

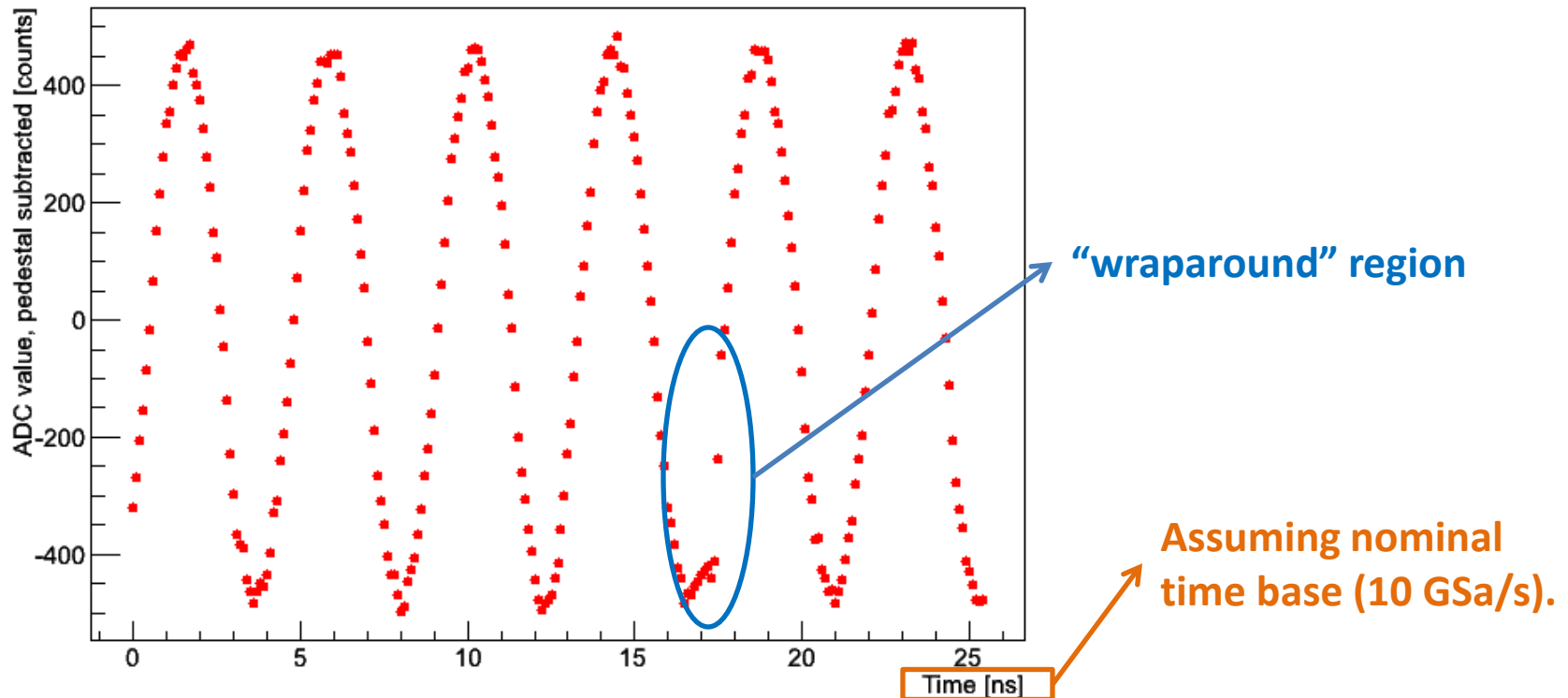
PSEC4 Timing Analysis

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Data Sample

- Data sent by Eric on September 29, 4 files:
 - All at ~ 10 GSa/s
 - CH3: 240 MHz sine wave input, 2000x2 events
 - CH4: 240 MHz sine wave input, 2000x2 events

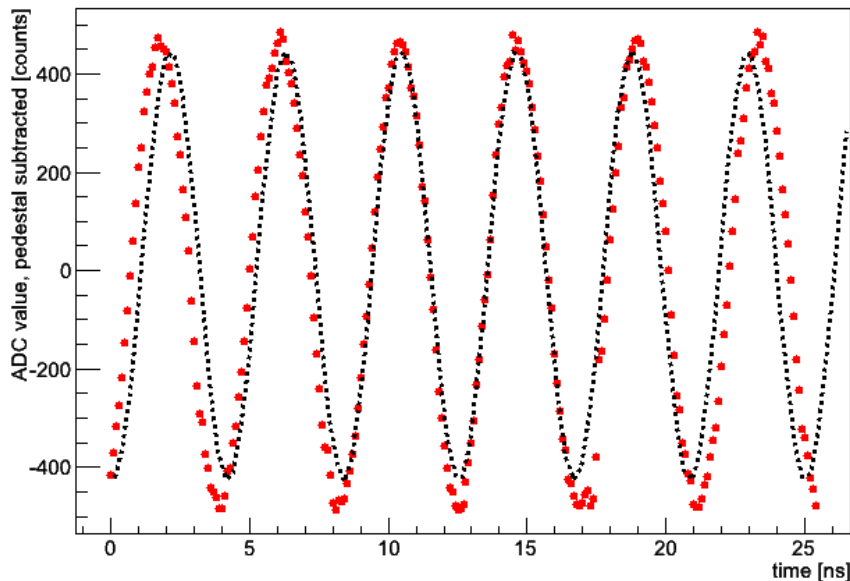
Example event:



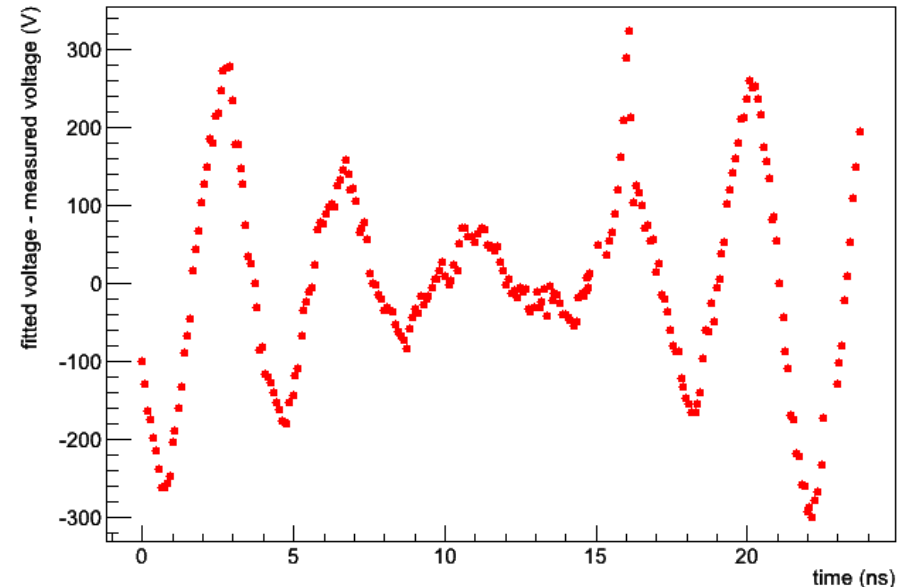
Raw Data, Fitted to Sine Wave

- Fit to a waveform with a sine wave:
 - Fixed: input frequency
 - Floated: DC offset, amplitude, phase

Example fit result:



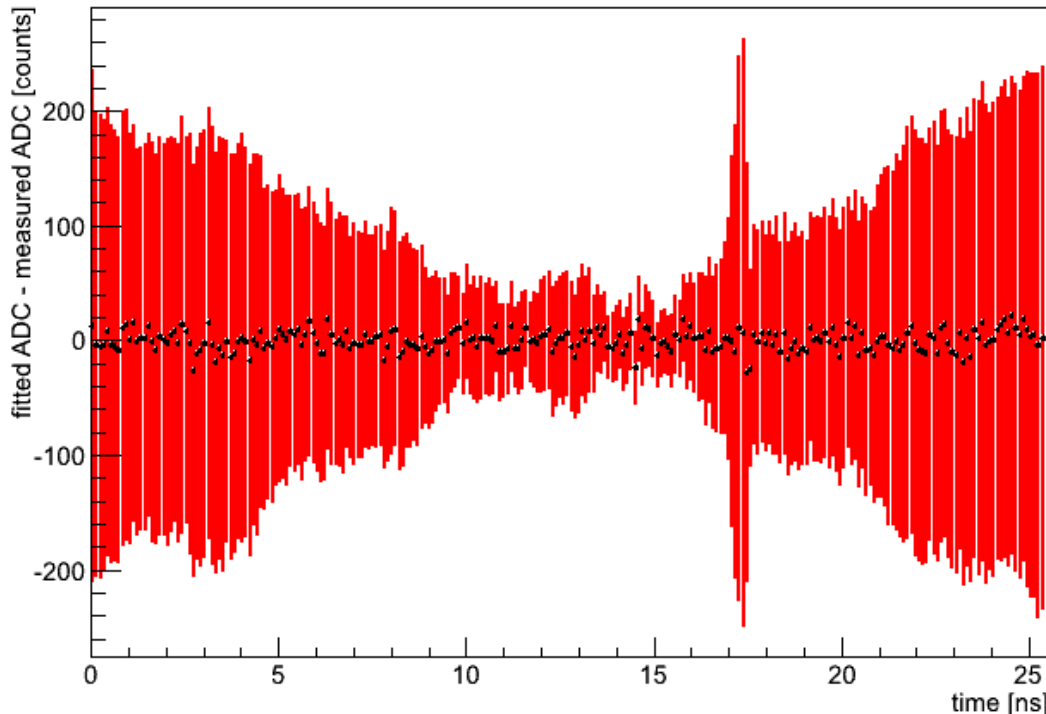
Example fit residuals:



Raw Data, Fitted to Sine Wave

- Fit to a waveform with a sine wave:
 - Fixed: input frequency
 - Floated: DC offset, amplitude, phase

Profile of fit residuals for 2000 events:



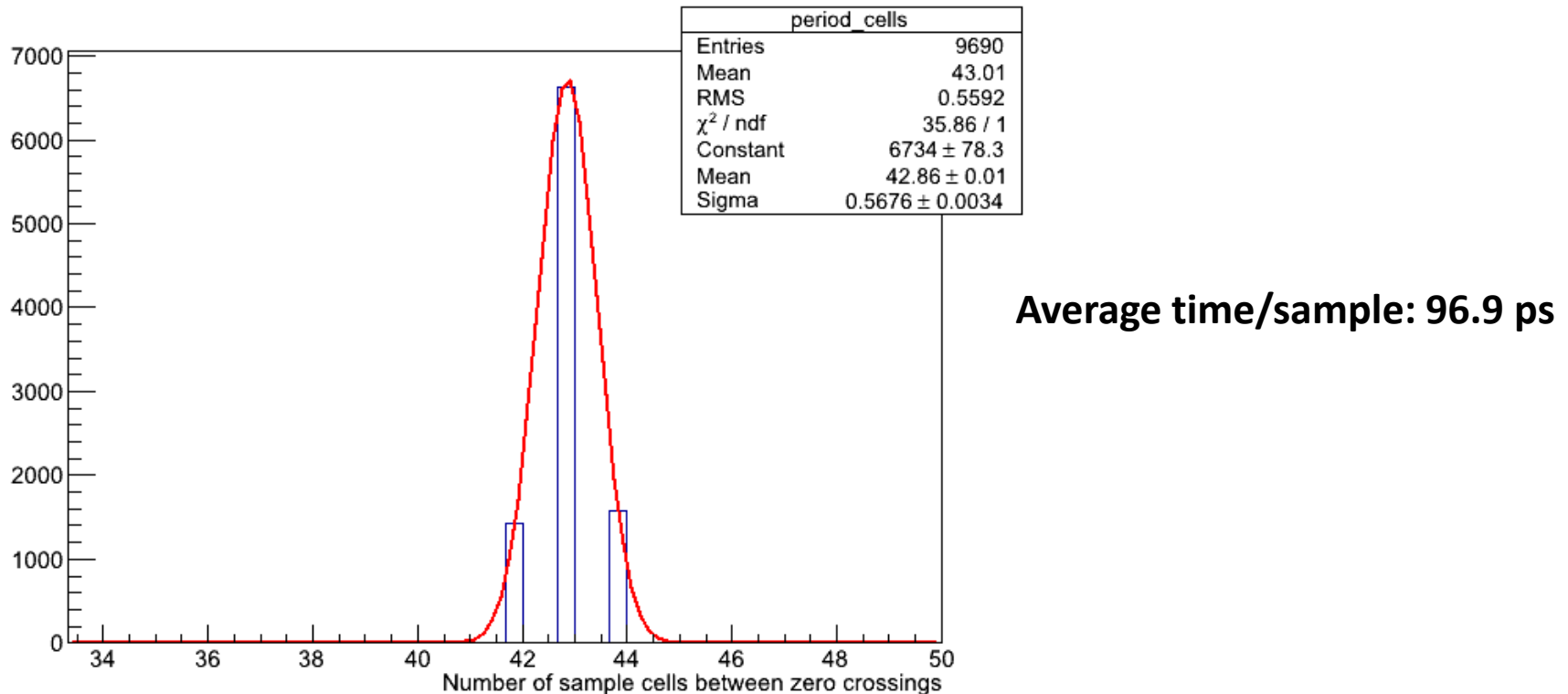
Black points – mean value of residuals over 2000 events
Red bars – RMS of residuals over 2000 events

Two observations:

1. Time base seems to require an overall correction.
2. Residuals near center of window seem better behaved.

Simple Time Base Calibration

- Count number of sample cells between zero crossings*. Calculate average time/sample.

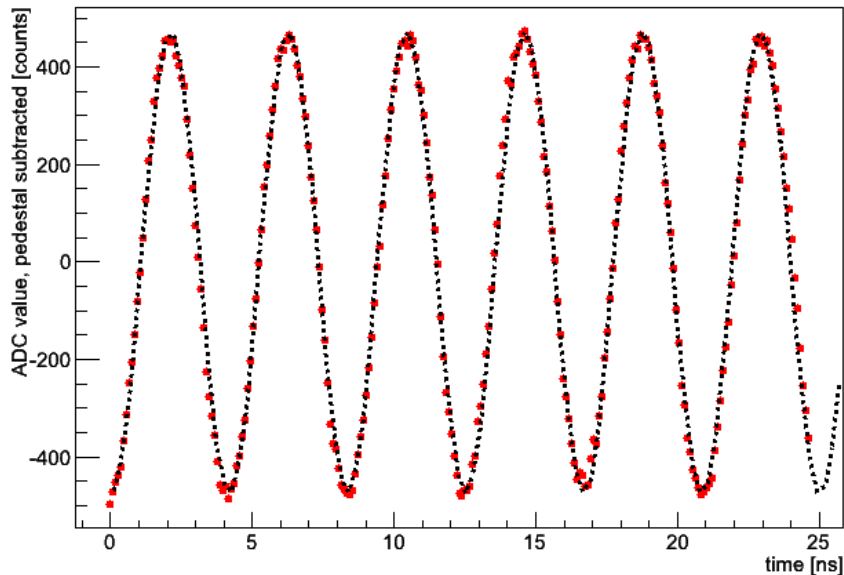


*Using samples 0-150 to avoid "wraparound" (average time/sample: 96.4 ps if all samples are used).

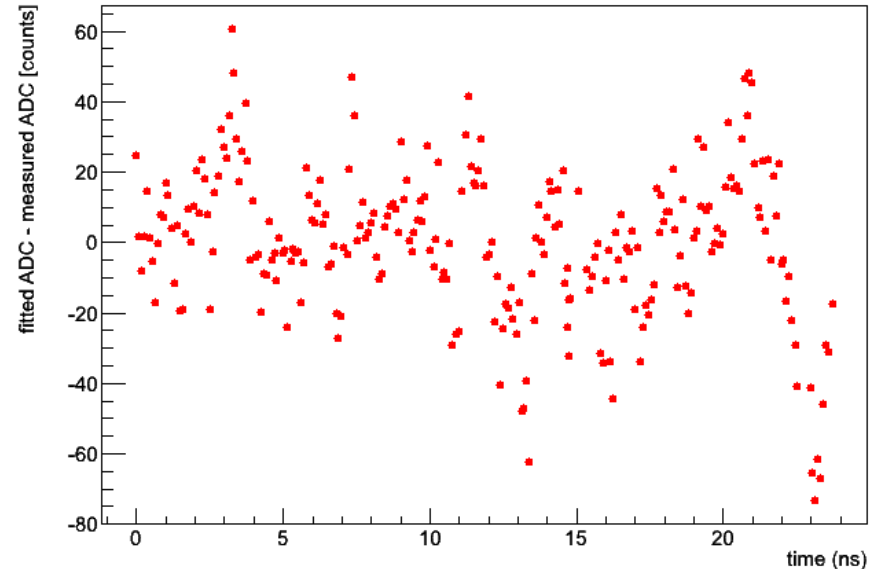
Sine Wave Fits w/ Corrected Time Base

- Refitted with new average Δt , no sample-to-sample corrections.

Example fit result:



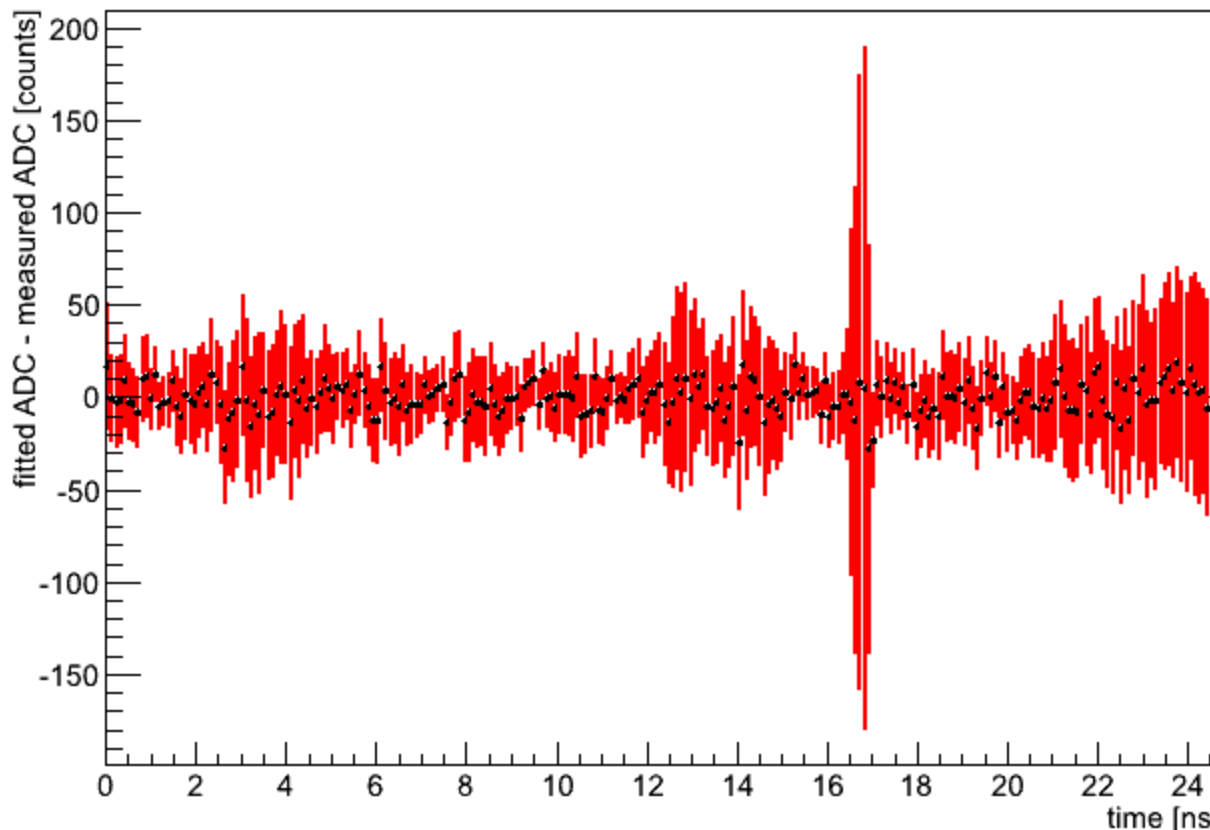
Example fit residuals:



Sine Wave w/ Corrected Time Base

- Refitted with new average Δt , no sample-to-sample corrections.

Profile of fit residuals for 2000 events:



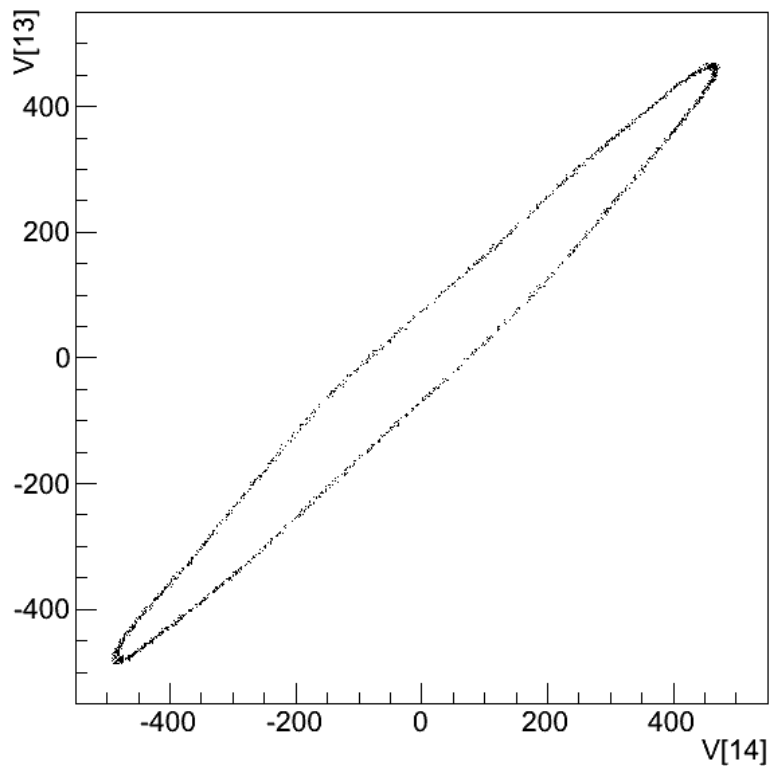
Black points – mean value of residuals over 2000 events
Red bars – RMS of residuals over 2000 events

Wraparound still fits poorly, as expected.
Some smaller scale structure persists as well.

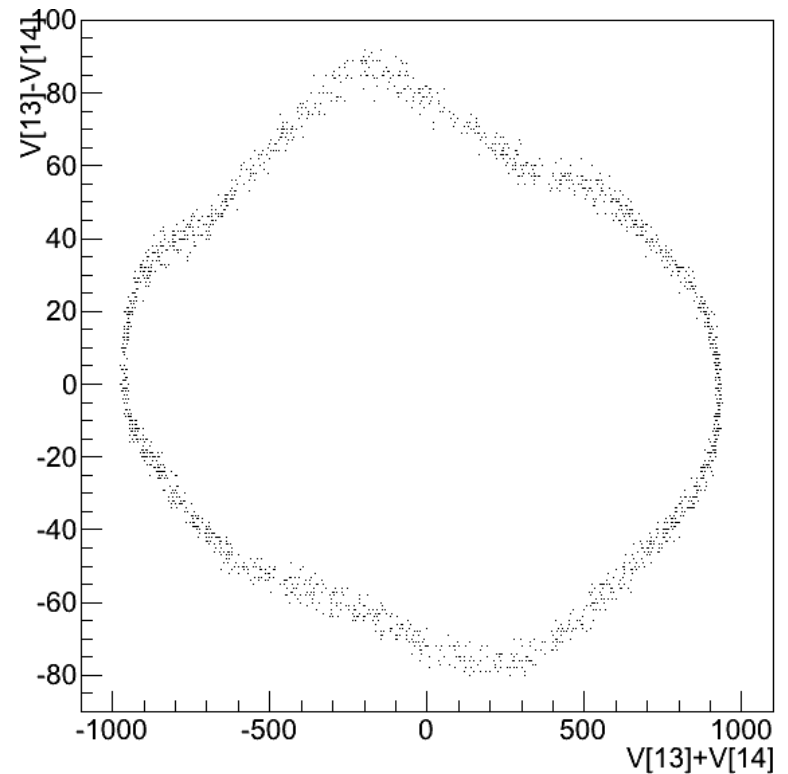
→ Overall significant improvement in fit quality!

What about ellipse method...?

- Ellipse fits w/ cells one-apart are complicated by strange contours:



Cell 13 vs. Cell 14

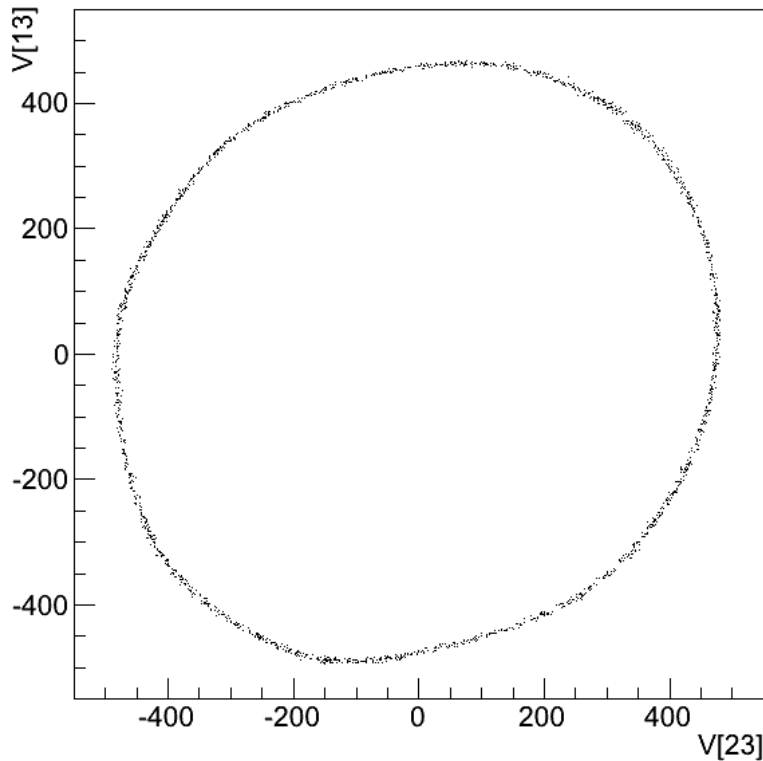


Cell 13, 14: difference vs. sum

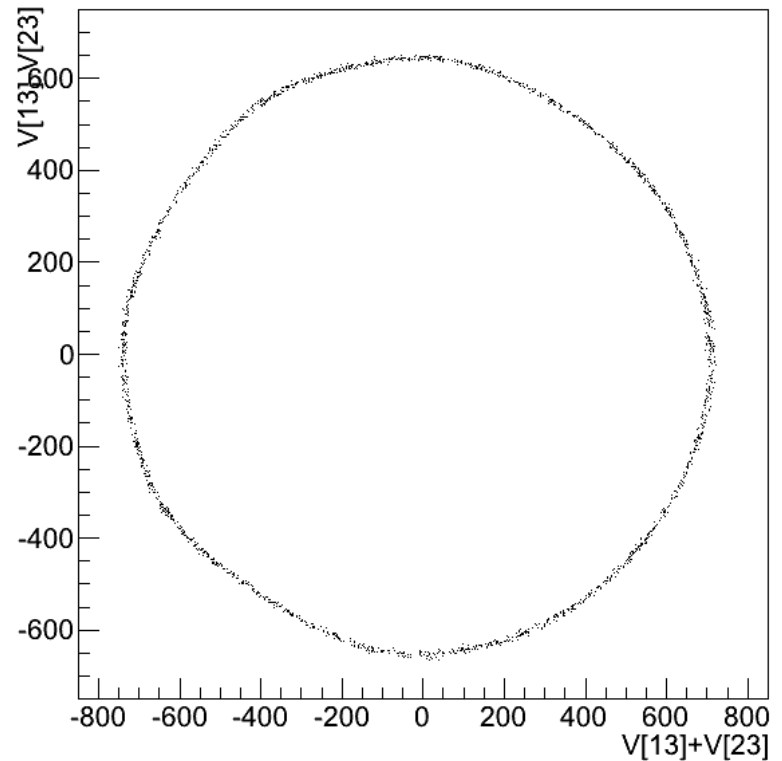
Is this nonlinearity? Correlated noise? Something else?
Can try applying nonlinearity correction, if available.

Ellipse Plots 9,10 Samples Apart

- Not as obviously misbehaved:



Cell 13 vs. Cell 23

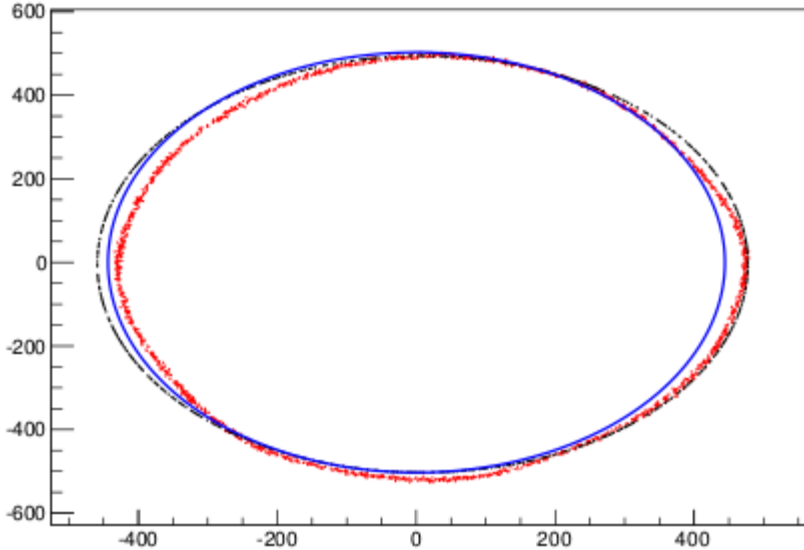


Cell 13, 23: difference vs. sum

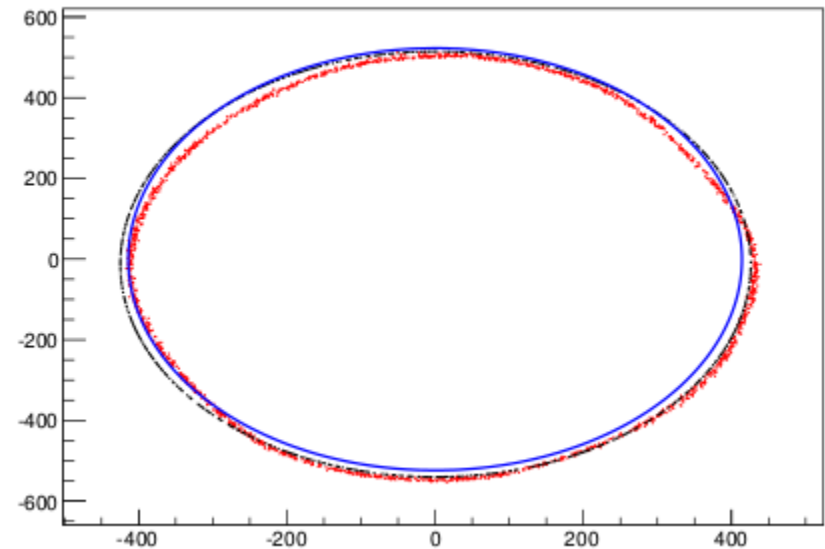
Fits to 9,10 Apart Contours

- Fits (assuming gain = 1 for all cells) are fairly well behaved (not too many fit failures), but they clearly miss some features of the data.

samples 35 and 45



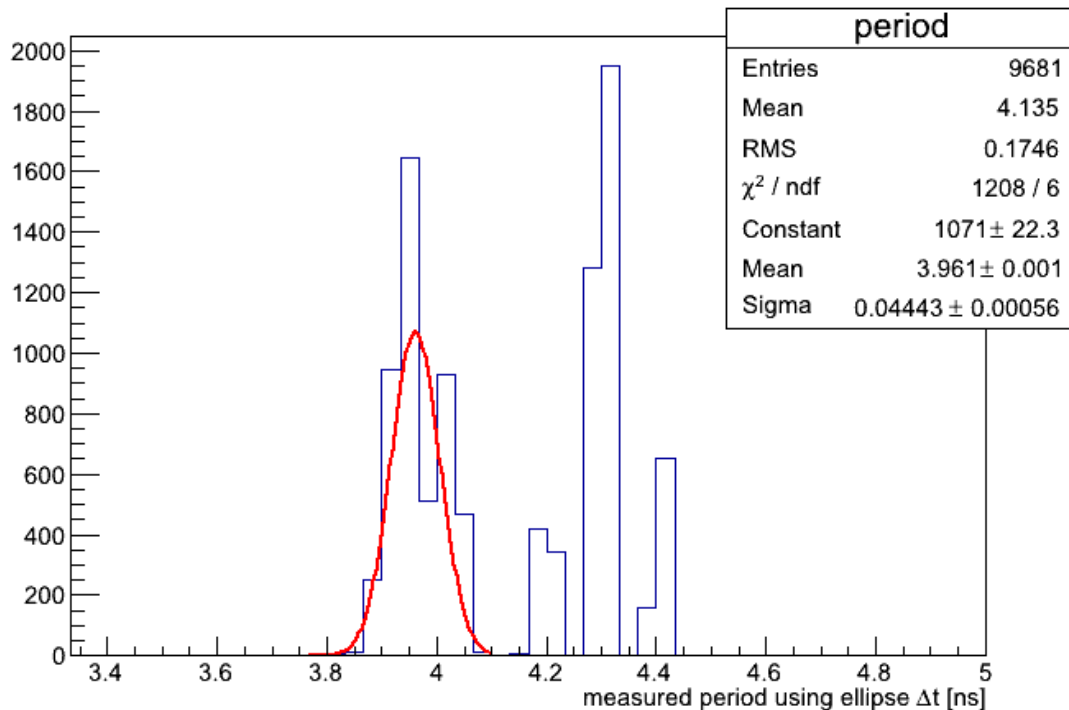
samples 36 and 45



Red – data points, **black** – nominal 10 GSa/s contour, **blue** – fitted contour

Cross Check on Ellipse Fits

- Measured period using Δt values from ellipse fits:



True T @ 240 MHz = 4.17 ns

Mean shows something near this value, but only due to some cancellation between different populations.

Summary & Plan

- Sampling rate appears to be slightly faster than 10 GSa/s.
 - Making this correction cleans up data quite a bit.
- For finer corrections, sample-to-sample Δt :
 - Can perform zero-crossing analysis, but may need more data. (Will try with what I have, first.)
 - Ellipse fitting will require better understanding of “kinks” in plots. As a first pass, I would like to try correcting out nonlinearity.