

# FASTER PARTON DISTRIBUTION EVALUATION IN MONTE CARLOS

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Based on ZS, [hep-ph/0403055](http://arxiv.org/abs/hep-ph/0403055) and new studies

## Contents:

1. How much time is spent retrieving CTEQ PDFs?  
(i.e. Why do we care?)
2. Speed enhancements to the evaluations:  
How do they work?  
How well do they work?
3. Potpourri (a few other irregularities to watch)

<http://home.fnal.gov/~zack/pdf/> has improved code and details.

## Why should we care?

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More time is spent retrieving PDFs than in any other routine in event generation (by a long way).

I first noticed this when profiling **ZTOP**, my fully differential NLO coding of s- and t-channel single-top-quark production, but it seems to be universal.

Fraction of time spent inside PDF functions  
using default CTEQ PartonX# on single-top

	CTEQ4/5	CTEQ6
<b>ZTOP</b>	90%	60%
<b>HERWIG</b>	70%	33%
<b>PYTHIA</b>	30%	16%

You should care because CTEQ5L and CTEQ5M1 are the default PDF choices for Run 2.

Why does **HERWIG** spend more time than **PYTHIA**?

**PYTHIA** calls **STRUCTM**  $\sim 100$  times/event requested.

**HERWIG** calls **STRUCTM**  $\sim 1100-1800$  times/event requested.

Each call of **STRUCTM** calls the PDF evaluation 8 times.

This means you can call the PDFs up to 13,000,000 times to get 1000 events using CTEQ5 and **HERWIG**!

Why is this so expensive? Why is CTEQ4/5 more expensive than CTEQ6?

CTEQ4/5 and 6 use completely different interpolation algorithms.

I don't want to change results, so how can I speed these up?

## **POLINT and CTEQ PDFs**

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### What is **POLINT**?

It is a routine from “Numerical Recipes” that performs a polynomial fit of degree  $n - 1$  to a set of  $n$  points based on *Neville’s algorithm*.

### Why is it used?

This is used by CTEQ to smoothly interpolate between values of  $x$  and  $Q^2$  that are read in form a best-fit table.

Most of the time spent in CTEQ4/5 occurs in **POLINT**, while CTEQ6 only uses **POLINT** at the endpoints of  $x$  or  $Q^2$ . (which is sometimes expensive).

### Why is **POLINT** so slow?

1. A line that is never called disables compiler optimizations.
2.  $n = 3(4)$  for CTEQ4/5(6). **POLINT**( $n$  points) is too general for most compilers to fully optimize.

Fix this by removing the unreachable line, and writing versions of **POLINT** for CTEQ4/5(6) that set  $n = 3(4)$ .  $\Rightarrow$   
ZS, hep-ph/0403055

There is still a lot of redundancy in **POLINT3**. Therefore, I’ve also written a version that removes all unnecessary calls.  $\Rightarrow$

```

SUBROUTINE POLINT (XA,YA,N,X,Y,DY)
IMPLICIT DOUBLE PRECISION (A-H, O-Z)
C Adapted from "Numerical Recipes"
PARAMETER (NMAX=10)
DIMENSION XA(N),YA(N),C(NMAX),D(NMAX)
NS=1
DIF=ABS(X-XA(1))
DO 11 I=1,N
    DIFT=ABS(X-XA(I))
    IF (DIFT.LT.DIF) THEN
        NS=I
        DIF=DIFT
    ENDIF
    C(I)=YA(I)
    D(I)=YA(I)
11 CONTINUE
Y=YA(NS)
NS=NS-1
DO 13 M=1,N-1
    DO 12 I=1,N-M
        HO=XA(I)-X
        HP=XA(I+M)-X
        W=C(I+1)-D(I)
        DEN=HO-HP
        IF(DEN.EQ.0.)PAUSE
        DEN=W/DEN
        D(I)=HP*DEN
        C(I)=HO*DEN
12 CONTINUE
    IF (2*NS.LT.N-M) THEN
        DY=C(NS+1)
    ELSE
        DY=D(NS)
        NS=NS-1
    ENDIF
    Y=Y+DY
13 CONTINUE
RETURN
END

```

```

cz This specialized recoding assumes N=3.
  SUBROUTINE POLINT3 (XA,YA,N,X,Y,DY)
  IMPLICIT DOUBLE PRECISION (A-H, O-Z)
C   Adapted from "Numerical Recipes"
  DIMENSION XA(3),YA(3),C(3),D(3)
  NS=1
  DIF=DABS(X-XA(1))
  DO 11 I=1,3
    DIFT=ABS(X-XA(I))
    IF (DIFT.LT.DIF) THEN
      NS=I
      DIF=DIFT
    ENDIF
    C(I)=YA(I)
    D(I)=YA(I)
11  CONTINUE
  Y=YA(NS)
  NS=NS-1
  DO 13 M=1,2
    DO 12 I=1,3-M
      HO=XA(I)-X
      HP=XA(I+M)-X
      W=C(I+1)-D(I)
      DEN=HO-HP
cz      IF(DEN.EQ.0.) PAUSE
      DEN=W/DEN
      D(I)=HP*DEN
      C(I)=HO*DEN
12  CONTINUE
    IF (2*NS.LT.3-M) THEN
      DY=C(NS+1)
    ELSE
      DY=D(NS)
      NS=NS-1
    ENDIF
    Y=Y+DY
13  CONTINUE
  RETURN
  END

```

```

cz This is a specialized recoding that
cz   assumes N=3.
cz Written by Z. Sullivan, 5/14/04
  SUBROUTINE POLINT3 (XA,YA,N,X,Y,DY)
C   Modified "Numerical Recipes" routine.
  IMPLICIT NONE
  DOUBLE PRECISION XA(3),YA(3),X,Y,DY
  DOUBLE PRECISION C1,HO,HP,HP2,W,D1,D2,DEN
  INTEGER N

  HO=XA(1)-X
  HP=XA(2)-X
  w=ya(2)-ya(1)
  DEN=HO-HP
  DEN=W/DEN
  D1=HP*DEN
  C1=HO*DEN

  HP2=XA(3)-X
  w=ya(3)-ya(2)
  DEN=HP-HP2
  DEN=W/DEN
  D2=HP2*DEN

  W=HP*DEN-D1
  DEN=HO-HP2

  if((x+x-xa(2)-xa(3)).gt.0d0) then
    y=ya(3)+d2+hp2*w/den
  elseif((x+x-xa(1)-xa(2)).gt.0d0) then
    y=ya(2)+d1+ho*w/den
  else
    y=ya(1)+c1+ho*w/den
  endif

  RETURN
  END

```

## **STRUCTM** and the Monte Carlos

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Both **PYTHIA** and **HERWIG** call PDFs by using the CERNLIB routine **STRUCTM**. This in turn loops over 8 uniquely defined PDFs:  $u_v, d_v, u_s, d_s, s, c, b, g$ .

Much of the code that sets up **POLINT**, or the CTEQ6 algorithm, is repeated with identical results between runs.

An obvious algorithmic improvement is to do 1 of 2 things:

1. **SAVE** the values of  $x$ ,  $Q^2$ , and the results of functions applied to them; and bypass that setup code unless  $x$  or  $Q^2$  change.
2. Write a custom **STRUCTM** to do the direct calls itself.

I won't show these here, but you can find them at:

<http://home.fnal.gov/~zack/pdf/>

How much do each of these changes help? . . .

## Benchmarks give a rough feeling

There are large discrepancies in the amount of execution time between different compilers and compiler flags.

1. **ifc** is ALWAYS faster than **g77** by a factor of 1–3.
2. Using **POLINT3** removes most of the difference between compiler flags.
3. Using **POLINT3**, CTEQ4/5 is  $\sim 20\%$  faster than CTEQ6.
4. Using **POLINT3**, CTEQ5 is faster and more accurate than the functional fit in **PYTHIA**.

Typical speed gains relative to **POLINT3/4** (looping over  $x$ )

Optimization	g77 3.1(2.95)		ifc 6.0	
	CTEQ4/5	CTEQ6	CTEQ4/5	CTEQ6
Default CTEQ	1/(1.1–1.2)	1.0	1/(1.5–2.7)	1/1.03
<b>POLINT3/4</b>	1.0	1.0	1.0	1.0
<b>POLINT3</b> (fast)	1.5	—	1.2	—
<b>SAVE</b> $x, Q^2$	1.26	2.6	1.12	2.4
<b>SAVE</b> $x, Q^2$ (fast)	2.3	—	1.9	—
fastest <b>STRUCTM</b>	3.1	3.1	4.6	2.7
fastest times:	40 s	50 s	17 s	35 s

While these numbers are a nice guide, a simple benchmark misses the fact that real programs interact in subtle ways with their routines. E.g., the CPU has more cache misses, memory alignment is different, . . .

We care about real code





## Speedup for NLO single-top code **ZTOP**

Typical speed gains relative to **POLINT3/4**

Optimization	<b>g77</b> 3.1(2.95)		<b>ifc</b> 6.0	
	<b>CTEQ4/5</b>	<b>CTEQ6</b>	<b>CTEQ4/5</b>	<b>CTEQ6</b>
Default CTEQ	1/1.2	1.0	1/(1.6–2.2)	1.0
<b>POLINT3/4</b>	1.0	1.0	1.0	1.0
<b>POLINT3</b> (fast)	1.3	—	1.25	—
<b>SAVE</b> $x, Q^2$	1.2	2.0	1.13	2.0
<b>SAVE</b> $x, Q^2$ (fast)	1.7	—	1.7	—
fastest <b>STRUCTM</b>	1.9	2.15	1.9, 2.7	2.15
fastest times:	86 s	98 s	60, 42 s	62 s

Using the fastest routines makes **ZTOP** run 2–5 times faster!

Publication-quality runs drop from over day to a few hours.

This is probably the biggest gain a theory calculation can achieve, but factors of 2–3 should be attainable.

I actually got another factor of 1.5 by hard-coding some calls to **CTQNPFD**. Removing unnecessary PDF calls can help.

## Speedup for HERWIG and PYTHIA

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Typical **HERWIG** speed gains relative to **POLINT3/4**

Optimization	<b>g77</b> 3.1(2.95)		<b>ifc</b> 6.0	
	CTEQ4/5	CTEQ6	CTEQ4/5	CTEQ6
Default CTEQ	1/1.12	1.0	1/(1.2–1.7)	1.0
<b>POLINT3/4</b>	1.0	1.0	1.0	1.0
<b>POLINT3</b> (fast)	1.25	—	1.1	—
<b>SAVE</b> $x, Q^2$	1.14	1.6	1.12	1.3
<b>SAVE</b> $x, Q^2$ (fast)	1.5	—	1.3	—
fastest <b>STRUCTM</b>	1.65	1.7	1.53	1.4
fastest times:	64 s	50 s	75 s	55 s

Strangely, **g77** was faster than **ifc**, and CTEQ6 was faster than CTEQ5. I do not know why yet, but it worries me (see next page).

Typical **PYTHIA** speed gains relative to **POLINT3/4**

Optimization	<b>g77</b> 3.1(2.95)		<b>ifc</b> 6.0	
	CTEQ4/5	CTEQ6	CTEQ4/5	CTEQ6
Default CTEQ	1/1.03	1.0	1/(1.15–1.25)	1.0
<b>POLINT3/4</b>	1.0	1.0	1.0	1.0
<b>POLINT3</b> (fast)	1.06	—	1.05	—
<b>SAVE</b> $x, Q^2$	1.04	1.13	1.05	1.13
<b>SAVE</b> $x, Q^2$ (fast)	1.1	—	1.1	—
fastest <b>STRUCTM</b>	1.14	1.15	1.18	1.15
fastest times:	55 s	58 s	42 s	43 s

**HERWIG** is 1.5–2.6 times faster.

**PYTHIA** is 1.15–1.5 times faster ( $\sim 3.3$  times faster than **HERWIG**).

## Some “gotchas” I’ve discovered

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I often use **PFTOPDG** rather than **STRUCTM** because it directly provides  $u, \bar{u}, d, \bar{d}, s, \bar{s}, c, \bar{c}, b, \bar{b}, t, \bar{t}, g$ .

**Warning:** The CERNLIB version just calls **STRUCTM**. New PDFs will have  $s \neq \bar{s}$ , so **PFTOPDG** will not return the correct PDFs as it is currently written.

We’ve already seen **HERWIG** calls the PDFs much too often.

**Before EVERY call to **STRUCTM**, **HERWIG** calls **PDFSET**!**

1. To use new PDFs, you must write your own **PDFSET** to fill the necessary common blocks. **HERWIG** mysteriously overwrites these common blocks, and so must call **PDFSET** to fix it’s mistake.
2. Switching to **LHAPDF** will require writing a CERNLIB-like **PDFSET** that secretly stores an extra copy of the variables, and refills the common blocks with each call.
3. **LHAPDF** currently includes the CTEQ6 evolution code (not CTEQ5). The code in EVLCTEQ is essentially **PFTOPDG**, so the same gains in speed can come from updating **evolvePDF**.

## Conclusions

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1. Retrieving PDFs is still the single most time consuming operation in **HERWIG** and theoretical codes (including those with fast detector simulations).

	CTEQ4/5	CTEQ6
<b>ZTOP</b>	90→42%	60→48%
<b>HERWIG</b>	70→30%	33→23%
<b>PYTHIA</b>	30→9%	16→9%

2. Additional improvements can still be made

- You can replace the binary search for  $x$  and  $Q^2$  with a hash  $\Rightarrow$  factor of 2 for that piece, but only **5%** overall.
- We may have reached the point of diminishing returns.

3. I've made the validated routines available at <http://home.fnal.gov/~zack/pdf/>

At least use **POLINT3**. I'll use the fastest version of **STRUCTM**.

4. Watch out for the interface bug in **PFTOPDG**.

5. It is more important to write correct code, than fast code.  
That does not mean it is unimportant to write fast code. . .