# Unfolding: the neutrino experiment experience 

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24 January 2019

## Introduction

- Who am I?
- MINOS, T2K, Minerva, DUNE; Neutrino-nucleus interaction measurement and fitting
- Which neutrino experiments use/don't use unfolding and why?
- What have we done in the past?
- What are we doing now?
- What are we likely to do in the future?


## Oscillation analyses don't (explicitly) use unfolding

- Measuring small number of parameters in a "well-known" model.
- But maybe future experiments will want to map out $P\left(\nu_{\alpha} \rightarrow \nu_{\beta}\right)$, say
- Implicit unfolding in "beam matrix" methods


NOvA, J. Wolcott, Nulnt18

## Unfolding is widely used in neutrino interaction analyses



- Approximate "effective" models used at each stage
- Want to measure identity and kinematics of final-state particles
- Renewed interest because of importance to oscillation analyses


## Some common features of neutrino interaction measurements




T2K, Phys Rev D 87, 092003
MINER $\nu$ A, Phys. Rev. D92, 092008 (2015)

- Measure flux-integrated distributions of kinematic variables to distinguish widely-varying models
- Often systematics-limited
- Largest systematic often flux: strong positive correlations


## What we've done in the past: D'Agostini, mostly

- $\mathrm{O}(1)$ iteration of D'Agostini on background-subtracted data
- "Unfolding error": 2nd iteration minus first
- Choice of $N_{\mathrm{it}}$ : warp MC to look like data. Iterate until bin contents "close" to truth
- Unfold using different models; add to systematic error


MINER A, Phys. Rev. Lett. 116, 071802 (2016)

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## What we've done in the past has some shortcomings

- Background subtraction breaks Poisson assumption in D'Agostini method
- Method of choosing $N_{\mathrm{it}}$ is ad hoc, doesn't consider stat error


## What we're doing now

- Providing unregularized results alongside Tikhonov-regularized ones (T2K)
- More details in Stephen Dolan's talk

- More careful checks with covariance matrix chi2 on data-driven warped distributions (Mineva)


## MINER $\nu$ A approach to D'Agostini unfolding: example




Trung Le, Fermilab JETP seminar, Sep 212018

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## MINER $\nu$ A approach to D'Agostini unfolding: example




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- "Feed-down" means wider migration matrix than usual. Procedure similar for all variables
- Reweight MC to look like data: unfold this warped fake data


## Two checks on number of iterations

1. When does the unfolded distribution closely approximate the (warped) truth?
2. When is the $\chi^{2} /$ dof $\sim 1$, averaged over many Poisson throws?
3. In this case, $\chi^{2} /$ dof $\sim 1$ with bins removed



## What we might do in the future

- L-curve-alike for D'Agostini unfolding: compare $N_{\text {it }}$ result to infinite iterations result. (More details from Stephen Dolan)
- Implement D'Agostini without background subtraction
- Forward folding: provide tools to compare theoretical models directly to reconstructed (smeared) data. Challenges:
- Publishing systematics
- Making the response matrix model independent
- Long-term sustainability (what if the tools are unmaintained?)


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## Fits to strongly-correlated data 1



MICROBOONE-NOTE-1045-PUB
$\phi$ covariance matrix


T2K, Phys Rev D 87, 092003

- Neutrino interaction data strongly positively correlated by flux uncertainties
- Well known that in such cases, the best fit can be well outside the data points
- "Peelle's Pertinent Puzzle" in nuclear physics. Several proposed interpretations/solutions: "International evaluation of neutron crosssection standards", IAEA (2007)

Fits to strongly-correlated data 2



$$
c_{i j}=\sum_{\text {universe } k}\left(y_{i}^{(k)}-y_{i}^{*}\right)\left(y_{j}^{(k)}-y_{j}^{*}\right)
$$

- "Multi-universe": throw random systematic universes, re-extract result

Fits to strongly-correlated data 2


$$
\chi^{2}=(\mathbf{D}-\mathbf{M})^{T} \mathbf{C}^{-1}(\mathbf{D}-\mathbf{M}) \quad C_{i j}=\sum_{\text {universe } k}\left(y_{i}^{(k)}-y_{i}^{*}\right)\left(y_{j}^{(k)}-y_{j}^{*}\right)
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- "Multi-universe": throw random systematic universes, re-extract result
- Empirically, $y \rightarrow \log (y)$, ameliorates the issue, $\Rightarrow$ log-normal uncertainties on $y(?)$
"Box-Cox transformation for resolving the Peelle's Pertinent Puzzle in curve fitting", Oh and Seo 2004
- Is this the best way to communicate our systematics?


## Some outstanding questions

- Can we unfold and quantify bias s.t. it's small enough to not matter?
- How can we adapt unfolding techniques from the literature to work with the multi-universe/multisim method used by Minerva and MicroBooNE?
- Evaluating unfolding bias by comparing to a model "warped like the data": what's the range of validity?
- How do we assign systematic uncertainties to "the unfolding technique" without double-counting?
- How do we deal with PPP? Or, what's the best way to preserve the features of our detailed systematic error estimates in a way that's digestible to users (theorists, other experiments)?


## Backup slides

## MINER $\nu \mathrm{A} \nu$ CCOpi $\chi^{2}$ vs number of iterations



- 4 iterations chosen


## More questions/thoughts

- Probable valuable: "bottom line tests" as in arXiv:1607.07038
- Covariance matrices:
- How to quote in data releases without numerical issues?
- How to approximate when large?
- How to make sure we have enough multisim throws?


## Some personal opinions

- Field is strongly "visual": we're always going to want something to look at to assess results
$\Rightarrow$ Unfolding will probably always be with us
- Tikhonov-regularized fits offer some clear advantages over iterative techniques
- Always show full data in reco space (in as many dimensions as the cross section). Not always done!
- Bias-variance tradeoff is a useful frame for thinking about unfolding. Links discussion to the literature. Makes clear pros and cons


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