Jet Substructure in Julia

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Initial Benchmarks

Comparing N2Plain Strategy: Julia and Python

In this strategy, every particle is compared with every other particle to determine the nearest neighbors according to the chosen distance metric (e.g., k_t , anti- k_t , or Cambridge/Aachen)

The adjoining plot shows that, similar to C++, the python bindings show poor scaling compared to Julia at higher particle density



Comparing N2Tiled Strategy: Julia and Python

In this strategy, the space in which the particles are distributed (usually the rapidity-azimuth plane) is divided into a grid of tiles. Each particle is assigned to a tile based on its coordinates. Instead of comparing every particle with every other particle, the algorithm only compares particles within the same tile and neighboring tiles. This reduces the number of distance calculations significantly.



It can be seen that like C++, there is small advantage of Julia over Python also.

Overall comparison: Julia and Python



From this graph, we see that, in terms of efficiency we have:

N2TiledJulia > N2TiledPython > N2PlainJulia > N2PlainPython

With N2Tiled Julia being the most efficient, for higher particle density.

Jet Substructure Modules

Jet Filtering

The jet filtering algorithm was implemented in Julia, and the corresponding result was compared with Python. In the adjoining plots, the y-axis represents the groomed parameters as obtained from Julia, and the x-axis represents the groomed parameters obtained from Python.



Figure: Comparing filtered pT

Figure: Comparing filtered eta

Jet Filtering: Efficiency

The adjoining plot compares the efficiency of the jet filtering algorithm implemented in Python and Julia.



Jet Trimming

A similar analysis for the jet trimming algorithm was done, and the following plots were obtained.



Jet Trimming: Efficiency



The adjoining plot compares the efficiency of the jet trimming algorithm implemented in Python and Julia. This follows a similar trend to that of jet filtering.

Filtering & Trimming: Codes

• Filter() (3/4)	
fastjet::Fite::#iter (double Rfit,	
Selector selector, double mis = 0.0	
)	
Same as the full constructor (see above) but just specifying the radius By default, Cambridge-Aachen is used If the jet (or all its pieces) is obtained with a non-default recombiner, that one will be used.	
Parameters Rfill the fibering radius	

Definition at line 124 of file Filter.hh.





Mass Drop Tagger

The MassDrop Tagging algorithm was similarly implemented in Julia and compared with the corresponding Python bindings, for the same set of data. The obtained results were concurrent with each other (shown in the plots below).



Figure: Comparing groomed pT

Figure: Comparing groomed mass

Mass Drop Tagger: Efficiency

As can be seen from the adjoining plot, the implemented Julia MassDrop Tagger is much more efficient compared to the corresponding Python bindings.



Mass Drop Tagger: Codes

```
PseudoJet MassDropTagger::result(const PseudoJet & jet) const{
    PseudoJet j = jet;
                                                                                                                                                                                                                                                                                                                                                                                      truct MassDropTagge
      unction apply massdropliet: Pseudolet, clusterseg: ClusterSequence, tag: MassDropTageer]
                                                                                                                                                                                                                                                                                                                                                                                              alliets = clustersen jets
      Pseudolet j1, j2:
bool had parents:
                                                                                                                                                                                                                                                                                                                                                                                              hist = clusterseq.history
      // we just ask that we can "walk" in the cluster sequence.
// appropriate errors will be thrown automatically if this is not
// the case
                                                                                                                                                                                                                                                                                                                                                                                                        had_parents, p1, p2 = has_parents(jet, hist)
                    c ((had_parents = j.has_parents(j1,j2))) {
  (j.#2() <= 0) {</pre>
                 if [had_parents]
                                                                                                                                                                                                                                                                                                                                                                                                                   parent1 = allJets[hist[p1].ietp index]
            // make parent1 the more massive jet
if (j1.m2() < j2.m2()) std::swap(j1,j2);</pre>
         // if we pass the conditions on the mass drop and its degree of
// if we pass the conditions on the mass drop and its degree of
// if (log)
// if
                                                                                                                                                                                                                                                                                                                                                                                                                            p1, p2 = p2, p1
                                                                                                                                                                                                                                                                                                                                                                                                                            parent1, parent2 = parent2, parent1
                                                                                                                                                                                                                                                                                                                                                                                                                   if (in2(parent1) < n2(iet)+tag.mu*2) 66 (kt_distance[parent1, parent2) > tag.vem2(iet)))
                 j = j1;
                                                                                                                                                                                                                                                                                                                                                                                                                                jet = parents
      if (!had parents)
    // no Higgs found, return an empty PseudoJet
    return PseudoJet();
    // create the result and its structure
Pseudolet result_local = j;
MassDropTaggerStructure = s = new MassDropTaggerStructure(result_local);
s => mu = j1.el) / j.nl;
s => mu = j1.el) / j.nl;
s => mu = j1.el) / j.nl;
      result_local.set_structure_shared_ptr(SharedPtr+PseudoJetStructureBase>(s));
      return result local:
```

Soft Drop Tagger

The SoftDrop Tagging algorithm was also implemented in Julia, but this time we compared it with the C++ FastJet Bindings, for the same set of data. The obtained results were concurrent with each other (shown in the plots below).



Figure: Comparing groomed pT

Figure: Comparing groomed mass

Plans Moving Forward

What More To Do

- Optimise and improve the built modules
- Add the remaining modules
- Build an analysis module
- Contribute to the already existing code base

I have added all the codes I have developed till now in this GitHub repository: Ojulia-JetSubstructure (https://github.com/sattwamo/julia-JetSubstructure)