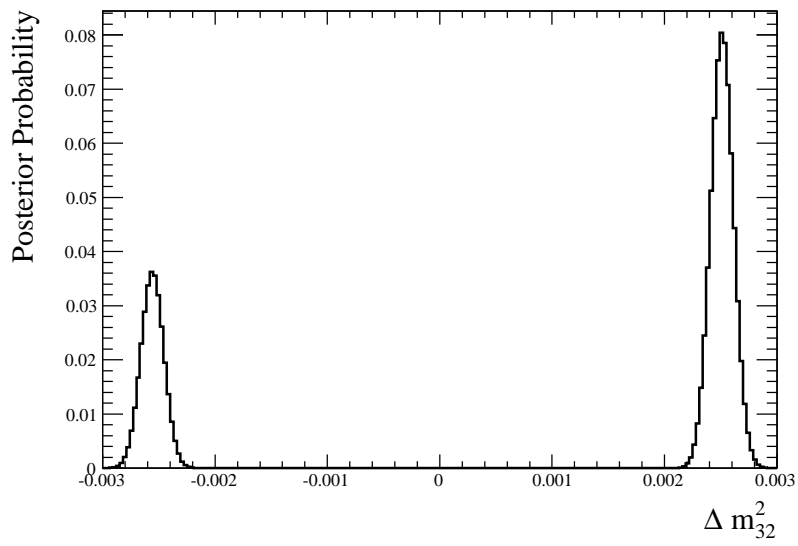


Figure 45: Marginalized Δm_{32}^2 posterior. Normal hierarchy is positive values and inverted hierarchy is negative values; 69.1% of the probability lies in the normal hierarchy.

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493

A Additional Fake Data Set Plots

494

A.1 T2K Only

Contour comparison with VALOR analysis.



plots/fds/fds_0_cont_th23dm23-eps-completed-to.pdf plots/fds/fds_0_cont_th13dcp-eps-converted-to.pdf

(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 46: Fake Data Set 0

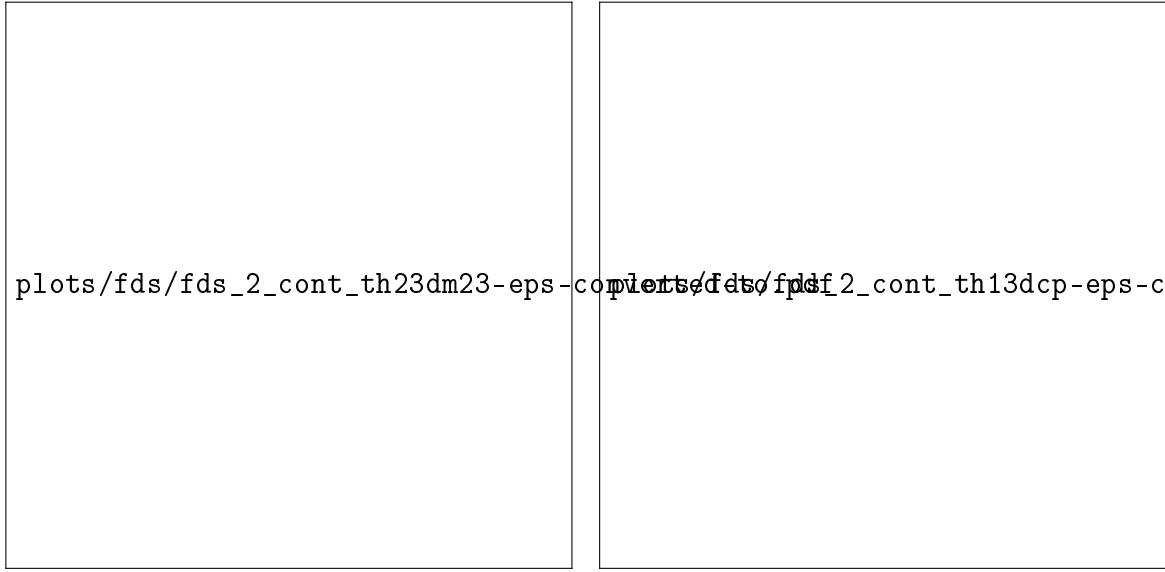
495

496

A.2 T2K with Reactor Constraint

497

MaCh3 only contours, both hierarchies.



(a) $1R\nu_\mu$

(b) $1R\nu_e$

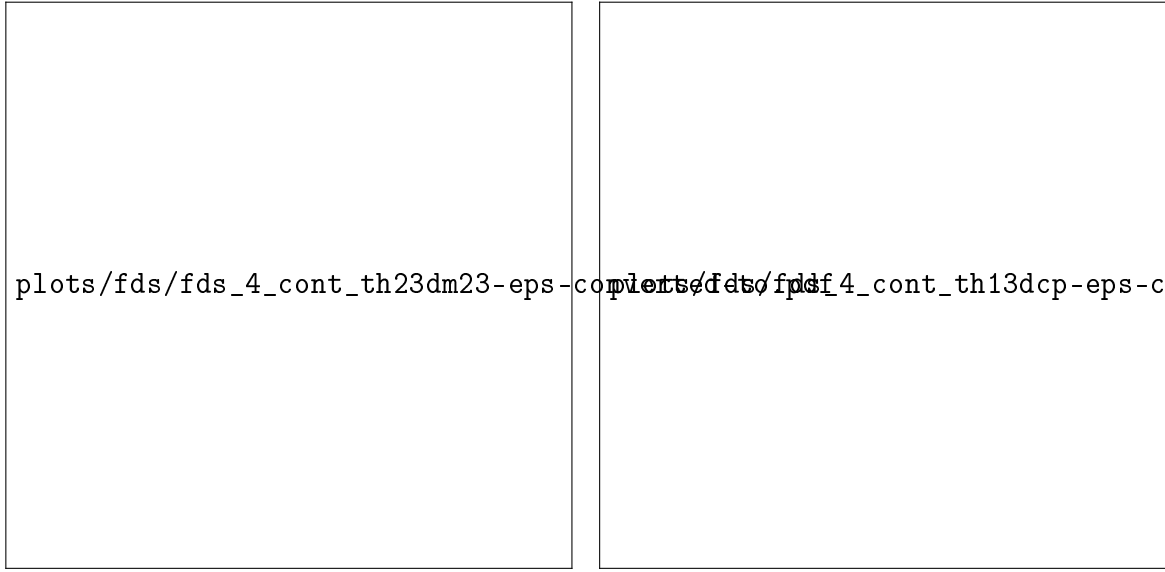
Figure 47: Fake Data Set 2



(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 48: Fake Data Set 3



(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 49: Fake Data Set 4



(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 50: Fake Data Set 5



(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 51: Fake Data Set 1



(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 52: Fake Data Set 2



(a) $1R\nu_\mu$

(b) $1R\nu_e$

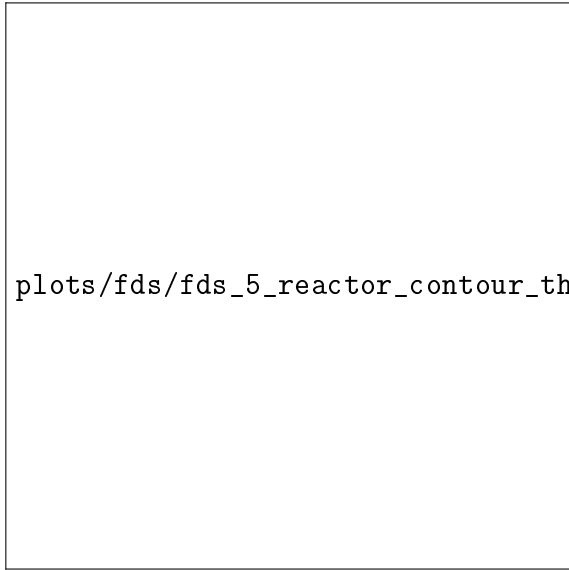
Figure 53: Fake Data Set 3



(a) $1R\nu_\mu$

(b) $1R\nu_e$

Figure 54: Fake Data Set 4



(a) $1R\nu_\mu$

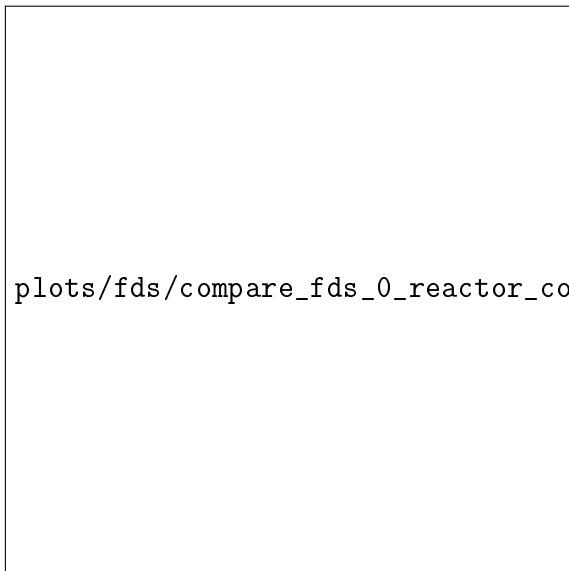


(b) $1R\nu_e$

Figure 55: Fake Data Set 5

498

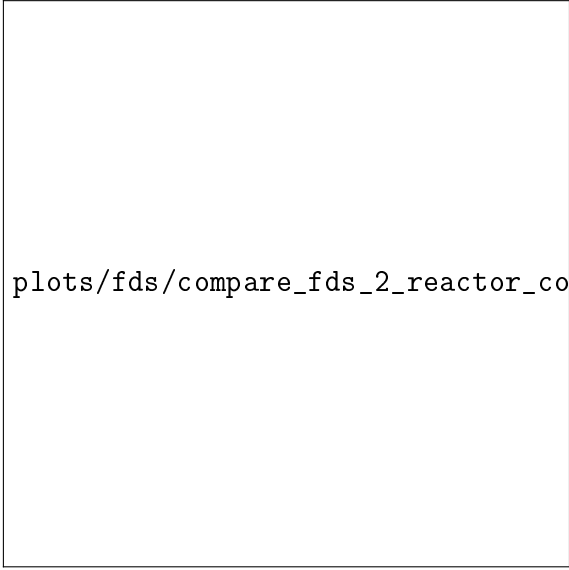
A.3 Comparison with VALOR



(a) Fake Data Set 0 with Reactor Constraint.



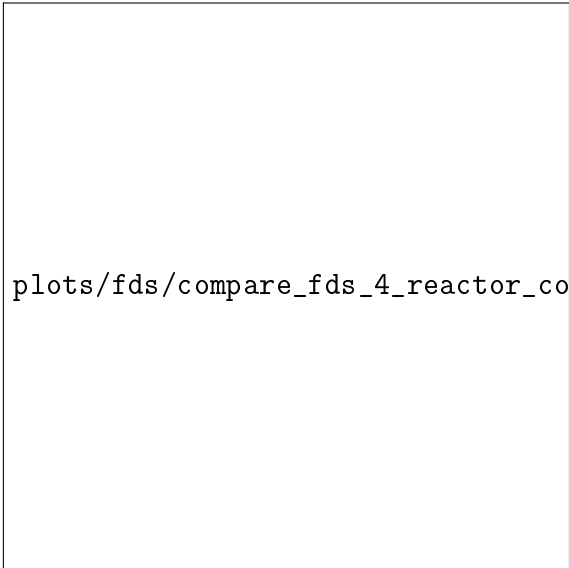
(b) Fake Data Set 1 with Reactor Constraint.



(c) Fake Data Set 2 with Reactor Constraint.



(d) Fake Data Set 3 with Reactor Constraint.



(e) Fake Data Set 4 with Reactor Constraint.



(f) Fake Data Set 5 with Reactor Constraint.