

Tools and Techniques for High-*p***^T Physics**

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Outline



From Raw Data to Physics Results

Monte-Carlo Event Generation

Physics Objects

Background Estimation

Advanced Signal Analysis





PHYSICS OBJECTS

Many classes: charged leptons, jets, b-jets, boosted objects ...

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Leptons: scale factors, e.g. via tag&probe method Jets: today...

Short Recap

Physics objects:

All objects must be properly calibrated:

- Lepton ID: multivariate discriminant, isolation

Tracks: **alignment** of tracking detectors (not shown)





What is a Jet?





www-cdf.fnal.gov

Jets can be defined on different technical levels:

- Parton level: for calculations in perturbative QCD ("theory jets")
- **Particle level**: jets reconstructed from stable hadrons
- Detector level: jets reconstructed from energy deposits in calorimeter and/or tracks in tracking detectors
- Design of successful jet algorithms:
 - Independent of technical level
 - Invariant under Lorentz boosts
 - Comparison with theory: infrared and collinear safe → find same jet even after emitting soft/collinear radiation

Infrared and Collinear Safety



Goal: jet definition on robust on all technical levels against additional radiation (low momentum or small angle)



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Sequential Recombination



- Jets at the LHC: sequential recombination
 - Infrared/collinear safe by construction
 - Define distance measure d_{ij} between particles i, j and distance of particle i to beam axis d_{iB}
 - LHC standard: "anti-kt algorithm"

$$d_{ij} = \min(k_{t,i}^{-2}, k_{t,j}^{-2}) \frac{\Delta R_{ij}^2}{R^2}$$
 $d_{iB} = k_{t,i}^{-2}$ LHC Run 2:
R = 0.4 ("ak4")

- Sequential recombination algorithm:
 - Compute *d_{ij}* for all pairs of (pseudo-)particles, combine pairs with *d_{ij}* < *d_{iB}* to new pseudo-particles
 - Termination condition: pseudo-particle \rightarrow jet if d_{iB} is the smallest distance d_{ij}



Sequential recombination



Jet Reconstruction





- Starting points for energy reconstruction in calorimeter:
 - **Calorimeter towers**: fixed grid grouping calorimeter cells, e.g. $\Delta\eta \times \Delta\phi = 0.1 \times 0.1$
 - Topological clusters ("topo clusters"): groups of cells with energy deposits
- Jet reconstruction strategies:
 - Pure calorimeter jets or track jets
 - **Combination** of calorimeter and tracker information
 - Particle flow: optimal combination of subdetectors for reconstruction of each particle type

Particle Flow



- Jet reconstruction with traditional calorimetry:
 - Reconstruct jet energies from ECAL and HCAL energy deposits
 - Approx. 70% from HCAL with rather poor energy resolution (≈ 50%/√E)
- Particle flow calorimetry: jet reconstruction exploiting each sub-detector optimally
 - Tracker: charged particle momentum
 - ECAL: photon energy and bremsstrahlung
 - HCAL: only neutral hadrons $(n, K_L) \rightarrow$ only 10% of jet energy reconstructed with rather bad resolution



Traditional Calorimetry





M. Thomson

Jet Calibration: CMS Method



Multi-stage jet energy corrections in CMS:

Applied to data –



- **Pileup** and electronic **noise** correction (MC-based + additional "residual correction" for data)
- **Response** correction: uniform response as a function of p_T and η (MC)
- **Residual** corrections for data in η (dijet events with one well-calibrated jet in the barrel region)
- **Residual** corrections for data in p_T (balance of Z or γ recoiling against jet)
- (optional) Jet flavor correction: different response from light quarks, gluons, heavy quarks (MC)

Jet Calibration: Results



- Typical uncertainties of jet energy corrections: 1–2%
- Uncertainties propagated into more complex observables, e.g. E^{miss}
- I Jet energy **resolution** in data worse than in $MC \rightarrow$ "smear" MC



B-Tagging



- B-tagging algorithms at hadron colliders based on:
 - Secondary vertices and tracks with large impact parameter: long B hadron lifetime (picoseconds)
 - **Soft leptons**: semileptonic decays $B \rightarrow \ell \nu X$
 - **Large b-quark mass:** wider jets, large relative p_T of lepton in $B \rightarrow \ell v X$
 - Hard b-quark fragmentation: B hadron carries most of b-quark energy
- LHC Run 2: above criteria combined in multivariate discriminant (increasingly based on deep learning)
- Relevant e.g. for t \rightarrow Wb, H \rightarrow bb, Z \rightarrow bb



B-Tagging Performance



Performance of b-tagging algorithms:

- b-jet tagging efficiency: fraction of true b-jets tagged as b-jets
- Misidentification probability ("mistag rate"): fraction of true lightflavor (uds), charm (c), or gluon (g) jets wrongly tagged as b-jets
- Depends on physics process considered, popular benchmark: tt
- Representation: receiveroperating characteristic (ROC)

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B-Tagging Calibration





Calibration of b-tagging algorithms:

Method: measure tagging efficiency and mis-ID probability for benchmark processes (e.g. QCD dijets, tī) with different methods in data and MC → correct MC with scale factor SF_b depending on jet kinematics
Several working points (= cuts on b-tagging discriminant) with fixed mis-

identification probability, e.g. 10% ("loose"), 1% ("medium"), 0.1% ("tight")

Very useful: calibration of full b-tagging discriminant shape let p_[GeV]

Boosted Objects

- Significant fraction of LHC events contains decays of heavy particles with large transverse momenta (p_T ≥ 200 GeV)
 - Decay products strongly collimated through large Lorentz boost
 - Need novel algorithms to reconstruct jet substructure and tag W/Z/H/top
 - Standard model example: associated tt
 H
 production with H
 → bb
 decay
 - BSM example: heavy Z' decay



Jet Substructure Algorithms



- Reconstruction of boosted objects: very active field at the LHC
 - General idea: reconstruct "fat jets" and study their substructure
 - More general sequential recombination jet algorithm:

$$d_{ij} = \min(k_{t,i}^{2n}, k_{t,j}^{2n}) \frac{\Delta R_{ij}^2}{R^2} \qquad d_{iB} = k_{t,i}^{2n}$$

(n = 0: Cambridge/Aachen, n = +1: k_t , n = -1: anti- k_t)

- Anti-kt jets: hard particle first clustered with surrounding soft particles → reversion of clustering steps ("declustering") not meaningful
- Jet clustering with Cambridge/Aachen or k_t algorithm \rightarrow declustering reveals substructure
- Typical fat-jet radius parameters: *R* = 0.8–1.5



Jet Substructure Algorithms



- Analysis of jet substructure:
 - Iterative declustering of fat jet
 - Grooming: removal of uncorrelated wide-angle soft emission from fat jet → better mass resolution, reduced pileup dependence
 - Jet shape algorithms, e.g. N-subjettiness
 - More involved algorithms used for tagging top and Higgs: combination of jet shape algorithms with grooming



CMS-PAS-B2G-17-001

Jet Substructure Landscape





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N-Subjettiness: Definition



- Jet-shape variable N-subjettiness τ_N : **energy flow** inside fat jets
- Jet with M particles \rightarrow deviation of energy flow from N subjet axes



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Short Summary



- Jet reconstruction:
 - Sequential recombination: infrared and collinear safe
 - Multi-stage calibration of energy scale
- Identification of b-jets: key to many process
 - Multivariate b-tagging algorithms
 - **Scale factors** for differences between data and MC
- **Boosted** jets: special treatment





BACKGROUND ESTIMATION

Sidebands

- General idea: extract information on background in phase space region that is signal-depleted but representative of signal region
- Sideband techniques:
 - Use case: signal = narrow invariant mass peak on large (often combinatorial) background (e.g. H → γγ)
 - Estimate background normalization (sometimes: also shape) from the same invariant mass spectrum **outside** peak



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- Use case: MC simulation of background in signal region unreliable (e.g. corner of phase space that is not well modeled or lacking statistics)
- Event selection for control region such that signal and control regions are mutually exclusive (jargon: "orthogonal"), e.g. by inverting certain selection criteria
- Measure background normalization (and shape) in control region, transfer to signal region (usually using MC simulation)

Control Regions: ABCD Method



Idea: measure background in three control regions → predict background in signal region

- Define four regions A, B, C, D in space of **uncorrelated** observables x and $y \rightarrow$ signal in A (small x AND small y)
- Measure background b in B, C, $D \rightarrow$ background in A given by

$$b(A) = \frac{b(B) \times b(C)}{b(D)}$$

Example: CMS $H \rightarrow \tau \tau$ analysis – QCD multijets in $H \rightarrow \tau_h \tau_\ell$

Observables: charge sign (same vs. opposite charge) and isolation of hadron and lepton (tight vs. relaxed)



Closure Test

- Background estimation from sidebands or control regions: consistency check required
 - **Closure test:** does the method "close" on simulated events (i.e. is known background process predicted accurately, are there biases)?
 - Often: amount of "non-closure" used as systematic uncertainty due to background estimation method
- Example: closure test of tau modeling method in $H \rightarrow \tau \tau$ ("embedding")





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In-Situ Background Determination



- So far: a-priori expectation of background normalization of shape in signal region from MC simulation or data-driven method
- Signal extraction in many analyses: **profile-likelihood fit**
 - Simultaneous fit of signal and control region(s)
 → background in control region(s)
 constrains background in signal region
 - Systematic uncertainties included as nuisance parameters (e.g. normalization and shape from ABCD = nominal value, uncertainty from closure test)
 - Assumptions: fit model adequate, correlations between signal and control regions well modeled



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Summary



- Various methods to determine backgrounds: MC simulation or data-driven methods
- Data-driven methods: Measure background in signal-depleted region in data → estimate background in signal-enriched region (closure test required)
- Various techniques, specific to analysis and background:
 - Combinatorial background underneath mass peak: fit to parametric signal and background model
 - Continuum background: control regions, ABCD method, …
 - If background is hard to model: embedding of MC objects in data events
 - Simultaneous profile-likelihood fits to signal and control regions





ADVANCED SIGNAL ANALYSIS

Differential Cross Section



- So far: reconstructed distributions of kinematic observables compared to expected distributions (from MC and/or data)
 - All physics effects forward-folded with detector effects (e.g. resolution)
 - Problem: distributions **cannot be compared** between experiments
- Way out: measurements presented as differential cross sections
 = cross sections as a function of one or more kinematic observable
 - Detector effects corrected by unfolding procedure
 - Typical result: fiducial differential cross section on level of stable particles
 - Differential distributions contain more information on physics processes than inclusive cross sections → more detailed comparison with theory

Unfolding Techniques



- Determine **true** distribution $f(\mathbf{y})$ from **reconstructed** distribution $g(\mathbf{x})$:
 - Relation: Fredholm integral equation

$$g(\mathbf{x}) = \int R(\mathbf{x}|\mathbf{y}) f(\mathbf{y}) \, \mathrm{d}\mathbf{y} + b(\mathbf{x}) = \int A(\mathbf{x}|\mathbf{y}) \, \epsilon(\mathbf{y}) \, f(\mathbf{y}) \, \mathrm{d}\mathbf{y} + b(\mathbf{x})$$

- **x**: observed (reconstruction-level) kinematics, **y**: "true" kinematics
- R(x|y) transfer function ("translation" from true to reconstructed kinematics), can be factorized in acceptance function A(x|y) and efficiency function e(y)
- b(x) background distribution

Unfolding Techniques



- Unfolding = solving integral equation for f(y)→ **ill-posed** mathematical problem, typical solutions:
 - First step: discretization (= histograms), **response/migration matrix** *R*

$$g_i = \sum_{j=1}^m R_{ij}f_j + b_i$$

If $R \sim$ diagonal: **bin-by-bin** correction factors c_i may be sufficient:

$$g_i = c_i f_j + b_i$$

• Matrix inversion of R: numerically unstable due to statistical fluctuations \rightarrow additional assumption: smooth distributions ("regularization")

Bin-by-bin Unfolding: $H \rightarrow ZZ \rightarrow 4\ell$



Correction Factors $H \rightarrow ZZ \rightarrow 4\ell$: clean signature, 1.4 high **purity** (i.e. response ATLAS Preliminary — Purity $1.2 - H \rightarrow ZZ^* \rightarrow 4I$ matrix almost diagonal) Unfolded $p_{T}(4\ell)$ Correction factor 13 TeV, 36.1 fb⁻¹ → **bin-by-bin** unfolding d $m o/d_{p_{T,4l}}$ [fb/GeV] ATLAS Preliminary Data 0.14 $H \rightarrow ZZ^* \rightarrow 4l$ Syst. uncertainties 0.8 13 TeV. 36.1 fb⁻¹ HRes k = 1.1. + XH0.12 NNLOPS k = 1.1, +XHReconstructed $p_T(4\ell)$ 0.6 MG5 FxFx k = 1.47, +XH0.1 thit XH = VBF+WH+ZH+ttH+bbH 0.4 3.5F Events / GeV p-value NNLOPS = 25% Data ATLAS Preliminary 0.08 Signal (m_=125 GeV) $H \rightarrow ZZ^* \rightarrow 4I$ p-value MG5 FxFx = 42% 0.2 Background ZZ* 3F 13 TeV. 36.1 fb⁻¹ p-value HRes = 21% Background tt+V. VVV 0.06 Background Z+jets, tt ______ 150 200 250 300 350 50 100 Uncertainty 2.5 0.04 $p_{_{\mathrm{T,4I}}}$ [GeV] 0.02 1.5 Data/Theory 2.5 1.5 0.5 0.5 120 200 350 250 0 10 15 20 30 60 80 200 300 350 50 100 150 $p_{T_{AI}}$ [GeV] $p_{T_{4I}}$ [GeV] ATLAS-CONF-2017-032

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Matrix Unfolding: $H \rightarrow WW$



Matrix-Element Method



- Entire parton-level kinematics of a process contained in squared scattering amplitude ("matrix element", ME)
- Matrix-element method (MEM): construct event-based likelihood discriminant that fully exploits all information from matrix element
 - Likelihood function for a given process contains hard-scattering matrix element for that process (→ next slide)
 - For each event: ratio of likelihood functions for observed set of kinematic variables x under signal hypothesis S and background hypotheses B_i

$$R(\mathbf{x}) = \frac{L(\mathbf{x}|S)}{L(\mathbf{x}|S) + \sum_{i} c_{i} L(\mathbf{x}|B_{i})}$$

Full discriminant: product of event-based discriminants for all events

Matrix Element and Phase Space



- Main ingredient of event-based likelihood: differential parton-level cross sections for signal and (main) backgrounds
 - Consider cross section for all processes $pp \rightarrow y$ with parton-level kinematics y that could have led to the reconstruction-level final state x with kinematics x

$$\sigma(pp \to y) = \sum_{jk}^{\text{partons}} \int_{0}^{1} dz_{j} dz_{k} \frac{f_{j}(z_{j}) f_{k}(z_{k})}{PDFs} \frac{(2\pi)^{4}}{z_{j} z_{k} s} \frac{|M(jk \to y)|^{2} d\Phi}{\text{matrix}}$$

- Approach uses **QCD factorization**: PDFs f_j , f_k , (squared) hard-scattering matrix element M, Lorentz-invariant phase space measure d Φ
- Current implementations: **LO matrix elements** (NLO in the works)
- Integration over all unobserved variables in the event: momentum fractions of colliding partons, phase space integral → often numerically expensive

Transfer Functions



- Translation of parton-level final state to **reconstruction level**:
 - Folding with **transfer functions** $W(\mathbf{x}|\mathbf{y})$ determined from MC simulation
 - W(x|y) accounts for limited detector resolution and combinatorics in matching parton level and reconstruction level objects (especially quarks/ gluons → jets)

$$\sigma(pp \to x) = \int \sigma(pp \to y) W(\mathbf{x}|\mathbf{y}) \, \mathrm{d}\mathbf{y}$$

Normalization to (fiducial) cross section of process *P* = *S*, *B*

$$L(\mathbf{x}|P) = \frac{\sigma(pp \to x)}{\sigma_{obs}^{P}} \quad \text{with} \quad \sigma_{obs}^{P} = \int \sigma(pp \to y) \ W(\mathbf{x}|\mathbf{y}) \cdot f_{acc}(\mathbf{x}) \, d\mathbf{x} \, d\mathbf{y}$$

and $f_{acc}(\mathbf{x}) = 0$;1 acceptance for single event with kinematics \mathbf{x}

MEM Application: $H \rightarrow ZZ \rightarrow 4\ell$

- $H \rightarrow ZZ \rightarrow 4\ell$ angular analysis \rightarrow Higgs-boson spin and parity
- Kinematics fully determined by
 - 2 masses *m*_{Z1}, *m*_{Z2}
 - Decay planes of $Z_{1,2}$ → 5 angles **Ω** = (θ^* , ϕ_1 , ϕ , θ_1 , θ_2)
 - Polar angle of Z bosons (θ^*)
 - Azimuthal angle of Z_1 plane (ϕ_1)
 - Azimuthal angle of Z₂ plane relative to Z₁ plane (φ)
 - Polar angles of leptons relative to $Z_{1,2}$ ($\theta_{1,2}$)





R. Wolf

MEM Application: MELA

- Application of MEM to angular analysis of $H \rightarrow ZZ \rightarrow 4\ell$
 - MELA: Matrix Element Likelihood Analysis (based on PRD 81 (2010) 075022) → already applied for CMS Higgs discovery analysis
 - Purely leptonic final state: no phase space integration and transfer functions required
 - MELA discriminant:



$$K_D = \frac{L(m_{Z1}, m_{Z2}, \Omega; m_{4\ell}|S)}{L(m_{Z1}, m_{Z2}, \Omega; m_{4\ell}|S) + L(m_{Z1}, m_{Z2}, \Omega; m_{4\ell}|B)}$$







Experimental particle physics: many tools and techniques, e.g.

- Simulation of collision processes: **Monte-Carlo** event generators
- Reconstruction, ID, calibration of all **physics objects**
- Treatment of **background** processes
- Many opportunities for you to dig deep into particle physics



Tutorial **TAG&PROBE EFFICIENCY**



- Tag&probe trigger efficiency definition: $\epsilon = \frac{N(\text{reference \&\& probe})}{N(\text{reference})} = \frac{N(\text{HLT_IsoMu20 \&\& HLT_PFJet500})}{N(\text{HLT_IsoMu20})}$
 - Motivate this definition by discussing the following questions:
 - What is the purpose of the reference trigger?

Cannot measure absolute efficiency in data.

Why do we require the muon of the reference trigger to be isolated? What could happen if one considers any muon (also non-isolated)?

Tag should be independent of probe \rightarrow muon. Non-isolated muon may be part of a jet \rightarrow not independent.



Tag&probe trigger efficiency definition:

 $\epsilon = \frac{N(\text{reference \&\& probe})}{N(\text{reference})} = \frac{N(\text{HLT_IsoMu20 \&\& HLT_PFJet500})}{N(\text{HLT_IsoMu20})}$

• Could we use a trigger that fires at random as the reference trigger? Yes, but the rate will be much too low \rightarrow see later.

Could we, instead of using the single-muon trigger, use a trigger that requires the presence of a jet with a p_T threshold lower than the threshold of HLT PFJet500 as the reference trigger, e.g. a trigger requiring a jet with $p_T > 300$ GeV?

Yes, but but reference trigger must be constant \rightarrow see later.



Inspect the two jet p_T histograms and discuss the following questions:

- Why does the number of entries per bin decrease towards large p_T? Differential cross sections of all processes decrease with increasing p_T
- What is the reason for the turn-on at low p_T ?

Events recorded with lepton trigger are mainly W+jets events, minimum momentum transfer \rightarrow minimal p_T of order of $m_w/2$



Inspect the trigger efficiency plot and answer the following questions:

- What is the efficiency of the HLT PFJet500 trigger path? Heavily p_T-dependent: zero efficiency → turn-on → plateau (if efficiency is quoted: plateau efficiency)
- Why is there a smooth turn-on region around 500 GeV where the efficiency gradually increases? Why does the trigger not reach its maximum efficiency instantly at $p_T = 500$ GeV?

Online reconstruction of p_T (in trigger) and offline reconstructed (for plotted p_T) different (\rightarrow resolution effect)



- In the light of the turn-on feature of a trigger efficiency, consider again the question:
 - Could we, instead of using the single-muon trigger, use a trigger that requires the presence of a jet with a lower p_T threshold than HLT PFJet500, e.g. a trigger that requires a jet with p_T > 300 GeV?
 - Reference trigger must be in plateau (not necessarily fully efficient!), i.e. beyond on turn-on, before turn-on of probe trigger starts
 - Which condition must be satisfied when a trigger with lower threshold is used as the reference trigger?

Need sufficient difference in thresholds. Exact conditions depend on the offline-vs-online resolution of objects, typically large for jets, small for leptons

Uncertainty of the Efficiency



- Have a look at the error bars in the efficiency plot produced in the previous exercise:
 - Are they reasonable? Not really, they extend below 0 and above 1.
 - How are they calculated? Error propagation of Poisson uncertainties.
- You can switch to using binomial uncertainties by adding the "B" option to the TH1::Divide method.
 - How do the error bars change? Error bars vanish for for 0 and 1.
 - Is this reasonable?
 Yes, except for 0 and 1.
- Adjust calculate eff.py to use Clopper–Pearson intervals as error bars: How do the error bars change? Is this reasonable? Yes, for entire efficiency interval [0,1].

When is Trigger "Fully Efficient"?



- Given the turn-on feature, we can assume the trigger efficiency to remain constant for large p_T far above the turn-on region. Why? Jet trigger and reconstruction stays fully efficient to very high p_T.
- A suitable function to fit the turn-on is

$$f(p_T; a_0, a_1, a_2) = \frac{1}{2} \cdot a_2 \cdot \left[\operatorname{erf}(\frac{1}{\sqrt{2}a_0}(p_T - a_1)) + 1 \right]$$

- How is the error function erf(x) defined and why is it suitable in this case? (Remember again what the reason for the turn-on feature was!) Cumulative distribution of the normal (Gaussian) distribution. Suitable because turn-on is a (Gaussian) resolution effect.
- What is the interpretation of the parameters a_i? (Which trigger threshold do you find? What is the efficiency of the trigger above the threshold?)

 a_0 : width of the Gaussian a_1 : turn-on point (50% of plateau efficiency) a_2 : normalization \rightarrow plateau efficiency

Efficiency of a Different Trigger



We want to use the tools developed above to measure the efficiency of a single-jet trigger with a different threshold. Adjust your calculate eff.py to measure the efficiency of the HLT PFJet60 trigger What do you observe?

This should be possible in principle (but not with the way the histograms in histos.root were prepared). However, certain triggers may have been pre-scaled (i.e. only every *n*-th event recorded) \rightarrow observable consequence: plateau much below 1