

Group Meeting

8/8/2007

Agenda

- Cell culture (C2C12 regular and 4R-CAT)
- Sonication
- ChIP preps and DNA concentration
- Solexa

Large Cell Culture Effort

**Goal: collect chromatin to be used in a
Solexa ChIP analysis**

Setup: 36x 15 cm plate and 30x 6 cm plate

15 cm plates harvested at 0, ~12, 24 and 60
hrs

6 cm plates fixed at 0, 6, 12, 18, 24, 36, 48,
60, 72, 96 hrs

More Cell Culture

As Biran and others have noted, cell density increases after the removal of serum and addition of differentiation medium.

To compensate, 2x 15 cm plates were used per sample for the 0 hrs timepoint

1x 15 cm plate was used per sample for all post-differentiation timepoints (for ease of comparison)

Even More Cell Culture

Actual timepoints harvested:

C2C12 (WT): 0, 15, 24, 61

C2C12 (4R): 0, 12, 25, 60

Small plates fixed as planned (but there was always a fix accompanying the collection of chromatin)

Sonication

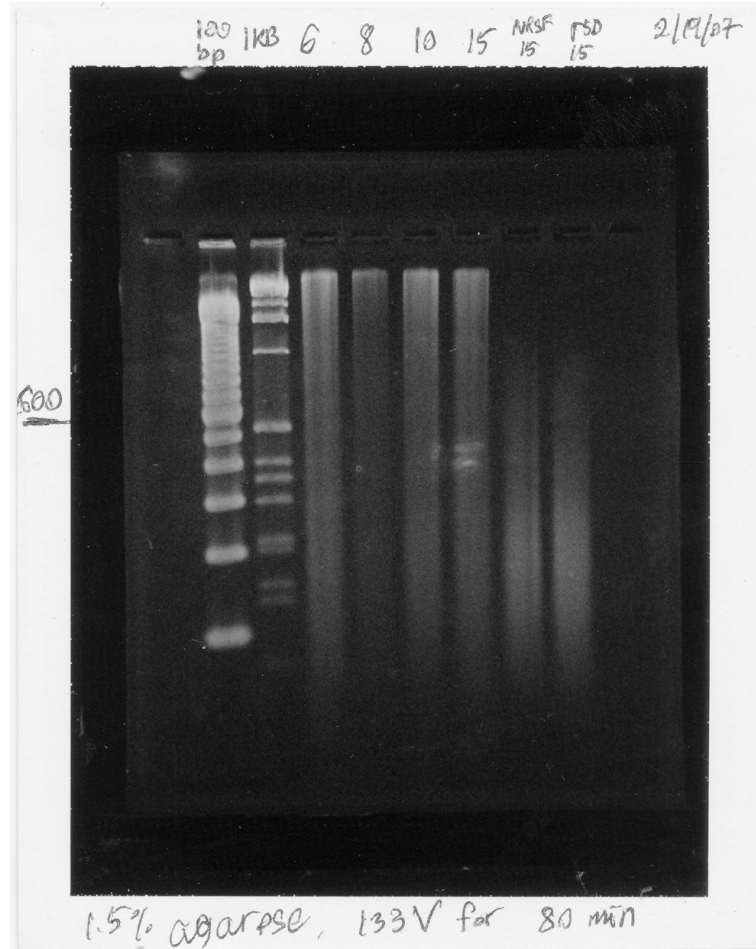
Which parameters are best?

Ali and Gordon get good results with 1 ml
volume

Brian gets good shearing with $\sim 200\text{-}300\ \mu\text{l}$
volume

What is a “good result”?

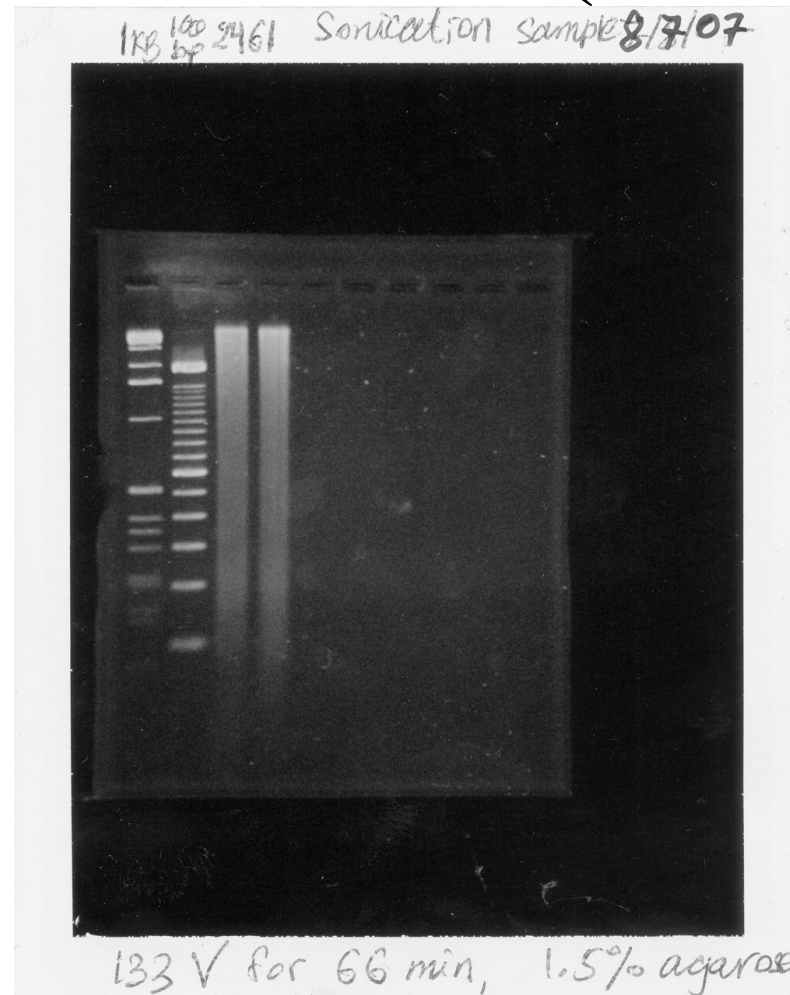
Sonication (5 months ago)



each lane = 1/2
of a 3x 10 cm
plate sample

Lanes 3-6 are sonicated using the number of cycles shown (2.0)
Lanes 7-8 are the product from lane 6 (15 cycles of sonication) that
was purified using the QIAquick purification kit / column.

Sonication (now)



Lanes 1-2 are the 1 KB and 100 bp ladders, respectively
Lanes 3-4 are sonication products of the 24 and 61 hrs diff
time points from the C2C12 (wt) t.c. (12 cycles @ 2.5, 300 μ l)

Sonication

Switching back to using 1 ml now after
having used 200-300 μ l for the last several
months

Ali and Gordon both use 1 ml

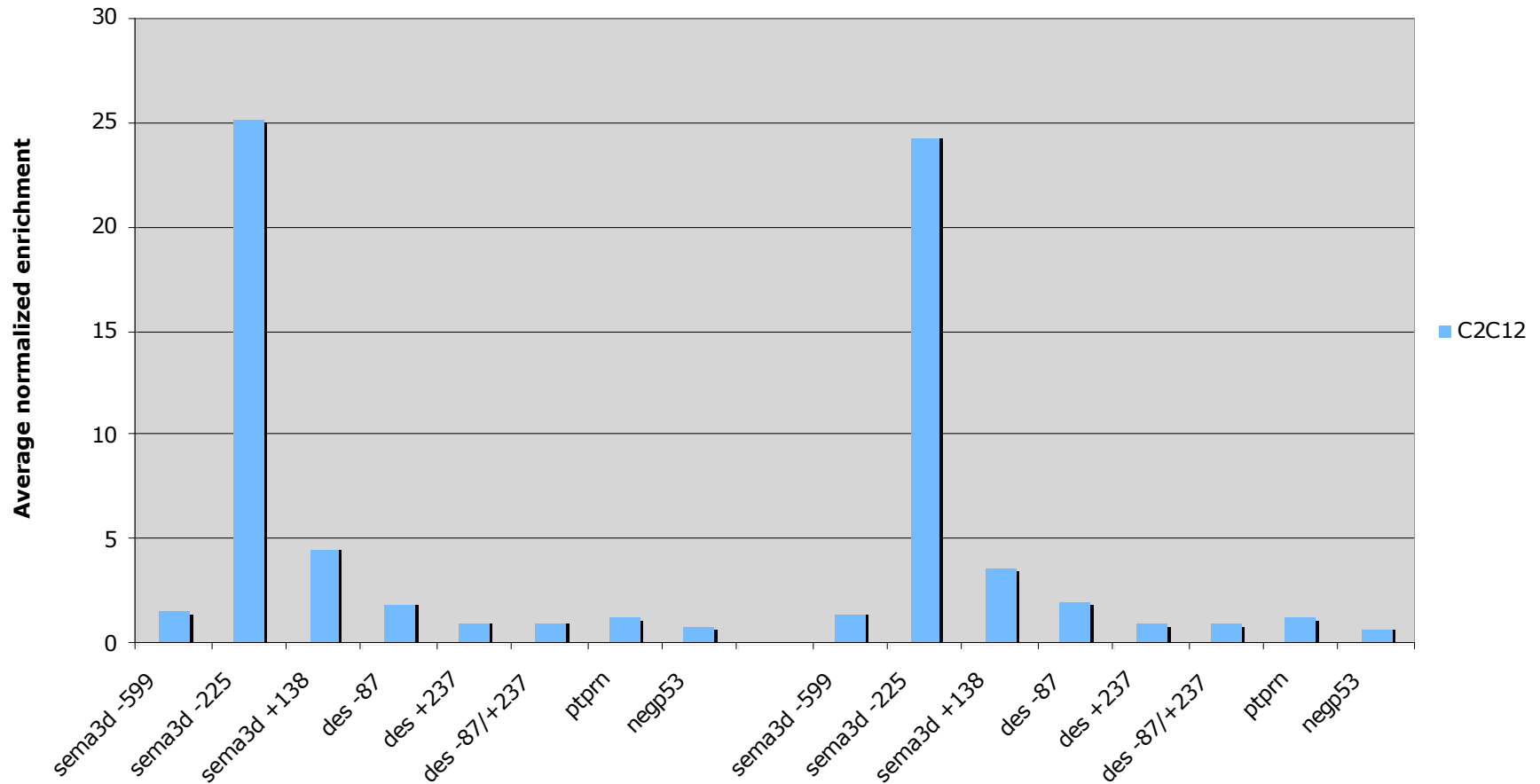
Did a batch of 8 sonications last night using
Gordon's parameters

Foaming is a problem

Results pending

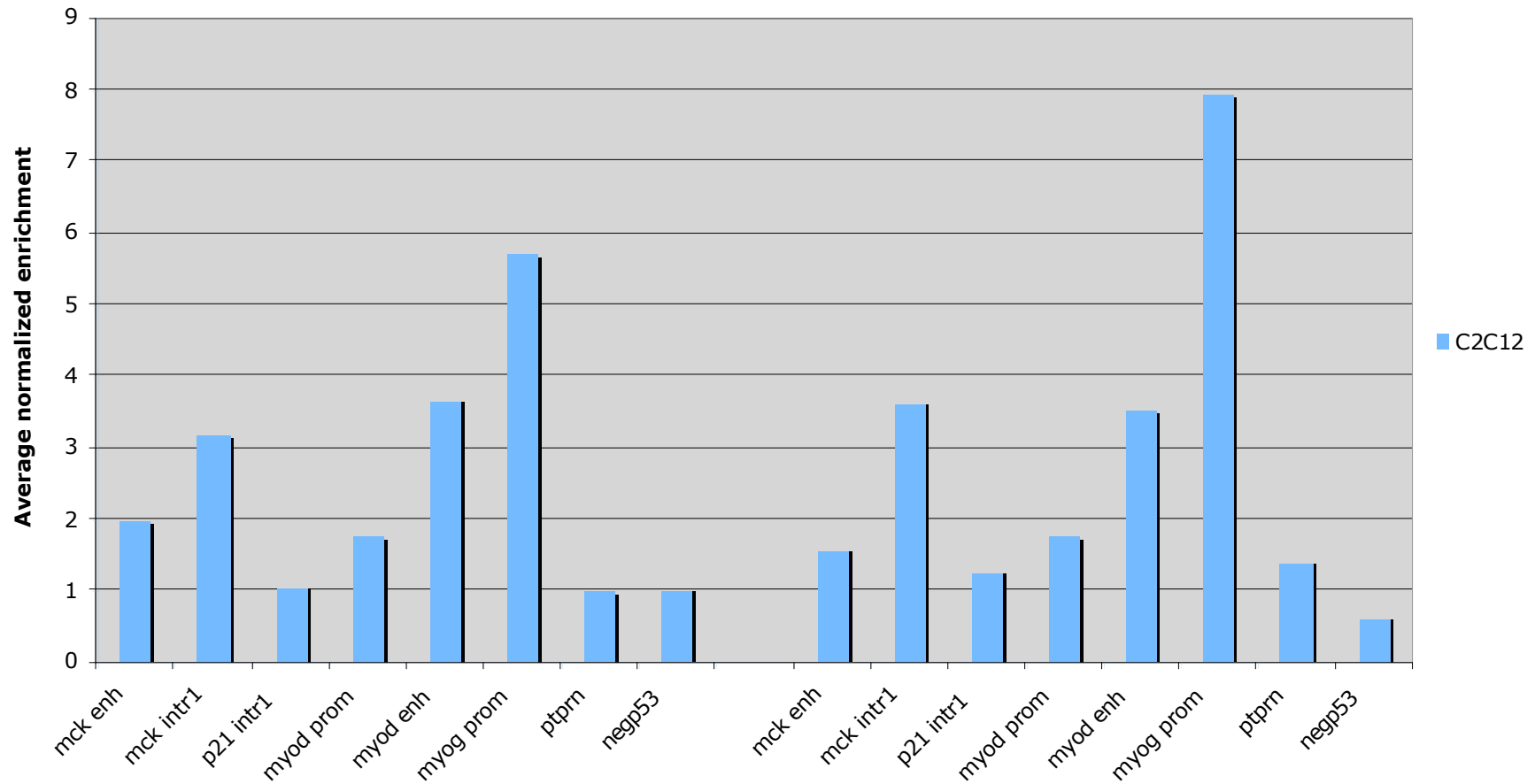
F5D ChIP (new primers)

F5D Santa Cruz ChIP 15 (3/23/2007)
100 hrs post differentiation C2C12s, 15 cycles of sonication, 100 ul beads,
1 ug antibody



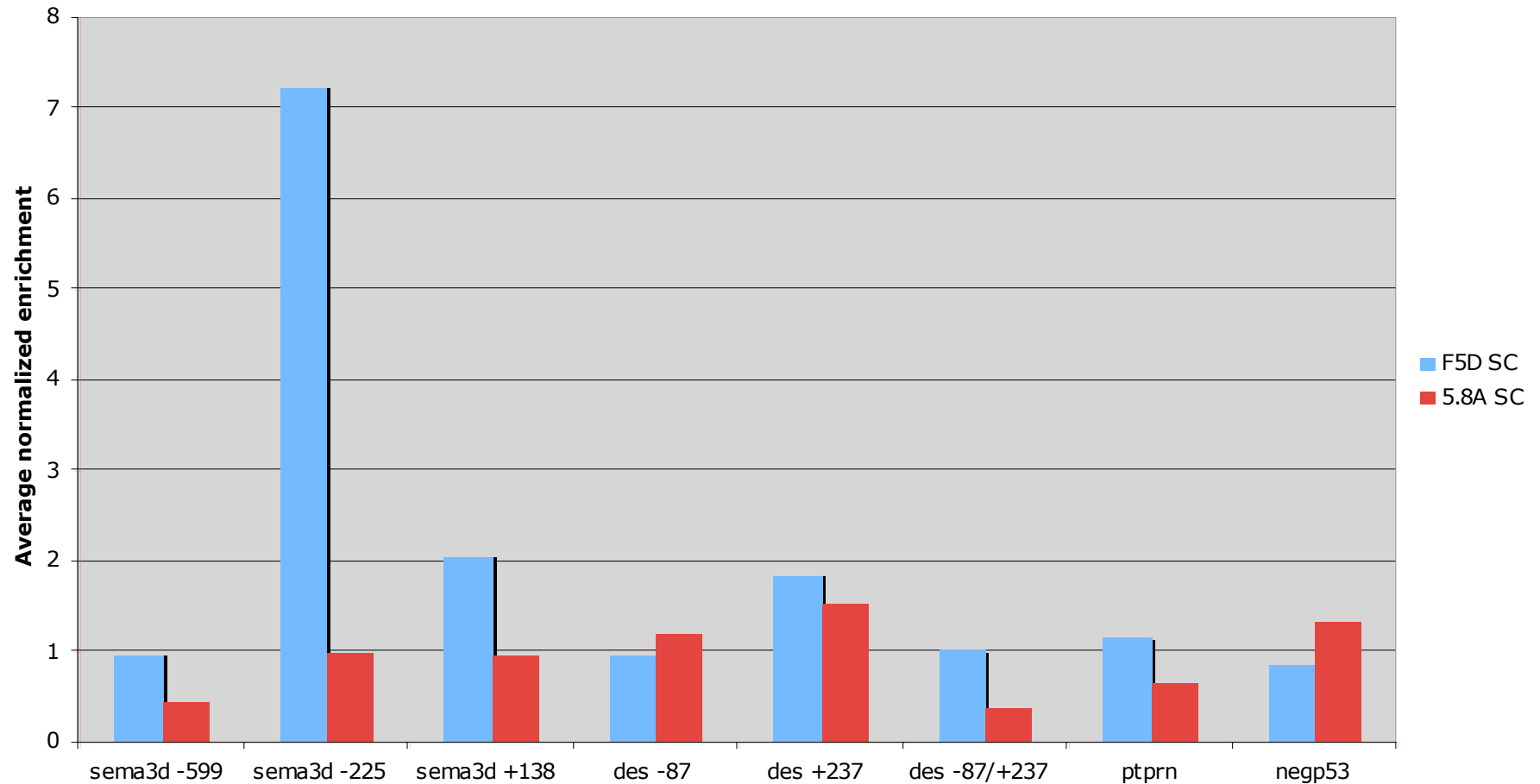
F5D ChIP (old primers)

F5D Santa Cruz ChIP 16 (3/23/2007)
100 hrs post differentiation C2C12s, 100 ul beads, 1 ug antibody, 15 cycles sonication,
Farnham + PI



F5D and 5.8A ChIP (new primers)

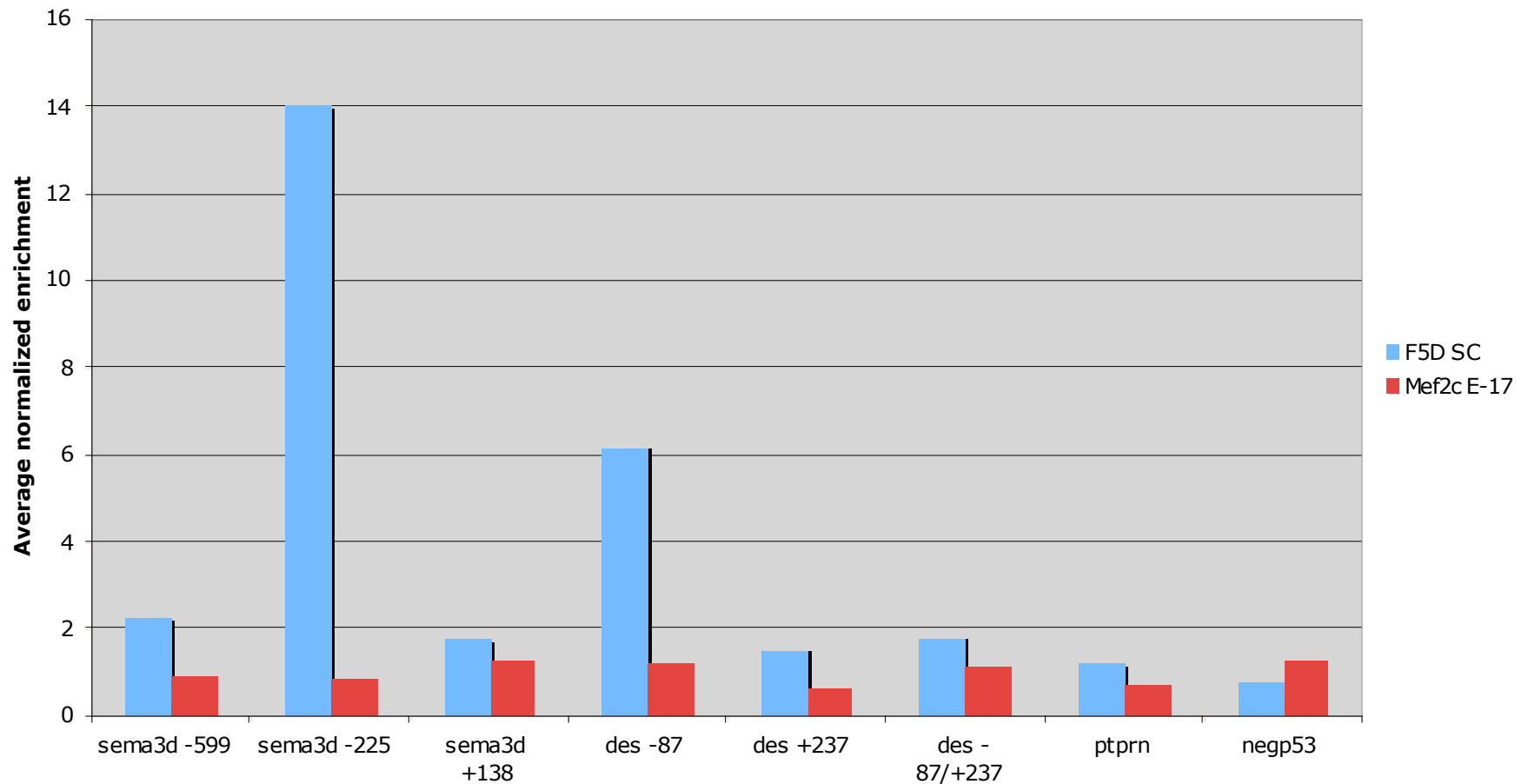
F5D SC and 5.8A ChIP 18 (4/23/2007)
63 hrs post differentiation C2C12s + ara-C, Farn + PI / 20 cycles of sonication, 100 ul
beads, 1 ug antibody



F5D and Mef2c - E17 ChIP

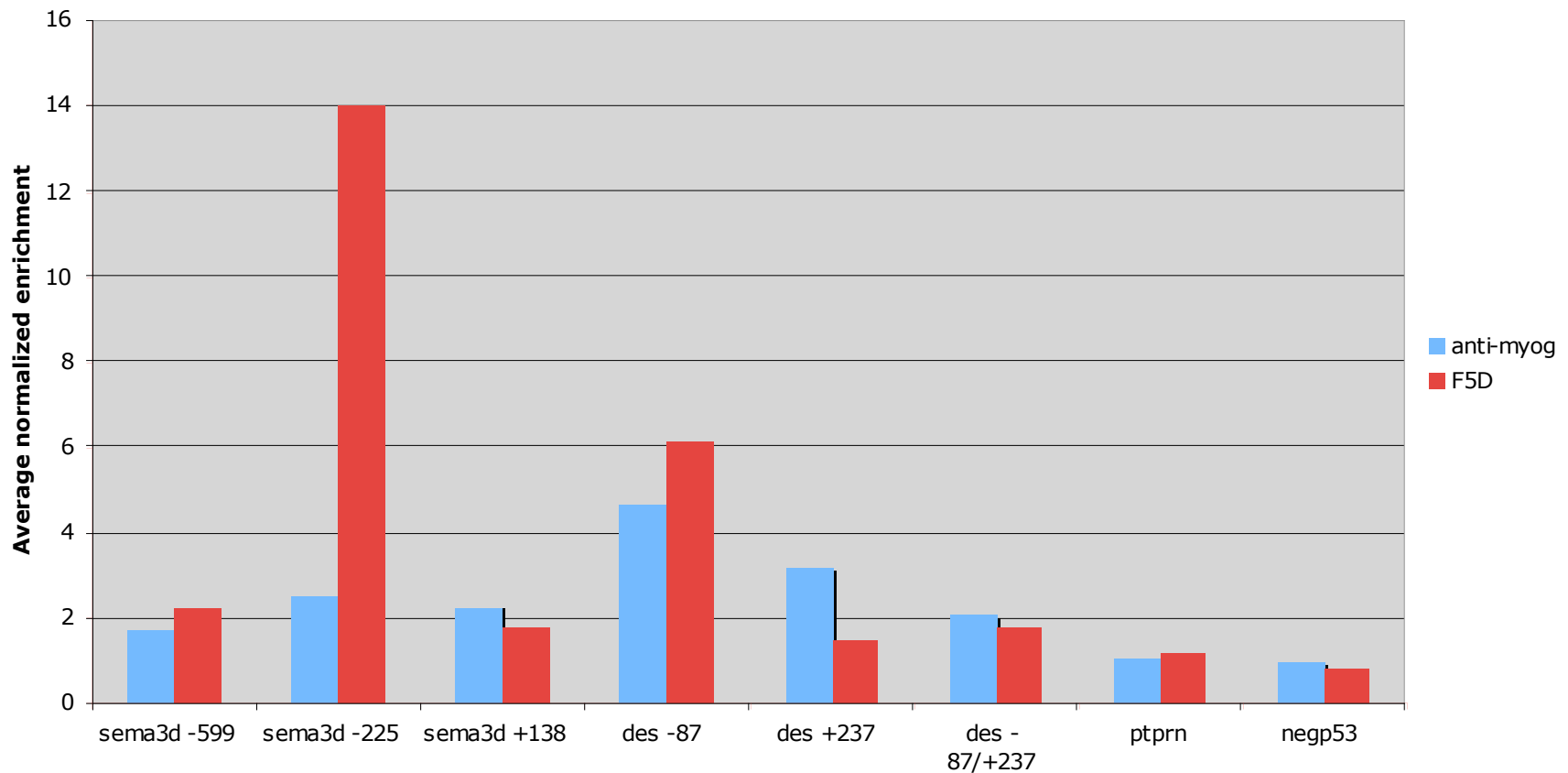
(new primers)

F5D Santa Cruz and Mef2c E-17 ChIP 19 (4/25/2007)
63 hrs post differentiation C2C12s + ara-C, Farn + PI / 20 cycles of sonication, 100 ul
beads, 1 ug antibody monoclonal / 4 ug polyclonal



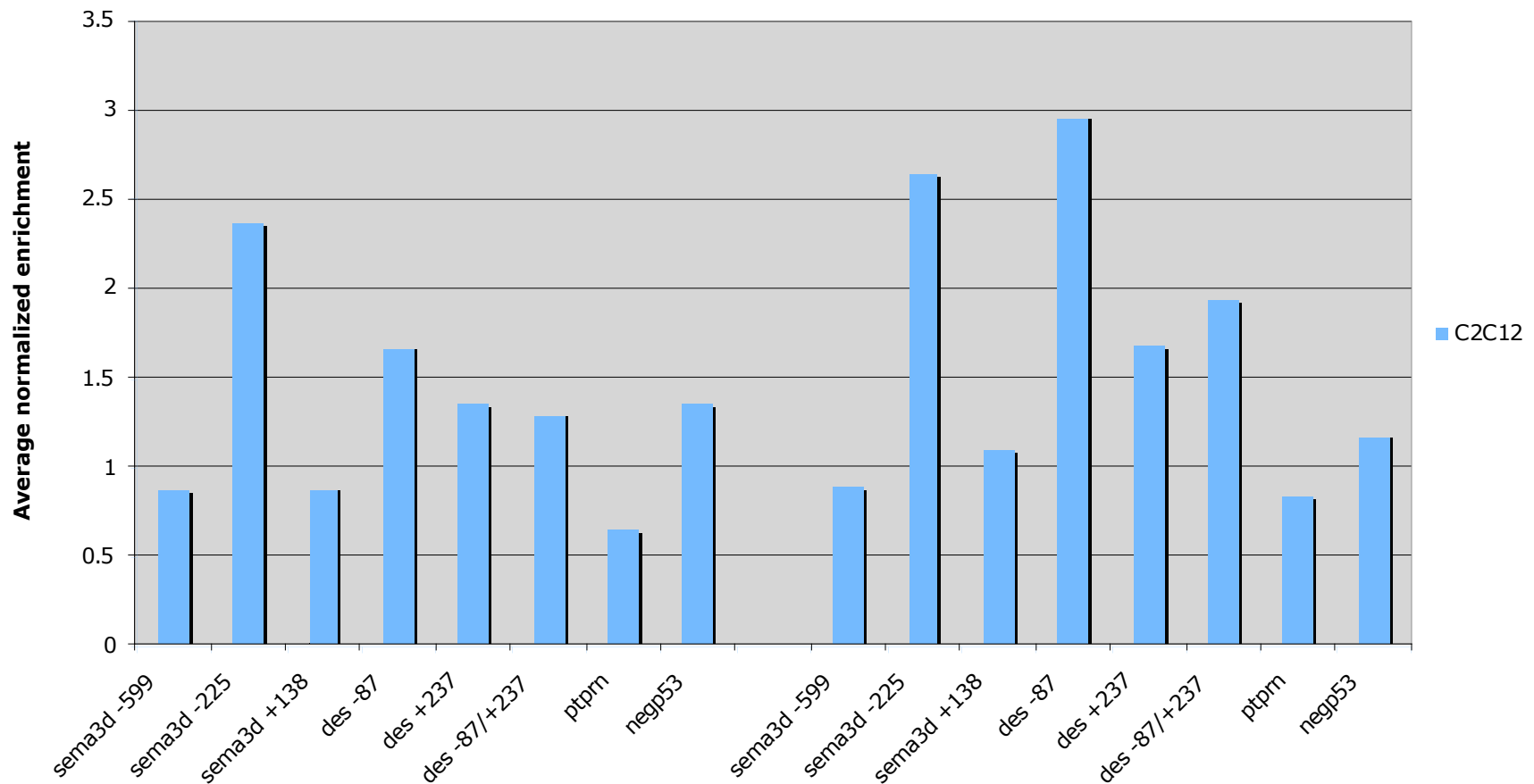
F5D SC and anti-myog polyclonal ChIP (new primers)

Polyclonal Rabbit anti-myog and F5D
63 hrs post differentiation C2C12s + ara-C, Farn + PI
15/20 cycles of sonication (respectively),
100 ul beads, 2 ug antibody polyclonal, 1 ug antibody monoclonal



F5D Home ChIP (new primers)

F5D "Home" ChIP 21 (8/5/2007)
24 and 61 hrs post differentiation C2C12s, no ara-C, Farn + PI,
12 cycles of sonication, 100 ul beads, 1 ug antibody



Cell Density/Growth

Cells (aka cell body) grow during differentiation → larger pellet at later timepoints

However, amount of DNA appears to increase as well

Will test the effect of ara-C on post-differentiation growth (results pending)

Cell Density/Growth

Density after 24 hrs diff: 197 ng/ μ l

Density after 61 hrs diff: 308 ng/ μ l

Extrapolated total chromatin:

- 24 hrs = 325 μ g
- 61 hrs = 508 μ g

Factors Affecting Final DNA Conc.

- Time
- Scraping efficiency
- Transfer efficiency during washes
- Phenol/chloroform extraction efficiency
- ara-C

Solexa

Myogenin Solexa run pending completion of:

- Test ChIPs
- Sonication tests
- Immunostaining

The End

RNA Timecourse Data (from Brian)

