

Deconvolution

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Method

required for single-bunch time resolution (Bingefors *et al.* 1993). The underlying notion is to use rather slow pulse shaping and then to apply a deconvolution algorithm to reconstruct the fast components of the input signal. For the sampled output of a *CR-RC* shaper with the step response

$$v(t) = \frac{t}{\tau} e^{-t/\tau} \quad (8.15)$$

this can be implemented by forming the weighted sum of three successive samples

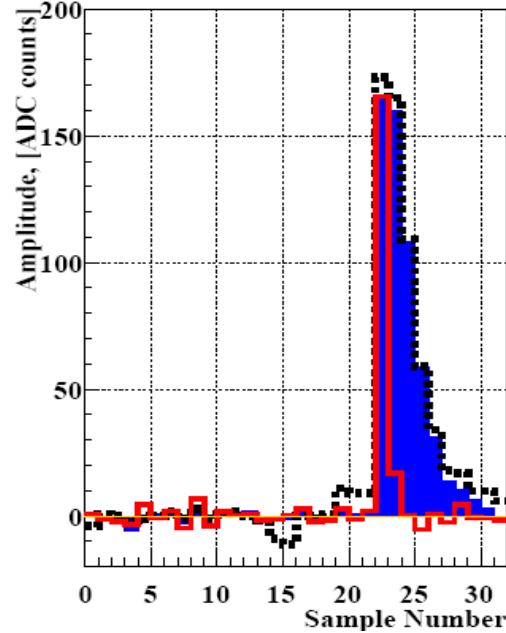
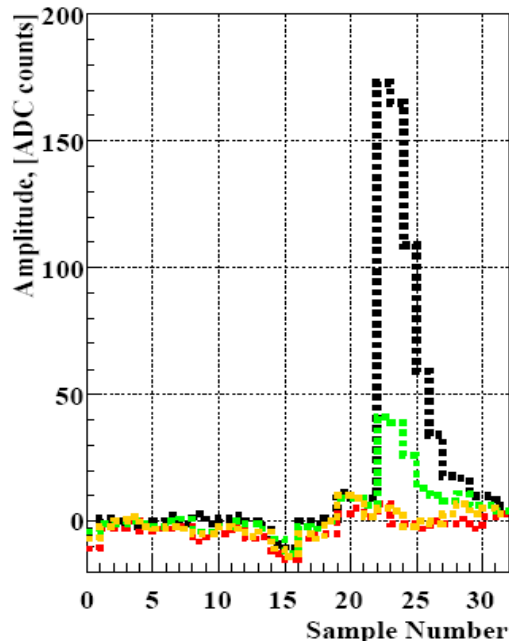
$$V_k = w_1 V_k + w_2 V_{k-1} + w_3 V_{k-2} \quad (8.16)$$

with the weights

$$w_1 = \frac{1}{x} e^{x-1}, \quad w_2 = -\frac{2}{x} e^{-1}, \quad w_3 = \frac{1}{x} e^{-(x+1)} \quad (8.17)$$

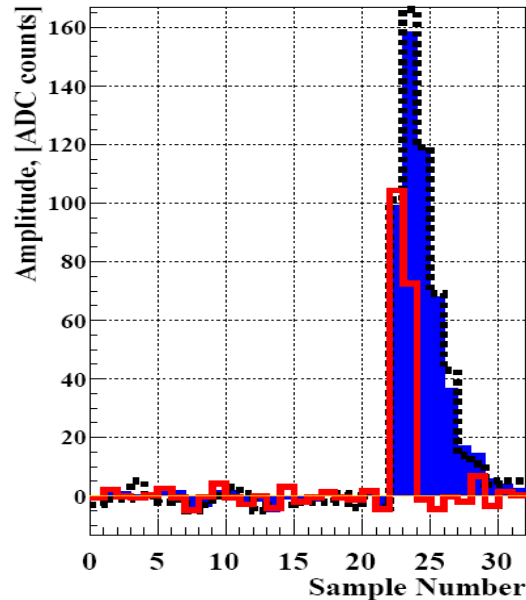
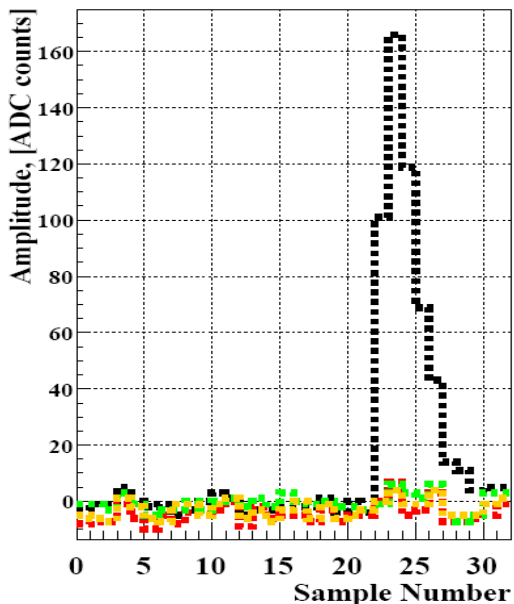
(Gadomski 1992, Bingefors *et al.* 1993). The weights depend on the sampling interval normalized to the shaping time constant $x = \Delta t / \tau$. For a step input the result of the deconvolution is zero. However, for a finite rise time the result is a short pulse with the duration of the rise time. The APV25 uses a time constant $\tau = 50$ ns in the *CR-RC* filter and samples the output at 40 MHz, so $x = 0.5$ and the weighting factors

Example



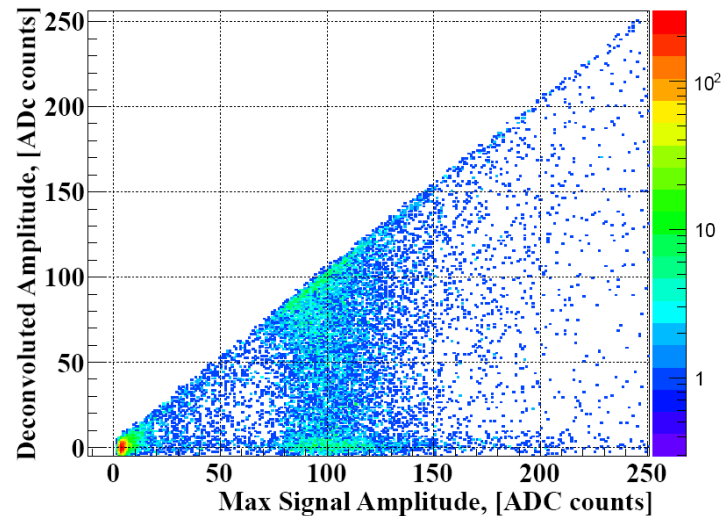
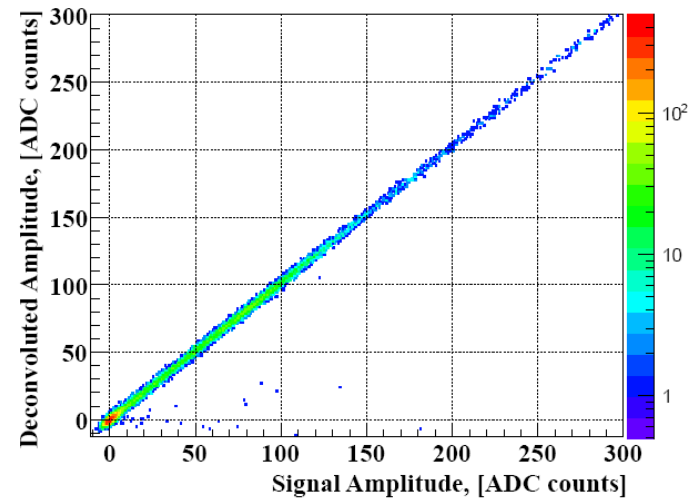
Signal Processing:

1. CMN calculation from all non hit channels of the same chip
2. CMN subtracted (blue histogram)
3. Deconvolution method applied (red histogram)
4. Different signals examples are shown.
5. Deconvolution method repeats first signal bin.
6. In case of synchronized data should work better



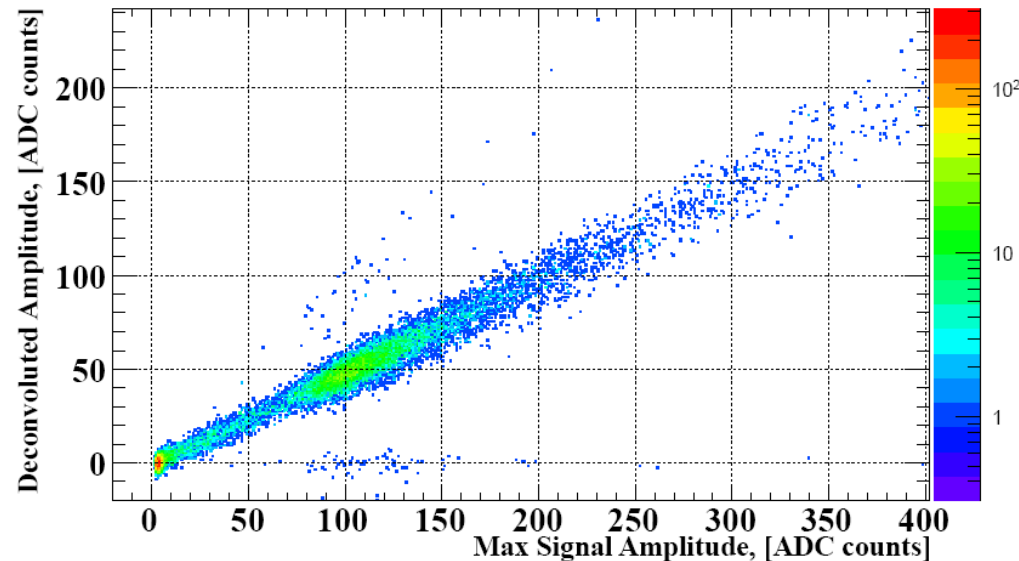
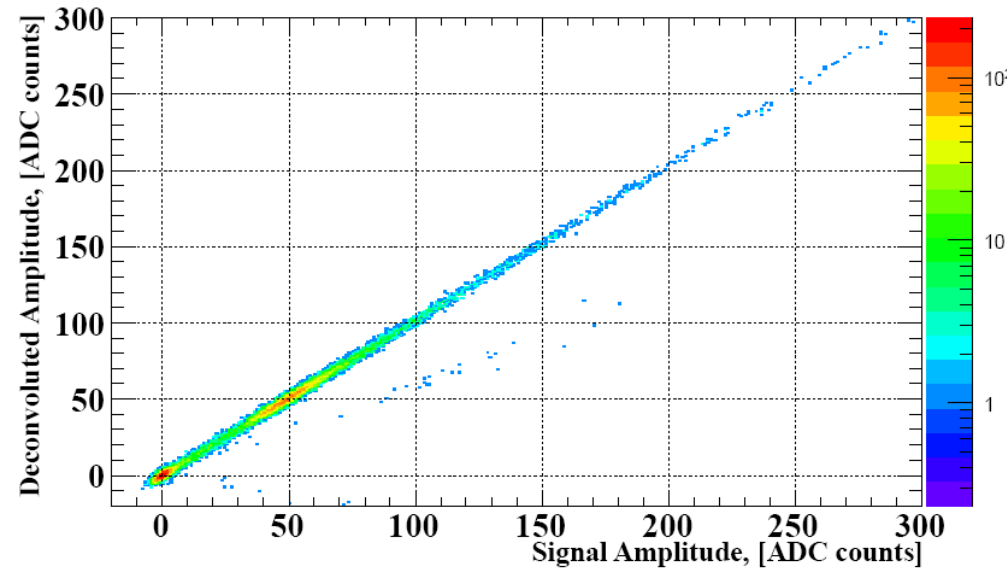
Correlations

- Bin with maximum deconvoluted value vs the same bin signal amplitude
 - Full proportionality
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- Bin with maximum deconvoluted value vs the maximal signal amplitude
 - Method depends on the first bin with the signal



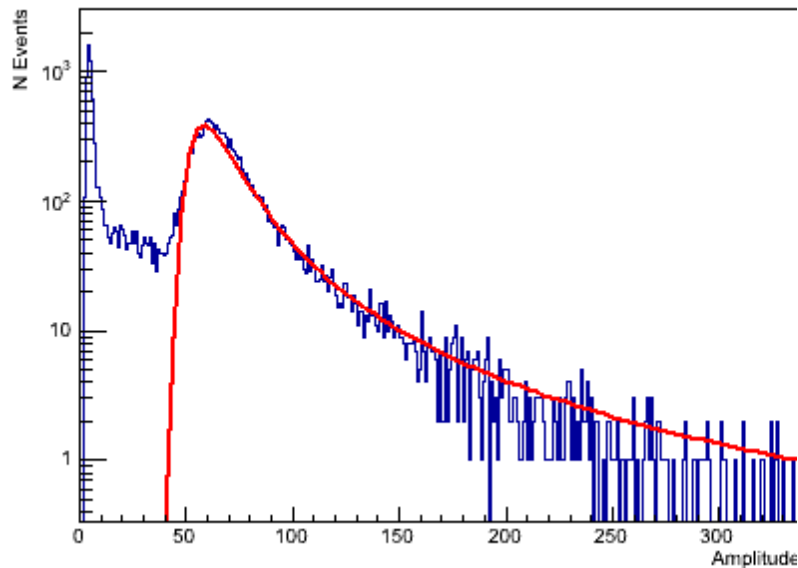
Synchronized data

- Bin with maximum deconvoluted value vs the same bin signal amplitude
- Full proportionality
- Bin with maximum deconvoluted value vs the maximal signal amplitude
- Method works better on synchronized data

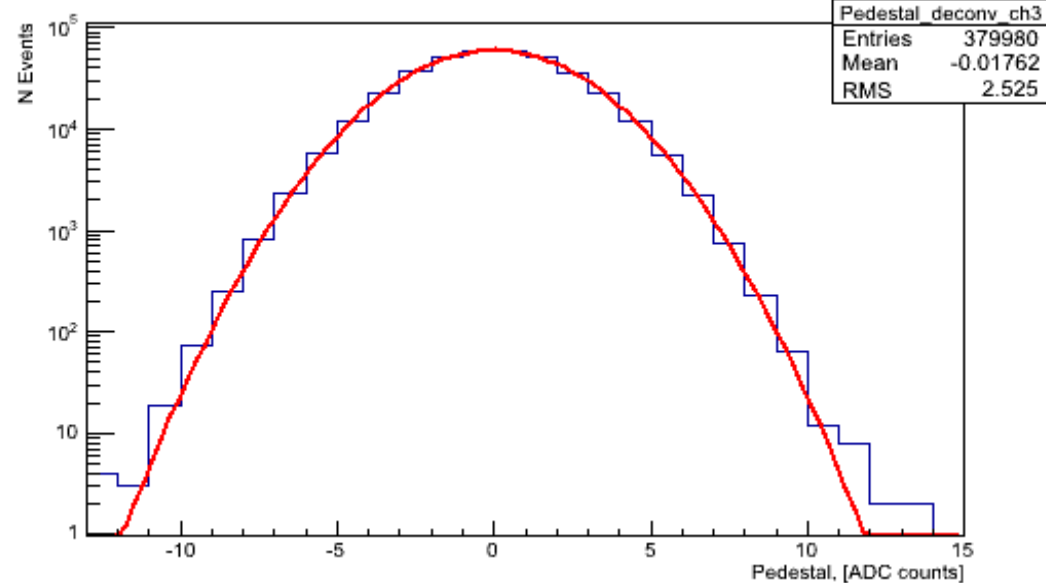


Conclusions

Amplitude values for each sample for ch3



Pedestals values for each sample for ch3



- Deconvolution Amplitude spectrum is distributed over Landau Gauss convolution
- Pedestal is calculated in the window before the signal coming (1-16 samples)
- $S/N = 59.6/2.5 = 24$

