

Monte Carlo reweighting for hadronization uncertainties

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Hadronization

Monte Carlo event generators simulate full particle collisions in three blocks:



Parameter uncertainties

In the string model, the probability f(z) to emit a hadron with a fraction of the string's longitudinal lightcone momentum z depends on parameters fixed from fits to experimental data, *e.g.*, r_b :

Reweighting

Generating simulation for each input parameter value is computationally expensive, especially when including detector effects. Reweighting calculates alternative weights for each event so that a single simulation can be used to estimate the effect of different param-

- 1. *hard process* initial high energy interaction (perturbative)
- 2. evolution perturbate to nonperturbative scale $\sim 1\,{\rm GeV}$
- 3. *hadronization* combine quarks and gluons into hadrons via phenomeno-logical models
- In the string model of hadronization (implemented in PYTHIA), partons are bound by the strong force into strings, which are iteratively split into hadrons:





This is conceptually straightforward but can be technically challenging.



Modified accept-reject

A probability distribution P(z) can be sampled when a reliable overestimate \widehat{P} is known. In the standard acceptreject algorithm, for a trial value of z,



Replicating distributions

Simulation generated with one set of hadronization parameter values can be reweighted to match another by implementing the modified accept-reject algorithm in PYTHIA, *e.g.*, values of r_b : $r_b=0.657$ $r_b=0.459$ $r_b=1.792$

Improving timing

The average time required to generate a single event using the modified accept-reject algorithm scales linearly as a function of the number of alternative values considered for parameter *a*:



 $P_{\text{accept}}(z) \equiv \frac{P(z)}{\widehat{p}} \leq 1.$

If an event generated with P(z) has weight w, the weight w' for an event generated with P'(z) can be calculated by tracking *all* the trial z values:





In this example, generating an additional value of a using the modified accept-reject algorithm is $\approx 6.5 \times$ faster than not using it; were detector simulation included, the speedup would be far greater, $\mathcal{O}(1000)$.

Publication

These findings have been submitted to SciPost, and the pre-print is available as arXiv:2308.13459.