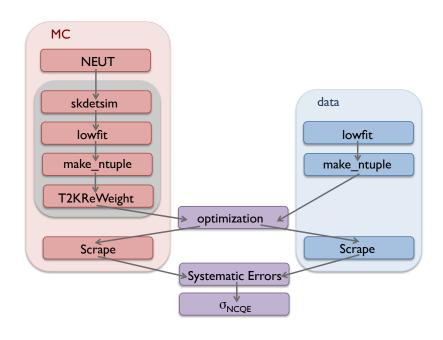
### ncgamma analysis tools



Corina Nantais group meeting 07 June 2017

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## Move from p2 to p3 of T2KReWeight\_v1r27

- Xiaoyue updating T2KReWeight from v1r27p2, because they found a bug
- and she made a mistake in the setup file regarding ROOT (maybe that's why everything just stopped working for me?)

Tag	Description and main features	Date	NEUT	GENIE	JReWeight	NIWGReWeight	GEANTReWeight	Change Log
v1r27p3	Patch the previous release for Summer 2017 OA.  Fixes from v1r27:  2p2h shape dial implemented with 3D(Enu, q0, q3) tables. Patched due to bugfix.  Improved and added validation scripts.  Updates from v1r27:  Include modern SK spline generation script.  Updated BANFF spline app  Update NEUT to include alternate form factor work by Patrick Stowell. Patch 3 completed validation with changes in NEUT at revision 301.  New Eb dial included, but is unstable and is biased in the forward region. We advise against using it for uncertainties.  5.3.3 MaQE bug fix in NEUT for hydrogen targets. Patch 4 (soon) for updated apps for BANFF/OA.  Note: "p1" also exists in CVS for both T2KReWeight and NIWGReWeight containing an additional set of SF->RFG reweighting tables that were removed because they did not resolve the underlying issue of mixing event-by-event reweighting with binned reweighting. They were discarded wholesale from this version because we do not want anyone using a "new" version that is not better than the old version, and has not gone through as much validation as the previous iteration of tables.	May 9th 2017	5.3.3 v1r27p3 maqefix (tarball) (this is also on SVN Revision at 405)	2.10.0 (>=2.8.0)	v1r13	v1r23p2	vlrl	

## Sophie's suggested example

- Sophie used this simpler executable
- T2K-SK 14a MC file
- doesn't need BANFF postfit

```
./genWeights_SK_SFRFG_RPA.exe

-s /disk01/sklb2/t2kmc/t2kmc_14c/t2k_14a_root/nu-
mode/numu_x_numu/root/numu_x_numu.h2o.sk.flux13a.neut_532_ap14a.000.999_fQ
v4r0.root
```

- IT WORKED!
- output file is testweights\_skexample\_nue.root

## Sophie's suggested example

#### testweights\_skexample\_nue.root

```
TFile**
            testweights_skexample_nue.root
 TFile*
           testweights_skexample_nue.root
 KEY: TTree weightstree:1 Tree storing TArray of weights and twkdials per event
[root [2] weightstree->Print()
:weightstree: Tree storing TArray of weights and twkdials per event *
*Tree
*Entries:
            984 : Total = 86352 bytes File Size = 11937 *
                : Tree compression factor = 7.51
*Br 0 :nweights : nweights/I
*Entries : 984 : Total Size= 4504 bytes File Size = 138 *
*Baskets: 1:Basket Size= 32000 bytes Compression= 29.12 *
*Br 1:weights : TArrayF
*Entries: 984: Total Size= 20255 bytes File Size = 6708 *
*Baskets: 1: Basket Size= 32000 bytes Compression= 2.95 *
*Br 2 :brNIWG2014a_SF_RFG : TArrayF
*Entries: 984: Total Size= 20310 bytes File Size = 1407 *
*Baskets :
           1 : Basket Size= 32000 bytes Compression= 14.06 *
   3 :brNIWG_rpaCCQE_norm : TArrayF
*Entries: 984: Total Size= 20315 bytes File Size = 1387 *
*Baskets: 1: Basket Size= 32000 bytes Compression= 14.26
   4 :brNIWG_rpaCCQE_shape : TArrayF
*Entries: 984: Total Size= 20320 bytes File Size = 1424 *
*Baskets: 1: Basket Size= 32000 bytes Compression= 13.89 *
```

### Use all 2015 files

```
./genWeights 2015.exe
-i /disk01/sklb2/t2kmc/t2kmc_14c/t2k_14a_root/nu-
mode/numu_x_numu/root/numu_x_numu.h2o.sk.flux13a.neut_532_ap14a.000.999_fQ
v4r0.root
-p postfit_banff_2015_data_20150417.root
-o weight_output.root
-horn I
-app 0
-t 5
--use-prefit
--drop-flux
IT WORKED!
```

18 branches

### want more recent executable

- ask Xiaoyue which genWeights
   → genWeights\_2016 written by Mark Scott in app
- had to fix bug in px, py, pz, E to E, px, py, pz
- Xiaoyue fixing another momentum bug
- can copy it over from p2 to p3

### Chatting with Mark Scott

#### like Xiaoyue said

- probably don't have to use NIWG, it's now included in NEUT reweight → ok?
- use genWeights\_SK\_2016
- maybe Alex edited src/T2KNeutReWeight → have to check
- best to try to add Alex's changes to genWeights\_SK\_2016
- would be able to use Alex's genWeights\_ncgamma if

```
rw.Systematics().Include(t2krew::kNIWG2014a_SF_RFG); → SF to RFG rw.Systematics().Include(t2krew::kNXSec_VecFFCCQE); → form factor maqe rw.Systematics().SetTwkDial(t2krew::kNIWG2014a_SF_RFG,1); rw.Systematics().SetTwkDial(t2krew::kNXSec_VecFFCCQE, 2);
```

- did he say set to 0?
- would also need 2012 banff postfit

#### most recent files

```
./genWeights_SK_2016.exe
-i /disk01/sklb2/t2kmc/t2kmc_14c/t2k_14a_root/nu-
mode/numu_x_numu/root/numu_x_numu.h2o.sk.flux13a.neut_532_ap14a.000.999_fQ
v4r0.root
-p BANFF_PostFit_170521.root
-o weight_2016_output.root
-horn I
-app 0
-t 5
--use-prefit
--drop-flux
IT WORKED!
```

16 branches

### now make it work with ncgamma files

- the input is after neut, skdetsim, lowfit, make\_ntuple
- I think it will take in lentuple/
- try from /disk01/usr4/cnantais/Attempts/7Jan2016/lemc\_jan7/lentuple/lentp\_numubar.ncgamma\_flux11a\_neut514.099.root
- root –l lentp\_numubar.ncgamma\_flux | la\_neut5 | 4.099.root
- TTree h1

/disk01/sklb2/t2kmc/	lentuple				
h1 DST	h1 T2KLowE Common Ntuple				
Br 373	Br 115				

### most recent files on 2016 ncgamma MC

```
./genWeights SK 2016.exe
-i lentp_numubar.ncgamma_flux | la_neut5 | 4.099.root
(copied from Attempts/7Jan2016/lemc_jan7/lentuple)
-p BANFF_PostFit_170521.root
-o weight_nc_output.root
-horn I
-app 0
-t 5
--use-prefit
--drop-flux
IT WORKED!
16 branches
```

### ProcessNCEL\_mc.sh

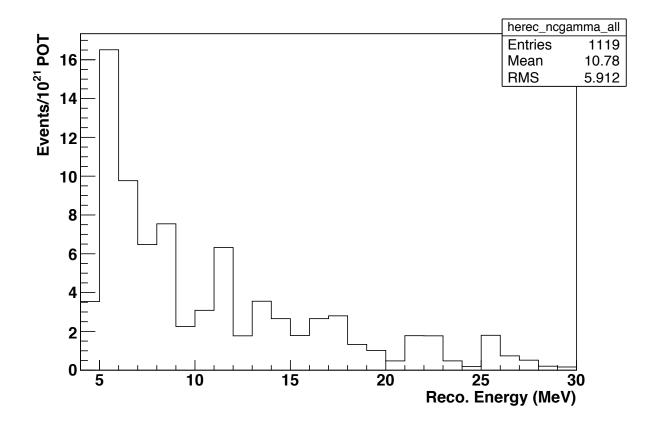
- ./ProcessNCEL\_mc.sh 001 numu
- IT WORKED!
- flux\_preffit/ 29 branches
- xsec\_prefit/16 branches
- compare to Attempts/7Jan2016/
- flux\_preffit/ 24 branches?
- xsec\_prefit/ 7 branches?
- → why are the number of branches of TArrayF different?

# Next in analysis flow

- next is Optimization
- can maybe do Alternate Processing instead
- .../ncgamma/Processing/
- \$ python SelectNCGamma.py —o ncgammamcRun4.root /disk/.../lemc/lentuple/lentp\_nu\*.root
- can try with just one file

# one MC file, looks good

e.g., herec\_ncgamma\_all



skip Optimization for now, until have all MC

# Do all MC Processing

- python PrepareNQS.py
- no T2KReWeight files???
- nothing in log
- try another single file, say numu 099
- problem probably in setup

- ./ProcessNCEL\_mc.sh 099 numu
- no shared library
- source my\_setup.csh
- make clean
- ./config.sh
- gmake
- ./ProcessNCEL\_mc.sh 050 numu
- IT WORKED!

# Moving on to Scrape step in the meantime

- runscrape.csh (and Scrapper.py) require OrderedDict and NumPy
- Hayato-san installed on sukap:
- numpy 1.7.1
- scipy 0.12.1
- ordereddict I.I.

~/ncgamma/Processing@sukap001[490]\_% ./runscrape.csh Getting flux weights from ~/ncgamma/SystematicErrors/beamweights/ No friend trees given Creating branch erec/D Creating branch pnu/D Creating branch inu/L Creating branch mode/L Creating branch isCC/L Creating branch isQE/L Creating branch appeared/L Creating branch onePi0/L some files totally missing (no candidates?) Creating branch onePipm/L Creating branch cosbeam/D some files have h1 Creating branch t2kposc/D Creating branch weight1/D some files don't have h1??? Creating branch weight2/D Creating branch weight3/D Creating branch weight4/D Creating branch nrun/L Creating branch nsub/L check 07 Jan 2016 Creating branch <u>nev/L</u> Creating branch mrrun/L lowedata/ missing? Creating branch Nn/L Creating branch Np/L Creating branch wallv/D Creating branch ng1/L Creating branch ng2/L check 17 Jan 2016 Creating branch toteg1/D Creating branch toteg2/D Creating branch wall/D Creating branch effwall/D Creating branch ovag/D Creating branch angle/D Creating branch anglepass/D Creating branch nompeak/L Creating branch dt0/D Creating branch dtbunchcenter/D Creating branch dt0passtight/L Creating branch dt0passloose/I Creating branch dt0beamunrelpre/L Creating branch dt0beamunrelpost/I Creating branch wallpass1/1 Creating branch effwallpass1/1 Creating branch ovagpass1/L Creating branch wallpass2/I Creating branch effwallpass2/I Creating branch ovaqpass2/1 Creating branch wallpass3/L Creating branch effwallpass3/I Creating branch ovaqpass3/L Creating branch wallpass4/L Creating branch effwallpass4/I Creating branch ovaqpass4/L 17 Begin processing 328 data files Error in <TChain::LoadTree>: Cannot find tree with name h1 in file /disk01/usr4/cnantais/lowedata/ntuple/data.lowfit.70620.merge.root

### Summary

- can use T2KReWeight on ncgamma mc
- problem with T2KReWeight setup when submitting many jobs
- then do Optimization
- then Scrape, must understand errors in output