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Proton spectrum analysis of 4-(ethylamino)-6-methyl-7-propylpyrido[3,4d]pyrimidin-8(7H)-one using 1-D NOE technique and Correlation spectroscopy methodology.

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Abstract: The power of Nuclear Magnetic Resonance spectroscopy (NMR) in structure elucidation derives in large part from its ability to establish bonding connectivity (via J- coupling interaction) or through space proximity (via dipolar coupling interactions) of nuclei the first is known as Correlation spectroscopy(COSY) and the second being Nuclear Overhauser effect spectroscopy(NOESY). The amount of time consumed in elucidating a structure depends on the rate at which these interaction can be detected by NMR and analyzed.1D NMR methods most often explore interactions between only few nuclei at a time: spin-spin decoupling measurements are used to demonstrate through-bond connectivity; and NOE measurements are used to probe inter-nuclear distances, 2-Dimensional NMR experiments provide much more structural information in a given time period.

Keywords: NMR; Nuclear Overhauser effect; Correlation spectroscopy; NOESY; decoupling

I. INTRODUCTION:

The key point in all is that magnetisation transfer occurs between coupled spins. To appreciate the outcome of this in the final COSY spectrum, consider the case of two J-coupled spins, A and X, with a coupling constant of J_{AX} and chemical shift offsets of \sqrt{A} and \sqrt{x} . The magnetisation associated with spin A will, after the initial 90⁰ pulse, precess during t1 according to its chemical shift offset, \sqrt{A} . The second 90⁰ pulse then transfers some part of this magnetisation to the coupled X spin, whilst some remains associated with the original spin A. That which remains with A will then precess in the detection period at a frequency \sqrt{A} just as it did during t1, so in the final spectrum, will produce a peak at \sqrt{A} in both dimensions, denoted (\sqrt{A} , \sqrt{A}). This peak is therefore equivalent to that observed for the uncoupled AX system and because it represents the same frequency in both dimensions, it sits on the diagonal of the 2D spectrum and is therefore referred to as a diagonal peak. In contrast, the transferred magnetisation will precess in t₂ at the frequency of the new 'host' spin X and will thus produce a peak corresponding to two different chemical shifts in the two dimensions (\sqrt{A} , \sqrt{X}). This peak sits away from the diagonal and is therefore referred to as an off-diagonal or, more commonly, a crosspeak This is the peak of interest as it provides direct evidence of coupling between spins A and X. The whole process operates in the reverse direction also, that is, the same arguments apply for magnetisation originally associated with the X spin, giving rise to a diagonal peak at (\sqrt{x}, \sqrt{x}) and a crosspeak at (\sqrt{x}, \sqrt{A}) . Thus, the COSY spectrum is symmetrical about the diagonal, with crosspeaks on either side of it mapping the same interaction.



Figure-1: Sample containing two uncoupled spins, A and X, of offsets \sqrt{A} and \sqrt{X} . Each produces a 2D peak at its corresponding chemical shift offset in both dimensions.

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Figure-2 : The COSY spectrum of a coupled, two-spin AX system. Diagonal peaks are equivalent to those observed in the 1D spectrum whilst crosspeaks provide evidence of a coupling between the correlated spins.

Theory of polarisation transfer in the INEPT experiment, it was shown that the basic requirement for the transfer of polarization was an anti-phase disposition of the doublet vectors of the source spin, which for INEPT was generated by a spin-echo sequence. Magnetisation components that were in-phase just before the second 90⁰ pulse would not contribute to the transfer, hence the Δ period was optimised to maximise the antiphase component. The same condition applies for magnetisation transfer between two protons as in the COSY experiment. This requires that the proton–proton coupling be allowed to evolve to give a degree of anti-phase magnetisation that may be transferred by the second pulse, whilst the in-phase component remains associated with the original spin. The coupling evolution period for COSY is the t₁ period so that the amount of transferred magnetisation detected in t₂ is also modulated as a function of t₁ (sin 180Jt₁). Likewise, the amplitude of the inphase, non-transferred component is also modulated in t₁ by the coupling (cos 180Jt₁), and this produces the coupling fine structure of the diagonal peak in f₁.



Figure-3: Coupling evolution during t₁ produces in-phase and antiphase magnetisation components. Only the anti-phase component contributes to magnetisation transfer and hence to crosspeaks in the 2D spectrum.



Figure-4: A schematic energy level diagram for the coupled two spin AX system.

As for 1D data, f_1 quadrature detection requires two data sets that differ in phase by 90⁰ to be collected, thus providing the necessary sine and cosine amplitude-modulated data. Since the f_1 dimension is generated artificially, there is strictly no reference rf to define signal phases, so it is the phase of the pulses that bracket t_1 that dictate the phase of the detected signal. Thus, for each t_1 increment, two data sets are collected, one with a 90_x preparation pulse (t_1 sine modulation) and the other with 90_y (t_1 cosine modulation), both stored separately These two sets are then equivalent to the two channel data collected with simultaneous acquisition, which produces the desired frequency discrimination when subject to a complex FT (also referred to as a

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hypercomplex transform in relation to 2D data). The rate of sampling in t_1 or, in other words, the size of the t_1 time increment, is dictated by the f_1 spectral width and is subject to the same rules as for the simultaneous sampling of one-dimensional data. This method is derived from the original work of States, Haberkorn and Ruben and is therefore often referred to in the literature as the States method of f_1 quad detection.



Figure-5 : The States method of f_1 quadrature detection requires two data sets to be acquired per increment to generate separate sine- and cosine modulated data sets.

II. METHODOLOGY :

The success of any NMR experiment is, of course, crucially dependent on the correct setting of the acquisition parameters. In the case of 2D experiments, one has to consider the parameters for each dimension separately, and we shall see that the most appropriate parameter settings for f_2 are rarely optimum for fl. Likewise, one has to give rather more thought to the setting up of a 2D experiment than is usually required for 1D acquisitions to make optimum use of the instrument time available and data storage space. spectral widths, which should be the same in both dimensions of the COSY experiment, should be kept to minimum values with transmitter offsets adjusted so as to retain only the regions of the spectrum that will provide useful correlations. It is usually possible to reduce spectral widths to well below the 10-ppm or so proton window observed in 1D experiments. The use of excessively large windows leads to poorer digital resolution in the final spectrum or requires greater data sizes, neither of which is desirable. The spectral widths in turn define the sampling rates for data in t₂, in exact analogy with 1D acquisitions, and the size of the t₁ increment, again according to the Nyquist criteria. The acquisition time (AQ_i) , and hence the digital resolution $(1/AQ_i)$, for each dimension is then dictated by the number of data points collected in each. For t_2 , this is the number of data points digitised in each FID, whilst for t₁ this is the number of FIDs collected over the course of the experiment. The appropriate setting of these parameters is a most important aspect to setting up a 2D experiment, and the way in which one thinks about acquisition times and digital resolution in a 2D data set is, necessarily, quite different from that in a 1D experiment. As an illustration, imagine transferring the typical parameters used in a 1D proton acquisition into the two dimensions of COSY. The acquisition time might be 4 s, corresponding to a digital resolution of 0.25 Hz/pt, with no relaxation delay between scans. On, for example, a 400-MHz instrument, with a 10 ppm spectral width, this digital resolution would require 32K words to be collected per FID. The 2D equivalent, with States quad detection in f_1 and with axial-peak suppression, requires four scans to be collected for each t_1 increment. The mean acquisition time for each would be 6 s (t_2 plus the mean t_1 value), corresponding to 24 s of data collection per FID. If 16K t_1 increments were to be made for the f_1 dimension (two data sets are collected for each t₁ increment remember), this would correspond to a total experiment time of about 4.5 days. Furthermore, the size of the resulting data matrix would be a little over 1000 million words, and with a typical 32-bit-per-word computer system, this requires some 4GB of disk space! We will agree that 4 days for a basic COSY acquisition is quite unacceptable, let alone the need for such disk space per experiment; so acquiring data with such high levels of digitisation in both dimensions is clearly not possible.

The key lies in deciding on what level of digitisation is required for the experiment in hand. The first point to notice is that adding data points to extend the t_2 dimension leads to a relatively small increase in the overall length of the experiment, so we may be quite profligate with these (although they will lead to a corresponding increase in the size of the data matrix). Moreover, adding t1 data points requires that a complete

FID of potentially many scans is required per increment, which makes a far greater increase to the total data collection time. Thus, one generally aims to keep the number of t_1 increments to a minimum, which is consistent with resolving the correlations of interest and increasing t_2 as required when higher resolution is necessary. For this reason, the digital resolution in f_2 is often greater than that in f_1 , particularly in the case of phase-sensitive data sets. The use of smaller AQt₁ is, in general, also preferred for reasons of sensitivity since FIDs recorded for longer values of t_1 will be attenuated by relaxation and so will contribute less to the overall signal intensity. The use of small AQt₁ is likely to lead to truncation of the t_1 data, and it is then necessary to apply suitable window functions that force the end of the data to zero to reduce the appearance of truncation artefacts.

For COSY in particular, one of the factors that limits the level of digitisation that can be used is the presence of intrinsically anti-phase crosspeaks, since too low a digitisation will cause these to cancel and the correlation to disappear The level of digitisation will also depend on the type of experiment and the data one expects to extract from it. For absolute-value COSY, one is usually interested in establishing where correlations exist, with little interest in the fine structure within these crosspeaks. In this case, it is possible to use a low level of digitisation consistent with identifying correlations. As a rule of thumb, a digital resolution of J to 2J Hz/pt (AQ of 1/J to 1/2J s) should enable the detection of most correlations arising from couplings of J Hz or greater. Thus for a lower limit of, say, 3 Hz, a digital resolution of 3–6 Hz/pt (AQ of \sim 300–150 ms) will suffice. The AQ_{t1} is typically half that for t₂ in this experiment, with one level of zero- filling applied in t1 so that the final digital resolution is the same in both dimensions of the spectrum (as required for symmetrisation).

For phase-sensitive data acquisitions, one is likely to be interested in using the information contained within the crosspeak multiplet structures, and a higher degree of digitisation is required to adequately reflect this, a more appropriate target being around J/2 Hz/pt or better (AQ of 2/J s or greater). Again, digitisation in t_2 is usually two or even four or eight times greater than that in t1. In either dimension, but most often in t_1 , this may be improved

by zero-filling, although one must always remember that it is the length of the time-domain acquisition that places a fundamental limit on peak resolution and the effective linewidths after digitisation, regardless of zero-filling. The alternative approach for extending the time domain data is to use forward linear prediction when processing the data. The rule as ever is that high resolution requires long data-sampling periods.

Having decided on suitable digitisation levels and data sizes, one is left to choose the number of scans or transients to be collected per FID and the repetition rates and hence relaxation delays to employ. The minimum number of transients is dictated by the minimum number of steps in the phase cycle used to select the desired signals. Further scans may include additional steps in the cycle to suppress artefacts arising from imperfections. Beyond this, further transients should be required only for signal averaging when sensitivity becomes a limiting factor. Since most experiments are acquired under 'steady-state' conditions, it is also necessary to include 'dummy' scans prior to data acquisition to allow the steady state to establish. On modern instruments that utilize double buffering of the acquisition memory, dummy scans are required only at the very beginning of each experiment to make a negligible increase to the total time required. On older instruments that lack this feature, it is necessary to add dummy scans for each t_1 increment, and these may then make a significant contribution to the total duration of the experiment. The repetition rate will depend upon the proton T_1s in the molecule, and since the sequence uses 90⁰ pulses, the optimum sensitivity is achieved by repeating every 1.3 T_1s .

Returning to the example of 400MHz acquisition discussed above, we can apply more appropriate criteria to the selection of parameters. Table 1 compares the result from above with more realistic data, and it is clear that under these conditions, COSY becomes a viable experiment, requiring only hours or even minutes to collect, rather than many days. The introduction of PFGs to high-resolution spectroscopy allows experiments to be acquired with only one transient per FID where sensitivity is not limiting, thus further reducing the total time required for data collection. The data storage requirements in these realistic examples are also well within the capabilities of modern computing hardware and are likely to become increasingly less significant as this develops further.

Experiment	Spectral width/ppm	$N(t_2)$	$N(t_1)$	Hz/pt (t ₂)	Hz/pt (t_1)	Experiment time	Raw data-set size
(a) Phase-sensitive	10×10	32K	32K	0.25	0.25	4.5 days	1000 M words
(c) Absolute value	6×6	2K 1K	256	4.6	4.6	22 min	0.25 M words

Table-1: Illustrative data tables for COSY experiments

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Scenario (a) transplants acquisition parameters from a typical 1D proton spectrum into the second dimension leading to unacceptable time requirements, whereas (b) and (c) use parameters more appropriate to 2D acquisitions. All calculations use phase cycles for f_1 quad detection and axial-peak suppression only and for (b) and (c), a recovery delay of 1 s between scans. A single zero-filling in f1 was also employed for (b) and (c). However when Nuclear Overhauser effect is considered ,resonance line intensity changes caused by dipolare cross relaxation from neighbouring spins with perturbed energy level populations. To understand the nature of the NOE, we have to look at a two-spin system I_1 and I_2 . Since NOE does not involves coherences, but merely polarization, i.e. population differences between the α and β states, we can use the energy level diagram here



Figure-6 Energy level diagram showing population differences between $\alpha \& \beta$ states

The possible transition for this two-spin system can be classified into three groups:

- W_1 transition involving a spin flip of only one of the two spins(either I_1 or I_2), corresponding merely to T_1 relaxation of the spin.

- a W₀ transition involving a simultaneous spin flip $\alpha \rightarrow \beta$ for one spin and $\alpha \rightarrow \beta$ for the other one (i.e., in summa a zero-quantum transition).

- a W_2 transition involving a simultaneous spin flip of both spins in the same direction, corresponding to a net double-quantum transition.

We are just contemplating spin state transitions here caused by relaxation, which do *not* involve a *coherent* process (like, e.g., I_x / I_y coherences, which require a phase coherent transition between two states that is generated by an r.f. pulse). If we perturb one spin, e.g., I_1 , i.e., change its populations of the α and β state (e.g., by saturating the resonance = creating *equal* population of both states), then relaxation will force I_1 back to the equilibrium BOLTZMANN distribution. With the W1 mechanism, spin I_1 will just relax without effecting spin I_2 . However, the other two mechanism *will* effect I_2 .



Figure-7 Energy level diagram showing population differences

With the I_1 polarization going back from saturation to the BOLTZMANN equilibrium, the W mechanism will cause the neighbouring (so far unperturbed) spin to deviate from its BOLTZMANN equilibrium towards a *decrease* in α , β population difference. After a 90° pulse, this will result in a *decrease* in signal intensity for I_2 — a "negative NOE effect". On the other hand, the W₂ mechanism will cause the population

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difference of the undisturbed spin I_2 to *increase*, corresponding to an *increase* in signal intensity: a "positive NOE effect". These effects can be directly observed in a very simple experiment, the 1D difference NOE sequence:



Figure-8 Figure showing the 1D difference NOE sequence

One spin is selectively saturated by a long, low-power CW (continuous wave) irradiation. As soon as the spin deviates from its BOLTZMANN population distribution, it starts with T_1 relaxation. Via the W_0 or W_2 mechanisms it causes changes in the population distribution of neighboring spins. After a 90° pulse, these show up as an increase or decrease in signal intensity. Usually, the experiment is repeated without saturation, giving the normal 1D spectrum. This is then subtracted from the irradiated spectrum, so that the small intensity changes from the NOE effects can be easier distinguished: spins with a positive NOE (i.e., higher intensity in the NOE spectrum than in the reference 1D) show a small positive residual signal, spins with a negative NOE yield a negative signal, spins without an NOE cancel completely.

RESULTS AND DISCUSSION:



Figure-9 Figure showing the 1H NMR spectra of the sample under study(source Bruker-AV400)



Figure-10 Figure showing expanded 1H NMR spectra of the sample under study (source Bruker-AV400)

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Figure-11 Figure showing COSY NMR spectra of the sample under study (source Bruker-AV400)



Figure-12 Figure showing expanded form of COSY NMR spectra of the sample under study (source Bruker-AV400)



Figure-13 Figure showing 1-D NOE NMR spectra of the sample under study (source Bruker-AV400) a clear correlation between CH₃(2.4ppm;s), CH₂(4.04ppm;q) & CH(6.2ppm;s) is observed.





IUPAC name of the sample under study is 4-(ethylamino)-6-methyl-7-propylpyrido[3,4-d]pyrimidin-8(7H)-one From the study of the compound spectra we were able to note the NMR spectrum as under: s(1H;8.6);q(2H;3.6);t(3H;1.3);s(1H;6.2);s(3H;2.4);t(2H;4.04);sextat(2H;1.7);t(3H;0.9); NH(1H;5.7;broad) the same has been shown in the figure-14 also.

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