# Fitting Polynomial Baselines to Complete Spectra

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### Syntax & parameters

fit.to: hyperSpec object with the spectra whose baselines are to be fitted.

apply.to: hyperSpec object giving the spectral range, on which the baselines should be evaluated.

If apply is NULL, a hyperSpec object with the polynomial coefficients is returned instead

of evaluated baselines.

poly.order: polynomial order of the baselines

npts.min: minimal number of data points per spectrum to be used for the fit.

npts.min defaults to the smaller of 3 times (poly.order + 1) or  $\frac{1}{20th}$  of the number of

data points per spectrum.

If  $npts.min \leq poly.order$ , a warning is issued and  $npts.min \leftarrow poly.order + 1$  is

ısed.

noise: a vector giving the amount of noise, see below.

short, user, date: are handed to logentry

# Specifying the spectral range

The polynomials are always fit to fit.to, but are evaluated on the the wavelengths of apply.to. Therefore it is possible to exclude spectral regions that do not contribute to the baseline from the fitting, while the baseline is used for the whole spectrum. The choice of the spectral range in fit.to influences the resulting baselines to a certain extent, as becomes clear from figure 1. Narrowing the spectral range in fit.to speeds up the fitting of the polynomials:

## Fitting polynomials of different orders

Figure 2 shows the resulting baseline polynomial of spc.fit.poly.below (chondro [1], poly.order = order) with order = 0 to 3 for the first spectrum of the chondro data set.

### The mechanisms of fitting the baselines

Appropriate spectral regions with supporting points for the baseline polynomials are calculated iteratively:

- 1. A polynomial of the requested order is fit to the considered spectral range.
- 2. Only the parts of the spectrum that lie below this polynomial plus the noise are retained as supporting points for the next iteration.

These two steps are repeated until either

- no further points are excluded, or
- the next polynomial would have less than npts.min supporting points.

The baselines and respective supporting points for each iteration of spc.fit.poly.below (chondro [1], poly.order = 1) are shown in figure 3.

### The noise level

Besides defining a minimal number of supporting points, a "noise level" may be given. Consider a spectral range consisting only of noise. The upper part of figure 4 illustrates the problem. As the baseline fitting algorithm cannot distinguish between noise and real bands appearing above the fitted polynomial, the resulting baseline (black) is too low if the noise parameter is not given.

Setting the noise level to 10 (2 standard deviations), the fitting converges immediately with a much better result. The resulting baselines for spc.fit.poly.below (chondro [1], poly.order = 1, noise = 12) of the whole spectrum are shown in the middle and lower part of figure 4

**noise** may be a single value for all spectra, or a vector with the noise level for each of the spectra separately.

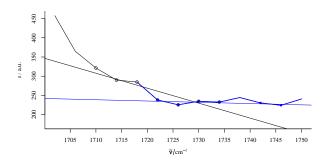


Figure 1: Influence of fit.to on the baseline polynomial. The black baseline is fit to the spectral range  $1700 - 1800 \,\mathrm{cm^{-1}}$ , the blue to  $1715 - 1800 \,\mathrm{cm^{-1}}$  only (dots & circles: supporting points).

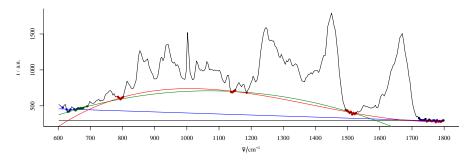


Figure 2: Baseline polynomial fit to the first spectrum of the chondro data set of order 0 (black), 1 (blue), 2 (green), and 3 (red). The dots indicate the points used for the fitting of the polynomial.

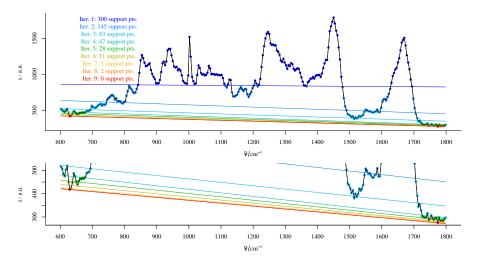


Figure 3: Iterative fitting of the baseline. The dots give the supporting points for the baselines in the same colour. The lower part is a magnification of the intensity axis.

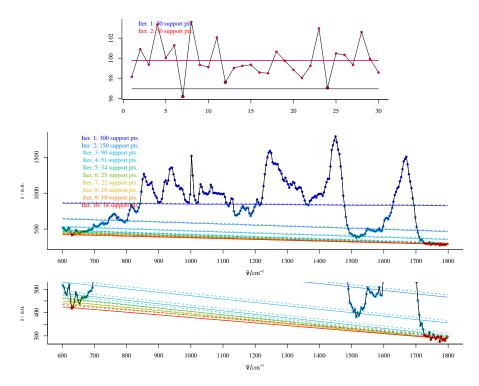


Figure 4: Iterative fitting of the baseline with noise level. Upper part: effects of the noise parameter on the baseline of a spectrum consisting only of noise and offset: without giving noise the resulting baseline (black) is clearly too low. A noise level of 10 results in the red baseline. The middle and lower part show the baseline fitting with noise level on the complete spectrum. Colour: iterations, dots/circles: supporting points for the respective baselines. Dashed: baseline plus noise. All points above this line are excluded from the next iteration.