# Statistical models for recasting of unfolded measurements and combination with searches

Andy Buckley University of Glasgow

**ECFA Higgs Factories, MC Generators Meeting 12 November 2021** 



## Executive briefing: unfolding

#### Unfolding has been around for decades

- General principle is to correct reco data for effects distorting our view of the physics of interest
- LHC in particular has refined this to mean unfolding to a *fiducial volume:* correct ~only for detector effects, to a well-defined final-state, cuts similar to reco
- ➢ Non-fiducial extrapolations, too in addition to fid xs
- Necessarily propagate uncertainties from both detector understanding, and the unfolding mechanism

## Main methods/tools

- Regularised inversion:
  - Iterative Bayes IBU (RooUnfold, pyunfold)
  - SVD/Tikhonov reg (RooUnfold, TUnfold)
- "Forward-propagation" with sampling/fitting:
  - TRExfitter, pyhf?
  - PyFBU, (hunfold)



## Unfolding vs search profiling

#### MC-driven unfoldings need the same things

- Binned data observable
- Binned MC truth and reco observables
  - Generally, truth/reco binnings (and even the observable!) don't need to match: *bin reco tighter*
- A response matrix of p(reco<sub>i</sub>|truth<sub>i</sub>) for truth, reco bins i, j
  - A good unfolding requires a "tight" response matrix, not a "diagonal" one
- > Lots of standard methods to assess unfolding non-closure and MC-model dependence

### Aim of unfolding is to extract fiducial, particle-level bin values & errors

> Note, not the true, integer bin-populations, but physics params e.g.  $d\sigma_i$  (differential xs in bin i)

 $\Rightarrow L_{\text{search}}(\text{obs}|\text{BSM params}) \text{ vs } L_{\text{meas}}(\text{obs}|\text{d}\sigma'\text{s})$ 

- > Different methods use different Poisson/multinomial internal models, plus inversion heuristics
- Fit methods not so different from a BSM-search fit/scan: POIs are bin-values, not model params
- > We don't often think of measurements as limit contours in a  $N_{\rm bin}$ -dim space, but they are!

### Systematic uncertainties

- > Need some mechanism to propagate systematic uncertainties
- ▶ Intrinsic/natural in fit/scan methods, just extend the param space to bin vals ⊕ nuisances
- More ad hoc in (most?) inversion tools updates coming in RooUnfold?

## Stat models from unfolding, and combination

#### Search-likelihood publication quite advanced now

- > Particularly via pyhf, path to common format for Poisson model in searches
- Don't want measurements to get left out, cf. e.g. Contur (and others') uses of LHC measurements for complementary BSM constraints

## Statistical models from unfolding tools

- > Obvious for sampling tools: Poisson likelihood/posterior are explicit
- Maybe less clear for inversion tools, but means and (correlated) uncertainties calculated ⇒ implicit b2b Gaussian pdf maybe templated wrt nuisances
- Obstacles to encoding unfolded likelihoods in pyhf?
- Alternatively, publish sampled point sets?

### Issues shared with searches

- elementary nuisances needed for fully coherent combination
- some standardisation and versioning of nuisance names needed and gauging degree of correlation between versions (nice problem to have...)
- Freq vs Bayes: combining likelihoods & posteriors?



## Summary/discussion

#### Input from toolkit authors

- Inversion and sampling methods developments, current and planned output data and formats.
- Possible to achieve technical compatibility with pyhf/etc. direct searches?
- In pyhf and similar tools: how best to encode Gaussian uncertainty models? Extend pyhf, build on pyhf, include one more stat distribution in common format?

### Experience of unfolding with model publication

- 2x public ATLAS top analyses, <u>1 total</u>, <u>1 differential</u>
- More?

#### Experience of combination (e.g. Convino)

- Need for standards / understanding of common & related systematics
- Combining limits from different stat paradigms

#### More?