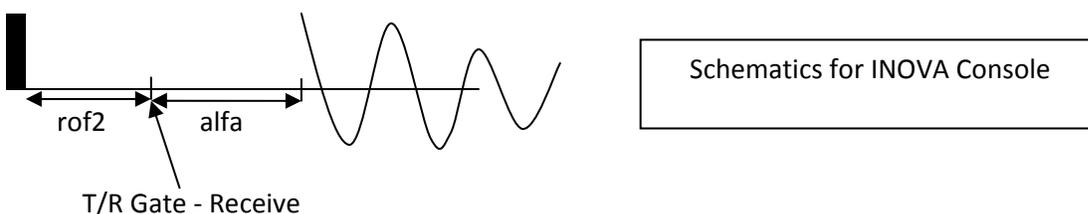


Getting Flat Baseline

A good flat baseline is an essential feature in a NMR spectrum for several reasons. A flat baseline ensures integrity of integrals, reduces t_1 noise in 2D spectrum and in indirect dimensions of nD spectrum. Baseline wobble in a nD dataset will lead to bad quality data.

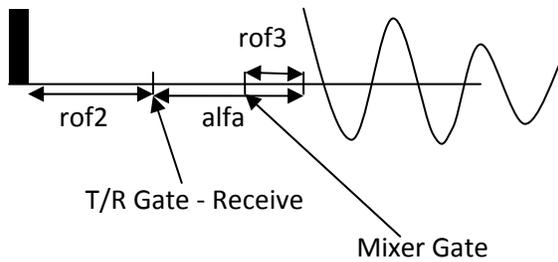
The baseline of a NMR spectrum depends on several parameters and the experiment. The protocol for controlling the baseline also depends on the vintage of the console. The steps involved for INOVA and VNMRS consoles are different.

There are few parameters that need to be set properly in order to get flat baseline: they are rof2 and alfa - common to INOVA and VNMRS, rof3 and ddrtc - only for VNMRS. The schematics below help understand these delays.



The first point of the FID is physically observed after a delay of $\text{rof2} + \text{alfa}$ after the pulse (all timing in μs) while ideally the first point should have been observed at the middle of the pulse. This in turn leads to linear phase difference for various resonances that is reflected as a non-zero l_p value when phasing the NMR spectrum. Larger the l_p value more the distortion of the resulting baseline. If we minimize the rof2 and alfa delays so that there is no large l_p then the rf breakthrough will distort the first few points of FID and will distort the NMR spectrum.

With INOVA spectrometers, a $\text{rof2} \sim 4$ to $5 \mu\text{s}$ and $\text{alfa} \sim 5$ to $6 \mu\text{s}$ will usually give zero l_p and flat baseline if $\text{dsp}='r'$ or $\text{dsp}='i'$. There are also macros crof2 or calfa that can be used after phasing a spectrum if there are large l_p values to get an estimate of rof2 and l_p values. Sometimes these macros will give a negative alfa and too short a rof2 value, so as a rule of thumb the above said values could be used. For cold probes - use $\text{dsp}='i'$ and $\text{qcomp}='y'$ along with the above said values. The dsp algorithm takes into account the total delay and constructs the required points of the FID to make it start from time $t=0$.



Schematics for VNMRS Console

With VNMRS console, there is a separate parameter `ddrtc` (μs) that controls the timing. This parameter doesn't exist by default, create it using the `create('ddrtc','pulse')` command. Usually

$\text{ddrtc} = \text{rof2} + \text{alfa} + \text{pw}/2$ (where `pw` is the pulse width) and $\text{rof2} \sim 28 \mu\text{s}$ and $\text{alfa} \sim 5$ to $6 \mu\text{s}$

The above expression is true only if the sequence is pulse acquire type. If there is a spin-echo type segment ($\tau - 180 - \tau$) just before acquisition then it is enough to set `ddrtc=alfa`. This, of course, presumes there is no `rof2` delay that is uncompensated in the echo segment. The parameter `dsp` has no effect in VNMRS console and can be set to anything ('r','n','i'). The default digital signal processing is done in three stages and there are quite many parameters control them. It is best to leave it in the default mode and worry about it only if extremely large spectral widths are to be acquired. The relatively large value of `rof2` is required for good baseline based on the dsp processing done by VNMRS and is used for both room temperature and cold probes (this value is large enough for cold probes also). The `rof3` value is by default $2 \mu\text{s}$ and is part of `alfa`. Thus `alfa` should be $\geq 2 \mu\text{s}$. Do not use the macros `crof2` or `calfa` with VNMRS console.