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Time-like Showers and Matching with Antennae

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Parton Showers

- ► The final answer depends on:
 - The choice of evolution variable
 - The splitting functions (finite/subleading terms not fixed)
 - The phase space map ($d\Phi_{n+1}/d\Phi_n$)
 - The renormalization scheme (argument of α_s)
 - The infrared cutoff contour (hadronization cutoff)
- Step 1, Quantify uncertainty: vary all of these (within reasonable limits)
- Step 2, Systematically improve: Understand the importance of each and how it is canceled by
 - Matching to fixed order matrix elements
 - Higher logarithms, subleading color, etc, are included
- Step 3, Write a generator: Make the above explicit (while still tractable) in a Markov Chain context → matched parton shower MC algorithm



VIRTUAL NUMERICAL COLLIDER WITH INTERLEAVED ANTENNAE

VINCIA

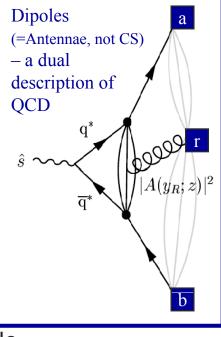
Gustafson, PLB175(1986)453; Lönnblad (ARIADNE), CPC71(1992)15. Azimov, Dokshitzer, Khoze, Troyan, PLB165B(1985)147 Kosower PRD57(1998)5410; Campbell,Cullen,Glover EPJC9(1999)245

Based on Dipole-Antennae

- Shower off color-connected pairs of partons
- Plug-in to PYTHIA 8.1 (C++)
- So far: Giele, Kosower, PS : hep-ph/0707.3652 + Les Houches 2007
 - Time-like QCD cascades (with massless quarks)
 - 2 different shower evolution variables:
 - pT-ordering (= ARIADNE ~ PYTHIA 8)
 - Dipole-mass-ordering (~ Thrust ≠ PYTHIA 6, SHERPA)
 - For each: an infinite family of antenna functions
 - Laurent series in branching invariants with arbitrary finite terms

Shower cutoff contour: independent of evolution variable → IR factorization "universal"

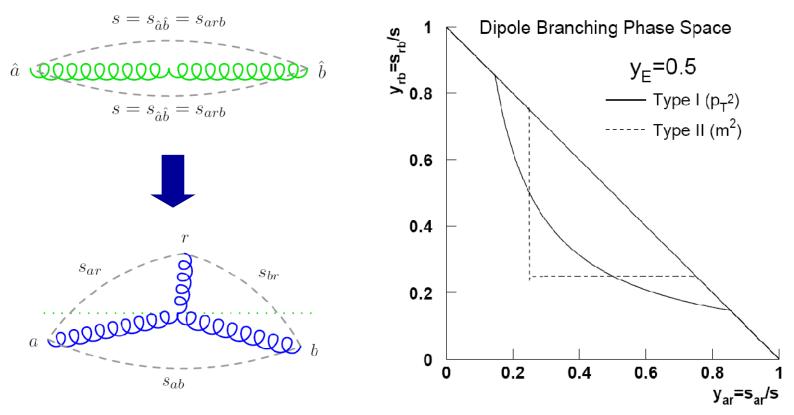
- Several different choices for α_s (evolution scale, p_T , mother antenna mass, 2-loop, ...)
- Phase space mappings: 2 different choices implemented
 - Antenna-like (ARIADNE angle) or Parton-shower-like: Emitter + longitudinal Recoiler





Dipole-Antenna Showers

Dipole branching and phase space



(\rightarrow Most of this talk, including matching by antenna subtraction, should be applicable to ARIADNE as well)

Giele, Kosower, PS : hep-ph/0707.3652



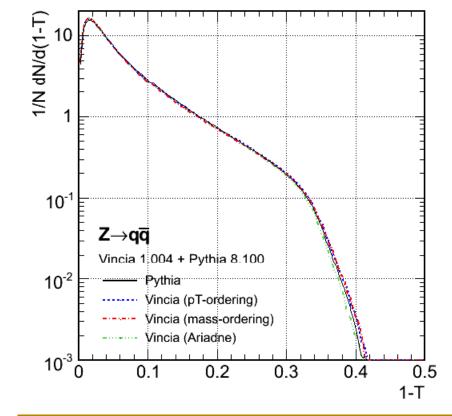
Example: Z decays

VINCIA and PYTHIA8 (using identical settings to the max extent possible)

 $\alpha_{s}(p_{T}),$ $p_{Thad} = 0.5 \text{ GeV}$ $\alpha_{s}(m_{Z}) = 0.137$ $N_{f} = 2$

Note: the default Vincia antenna functions reproduce the $Z \rightarrow 3$ parton matrix element;

Pythia8 includes matching to $Z \rightarrow 3$



Frederix, Giele, Kosower, PS : Les Houches Proc., in preparation



Example: Z decays

- Why is the dependence on the evolution variable so small?
 - Conventional wisdom: evolution variable has huge effect
 - Cf. coherent vs non-coherent parton showers, mass-ordered vs p_T -ordered, etc.
- Dipole-Antenna showers resum radiation off pairs of partons
 - \rightarrow interference between 2 partons included in radiation function
 - If radiation function = dipole formula \rightarrow intrinsically coherent
 - \rightarrow angular ordering there by construction Gustafson, PLB175(1986)453
 - Remaining dependence on evolution variable much milder than for conventional showers
- The main uncertainty in this case lies in the choice of radiation function away from the collinear and soft regions
 - \rightarrow dipole-antenna showers under the hood ...



Dipole-Antenna Functions

Giele, Kosower, PS : hep-ph/0707.3652

Starting point: "GGG" antenna functions, e.g., gg→ggg:

$$Gehrmann-De Ridder, Gehrmann, Glover, JHEP 09 (2005) 056$$

$$f_3^0(p_a, p_r, p_b) = \frac{1}{s^{[i]}} \left[(1 - y_{ar} - y_{rb}) \left(\underbrace{\frac{2}{y_{ar}y_{rb}}}_{\text{"soft"}} + \underbrace{\frac{y_{ar}}{y_{rb}}}_{\text{"collinear"}} + \underbrace{\frac{y_{rb}}{y_{ar}}}_{\text{"collinear"}} \right) + \frac{8}{3} \right] \qquad \underbrace{y_{ar} = \frac{s_{ar}}{s_i}}_{s_i = \text{ invariant}}$$

 $s_i = invariant$ mass of i'th dipole-antenna

• Generalize to arbitrary Laurent series:

$$A(s_{ar}, s_{rb}) = \frac{4\pi\alpha_s(\mu_R)N_C}{s_{arb}} \sum_{\alpha,\beta \ge -1} C_{\alpha\beta} \frac{s_{ar}^{\alpha} s_{rb}^{\beta}}{s_{arb}^{\alpha+\beta}} \qquad \begin{array}{l} \text{Singular parts fixed,} \\ \text{finite terms arbitrary} \end{array}$$

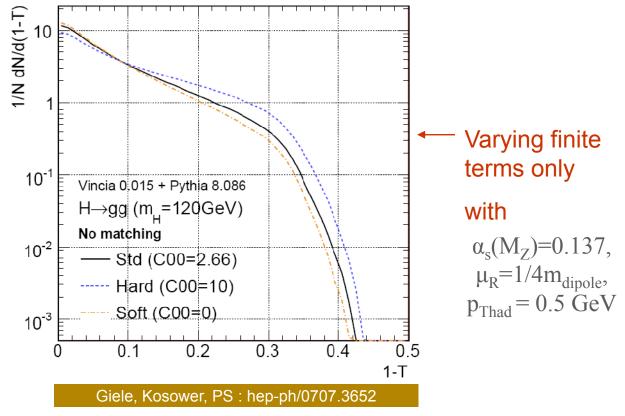
→ Can make shower systematically "softer" or "harder"

- Will see later how this variation is explicitly canceled by matching
- → quantification of uncertainty
- \rightarrow quantification of improvement by matching

Quantifying Matching

► The unknown finite terms are a major source of uncertainty

- DGLAP has some, GGG have others, ARIADNE has yet others, etc...
- They are arbitrary (and in general process-dependent)





Tree-level matching to X+1

1. Expand parton shower to 1st order (real radiation term)

$$\mathbf{PS} : \int \mathrm{d}\Phi_X |M_{X+0}^{(0)}|^2 \int_{t_{X+0}}^{t_{\mathrm{had}}} \mathrm{d}t_{X+1} \sum_i \int \frac{\mathrm{d}\Phi_{X+1}^{[i]}}{\mathrm{d}\Phi_X} \,\delta(t_{X+1} - t^{[i]}(\{p\}_{X+1})) A_i(\ldots) \delta\left(\mathcal{O} - \mathcal{O}(\{p\}_{X+1})\right)$$

$$\mathbf{ME} : \int_{t < t_{had}} \mathrm{d}\Phi_{X+1} |M_{X+1}^{(0)}|^2 \delta\left(\mathcal{O} - \mathcal{O}(\{p\}_{X+1})\right) \qquad \qquad \text{map} \sim \text{clustering} \\ \{\hat{p}_i\}_X = \{\kappa_i^{-1}(\{p\}_{X+1})\}_X$$

→ Matching Term:

MT :
$$w_{X+1}^{(R)} = |M_{X+1}^{(0)}|^2 - \sum_{i \in X \to X+1} A_i(...) |M_{X+0}^{(0)}(\{\hat{p}_i\}_{X+0})|^2$$

- \rightarrow variations in finite terms (or dead regions) in A_i canceled (at this order) ۲
- (If A too hard, correction can become negative \rightarrow negative weights) ٠

$$= \frac{8\pi\alpha_s(\mu_R)N_c}{M_H^2} \Big(8 - 3C_{00} - (C_{10} + C_{01}) - C_{11}\left(y_{12}y_{23} + y_{23}y_{31} + y_{31}y_{12}\right) + \cdots \Big) |M_2^{(0)}|^2$$

Giele, Kosower, PS : hep-ph/0707.3652

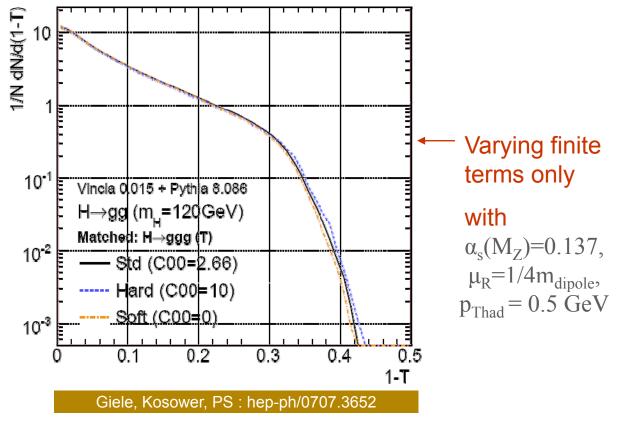


Inverse phase space

0

Quantifying Matching

- ► The unknown finite terms are a major source of uncertainty
 - DGLAP has some, GGG have others, ARIADNE has yet others, etc...
 - They are arbitrary (and in general process-dependent)





1-loop matching to X

- $\blacktriangleright \text{ NLO "virtual term" from parton shower (= expanded Sudakov: exp=1 ...)}$ $PS : -\int d\Phi_X |M_X^{(0)}|^2 \delta \left(\mathcal{O} \mathcal{O}(\{p\}_X) \int_{t_X}^{t_{\text{had}}} dt_{X+1} \sum_i \int \frac{d\Phi_X^{[i]}}{d\Phi_X} \delta(t_{X+1} t^{[i]}(\{p\}_{X+1})) A_i(...)$
- Matrix Elements (unresolved real plus genuine virtual)

$$\mathbf{ME} : \int_{t>t_{had}} \mathrm{d}\Phi_{X+1} |M_{X+1}^{(0)}|^2 \delta\left(\mathcal{O} - \mathcal{O}(\{p\}_{X+1})\right) + \int \mathrm{d}\Phi_X 2\operatorname{Re}[M_X^{(0)}M_X^{(1)*}]\delta\left(\mathcal{O} - \mathcal{O}(\{p\}_X)\right)$$

Matching condition same as before (almost):

$$\begin{aligned} \mathsf{MT:} \quad w_X^{(V)} &= 2\operatorname{Re}[M_X^{(0)}M_X^{*(1)}] + |M_X^{(0)}|^2 \sum_i \int_{\mathrm{all}\,t} \frac{\mathrm{d}\Phi_{X+1}^{[i]}}{\mathrm{d}\Phi_X} A_i(\ldots) + \int_{t>t_{\mathrm{had}}} \mathrm{d}\Phi_{X+1} w_{X+1}^{(R)} \\ &= 2\operatorname{Re}[M_X^{(0)}M_X^{*(1)}] + |M_X^{(0)}|^2 \sum_i \int_{\mathrm{all}\,t} \frac{\mathrm{d}\Phi_{X+1}^{[i]}}{\mathrm{d}\Phi_X} A_i(\ldots) + \mathcal{O}(t_X/t_{\mathrm{had}}) \,, \end{aligned}$$

• (This time, too small $A \rightarrow$ correction negative)

Tree-level matching just corresponds to using zero

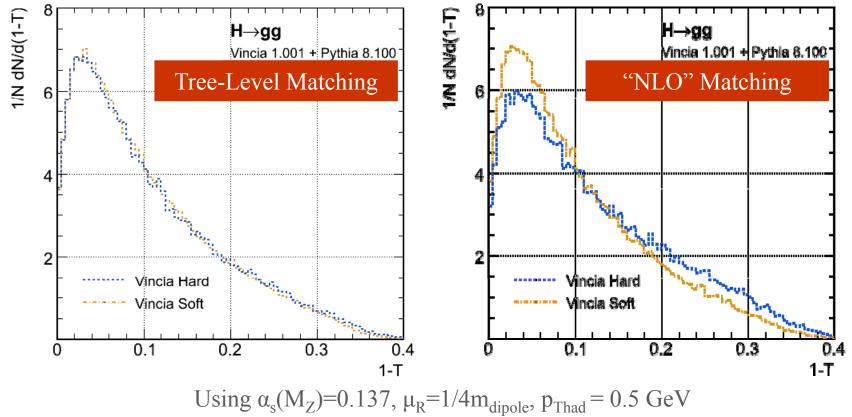
You can choose anything for A_i (different subtraction schemes) as long as you use the same one for the shower

Giele, Kosower, PS : hep-ph/0707.3652



Note about "NLO" matching

- Shower off virtual matching term → unmatched O(α²) contribution to 3-jet observables (only canceled by 1-loop 3-parton matching)
- ► While normalization is improved, most shapes are not





What happened?

- Brand new code, so bear in mind possibility of bugs
- Naïve conclusion: tree-level matching "better" than NLO?
 - No, first remember that the shapes we look at are not "NLO"
 - E.g., 1-T appears at $O(\alpha)$ below 1-T=2/3, and at $O(\alpha^2)$ above
 - An "NLO" thrust calculation would have to include at least 1-loop corrections to $Z \rightarrow 3$
 - So: both calculations are LO/LL from the point of view of 1-T
- What is the difference then?
 - Tree-level $Z \rightarrow 3$ is the same (LO)
 - The O(α) corrections to Z \rightarrow 3, however, are different
 - The first non-trivial corrections to the shape!
 - So there *should* be a large residual uncertainty → the 1-loop matching is "honest"
 - The real question: why did the tree-level matching not tell us?
 - I haven't completely understood it yet ... but speculate it's to do with detailed balance
 - In tree-level matching, unitarity \rightarrow Virtual = Real \rightarrow cancellations. Broken at 1 loop
 - + the tree-level curves had different normalizations (covering the shape uncertainty?)



What to do next?

Further shower studies

- Interplay between shower ambiguities and hadronization "tuning"
- Shower+hadronization tuning in the presence of matching
- Go further with tree-level matching
 - Demonstrate it beyond first order (include H,Z \rightarrow 4 partons)
 - Automated tree-level matching (automated subtractions)
- Go further with one-loop matching
 - Demonstrate it beyond first order (include 1-loop H,Z \rightarrow 3 partons)
 - Should start to see genuine stabilization of shapes as well as normalizations
- Extend to the initial state
- Extend to massive particles
 - Massive antenna functions, phase space, and evolution

We're only a few people working part-time, so plenty of room for collaboration

Extra Material

Frederix, Giele, Kosower, PS : Les Houches Proc., in preparation

- Number of partons and number of quarks
 - N_a shows interesting dependence on ordering variable

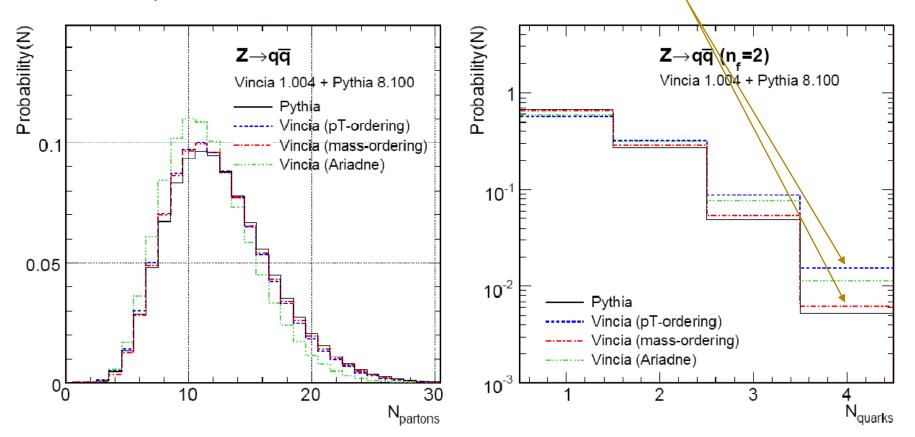


Fig. 3: Number of partons (left) and number of quarks (right) at shower termination, with 2 massless quark flavors.



Comparison $A(s_{ar}, s_{rb}) = \frac{4\pi\alpha_s(\mu_R)N_C}{s_{arb}} \sum_{\alpha,\beta\geq-1} C_{\alpha\beta} \frac{s_{ar}^{\alpha}s_{rb}^{\beta}}{s_{arb}^{\alpha+\beta}}$

			-						-		
		C_{-1-1}	C_{-10}	C_{0-1}	C_{-11}	C_{1-1}	C_{-12}	C_{2-1}	C_{00}	C_{10}	C_{01}
(GGG										
Q	q ar q o q g ar q	2	-2	-2	1	1	0	0	0	0	0
Ģ	$qg \rightarrow qgg$	2	-2	-2	1	1	0	-1	$\frac{5}{2}$	-1	$\frac{3}{2}$
g	$gg \rightarrow ggg$	2	-2	-2	1	1	-1	-1	5 28 3	-1	-1
Q	$qg ightarrow q \overline{q}' q'$	0	0	$\frac{1}{2}$	0	-1	0	1	$-\frac{1}{2}$	1	0
g	$gg \to g\overline{q}q$	0	0	$\frac{\frac{1}{2}}{\frac{1}{2}}$	0	-1	0	1	-1	1	$\frac{1}{2}$
	ARIADNE		I	-					I		_
	$q\bar{q} ightarrow qg\bar{q}$	2	-2	-2	1	1	0	0	0	0	0
4	$qg \rightarrow qgg$	2	-2	-3	1	3	0	-1	0	0	0
g	$gg \rightarrow ggg$	2	-3	-3	3	3	-1	-1	0	0	0
Ģ	$qg \rightarrow qq'q'$	0	0	$\frac{1}{2}$	0	-1	0	1	-1	1	$\frac{1}{2}$
	$gg \to g\bar{q}q$	0	0	$\frac{\overline{1}}{2}$	0	-1	0	1	-1	1	$\frac{\frac{1}{2}}{\frac{1}{2}}$
	ARIADNE2 (reparametrization of ARIADNE functions à la GGG, for comparison)										
Ģ	$q \overline{q} o q g \overline{q}$	2	-2	-2	1	1	0	0	0	0	0
Q	$qg \rightarrow qgg$	2	-2	-2	1	1	0	-1	-1	0	0
9	$gg \rightarrow ggg$	2	-2	-2	1	1	-1	-1	$-\frac{4}{3}$	-1	-1

Table 1: Laurent coefficients for massless LL QCD antennae $(\hat{a}\hat{b} \rightarrow arb)$. The coefficients with at least one negative index are universal (apart from a reparametrization ambiguity for gluons). For "GGG" (the defaults in VINCIA), the finite terms correspond to the specific matrix elements considered in [4]. In particular, the $q\bar{q}$ antenna absorbs the tree-level $Z \rightarrow qg\bar{q}$ matrix element [5] and the qq antennae absorb the tree-level $h^0 \rightarrow qq \rightarrow qqq$ and $h^0 \rightarrow qq \rightarrow q\bar{q}q$ matrix elements [6]. The

MC4BSM



Monte Carlo Tools for Beyond the Standard Model Physics

ORGANIZERS: mc4bsm.AT.phys.ufl.edu

RESOURCES:

- <u>Video Lectures on</u> <u>Monte Carlo for the</u> <u>LHC</u>
- BSM tool repository
- Summary of

3rd workshop: MARCH 10-11, 2008 (CERN)

Note: both days will be full days. Participants who don't want to miss anything should plan on arriving Mar 9 and leaving Mar 12.

Organizing committee: Georges Azuelos, Christophe Grojean, Jay Hubisz, Borut Kersevan, Joe Lykken, Fabio Maltoni, Konstantin Matchev, Filip Moortgat, Steven Mrenna, Maxim Perelstein, Peter Skands : mc4bsm.AT.phys.ufl.edu

Main Focus: Alternative (non-MSSM) TeV scale physics models

http://theory.fnal.gov/mc4bsm/

