### Jet Substructure

4/27/2016 Physics 290E Brian Amadio As the center-of-mass energy of the LHC increases, it becomes harder to resolve the decay products of increasingly boosted SM particles.



### For 1.6 TeV Z'→ tt, very few of the top quarks can be fully resolved with R=0.4 jets



### Collider energies and heavy resonance limits are going up → The future of jets lies in jet substructure!



From BOOST 2015 Summary Talk

# Many BSM searches benefit from jet substructure methods.







## Jet substructure is a wide, active, and rapidly expanding field.



# The most straightforward jet substructure variable is mass.

$$m^2 = \left(\sum E_i\right)^2 - \left(\sum \vec{p_i}\right)^2$$

Boosted resonance:  $m_{jet} \approx m_{resonance}$ 

QCD jet: ???

### QCD jets are not massless!

### For finite-radius jets, the mass is sensitive to unassociated radiation in the event.



Trimming is one of several jet grooming methods used to reduce this sensitivity.



Unassociated radiation should be ≈evenly distributed in the event, while radiation from parent partons is clustered near the parent. Use k<sub>T</sub> for reclustering to keep more FSR.

Mass-drop/filtering takes advantage of the fact that large changes in mass are exponentially suppressed in QCD showers.

Sudakov form factor 
$$\Delta(t) = exp\left[-\int_{t_0}^t \frac{dt'}{t'} dz \frac{\alpha_s}{2\pi} \mathcal{P}(z)\right]$$
  
$$\Delta(t) \propto \left(\frac{t_0}{t}\right)^p$$

But when a heavy particle decays, large changes in mass are guaranteed!

Mass-drop/filtering attempts to optimize the reclustering scale for a specific decay topology, originally h→bb.

Algorithm: unwind C/A jets and require:

- 1. large mass drop  $max(m_1, m_2) < \mu m_J$ and  $\cdot (2 - 2)$
- 2. symmetric splitting

$$\frac{\min\left(p_{T1}^2, p_{T2}^2\right)}{m_J^2} \Delta R_{1,2}^2 > y_{cut}$$

If either requirement fails, throw out softer subjet and unwind harder subjet with same method. Continue until a splitting passes both requirements to obtain  $R_{filt} = min(0.3,\Delta R/2)$  Then recluster N subjets at  $R_{filt}$ .

Pruning, inspired by filtering, explicitly vetoes soft, wide angle radiation.

Algorithm: unwind C/A jets and require:

 hard splitting or
 small angle

$$\frac{\min\left(p_{Ti}, p_{Tj}\right)}{p_{TJ}} > z_{cut}$$
$$\Delta R_{ij} < D_{cut}$$

At each step, if both checks fail, drop the softer constituent and unwind the harder. Stop when either 1 or 2 are satisfied.

## All grooming methods come with parameters that have to be tuned.



# Grooming can significantly reduce pileup sensitivity.



# Grooming improves the jet mass resolution for signal and reduces the QCD background.



(a) dijets, 500–600 GeV

(b)  $t\bar{t}$ , 500–600 GeV

Pruning most aggressive at small mass, as D scales like m/p<sub>T</sub>, whereas R<sub>sub</sub> is fixed. Filtering least aggressive, as it keeps a fixed number of subjets - in this case 3.

## N-subjettiness quantifies the degree to which a jet is comprised of N subjets.

Given N axes  $\hat{n}_k$ ,  $\tau_N = \frac{\sum_{i \in J} p_{T,i} min\left(\Delta R_{ik}\right)}{\sum_{i \in J} p_{T,i} R_0}$ 

 τ<sub>N</sub> small: jet is well
 described by N or
 fewer subjets

 τ<sub>N</sub> large: more subjets
 will likely describe jet
 better



# Ratios of N-subjettiness provide more discriminating power.









The kT splitting scale can also be used to separate QCD jets from boosted resonance decay jets.

Recluster jets using k<sub>T</sub> algorithm  $\sqrt{d_{ij}} = min(p_{Ti}, p_{Tj}) \Delta R$  $\left< d_{23} \right> \approx m_W / 2$  $\left( d_{12} \right) pprox m_t / 2$ For QCD jets:  $d_{34} \ll d_{23} \ll d_{12}$ 

## To study top-tagging performance, we use a semileptonic ttbar selection.



### Selection

1 lepton, 2 b-tagged jets.

#### **Truth-matching**

large-R jet within  $\Delta R = 0.75$  of hadronicallydecaying top at decay vertex.

# Substructure-based top-tagging can combine mass, n-subjettiness, and k<sub>T</sub> splitting scales.



(c)

(d)

## The signal-to-background is improved as compared to mass cuts alone.



Tagger	Top-tagging criterion
Substructure tagger I	$\sqrt{d_{12}} > 40 { m GeV}$
Substructure tagger II	m > 100  GeV
Substructure tagger III	$m > 100 \text{ GeV}$ and $\sqrt{d_{12}} > 40 \text{ GeV}$
Substructure tagger IV	$m > 100 \text{ GeV}$ and $\sqrt{d_{12}} > 40 \text{ GeV}$ and $\sqrt{d_{23}} > 10 \text{ GeV}$
Substructure tagger V	$m > 100 \text{ GeV}$ and $\sqrt{d_{12}} > 40 \text{ GeV}$ and $\sqrt{d_{23}} > 20 \text{ GeV}$
W' top tagger	$\sqrt{d_{12}}$ > 40 GeV and 0.4 < $\tau_{21}$ < 0.9 and $\tau_{32}$ < 0.65

Table 2: Top taggers based on substructure variables of trimmed anti- $k_t R = 1.0$  jets.

### Performance is studied in MC, using Z'→ ttbar for signal and multijets for background.



### Additional Material

#### HEPTopTagger

#### Use Cambridge-Aachen, R = 1.5 jets Algorithm

- unwind the jet and require max(m<sub>1</sub>,m<sub>2</sub>) > 0.8 m<sub>jet</sub> to keep both constituents, otherwise keep higher mass
- keep unwinding all subjets as long as m > 30 GeV. Keep all subjets that haven't been discarded by requirements in step 1

cluster subjets with  $R_{filt} = min(0.3, \Delta R_{jk}/2)$ calculate mass of 5 hardest constituents in each subjet

choose pairings of subjets to obtain total mass closest to  $\ensuremath{m_t}$ 

### Because of changes in q/g, $\alpha_s$ , and pdfs, QCD jet mass is not quite linear in $p_T$



Better than 25%