

Line Broadening and Negative Line Broadening

Some hints and tips

Maths time!

- NMR signal is (roughly)

$$S(t) = \exp(i\omega t) \cdot \exp(-t/T_2) \cdot \exp(-t/T_1)$$

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- An oscillation

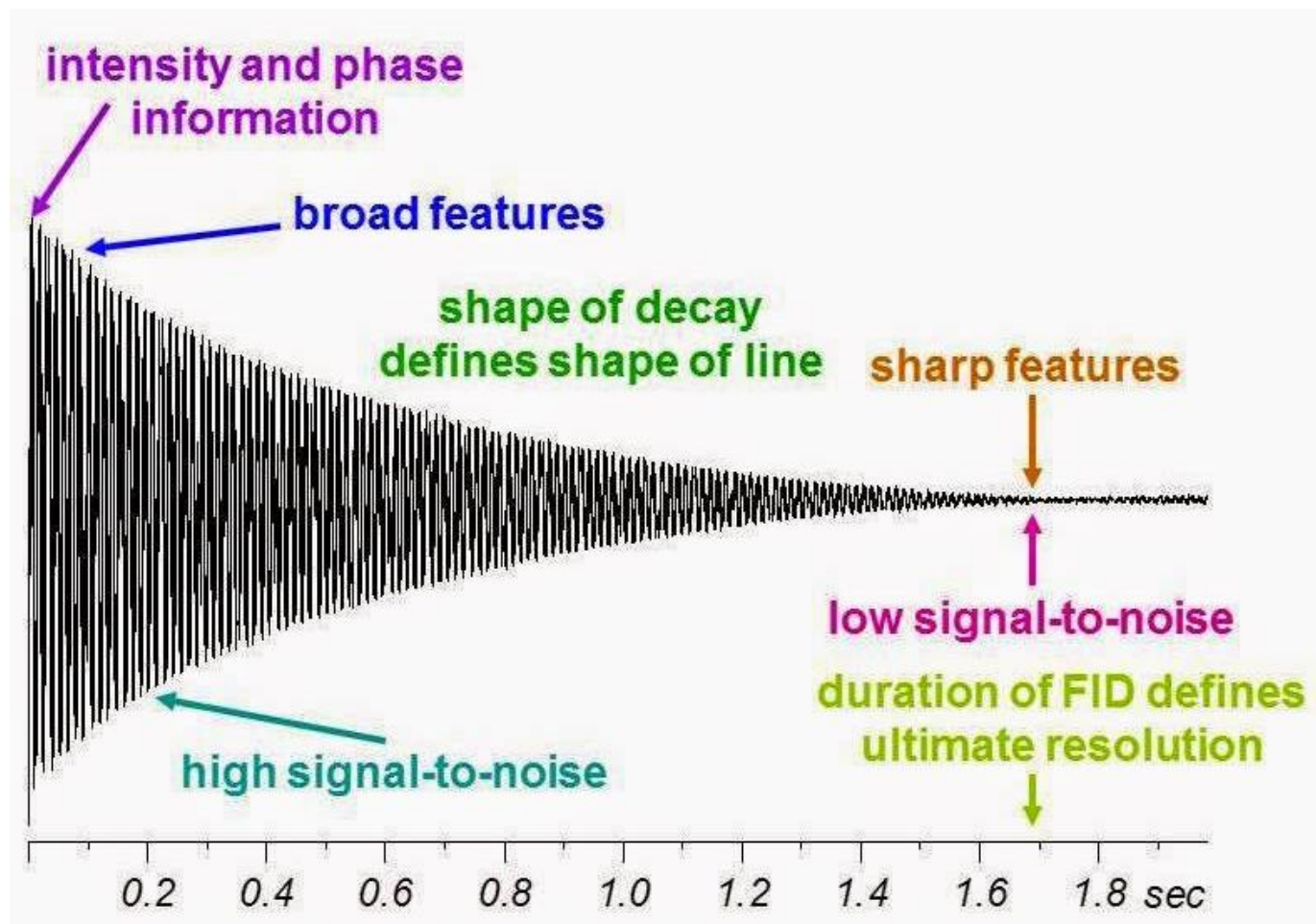
Maths time!

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$$S(t) = \exp(i\omega t) \cdot \exp(-t/T_2) \cdot \exp(-t/T_1)$$

- An oscillation
- And a decay
 - In liquids $T_1 \approx T_2$
 - In solids $T_1 \gg T_2$

Anatomy of a FID



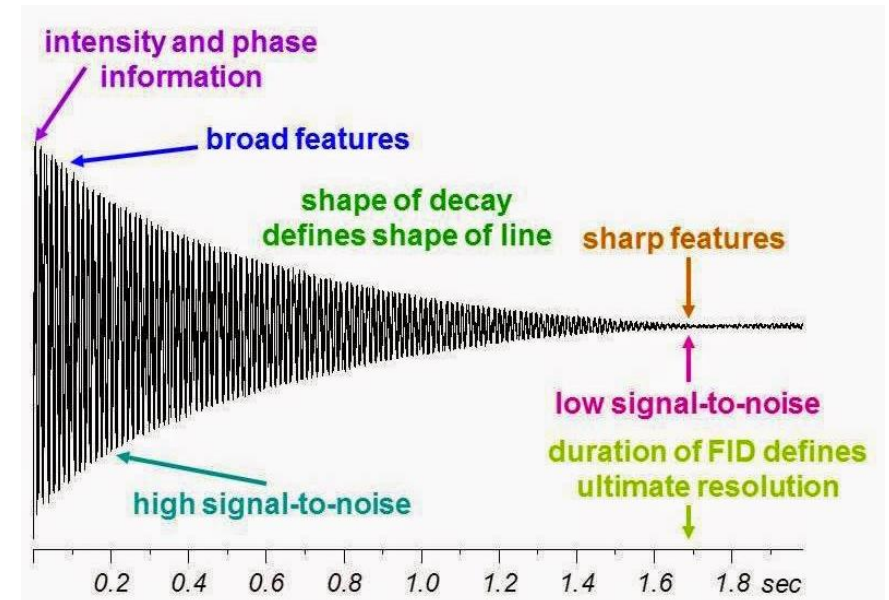
Shamelessly stolen from Glenn Facey's blog

Line broadening

- When we use line broadening ($LB = x$, WDW = EM, efp) we introduce another decay

$$S(t) = \exp(i\omega t) \cdot \exp(-t/T) \cdot \exp(-t/kx)$$

- Reduces low SNR end of the FID, increases the high SNR end
- Enhances broad features, smooths out sharp features
- In signal processing, the concept of the “matched filter” says that the best value of x is to match the linewidth

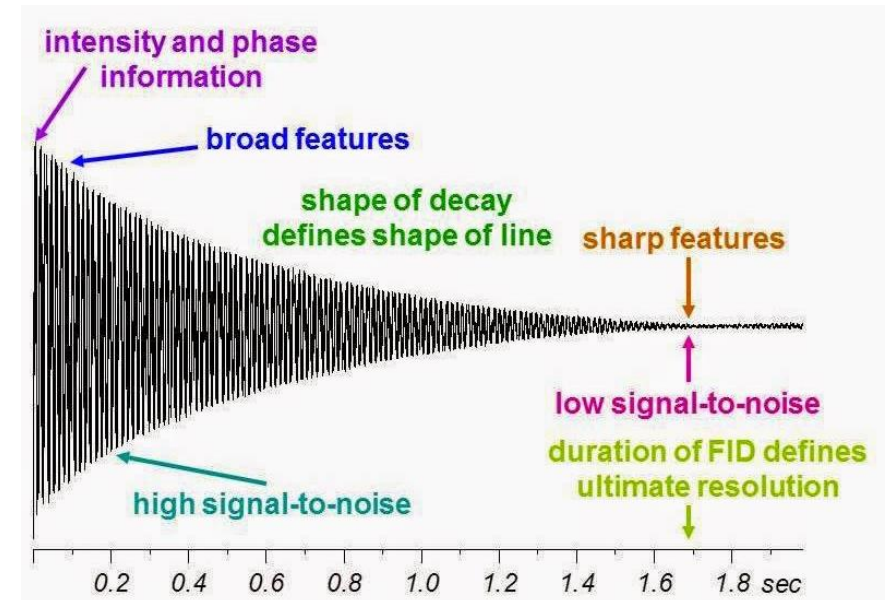


Negative line broadening

- But what if we care about the sharp features? What if we have more than enough broad features already and we want rid of them?
- How do you emphasise the other end of the FID?
- Use an exponential growth instead of a decay! (LB = -x)

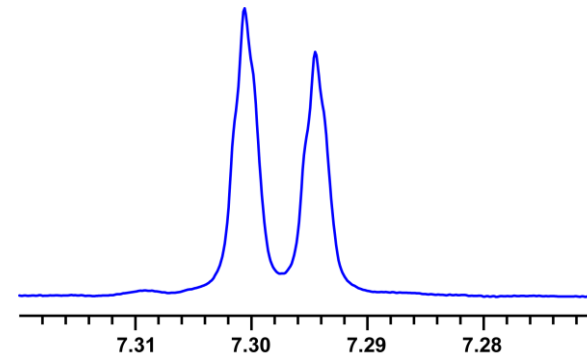
$$S(t) = \exp(i\omega t) \cdot \exp(-t/T) \cdot \exp(-t/-kx)$$

- Reduces high SNR end of the FID, increases the low SNR end
- Enhances sharp features and noise

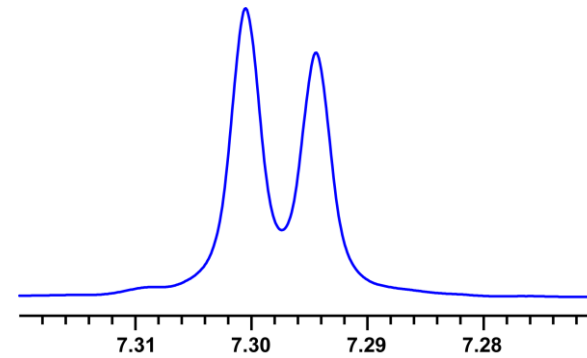


Practical example

“raw” data: LB = 0; SINO = 787; inherent linewidth 0.41 Hz



matched filter: LB = 0.4 Hz; SINO = 2195



resolution enhanced: LB = -0.2; SINO = 82

