

A Three-Dimensional NMR Experiment for the Separation of Aliphatic Carbon Chemical Shifts via the Carbonyl Chemical Shift in ^{15}N , ^{13}C -Labeled Proteins

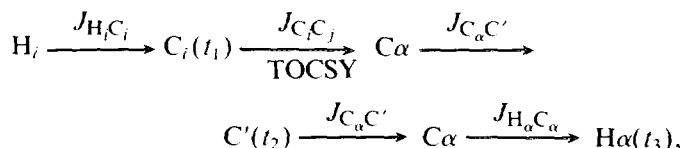
LEWIS E. KAY

Protein Engineering Network Centers of Excellence and Departments of Medical Genetics, Biochemistry and Chemistry,
Medical Sciences Building, University of Toronto, Toronto, Ontario, Canada M5S 1A8

Received September 24, 1992

The assignment of the proton spectrum of a protein is the first step in its structure determination by NMR methods (1). For small proteins, with molecular weights less than approximately 10 kDa, this can be successfully accomplished using homonuclear 2D experiments. For larger proteins, NOE-based assignment strategies often fail due to spectral overlap, while experiments which correlate chemical shifts via ^1H – ^1H scalar couplings become insensitive. The development of triple-resonance multidimensional spectroscopy has significantly extended the molecular weight limit of proteins that can be studied with NMR techniques (2, 3). Using a variety of 3D (2–7) and 4D (8–10) experiments, it is possible to assign the backbone resonances in ^{15}N , ^{13}C uniformly labeled proteins using only through-bond correlations. The assignment of side-chain chemical shifts is accomplished using the HCCH–COSY (11, 12) and the HCCH–TOCSY (13, 14) experiments, which transfer magnetization via the large and uniform one-bond ^{13}C – ^{13}C couplings. Recently, experiments have been developed which separate side-chain correlations via the backbone carbonyl (C') chemical shift (15). One such experiment, the HCACO–COSY, provides correlations linking the C' , $\text{H}\alpha$, and $\text{H}\beta$ chemical shifts, while the HCACO–TOCSY experiment links the C' chemical shift with all the ^1H shifts along a given side chain. A third experiment correlates the C' , $\text{C}\beta$, and $\text{H}\beta$ as well as the C' , $\text{C}\alpha$, and $\text{H}\alpha$ chemical shifts. Grzesiek and Bax have developed experiments which link ^{15}N and NH chemical shifts with intra-residue and sequential $\text{C}\alpha$ and $\text{C}\beta$ chemical shifts (16, 17). In this Communication, an experiment is presented which separates side-chain carbon chemical shifts via the intra-residue C' chemical shift. This experiment is closely related to the HCACO–TOCSY scheme (15), and these two experiments together provide a powerful approach for the assignment of side-chain ^1H and ^{13}C resonances.

Figure 1 illustrates the pulse sequence used to correlate the carbon side chain and C' chemical shifts. Magnetization transfer follows the pathway



where H_i and C_i are side-chain proton and carbon spins, respectively, and the active couplings responsible for magnetization transfer are indicated above the arrows. The pulse sequence can be understood in more detail by considering a product-operator description (18) of the magnetization-transfer process. In what follows such a description is provided. For simplicity, the effects of relaxation are neglected, only the terms of interest are retained, and numerical factors preceding these terms are omitted. The side-chain ^1H and ^{13}C spins are denoted by I' and S' , respectively, the $\text{H}\alpha$, $\text{C}\alpha$, and C' spins are denoted by I , S , and C' , and J_{ij} is the scalar coupling constant between spins i and j . ^1H longitudinal magnetization at time a is transferred via an INEPT transfer (19) to the side-chain carbons. At time b in the sequence, the magnetization is given by

$$\rho_b = \text{I}'_z \text{S}'_y \sin(2\pi J_{\text{I}'\text{S}'}\tau_a). \quad [1]$$

During τ_b , carbon magnetization which is antiphase with respect to the directly coupled proton spin(s) refocuses, and the carbon chemical shift is recorded in a constant-time manner (20, 21) during the period $2T$. Because each carbon spin can be attached to one, two, or three protons, a compromise value of $\tau_b \sim 0.3/J_{\text{I}'\text{S}'}$ is chosen. In order to enhance the sensitivity of the experiment, carbon magnetization is maintained in-phase with respect to the directly coupled proton spin(s) by the application of coherent decoupling on the proton channel (5, 6). The effects of C' – C and N – C scalar couplings are refocused by the application of C' and N 180° pulses at $t_1/2$. At time c , the magnetization of interest is given by

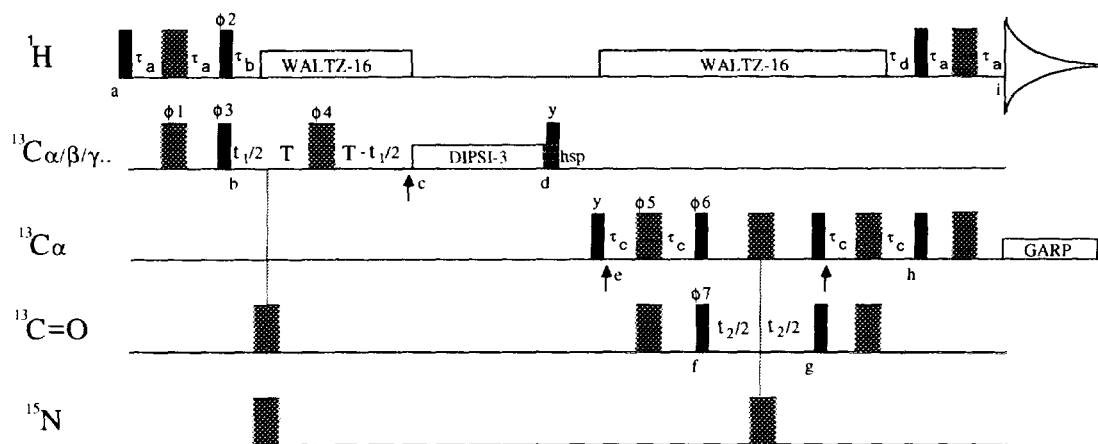


FIG. 1. Pulse sequence used to correlate aliphatic carbon shifts with the intrasidue C' shift. All narrow pulses have a flip angle of 90° with wider pulses having a flip angle of 180° . Pulses for which the phases are not indicated are applied along the x axis. The $C\alpha/\beta/\gamma \dots$ pulses are applied at 43 ppm with an RF field strength of 4.4 kHz for the 90° pulses and a field strength of 9.8 kHz for the 180° pulses. The $C\alpha$ pulses are applied at 54 ppm with an RF field strength of 4.0 kHz for the 90° pulses and a field strength of 9.0 kHz for the 180° pulses. This ensures that the application of the aliphatic pulses produces minimal effects in the C' region of the spectrum and vice versa. All C' pulses are implemented as phase-modulated pulses. A DIPSI-3 mixing scheme (22) of 16 ms is employed, preceded by a 1 ms trim pulse and followed by a trim pulse of 0.5 ms. Because the carbon synthesizer on our spectrometer does not maintain a constant phase during a frequency jump, magnetization is transferred to the z axis prior to shifting the frequency from 43 to 54 ppm. A 10 ms homospoil (hsp) pulse is applied to eliminate magnetization not along the z axis during this interval. Carbon decoupling during acquisition is achieved via GARP decoupling (27) with a 2 kHz RF field. A WALTZ (28) decoupling field of 6 kHz was employed to maintain in-phase carbon magnetization throughout the course of much of the sequence. The delays used are $\tau_a = 1.78$ ms, $\tau_b = 2.2$ ms, $\tau_c = 3.6$ ms, $\tau_d = 3.5$ ms, and $T = 3.0$ ms. The phase cycle used is $\phi_1 = x, -x$; $\phi_2 = y, -y$; $\phi_3 = x$; $\phi_4 = 2(x), 2(y), 2(-x), 2(-y)$; $\phi_5 = 8(x), 8(-x)$; $\phi_6 = 8(x), 8(-x)$; $\phi_7 = 16(x), 16(-x)$; Rec = $x, 2(-x), 2(x), 2(-x), x, -x, 2(x), 2(-x), 2(x), -x$. The receiver is inverted for the last 16 scans. Quadrature in t_1 and t_2 is obtained using the STATES-TPPI method (29) by incrementing the phases ϕ_3 and ϕ_7 . The arrows indicate the points at which the C' compensation 180° pulses are applied.

$$\rho_c = S'_x \cos(\omega_S t_1) \sin(\pi J_{1'S} \tau_b) \times \cos^n(\pi J_{1'S} \tau_b) \prod_{K \neq S'} \cos(2\pi J_{S'K} T), \quad [2]$$

where ω_S is the Larmor frequency of spin S' , $n = 0, 1, 2$ for methine, methylene, and methyl carbons, respectively, and $J_{S'K}$ is a homonuclear $^{13}\text{C}-^{13}\text{C}$ coupling constant. Subsequently ^{13}C magnetization is relayed among aliphatic side-chain carbons during the DIPSI-3 (22) mixing scheme. Magnetization that resides on the $C\alpha$ carbon after the DIPSI mixing period (point d) is then transferred to the coupled C' spin to give magnetization at time f of

$$\rho_f = S'_x C'_z \sin(2\pi J_{SC'} \tau_c) \cos(2\pi J_{SS'} \tau_c). \quad [3]$$

After the simultaneous 90° pulses on the $C\alpha$ and C' spins, the C' shift is recorded during t_2 while the $C'-C\alpha$ and $C'-N$ scalar couplings are refocused by the application of $C\alpha$ and N 180° pulses in the middle of the t_2 evolution. At time g , the magnetization is given by

$$\rho_g = S'_x C'_z \cos(\omega_C t_2). \quad [4]$$

The magnetization is subsequently transferred from the C' spin to the $C\alpha$ spin. The $C\alpha$ magnetization is refocused due

to the $C'-C\alpha$ scalar coupling during $2\tau_c$ as it dephases due to the $H\alpha-C\alpha$ coupling during the period τ_d . Immediately prior to detection, the magnetization is given by

$$\rho_g = I_x \cos(\omega_S t_1) \cos(\omega_C t_2) \prod_{K \neq S'} \cos(2\pi J_{S'K} T) \times \sin(2\pi J_{1'S} \tau_a) \sin(\pi J_{1'S} \tau_b) \cos^n(\pi J_{1'S} \tau_b) \Gamma(\text{DIPSI}) \times \sin^2(2\pi J_{SC'} \tau_c) \cos^2(2\pi J_{SS'} \tau_c) \times \sin(\pi J_{IS} \tau_d) \cos^k(\pi J_{IS} \tau_d) \sin(2\pi J_{IS} \tau_a), \quad [5]$$

where $\Gamma(\text{DIPSI})$ denotes the efficiency of magnetization transfer from spin S' to spin S , $k = 1$ for glycine, and $k = 0$ for all other residues. Bax and co-workers have presented figures showing the transfer of magnetization between particular carbons for different amino acid side chains as a function of mixing time (13, 23). Fourier transformation of the resulting signal gives rise to peaks at coordinates $(\omega_{S'}, \omega_{C'}, \omega_1)$. Therefore, the side-chain carbon chemical shifts are displayed along a line which intersects the F_3 axis at the intrasidue $H\alpha$ chemical shift.

The carbon pulse widths in this experiment are adjusted so that the application of a C' pulse produces minimal effects in the $C\alpha$ (54 ppm) or $C\alpha/\beta/\gamma \dots$ (43 ppm) regions of the spectrum and vice versa. The C' pulses are applied 15.6 kHz

(17.0 kHz) downfield from the $C\alpha$ ($C\alpha/\beta/\gamma \dots$) carrier by phase modulation of the RF source. Despite careful adjustments of the pulse widths, the off-resonance effects of the pulses cannot be neglected (24, 25). For example, consider the evolution of aliphatic γ magnetization during the application of a C' x pulse centered ν_0 hertz downfield from the carbon carrier. It can be shown that

$$S'_y \rightarrow S'_y \cos 2\pi\{\Delta + 0.5(\nu_1^2/\nu_0)\}\tau_p - S'_x \sin 2\pi\{\Delta + 0.5(\nu_1^2/\nu_0)\}\tau_p, \quad [6]$$

where S'_x and S'_y are the x and y components of aliphatic magnetization, ν_1 is the strength of the applied RF field, τ_p is the duration of the pulse, and Δ is the chemical-shift offset of the aliphatic magnetization from the carbon carrier. In the derivation of Eq. [6] it is assumed that $\nu_1/\nu_0 \ll 1$ and that $\Delta \ll \nu_0$. At a spectrometer frequency of 500 MHz and for $C\alpha$ magnetization with the carbon carrier centered at 54 ppm, $|\Delta| < \sim 1.3$ kHz, $\nu_1 = 4.0$ kHz, and $\nu_0 = 15.6$ kHz.

Equation [6] indicates that the C' 180° pulse applied between points b and c in Fig. 1 produces a phase shift, $\phi \sim \pi(\nu_1^2/\nu_0)\tau_p$, of the transverse aliphatic magnetization that cannot be corrected by simply inserting a delay equal to the C' pulse width on the opposite side of the aliphatic 180° pulse (ϕ^4). Following the lead of Bax and co-workers (16, 17, 26), the phase error introduced by the application of the C' 180° pulse is compensated by the application of a second C' 180° pulse immediately prior to the DIPSI mixing period. This ensures that carbon magnetization is modulated by a pure cosine (or sine) function in t_1 . In a similar fashion, the application of the C' 180° pulse between points e and f and between points g and h results in a phase shift of the transverse $C\alpha$ magnetization. Although not indicated in the figure, in both cases the C' 180° pulse is applied immediately after the $C\alpha$ 180° pulse. Unless properly corrected, the off-resonance effects of these C' pulses result in a decrease in the overall sensitivity of the experiment since the $C\alpha$ 90° pulses that follow the $C\alpha/C'$ 180° pulse pairs require a particular phase of the magnetization for optimal transfer. In the present experiment, the C' 180° pulses are applied with a strength of 4.0 kHz for a duration of 124.1 μ s. This creates an off-resonance phase shift of $\sim 25^\circ$, giving rise to a decrease in sensitivity of $\sim 10\%$ for each occasion where the C' pulse is compensated only by a delay equal to its length. The arrows in Fig. 1 indicate the points at which the C' compensation pulses are applied.

The sequence described is illustrated on a 1.5 mM sample of ^{15}N , ^{13}C -labeled calmodulin from *Xenopus laevis*, complexed with four molar equivalents of calcium, 0.1 M KCl, pH 6.3 (uncorrected), and dissolved in 0.5 ml D_2O . The experiment was performed on a Varian UNITY-500 spectrometer at 37°C . The spectrum was obtained from a $42 \times 42 \times 512$ complex matrix with acquisition times of 6.0 ms

(t_1), 28.0 ms (t_2), and 85.0 ms (t_3). A 32-step phase cycle was used with a repetition delay of 0.8 s, giving rise to a total acquisition time of 2.5 days.

Figures 2 and 3 illustrate the quality of the data obtained. For the 16 ms mixing time employed, many of the side chains show complete correlations. For example, the $C\alpha$, $C\beta$, $C\gamma$, and $C\delta$ peaks of R106 are clearly illustrated in the plane with a C' shift of 178.9 ppm and, although the maximum intensities for the cross peaks from K115 are on an adjacent slice, it is still possible to obtain all the carbon side-chain chemical shifts from the slice at 176.1 ppm.

In summary, a pulse sequence has been presented in this Communication for the correlation of the side-chain carbon chemical shifts with the backbone C' and $H\alpha$ chemical shifts. A related set of useful experiments consists of those which separate side-chain chemical shifts via the backbone ^{15}N and NH shifts (16, 17). The motivation for separating side-chain shifts based on the backbone C' resonances in this and other studies (15) is based on the fact that experiments which correlate both side-chain and backbone shifts are, in general, less sensitive than the corresponding experiments which correlate side-chain or backbone resonances exclusively. In the

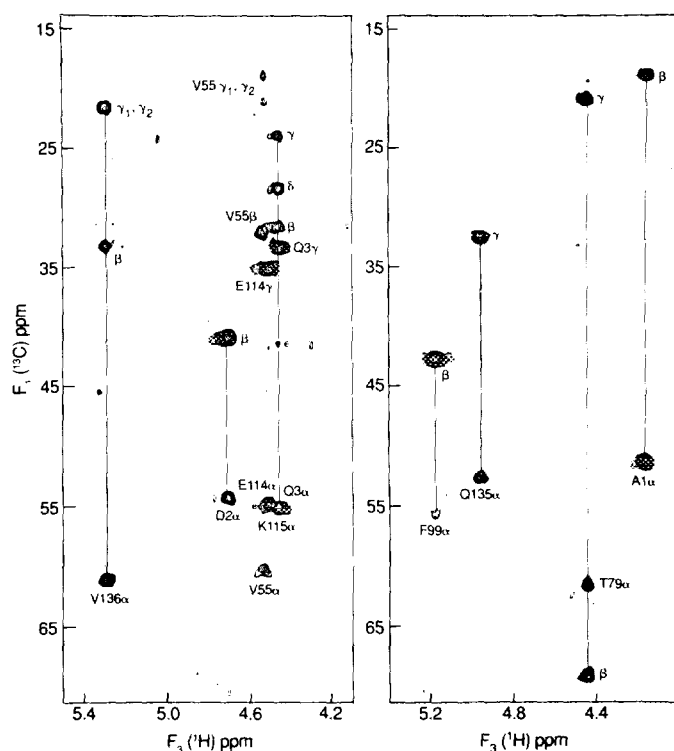


FIG. 2. Sections of slices (F_1/F_3) from the C' -edited ^{13}C - ^{13}C TOCSY experiment at C' shifts of 174.9 and 176.1 ppm. The experiment was recorded on a 1.5 mM sample of *Xenopus laevis* calmodulin at 500 MHz, 37°C , pH 6.8. The spectrum was recorded as a $42 \times 42 \times 512$ complex data matrix. After processing, including linear prediction of the t_1 time domain, the absorptive part of the final 3D spectrum was $128 \times 128 \times 1024$. (Left) $F_2(^{13}\text{C}') = 176.1$ ppm; (right) $F_2(^{13}\text{C}') = 174.9$ ppm.

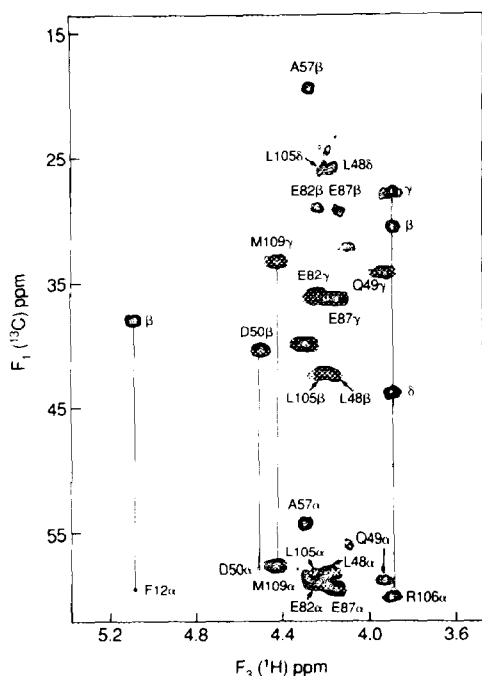


FIG. 3. Expanded region of the F_1/F_3 slice of the C' -edited TOCSY experiment at a C' shift of 178.9 ppm. $F_2(^{13}\text{C}) = 178.9$ ppm.

construction of backbone-edited side-chain experiments it is important, therefore, to ensure that each magnetization-transfer step is as efficient as possible. In this regard, HCACO-based side-chain experiments are a particularly good choice because of the very high intrinsic sensitivity of the HCACO experiment. The experiment presented here, when used in conjunction with the HCCH-COSY/TOCSY and HCACO-COSY/TOCSY schemes, should enable the assignment of aliphatic side-chain ^1H and ^{13}C shifts in a straightforward fashion.

ACKNOWLEDGMENTS

The author is grateful to Dr. Mitsu Ikura, Ontario Cancer Institute, Toronto, for kindly providing a sample of ^{15}N , ^{13}C -labeled calmodulin and for helpful discussions. Discussions with Drs. Ad Bax, NIH, Bethesda, and Mark McCoy, Eastman Kodak, Rochester, are also acknowledged. This research was supported through a grant from the Natural Sciences and Engineering Research Council of Canada.

Note added in proof. Slightly better sensitivity can be obtained by setting $\tau_c \sim 2.7$ ms. In this case, the second ^1H WALTZ-16 decoupling period should end at position g in the sequence and an additional ^1H 180° pulse is inserted 1.8 ms after position g . In this case τ_d has no meaning.

REFERENCES

1. K. Wüthrich, "NMR of Proteins and Nucleic Acids," Wiley, New York, 1986.
2. M. Ikura, L. E. Kay, and A. Bax, *Biochemistry* **29**, 4659 (1990).
3. L. E. Kay, M. Ikura, R. Tschudin, and A. Bax, *J. Magn. Reson.* **89**, 496 (1990).
4. A. G. Palmer III, W. J. Fairbrother, J. Cavanagh, P. E. Wright, and M. Rance, *J. Biomol. NMR* **2**, 103 (1992).
5. B. T. Farmer II, R. A. Venters, L. D. Spicer, M. G. Wittekind, and L. Mueller, *J. Biomol. NMR* **2**, 195 (1992).
6. S. Grzesiek and A. Bax, *J. Magn. Reson.* **96**, 432 (1992).
7. R. T. Clubb, V. Thanabal, and G. Wagner, *J. Magn. Reson.* **97**, 213 (1992).
8. L. E. Kay, M. Ikura, G. Zhu, and A. Bax, *J. Magn. Reson.* **91**, 422 (1991).
9. W. Boucher, E. D. Laue, S. Campbell-Burk, and P. J. Dommelle, *J. Am. Chem. Soc.* **114**, 2262 (1992).
10. L. E. Kay, M. Wittekind, M. A. McCoy, M. S. Friedrichs, and L. Mueller, *J. Magn. Reson.* **98**, 443 (1992).
11. L. E. Kay, M. Ikura, and A. Bax, *J. Am. Chem. Soc.* **112**, 888 (1990).
12. A. Bax, G. M. Clore, P. C. Driscoll, A. M. Gronenborn, M. Ikura, and L. E. Kay, *J. Magn. Reson.* **87**, 620 (1990).
13. A. Bax, G. M. Clore, and A. M. Gronenborn, *J. Magn. Reson.* **88**, 425 (1990).
14. S. W. Fesik, H. L. Eaton, E. T. Olejniczak, E. R. P. Zuiderweg, L. P. McIntosh, and F. W. Dahlquist, *J. Am. Chem. Soc.* **112**, 886 (1990).
15. L. E. Kay, M. Ikura, A. A. Grey, and D. R. Muhandiram, *J. Magn. Reson.* **99**, 652 (1992).
16. S. Grzesiek and A. Bax, *J. Magn. Reson.* **99**, 201 (1992).
17. S. Grzesiek and A. Bax, *J. Am. Chem. Soc.* **114**, 6291 (1992).
18. O. W. Sørensen, G. Eich, M. H. Levitt, G. Bodenhausen, and R. R. Ernst, *Prog. NMR Spectrosc.* **16**, 163 (1983).
19. G. A. Morris and R. Freeman, *J. Am. Chem. Soc.* **101**, 760 (1979).
20. A. Bax and R. Freeman, *J. Magn. Reson.* **44**, 542 (1981).
21. O. W. Sørensen, *J. Magn. Reson.* **90**, 433 (1990).
22. A. J. Shaka, C. J. Lee, and A. Pines, *J. Magn. Reson.* **77**, 274 (1988).
23. G. M. Clore, A. Bax, P. C. Driscoll, P. T. Wingfield, and A. M. Gronenborn, *Biochemistry* **29**, 8172 (1990).
24. M. A. McCoy and L. Mueller, *J. Magn. Reson.* **99**, 19 (1992).
25. M. A. McCoy and L. Mueller, *J. Magn. Reson.* **98**, 674 (1992).
26. G. W. Vuister and A. Bax, *J. Magn. Reson.* **98**, 428 (1992).
27. A. J. Shaka, P. B. Barker, and R. Freeman, *J. Magn. Reson.* **64**, 547 (1985).
28. A. J. Shaka, J. Keeler, T. Frenkiel, and R. Freeman, *J. Magn. Reson.* **52**, 335 (1983).
29. D. Marion, M. Ikura, R. Tschudin, and A. Bax, *J. Magn. Reson.* **85**, 393 (1989).