

Rivet analysis preservation & recasting

including observations from Gambit, TopFitter, Les Houches, LPCC forum...

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ATLAS data-reinterpretation workshop,
CERN, 21 July 2017



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SOCIETY



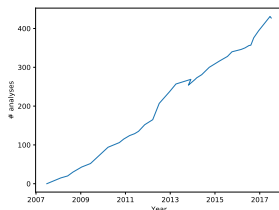
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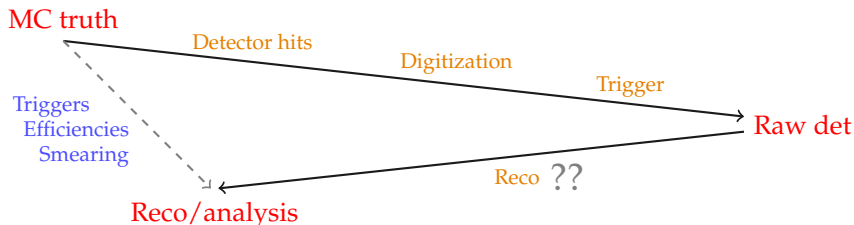
Rivet background

Rivet is an analysis system for MC events, and *lots* of analyses

- ▶ Easy & powerful tool to get many physically meaningful plots from many MC gens
- ▶ “Physicist-friendly” code interface
- ▶ LHC standard for archiving *unfolded* data analyses
- ▶ Well-established in ATLAS: 172 analyses and official support manpower
- ▶ Athena interface and standalone tools in ATLAS software
- ▶ Since version 2.5.0, includes [detector-smearing tools](#) for BSM preservation/recasting



Detector effects in ‘fast-sim’ vs. ‘smearing’



- ▶ Explicit fast-sims work hard to get to the “Raw det” stage, but the big reco step is not known in detail
- ▶ In practice, kinematic smearing is subleading: we calibrate well! (And where it matters, is a fast-sim accurate enough?!)
- ▶ ~All det effects in search analyses are lepton/tagging efficiencies
- ▶ \Rightarrow Rivet takes the “short route” of simple smearing, plus tabulated efficiencies cf. Delphes (+ CP notes & analysis papers)
- ▶ Effs (and resolutions) can be *analysis-specific*: many different WPs/effs of taggers, calibration & isolation across analyses/runs

Detector sim in Rivet

- ▶ Rivet smearing/efficiency approach based on a refinement of the GAMBIT system, with less “code noise” and more scope for user-defined smearing functions
- ▶ Analysis-specific efficiencies and smearings are more precise and allow use of **multiple jet sizes, tagger & ID working points, isolations, ...** ⇒ **many variations in real analyses**
- ▶ Smearing uses established “projection” mechanism: familiar and efficient. Smearred objects are just “wrappers” around truth-level definitions
- ▶ BSM developments also produced powerful filtering tools: Rivet `cut` objects and generalised C++ functions can all be used to apply complex selections. Cut-flow tools are included.

Rivet smearing examples

Leptons:

```
// Definition
FinalState es1(Cuts::abseta < 3.2 && Cuts::abspid == PID::ELECTRON);
SmearredParticles es2(es, ELECTRON_EFF_ATLAS_RUN2, ELECTRON_SMEAR_ATLAS_RUN2);
declare(es2, "Elecs");
...
// Usage
Particles elecs = apply<ParticleFinder>(event, "Elecs").particles(10*GeV);
```

Jets:

```
// Definition
FastJets js1(FinalState(Cuts::abseta < 4.9), FastJets::ANTIKT, 0.4);
SmearredJets js2(js1, JET_SMEAR_PERFECT, JET_BTAG_EFFS(0.7, 0.12, 0.02));
declare(js2, "Jets");
...
// Usage
Jets jets = apply<JetAlg>(event, "Jets").jetsByPt(30*GeV);
```

Also a SmearredMET ...

Standard global functions, plus user-defined. C++11 lambda fns etc. are allowed. Rivet 2.6 allows *chaining* of smearings and efficiencies.

From analysis preservation to reinterpretation

Description	Rivet			MadAnalysis5			CheckMATE
	#evt	tot.eff	rel.eff	#evt	tot.eff	rel.eff	tot.eff
2jm cut-flow:							
Pre-sel+MET+pT1	31250	100%	-	32150	100%	-	100%
Njet	28472	91%	91%	28478	91%	91%	91%
Dphi_min(j,MET)	28472	91%	100%	28477	91%	100%	91%
pT2	22950	73%	81%	22889	73%	80%	73%
MET/sqrtHT	22950	73%	100%	22889	73%	100%	73%
m_eff(incl)	10730	34%	47%	10710	34%	47%	33%
	10630	34%	99%	10609	34%	99%	32%

Prelim
results from
LH2017 BSM
(AB,
Grellscheid,
Fuks, Desai)

Current Les Houches benchmarking study: Rivet analyses reproduce published full-sim & custom-config Delphes fast-sim within a few %

But signal-region counts are just the beginning. Plan to augment Rivet with a statistics suite to turn SR counts into BSM limits: Rivet CL_s implementations in CONTUR and in Rivet contrib — potential for limit setting with combined SM+BSM data!

To streamline, we need to *standardise* distribution of not just observed counts, but also **SM background expectations, efficiency tables, cut-flows + other data for signal validation, and correlations.**
Obvious route is HepData. Interest in ATLAS top, CMS, BSM pheno...

Correlations and simplified likelihoods

Without correlations, reinterpretations have to be conservative: only use the single best-expected- Δ LL SR from each correlated group.

Better: full likelihoods or *simplified likelihoods* cf. CMS

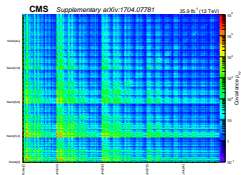
$$\mathcal{L}_S(\mu, \theta) = \prod_{i=1}^N \frac{(\mu \cdot s_i + b_i + \theta_i)^{n_i} e^{-(\mu \cdot s_i + b_i + \theta_i)}}{n_i!} \cdot \exp\left(-\frac{1}{2} \theta^T \mathbf{V}^{-1} \theta\right)$$

$$V_{ij} = E[\theta_i \times \theta_j]$$

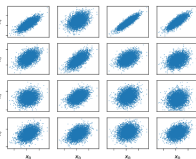
See CMS NOTE-2017/001

with averaging over elementary bkg nuisance distributions

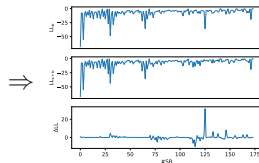
SL cov



Corr sampling



Δ LLs



Technical discussion / implementation needed on whether separated systematic cov matrices, simplified cov matrices, or nuisance param forms best. How to identify dataset types, and match cov indices across observables? Etc.