

## Solution 2: Nonlinear regression (Lmfit package)

The second approach that could be used for the luminescence decay computing is nonlinear regression. The measured output signal  $y(t)$  comprises convolution of luminescence  $f(t)$  and excitation impulse  $g(t)$ :

$$y(t) = f(t) * g(t), \text{ or } y(t) = \int_0^t f(\tau) g(t-\tau) d\tau.$$

If we know the hypothesis model for  $f(t)$  with set of some parameters, the corresponding cost function may be built and its minimization procedure could be performed. In our particular case

$$f_{\text{model}}(t) = I_0 \exp(-t/t_0), \text{ with two parameters } I_0 \text{ and } t_0.$$

Therefore, the model for the output signal will be:

$$y_{\text{model}}(t) = \int_0^t I_0 \exp(-\tau/t_0) g(t-\tau) d\tau.$$

The objective is to derive parameters  $I_0$  and  $t_0$  in order to match  $y_{\text{model}}(t)$  to  $y(t)$ . This is typical nonlinear regression problem. The benefit of this approach is obvious: avoiding noise and regularization like in Fourier transform case. Since no signal processing is performed, a considerable gain in accuracy is achieved.

**Lmfit** is a special extension package for Python – a high-level interface to non-linear optimization and curve fitting problems. It allows computing our problem by minimizing residual (i.e. data-model) array  $\{y(t) - y_{\text{model}}(t)\}$ .

More information could be found in [lmfit github repository](#).

Unlike [FFT/inverse FFT method](#), where full profile of the  $f(t)$  curve is computed, regression works with exponential model  $f_{\text{model}}(t)$  only (Fig. 1).

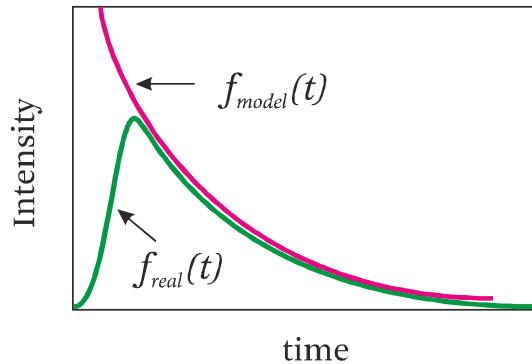


Fig. 1 – The real and modeled luminescence decay curves

This causes some inconveniences with time offsets. Simplified exponential model represents only decay and not the full profile of  $f(t)$ . Unless the entire curve is obtained, the input data must be preprocessed.

Time-wise, all the signals are biased. Attack times of  $y(t)$  and impulse  $g(t)$  are different and they are both measured at different times. It could be irrelevant for the conditions, mentioned above. However, due to a simplified model  $f_{\text{model}}(t)$ , one have to match all the signals to the time scale so that they all have the same starting point at zero seconds (Fig. 2).

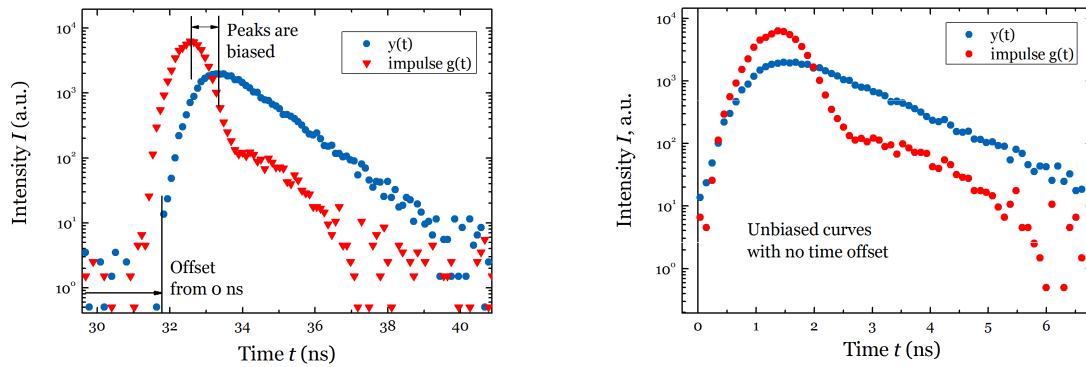


Fig. 2 – Measured (left) and preprocessed (right) input data

Another common problem in nonlinear regression is estimating errors. Asymptotic Standard Errors (ASE) *should not be used for nonlinear models*, since they underestimate the magnitude of the parameter uncertainties. The ASE are based on the “information matrix” and they ignore the off-diagonal elements. Probably the best alternative is using the [Bootstrap approach](#). It provides multiple curve-fitting, seeding the data with randomly sampled residuals  $\{y(t) - y_{\text{model}}(t)\}$ . Histograms of the parameters are obtained as the result.

Finally, mathematical model  $f_{\text{model}}(t)$  (or any arbitrary one) could be verified by performing the [Runs Test](#). It does not validate the model itself, it simply says if the data and/or assumptions about the data are not consistent with the model.

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The input data file “*preprocessed\_BaF2\_78nm.dat*” contains of 3 columns: time  $t$ , output  $y(t)$  and excitation impulse  $g(t)$ . The script file performs nonlinear regression on the arbitrary file as **Nonlinear\_regression\_BaF2.py** <file name>, including Bootstrap, Runs Test, plotting curves, residuals, histograms of parameters  $I_0$  and  $t_0$ , as well as printing the results to stdout.