



## Heteronuclear decoupling by multiple rotating frame technique

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### ABSTRACT

The paper describes the multiple rotating frame technique for designing modulated rf fields, that perform broadband heteronuclear decoupling in solution NMR spectroscopy. The decoupling method presented here is understood by performing a sequence of coordinate transformations, each of which demodulates a component of the rf field to a static component, that progressively averages the chemical shift and the dipolar interaction. We show that by increasing the number of modulations in the decoupling field, the ratio of dispersion in the chemical shift to the strength of the static component of the rf field is successively reduced in the progressive frames. The known decoupling methods like continuous wave decoupling, TPPM, etc., can be viewed as special cases of this method and their performance improves by adding additional modulations in the decoupling field. The technique is also expected to find use in design of broadband excitation, inversion and mixing sequences and broadband experiments in solid state NMR.

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### 1. Introduction

Heteronuclear decoupling methods have a long history in NMR spectroscopy [1–13,15–27,29,31–33,49–58,61–63] and *in vivo* applications [34–39]. The goal of broadband heteronuclear decoupling sequences is to observe a spin *S* by irradiating spin *I* that is coupled to *S* in order to simplify the spectra and to increase the signal-to-noise ratio. At the same time, the decoupling sequence should introduce only a minimal amount of artifacts, such as decoupling sidebands. Furthermore, in order to avoid undesirable sample heating or damage to the probe, the radio frequency (rf) power of the decoupling sequence should be as small as possible. This is of particular importance in medical imaging or *in vivo* spectroscopy of humans [34–39].

The earliest heteronuclear decoupling methods were based on CW irradiation [10] and noise decoupling [12]. Significantly improved decoupling sequences were found based on composite [8,9,13,15,17–20] or shaped [25–27,29,31,33] inversion pulses in combination with highly compensated cycles and supercycles [8,9,17,40–46]. Theoretical approaches that have been used for the analysis and design of decoupling sequences include average Hamiltonian [16,47,48] and Floquet [32,33] theory.

In this paper, we describe the multiple rotating frame technique for design of heteronuclear decoupling sequences.

In the absence of chemical shifts, the most straightforward way to decouple two heteronuclear spins *I* and *S* is to just irradiate with

full power on resonance to spin *I*. However, in the presence of large chemical shifts, one creates an effective field that is not perpendicular to the coupling interaction and therefore part of the coupling interaction parallel to this field is not averaged. We show that it is possible to design a multiply-MODulated rf field (MODE sequence) for broadband decoupling, whose effect is best understood by performing a sequence of coordinate transformations, *specific to each chemical shift*. Each transformation demodulates a component of the multiply-modulated rf field to a static component. The subsequent transformation is in the rotating frame defined by new static component and the residual chemical shift. Each transformation averages part of the coupling and reduces the ratio of chemical shift interaction to static rf field, progressively making the effective field perpendicular to the coupling interaction. We show that by increasing the number of modulations in the decoupling field, we can significantly improve the decoupling. Methods to make this method robust to rf inhomogeneity are discussed.

### 2. Heteronuclear decoupling

Consider two heteronuclear spins *I* and *S*. The Hamiltonian of the spin system takes the form

$$\mathcal{H}_0 = \omega_I I_z + 2\pi J_{Iz} S_z + \omega_S S_z, \quad (1)$$

where  $\omega_I$  and  $\omega_S$  are the chemical shifts of spin *I* and *S*. Assuming spin *S* is being observed, we can without loss of generality take the chemical shift of the spin *S* to be zero and denoting  $\omega_I = \omega$ , consider the natural Hamiltonian

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$$\mathcal{H}_0 = \underbrace{\omega I_z}_{H_{cs}} + \underbrace{2\pi J I_z S_z}_{H_c} \quad (2)$$

where  $\omega \in [-c_0, c_0]$ . Now, consider the rf Hamiltonian

$$H'_0 = w_0 I_x + (w_1 \sin v_1 t + w_2 \cos v_1 t \sin v_2 t + \dots) I_y. \quad (3)$$

A more general form of the rf Hamiltonian is then

$$H'_0 = w_0 I_x + \left( \underbrace{\sum_{k=1}^N w_k \prod_{j=1}^{k-1} \cos v_j t \sin v_k t}_{A_0(t)} \right) I_y, \quad (4)$$

where  $v_k > v_{k+1}$ . We adopt the notation  $\bar{w}_k = 2^{-k} w_k$ , and rewrite the total Hamiltonian  $H_0 = \mathcal{H}_0 + H'_0$  as

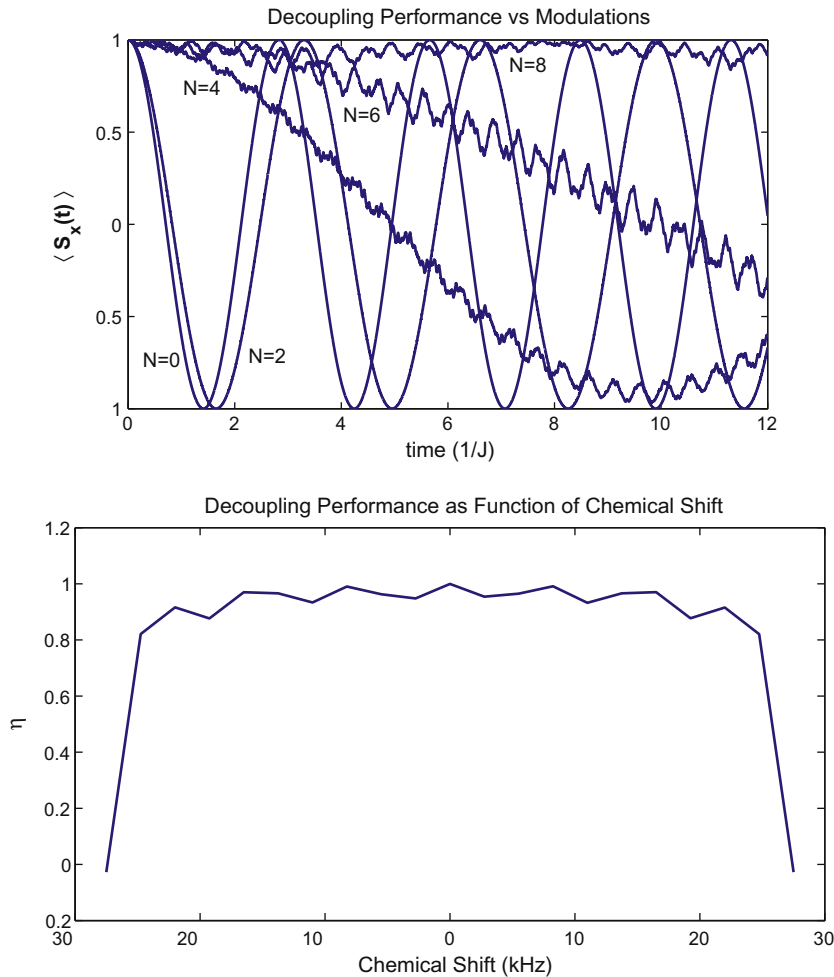
$$H_0 = \tilde{\omega} I_{z_1(\omega)} + A_0(t) I_y + 2\pi J I_z S_z, \quad (5)$$

where  $z_1(\omega)$  is the unit vector along the direction  $\omega \hat{z} + w_0 \hat{x}$  and define  $\theta_1(\omega) = \tan^{-1}(\frac{w_0}{\omega})$ , the spread of frequencies  $\tilde{\omega} \in [w_0, \sqrt{w_0^2 + c_0^2}]$ . Note,  $z_1(\omega)$  is different for each  $\omega$ . Now, by choosing  $v_1$  in the first transformation as exactly the center of this spread and transforming into a frame rotating around  $I_{z_1(\omega)}$ , with frequency  $v_1$ , we get the Hamiltonian

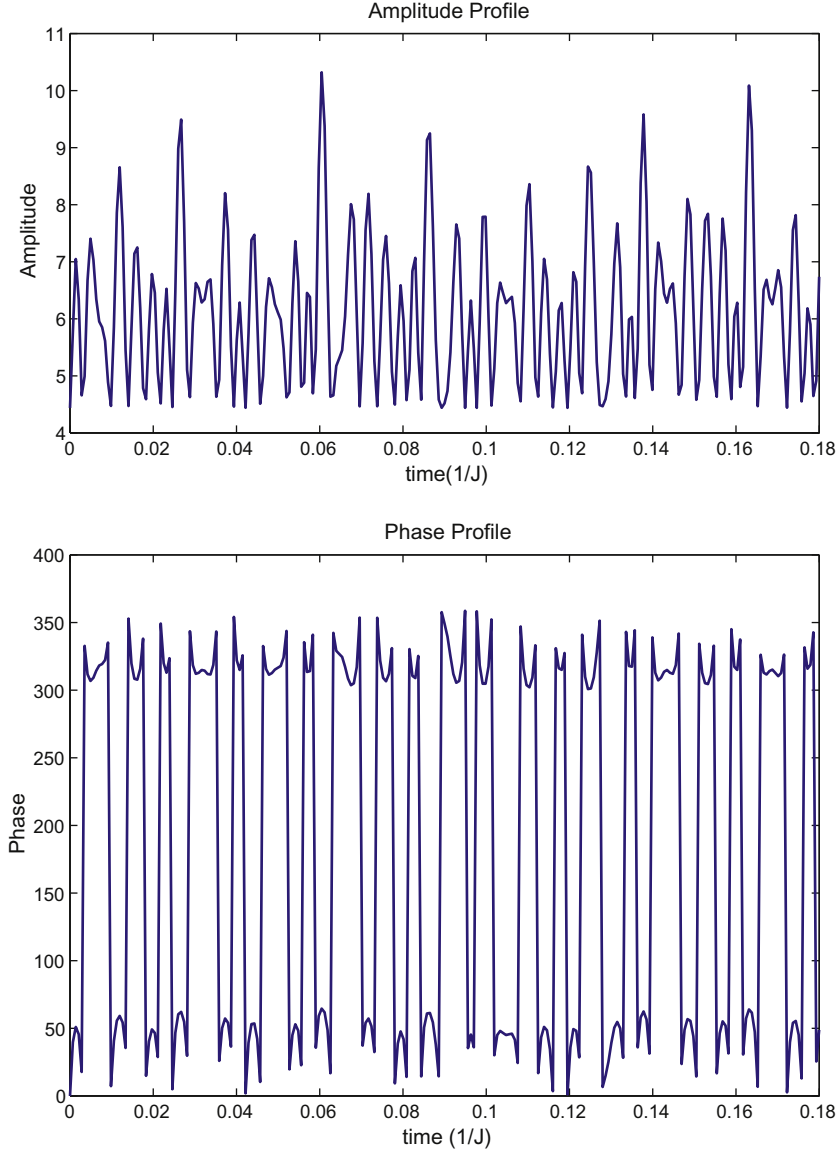
$$H_1 = \underbrace{(\tilde{\omega} - v_1)}_{f_1(\omega)} I_{z_1(\omega)} + 2\pi J \cos \theta_1(\omega) I_{z_1(\omega)} S_z + \underbrace{\bar{w}_1 I_{x_1} + A_1(t) I_y}_{H'_1} + H''_1(t) + H'''_1(t), \quad (6)$$

where  $H'_1$  is the demodulated part of the rf Hamiltonian  $H'_0$ , in the interaction frame of  $I_{z_1(\omega)}$ . The form of  $H'_1$  resembles  $H'_0$ .  $H''_1(t)$  and  $H'''_1(t)$  are the fast oscillating parts of the rf and coupling Hamiltonian that we ideally want to average out and we neglect these terms for now.  $H''_1(t)$  is simply the part of the coupling perpendicular to the effective field direction  $I_{z_1(\omega)}$ , that oscillates with frequency  $v_1$ . The new frequency  $f_1(\omega) \in [-c_1, c_1]$ , where  $c_1 < c_0$ .  $A_1$ ,  $H''_1$  and  $H'''_1$  are written in their general form below. The system obtained after first coordinate transformation has the desired feature that the ratio of chemical shift spread to static component of the rf field,  $\alpha_1 = \frac{c_1}{w_1}$  is reduced over  $\alpha_0 = \frac{c_0}{w_0}$  for the original system. For example, if  $c_0 = w_0$ , and  $w_1 = w_0$ , then  $\frac{c_1}{w_1} = \sqrt{2} - 1$ .

We can now iterate the above construction. We go into successive rotating frames around axis  $I_{z_k(\omega)}$  with frequency  $v_k$ . Unit vectors  $(x_k(\omega), y_k(\omega), z_k(\omega))$  define the  $k$ th frame, where we suppress the argument  $\omega$  subsequently.  $f_k(\omega)$  is the chemical shift in the  $k$ th rotating frame, starting with  $\omega$  in  $H_0$ .  $c_k$  represents the limit of the chemical shifts in the  $k$ th coordinate frame and  $\bar{w}_k = 2^{-k} w_k$  is the strength of the rf field along the direction  $x_k$ .



**Fig. 1.** The top figure shows how the coherence  $S_x(t)$  evolves as a function of time in units of  $J^{-1}$  as  $N$  the number of modulations in the rf field keeping  $A_{eff}$  constant. The simulations correspond to  $J = 140$  Hz,  $A_{eff} = 6.27$  kHz and  $\frac{c_0}{2\pi} = 27.5$  kHz corresponding to  $\frac{c_0}{2\pi A_{eff}} = 4.39$  and the chemical shift value  $\frac{\omega_0}{2\pi A_{eff}} = 1$ . The bottom figure shows the decoupling efficiency  $\eta = \frac{1}{T} \int_0^T \langle S_x(t) \rangle dt$ , as function of  $\omega_0$ . Here  $T = \frac{1}{J} = 85.7$  ms.



**Fig. 2.** The top figure shows the amplitude (in kHz) of the rf field as a function of time in units of  $J^{-1}$ , corresponding to  $N = 8$  modulations, with  $\frac{\omega_0}{2\pi} = 4.44$  kHz. This corresponds to  $A_{\text{eff}} = 6.27$  kHz. The bottom panel shows the phase (in degrees) as function of time. For  $J = 140$  Hz, the time axis corresponds to  $\frac{1}{J} = 7.1$  ms.

$$H_k = f_k(\omega) I_{z_k(\omega)} + \frac{W_k}{2^k} I_{x_k} + 2\pi J_k I_{z_k} S_z + A_k(t) I_y + H_k''(t) + H_k'''(t), \quad (7)$$

$$A_k(t) = \frac{1}{2^k} \left\{ \sum_{m=k+1}^n w_m \prod_{i=k+1}^{m-1} \cos(v_i t) \sin(v_m t) \right\} I_y, \quad A_n(t) = 0, \quad (8)$$

$$H_k''(t) = \exp(i2v_k I_{z_k} t) \left( -\frac{W_k}{2^k} I_{x_k} + A_k(t) I_y \right) \exp(-i2v_k I_{z_k} t), \quad (9)$$

$$H_k'''(t) = J_k^\perp \exp(iv_k I_{z_k} t) I_{x_k} S_z \exp(-iv_k I_{z_k} t), \quad (10)$$

$$c_{k+1} = \frac{\sqrt{c_k^2 + \bar{w}_k^2} - \bar{w}_k}{2}; \quad v_{k+1} = \frac{\sqrt{c_k^2 + \bar{w}_k^2} + \bar{w}_k}{2}; \quad \tan \theta_k(\omega) = \frac{\bar{w}_k}{f_k(\omega)}, \quad (11)$$

$$J_k = \prod_{j=1}^k \cos \theta_j(\omega) J, \quad (12)$$

$$J_k^\perp = - \prod_{j=1}^{k-1} \cos \theta_j(\omega) \sin \theta_k(\omega) J. \quad (13)$$

As discussed in the next section the iterated relations on  $c_k$  ensures that the ratio  $\alpha_k = \frac{c_k}{w_k}$  is decreasing and  $\theta_k \rightarrow \frac{\pi}{2}$ . This ensures that  $J_k \rightarrow 0$ .

### 3. Scaling

To fix ideas, we choose  $w_k = w_0$ , i.e.,  $\bar{w}_k = 2^{-k} w_0$ . From Eq. (11), we have the relation,

$$\alpha_{k+1} = \sqrt{1 + \alpha_k^2} - 1 < \alpha_k. \quad (14)$$

This ensures that  $\alpha_k$  is decreasing. We explore two limits, for  $\alpha_k \ll 1$ , we obtain,

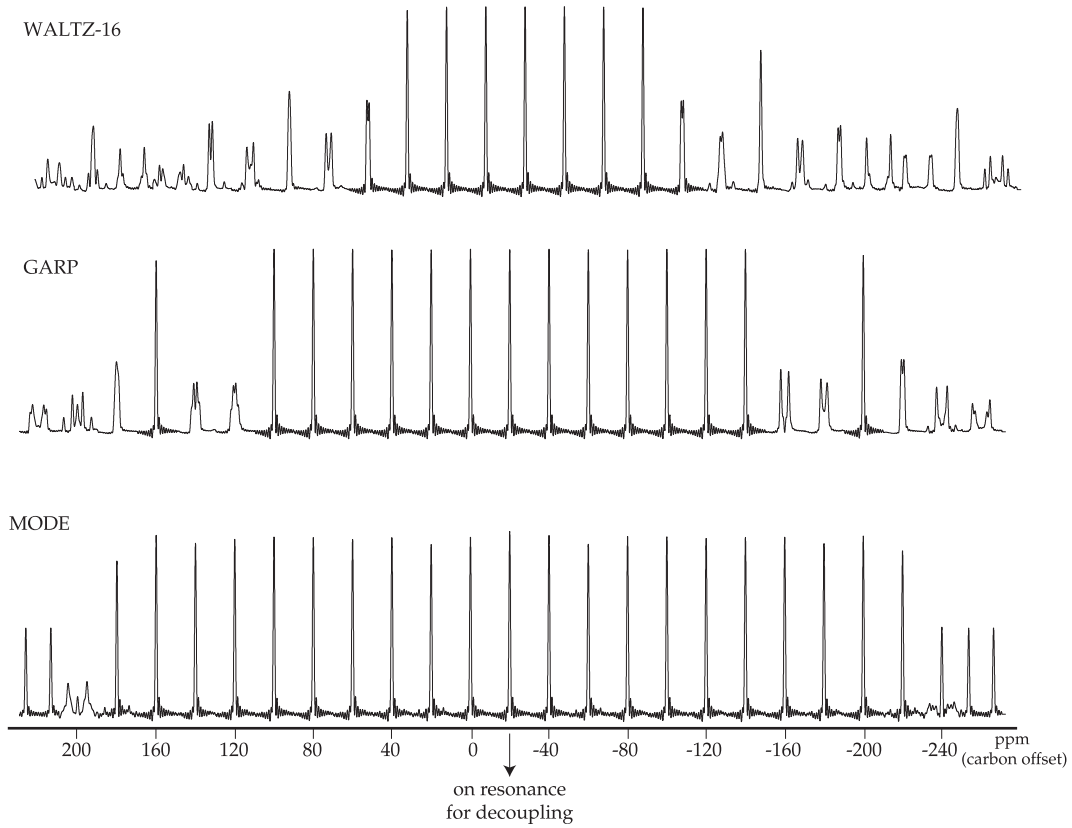
$$\frac{\alpha_{k+1}}{\alpha_k} \sim \frac{\alpha_k}{2}, \quad (15)$$

$\alpha_k \gg 1$ , we have

$$\alpha_{k+1} \sim \alpha_k - 1. \quad (16)$$

Since  $\alpha_k \rightarrow 0$ , we have  $\theta_k \rightarrow \frac{\pi}{2}$ . This ensures that  $J_k \rightarrow 0$ .

We evaluate the root mean square amplitude ( $A_{\text{eff}} = \sqrt{\frac{1}{T} \int_0^T A^2(\tau) d\tau}$ ) for this rf field as  $N \rightarrow \infty$



**Fig. 3.** The above figure show the experimental proton spectra obtained at 500 MHz for a  $^{13}\text{C}$ -iodomethane sample, dissolved in D-chloroform (D, 99.8%, Cambridge Isotope Laboratory, Inc.) The top two panels show performance of WALTZ-16 [17] and GARP [19] pulse sequences as function of the offset. The rf field for MODE corresponds to the simulation parameters with  $N = 8$  (bottom trace). The  $A_{\text{eff}}$  for all the decoupling sequence is at 6.27 kHz. The MODE sequence is implemented as small  $M = 17,136$  flip angle pulses with duration  $\Delta = 5 \mu\text{s}$ .

$$2\pi A_{\text{eff}} = w_0 \sqrt{\sum_{k=0}^N 2^{-k}} \sim \sqrt{2} w_0. \quad (17)$$

The ratio  $\frac{2v_k}{w_k}$  describes how well the oscillating component

$$\bar{w}_k \exp(j2v_k I_{z_k}) I_{x_k} \exp(-j2v_k I_{z_k}), \quad (18)$$

is averaged. This ratio

$$\frac{2v_k}{w_k} = 2 \left( \sqrt{\alpha_{k-1}^2 + 1} + 1 \right) > 4, \quad (19)$$

$$\frac{v_k}{v_{k+1}} = \frac{\bar{w}_k}{\bar{w}_{k+1}} \frac{(\sqrt{\alpha_k^2 + 1} + 1)}{(\sqrt{\alpha_{k+1}^2 + 1} + 1)} > 2. \quad (20)$$

#### 4. Non-resonant conditions

We now check whether all the oscillating terms captured by Hamiltonians  $H_k''$  and  $H_k'''$  that were neglected result in an effective coupling. If this were to happen, in the modulation frame of the rf field and chemical shifts, we will see a net coupling evolution. Therefore, we evaluate the evolution of the couplings in the modulation frame defined by the Hamiltonian  $H_m = H_{\text{CS}} + H_0$ . The evolution of this frame takes the form

$$U(\omega, t) = \exp(-iv_1 I_{z_1(\omega)} t) \dots \exp(-iv_k I_{z_k(\omega)} t) \dots \exp(-iv_N I_{z_N(\omega)} t) \Theta_N(\omega, t),$$

where number of modulations is  $N - 1$  and

$$\dot{\Theta}_N(\omega) = -i \{ f_N(\omega) I_{z(\omega)} + \tilde{H}(t) \} \Theta_N(\omega), \quad (21)$$

such that  $|f_N(\omega)| \ll c_0$ . Here,  $f_N(\omega)$  represents the chemical shift in the  $N$ th frame. Where

$$\tilde{H}(t) = \sum_{k=1}^{N-1} V_{k+1}^\dagger H_k'' V_{k+1}, \quad (22)$$

where

$$V_k(t) = \exp(-iv_k I_{z_k} t) \dots \exp(-iv_N I_{z_N} t). \quad (23)$$

In this notation,

$$U(\omega, t) = V_1(t) \Theta_N(\omega, t), \quad (24)$$

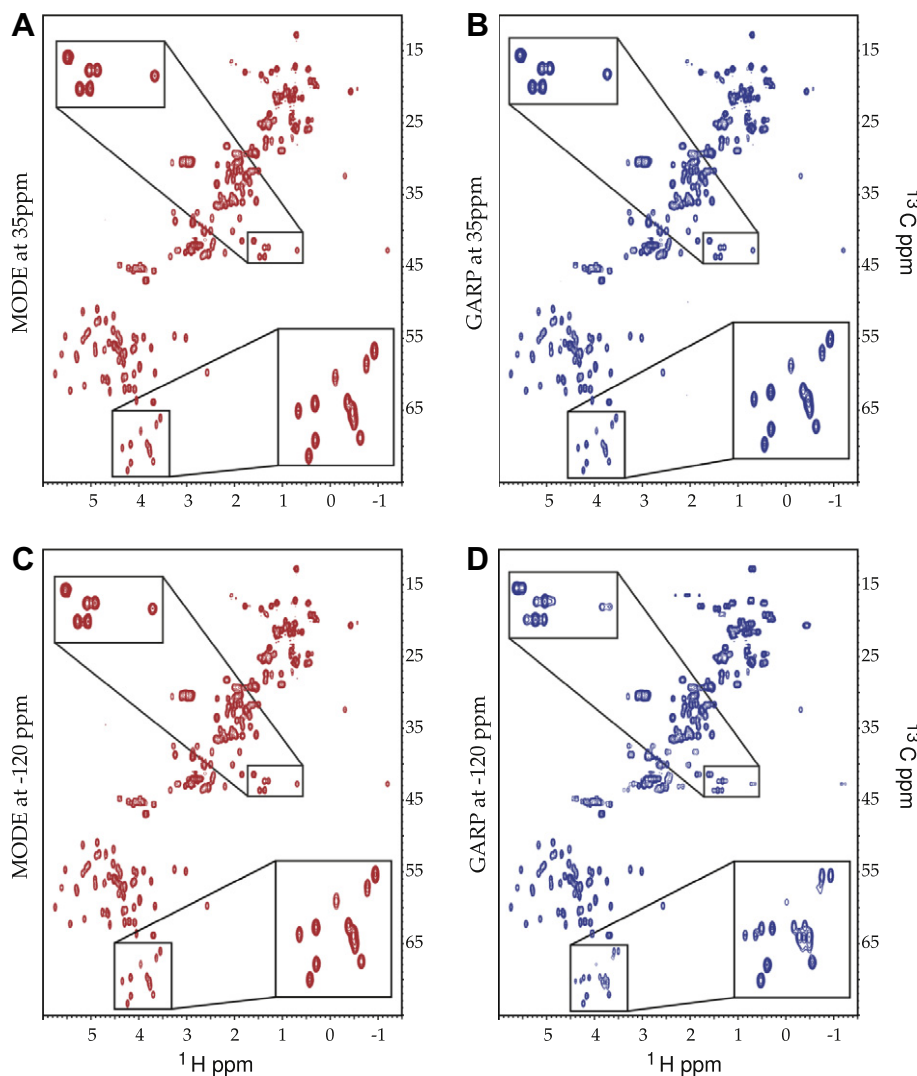
where

$$\Theta_N(\omega, t) = I - i \int_0^t \tilde{H}(\tau) d\tau - \int_0^t \int_0^\tau \tilde{H}(\tau) \tilde{H}(\sigma) d\tau d\sigma + \dots \quad (25)$$

Then in the frame of the Hamiltonian  $H_m(t)$ , the evolution takes the form

$$I - i2\pi J \int_0^t V_1' I_{z_2} S_z V_1 d\tau - 2\pi J \underbrace{\int_0^t \left[ V_1' I_{z_2} S_z V_1, \int_0^\tau \tilde{H}(\sigma) d\sigma \right] d\tau}_{J(\tau)} + \dots \quad (26)$$

where, the first integral is averaged out in the subsequent frames as  $\alpha_k \rightarrow 0$  and  $v_k > 2v_{k+1}$  in Eq. (20) prevents generation of static components from the oscillating parts. We evaluate the second integral in the series above. If  $J(\tau)$  has any static components this would reflect residual couplings in the system. We use the notation  $H_k''(f)$  to denote the discrete set of frequencies present in oscillating Hamiltonian  $H_k''(t)$ . The discrete frequencies is the set



**Fig. 4.** The above figures show the HSQC spectra on a sample of the protein GB1 obtained at 500 MHz. In the indirect dimension, the proton is decoupled by a refocusing  $\pi$  pulse. In the direct dimension, the carbon is decoupled by application of GARP and MODE decoupling sequences with mean rf amplitude of 6.27 kHz. The MODE sequence used is the same as described earlier in the section. A  $^{15}\text{N}/^{13}\text{C}$  uniformly labeled sample of protein GB1 was prepared to a final concentration of 1 mM in phosphate buffer (20 mM, pH = 6.5) with 150 mM NaCl and 1 mM EDTA. A comparison of decoupling with GARP and MODE with decoupling offsets at 35 ppm and  $-120$  ppm is shown.

$$\{\exp(-i\omega I_x)H_k''\exp(i\omega I_x)\}(f) = \{\omega \pm H_k''(f), \pm H_k''(f)\}, \quad (27)$$

$$(V_{k+1}'H_k''V_{k+1})(f) = a_k v_k + \sum_{j>k} b_j v_j, \quad a_k \in \{\pm 2\}, \quad b_j \in \{0, \pm 1, \pm 2\}$$

Similarly, we have

$$V_1' I_z S_2 V_1(f) = \sum_{k=1}^N c_k v_k,$$

where  $c_k \in \{0, \pm 1\}$ . We compute the overlap of the two set of frequencies, we find the smallest value of  $|\Delta|$  satisfying

$$\sum_{j=1}^k c_j v_j + a_k v_k + \sum_{j=k+1}^N d_j v_j = \Delta, \quad (28)$$

where  $d_j \in \{0, \pm 1, \pm 2, \pm 3\}$ . If  $|\Delta| > 0$ , we avoid a resonance condition, i.e., the second integral in Eq. (26) has no static components.

#### 4.1. Simulation and experiments

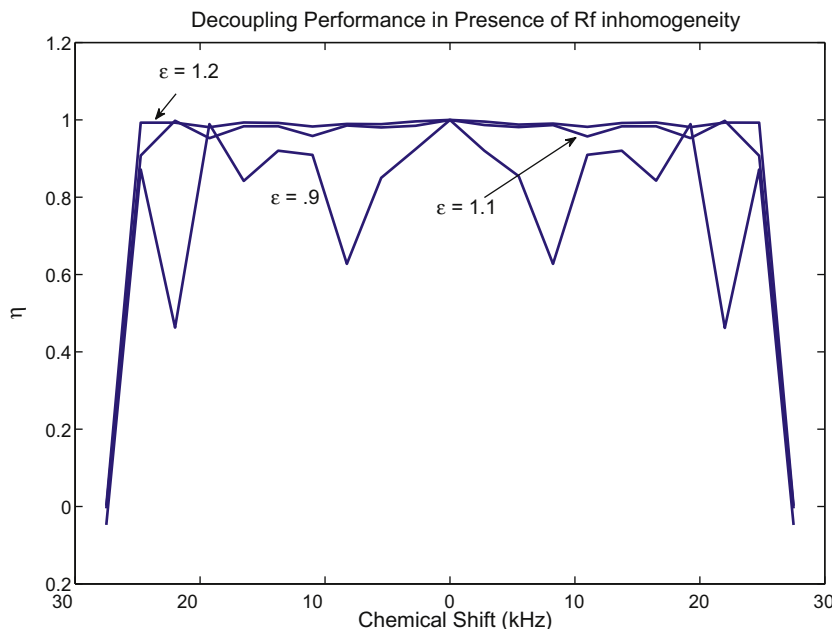
Simulations were carried for a carbon–proton ( $IS$ ) spin system, with a coupling constant  $J = 140$  Hz in Eq. (1). The carbon chemical

shift range was  $[-c_0, c_0]$ , where  $c_0(2\pi)^{-1} = 27.5$  kHz, which corresponds to a 244 ppm carbon chemical shift range at 900 MHz proton frequency. At 6.25 kHz rf power, the  $\frac{\pi}{2}$  pulse corresponds to 40  $\mu\text{s}$ . We choose  $w_k = w_0$ , and calculate  $w_0$  that corresponds to  $A_{\text{eff}} \sim 6.25$  kHz. For  $N = 8$ , this corresponds to  $\frac{w_0}{2\pi} = 4.44$  kHz (for  $n \geq 4$ ,  $\frac{w_0}{2\pi} \sim \frac{A_{\text{eff}}}{\sqrt{2}}$ ).  $w_0$  may be adjusted if the resulting rms power  $A_{\text{eff}}$ , is not exactly  $A_{\text{eff}}$ . For  $N = 8$ , the parameter  $\alpha_N = .1$  and the sequence is broadband over the given bandwidth, see Fig. 1. The resulting  $A_{\text{eff}} = 6.27$  kHz. Fig. 2 shows the amplitude and phase profile of the MODE sequence. Fig. 1 shows evolution of initial magnetization  $\langle S_x(0) \rangle = 1$  under the MODE sequence for various values of  $N$  at  $A_{\text{eff}} = 6.27 \pm .02$  kHz, for a period of  $\sim 12 J^{-1}$ . Fig. 1 also shows the decoupling efficiency as time averaged  $x$ -magnetization on spin  $S$  [59],

$$\eta = \frac{1}{T} \int_0^T \langle S_x(t) \rangle dt, \quad (29)$$

for various offsets.

The parameters of the MODE sequence for  $N = 8$ ,  $w_0 = w_k = 4.44$ ,  $c_0 = 27.5$ ,  $c_1 = 11.7$ ,  $c_2 = 4.85$ ,  $c_3 = 1.93$ ,  $c_4 = .73$ ,  $c_5 = .25$ ,  $c_6 = .07$ ,  $c_7 = .016$  and  $c_8 = .002$ , all in units of  $2\pi$  kHz. The modulation



**Fig. 5.** The above figure shows the decoupling efficiency  $\eta$ , as function of  $\omega_0$ . Here  $T \sim \frac{12}{J} \sim 85.7$  ms for rf inhomogeneity values corresponding to  $\epsilon = .9$ ,  $\epsilon = 1.1$  and  $\epsilon = 1.2$  with  $\delta = -.1$ ,  $\delta = .1$  and  $\delta = .2$  respectively. The mode sequence is designed with  $N = 8$ ,  $\frac{c_0}{w_0} = 4.39$  and  $c_0 = 27.5$  kHz.

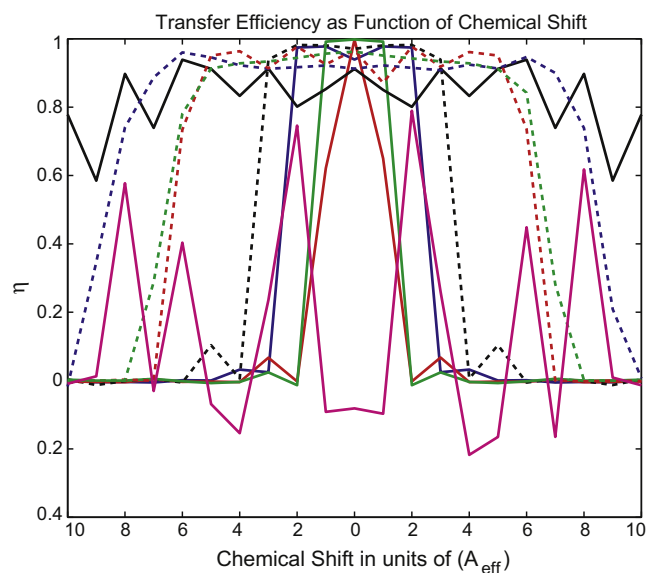
frequencies are  $v_1 = 16.15$ ,  $v_2 = 7.07$ ,  $v_3 = 3.04$ ,  $v_4 = 1.28$ ,  $v_5 = 0.53$  and  $v_6 = 0.21$ ,  $v_7 = .085$  and  $v_8 = .0364$  all in units of  $2\pi$  kHz. The ratio  $\frac{c_0}{w_0} = 6.2$ ,  $\frac{c_1}{w_1} = 5.28$ ,  $\frac{c_2}{w_2} = 4.37$ ,  $\frac{c_3}{w_3} = 3.48$ ,  $\frac{c_4}{w_4} = 2.62$ ,  $\frac{c_5}{w_5} = 1.81$ ,  $\frac{c_6}{w_6} = 1.07$ ,  $\frac{c_7}{w_7} = .46$ , and  $\frac{c_8}{w_8} = .10$ . In the last frame, the ratio of chemical shift to effective control is significantly reduced. The smallest frequency in Eq. (28) is .001 Hz, for the combination  $(v_1 - 2v_2 + v_3 - 3v_4 - 2v_5 - v_6 + v_7 - 2v_8 + 3v_9)$ , where  $v_9 = .0175$ , in the units of  $2\pi$  Hz. See Supplementary material for another design of a MODE sequence with  $N = 6$ .

The decoupling efficiency of the MODE decoupling sequence was studied experimentally using a  $^{13}\text{C}$ -iodomethane sample, dissolved in deuterated-chloroform. The experiments were carried out on a 500 MHz Bruker spectrometer at room temperature. The MODE pulse sequence can be implemented on a spectrometer, as sequence of small flip angle pulses with defined amplitude and phases. The MODE sequence of duration  $T = \frac{12}{J} = 85.7$  ms is finely discretized in  $M$  steps to capture all its modulations. The discretization used corresponds to  $\Delta \sim \frac{51}{v_1}$  and  $M = 17,136$  (a finer discretization may be used), and  $A_{\text{eff}}$  for this discretized sequence is 6.27 kHz. The closed form formula for the MODE sequence (4) is in terms of the amplitude of the  $x$  and  $y$  component of the rf-Hamiltonian, which allows the computation of the amplitude  $A_k = (2\pi)^{-1} \sqrt{A_0^2(k\Delta) + w_0^2}$  and the phase  $\tan \phi_k = \frac{A_0(k\Delta)}{w_0}$ , for the  $k$ th pulse. The MODE sequence was implemented as a table of amplitude and phases, where the amplitudes are specified as percentage of the peak amplitude.

Proton resonances were observed with carbon decoupling during acquisition. A thousand two hundred points were collected with a dwell-time of 71.4  $\mu\text{s}$  and a total acquisition time of 85.7 ms. Sixteen scans were averaged with a recycle delay of 15 s between them. The carbon resonance of iodomethane occurs at  $-20.7$  ppm. The offset on the carbon channel (decoupling offset) was varied from  $-240$  to 220 ppm to study the effect of the various decoupling sequences.

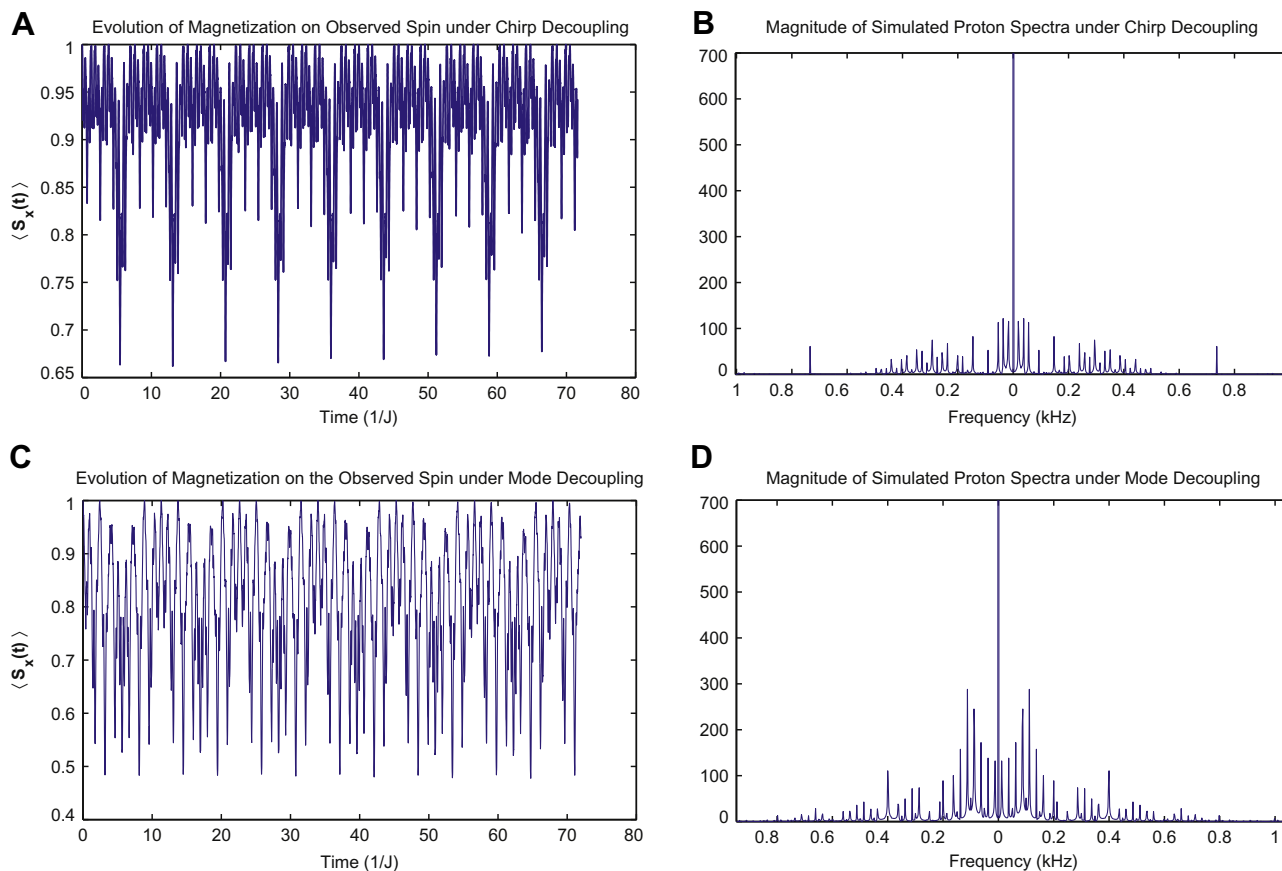
Fig. 3 shows the experimental spectra obtained on a system of methyl iodide with MODE sequence. The experimental parameters of the system are the same as in simulations. Comparison of conventional decoupling sequences with MODE shows it is more broadband for the same root mean square rf power.

Fig. 4 shows the 2D  $^{13}\text{C}$  HSQC spectra of the protein GB1, comparing GARP and MODE decoupling sequences on  $^{13}\text{C}$  during direct detection.  $^1\text{H}$ - $^{13}\text{C}$  HSQC experiments were collected with 128 complex points in the indirect ( $^{13}\text{C}$ ) dimension and 1200 points in the



**Fig. 6.** The above figure shows performance of the state of the art decoupling sequences as a function of the chemical shifts. MODE (black), SUSAN (dashed black) [60], GARP (blue) [19], WALTZ-16 (green) [17], MLEV-16 (red) [14], CHIRP (dashed blue) [30], STUD (dashed red) [26], MPPF9 (dashed green) [24], WURST (magenta) [28]. The acquisition time of  $72 J^{-1}$ , which for a coupling of 140 Hz translates to  $\sim 510$  ms. The performance parameter  $\eta$  is plotted as a function of chemical shift in units of root mean square amplitude  $A_{\text{eff}}$ . We use  $A_{\text{eff}} = 2.25$  kHz such that a total bandwidth of  $20A_{\text{eff}}$  corresponds to 45 kHz. The various pulse sequences were implemented exactly the way described in their respective references. The coupling and the acquisition time was scaled so that at the scaled rms amplitude of 2.25 kHz, all pulse sequences see a coupling of 140 Hz and acquisition time of  $\sim 510$  ms.





**Fig. 7.** Comparison of CHIRP (top) and MODE decoupling (bottom). Panel A shows simulation for time evolution of proton magnetization under CHIRP decoupling [30], for a duration of  $72J^{-1}$ , which for a coupling of 140 Hz, translates to  $\sim 510$  ms with  $A_{\text{eff}} = 2.25$  kHz. Panel C shows the same evolution under MODE decoupling. The parameters of the MODE decoupling sequences are described as above and the CHIRP decoupling sequence is as in [30]. B and D shows the Fourier transform of this time evolution. In both cases we take approx 7200 points, which corresponds to a spectral width of  $\sim 14$  kHz. The offset for the irradiated  $^{13}\text{C}$  is taken to be  $4A_{\text{eff}}$ , i.e. 9 kHz. Limits on vertical axis in B and D are chosen to magnify any modulation sidebands. The magnitude of the primary resonance is outside the scale limits.

direct ( $^1\text{H}$ ) dimension with eight scans for every indirect point with a recycle delay of 3 s. The carrier was centered at 4.7 ppm on the proton channel and 35 ppm on the carbon channel. The carrier on the carbon channel is shifted before acquisition/decoupling to study the broadband performance of decoupling sequences. With the MODE decoupling we observe H-alpha protons that are attached to the down field carbon C-alpha (50–60 ppm) to be decoupled even when the carbon carrier was shifted to  $-120$  ppm. In case of the GARP decoupling, we observe minor imperfections in decoupling (highlighted and zoomed in boxed regions) under similar conditions. These experiments also show that MODE sequences can be readily incorporated in standard NMR experiments.

Fig. 5 shows simulations of the performance of the MODE decoupling sequence as function of rf inhomogeneity, which is captured by the parameter  $\epsilon = (1 + \delta)$ , such that the actual amplitude of the rf field is  $\epsilon A$  where  $A$  is the nominal amplitude.

In the absence of inhomogeneity, for all  $\omega \in [-c_0, c_0]$ ,  $f_k(\omega) \leq f_k(c_0)$  and the ratio  $\frac{c_k}{\bar{w}_k}$  is constantly decreasing. In the presence of rf inhomogeneity, for our simulation with  $\delta = .1$  and  $\delta = .2$ , it is observed that  $\frac{f_k(c_0)}{\bar{w}_k}$  begins to increase. For  $\delta = -1$ , we observe that  $f_k(\omega) \not\leq f_k(c_0)$ . The performance of the MODE sequence in the presence of rf inhomogeneity for  $\delta > 0$  is comparable to the ideal case. For  $\delta = -1$ , offset frequencies are affected by rf inhomogeneity. The angle  $\prod_k \cos(\theta_k)$  in (12) (measuring residual coupling) is largest for the offset at  $\pm 8.25$  kHz which is close to resonance rather than at the extreme. This may be attributed to  $f_k(\omega) \not\leq f_k(c_0)$ . These simulations suggest that operating the MODE sequence

slightly above the nominal value can alleviate the performance of MODE for rf values below nominal.

Alternatively, in the presence of rf inhomogeneity, we can redefine our iterative procedure for computing the modulation frequencies  $\nu_k$ . In presence of inhomogeneity, the spread of effective shifts after the first frame transformation is from  $\left[ (1 - \delta)w_0, \sqrt{(1 + \delta)^2 w_0^2 + c_0^2} \right]$ , the left limit corresponds to zero offset and smallest rf amplitude. The right limit corresponds to largest rf amplitude and chemical shift. We choose  $\nu_1$  as center of this spread. Following this we obtain that

$$\begin{aligned} \nu_{k+1} &= \frac{\sqrt{c_k^2 + \bar{w}_k^2(1 + \delta)^2} + \bar{w}_k(1 - \delta)}{2}; \\ c_{k+1} &= \frac{\sqrt{c_k^2 + \bar{w}_k^2(1 + \delta)^2} - \bar{w}_k(1 - \delta)}{2}. \end{aligned} \quad (30)$$

This gives

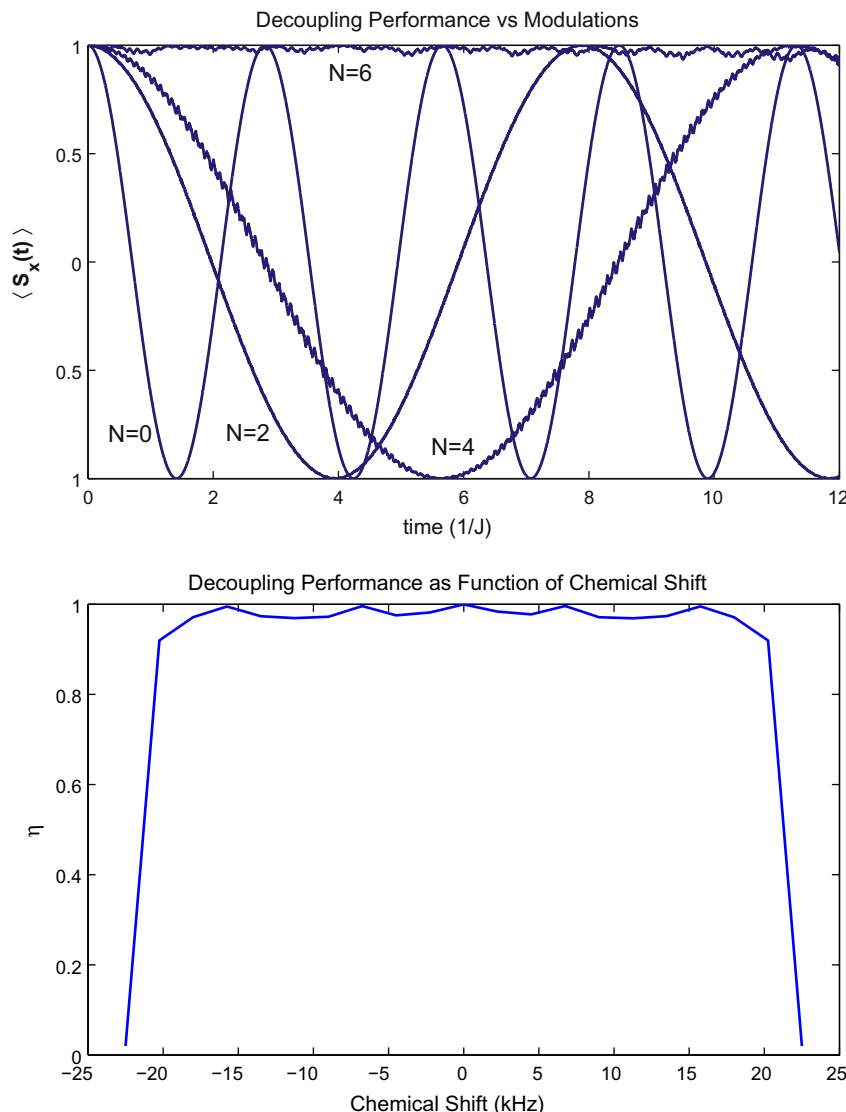
$$\alpha_{k+1} = \sqrt{\alpha_k^2 + (1 + \delta)^2} - (1 - \delta). \quad (31)$$

Note

$$\sqrt{\alpha_k^2 + (1 + \delta)^2} - (1 - \delta) \leq \alpha_k, \quad (32)$$

with equality when

$$\alpha_k = \alpha_p = \frac{2\delta}{1 - \delta}.$$



**Fig. 8.** The top figure shows how the coherence  $S_x(t)$  evolves as a function of time in units of  $J^{-1}$  as  $N$  the number of modulations in the rf field keeping  $A_{eff}$  constant. The simulations correspond to  $J = 140$  Hz,  $A_{eff} = 6.25$  kHz and  $\frac{\omega_0}{2\pi} = 22.5$  kHz corresponding to  $\frac{c_0}{2\pi A_{eff}} = 3.6$  and the chemical shift value  $\frac{\omega_0}{2\pi A_{eff}} = 1$ . The bottom figure shows the decoupling efficiency  $\eta = \frac{1}{T} \int_0^T \langle S_x(t) \rangle dt$ , as function of  $\omega_0$ . Here  $T = \frac{1}{J} = 85.7$  ms.

The  $\alpha_k$  values decrease until they reach  $\alpha_p$ . For  $\delta = .1$ , we have  $\alpha_p \sim .22$ , which is good enough as the ratio of chemical shift to rf strength is small. More generally, relaxing the constraint,  $w_{k+1} = w_k$ , we can move this fixed point close to zero.

## 5. Discussion and conclusion

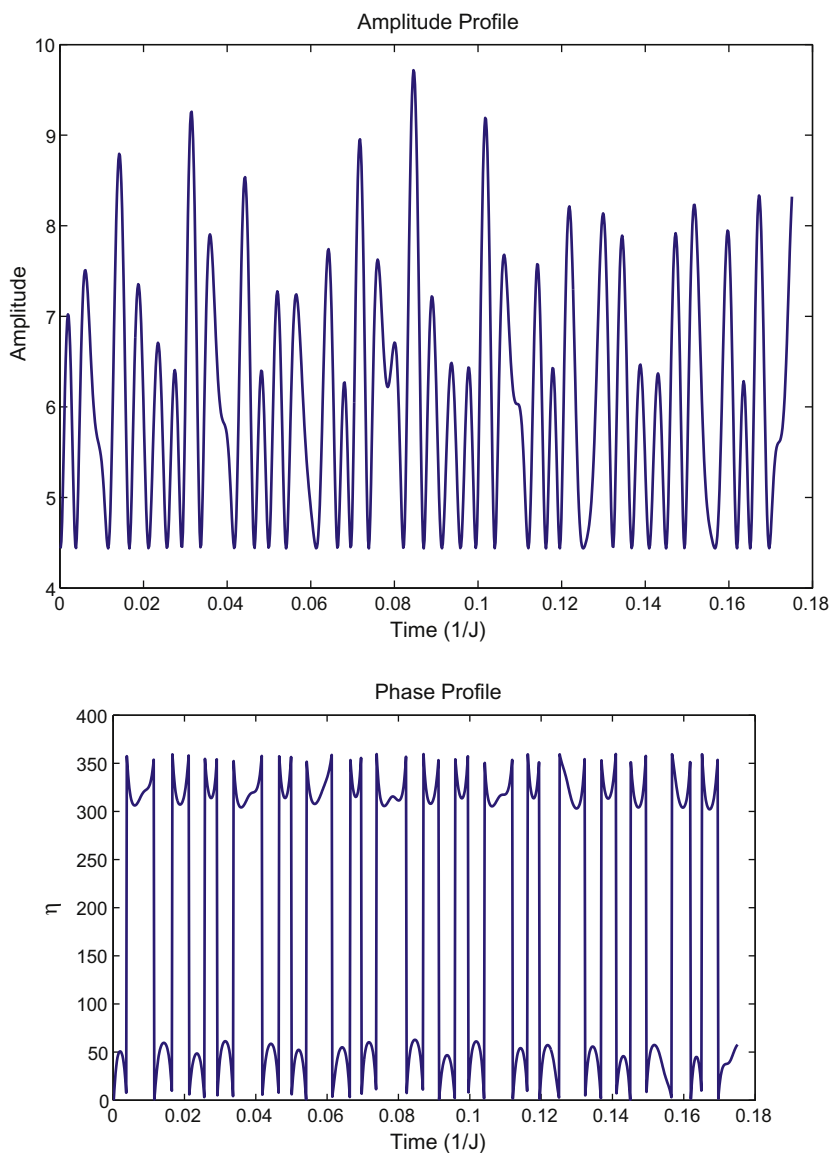
The Fig. 6 simulates performance of the state of the art decoupling pulse sequences for an acquisition time of  $72 J^{-1}$ , which for a coupling of 140 Hz translates to 510 ms. The performance parameter  $\eta$  is plotted as a function of chemical shift in units of rms amplitude  $A_{eff}$ . We use  $A_{eff} = 2.25$  kHz, such that a total bandwidth of  $20 A_{eff}$ , corresponds to 45 kHz, the full 200 ppm carbon spectra on a 900 MHz spectrometer. The various pulse sequences were implemented exactly the way described in their respective references. The coupling and the acquisition time was scaled so that at the scaled rms amplitude of 2.25 kHz, all pulse sequences see a coupling of 140 Hz and acquisition time of  $\sim 510$  ms. The parameters for the MODE pulse sequence are  $N = 9$ ,  $w_0 = 0.2250$ ,  $w_1 = 0.4460$ ,  $w_2 = 0.8762$ ,  $w_3 = 1.6916$ ,  $w_4 = 3.1565$ ,  $w_5 = 5.5235$ ,  $w_6 = 8.6296$ ,  $w_7 = 11.3585$ ,  $w_8 = 12.4943$ ,  $w_9 = 6.3103$ , all in units of  $2\pi$  kHz. This corresponds to a

rms amplitude of  $A_{eff} = 2.25$  kHz. The chemical shifts in the various frames is  $c_0 = 25.0150$ ,  $c_1 = 12.3955$ ,  $c_2 = 6.0873$ ,  $c_3 = 2.9361$ ,  $c_4 = 1.3661$ ,  $c_5 = 0.5915$ ,  $c_6 = .2218$ ,  $c_7 = .0624$ ,  $c_8 = .0099$ ,  $c_9 = .0005$ , all in units of  $2\pi$  kHz. The ratio of the chemical shifts to static rf control in each frame is  $\frac{c_0}{w_0} = 111.1779$ ,  $\frac{c_1}{w_1} = 55.5867$ ,  $\frac{c_2}{w_2} = 27.7889$ ,  $\frac{c_3}{w_3} = 13.8854$ ,  $\frac{c_4}{w_4} = 6.9248$ ,  $\frac{c_5}{w_5} = 3.4268$ ,  $\frac{c_6}{w_6} = 1.6448$ ,  $\frac{c_7}{w_7} = 0.7027$ ,  $\frac{c_8}{w_8} = 0.2020$ ,  $\frac{c_9}{w_9} = 0.0400$ . In the last frame, the ratio of chemical shift to effective control is significantly reduced. The various modulation frequencies are  $\nu_1 = 12.6205$ ,  $\nu_2 = 6.3103$ ,  $\nu_3 = 3.1551$ ,  $\nu_4 = 1.5776$ ,  $\nu_5 = 0.7888$ ,  $\nu_6 = 0.3944$ ,  $\nu_7 = .1972$ ,  $\nu_8 = 0.0986$ ,  $\nu_9 = .0493$  in units of  $2\pi$  kHz. Note, since we are using low rf power, the starting value  $\alpha_0$  is large. Eq. (16) entails that in this limit  $\alpha_{k+1} \sim \alpha_k - 1$ . This will require large number of modulations  $N$  to reduce  $\alpha_N$  below a desired threshold. To achieve a faster decrease in the value of  $\alpha_k$ , we start with an acceptable value of  $\alpha_N$  in the final frame which is chosen to be  $\alpha_N = .04$  and  $\nu_N = 4\bar{w}_N$ . We solve the forward equations

$$c_{k+1} = \frac{\sqrt{c_k^2 + \bar{w}_k^2} - \bar{w}_k}{2}; \quad \nu_{k+1} = \frac{\sqrt{c_k^2 + \bar{w}_k^2} + \bar{w}_k}{2}. \quad (33)$$

backwards, by choosing  $\nu_k = 2\nu_{k+1}$ , and





**Fig. 9.** The top figure shows the amplitude (in kHz) of the rf field as a function of time in units of  $J^{-1}$ , corresponding to  $N = 6$  modulations, with  $\frac{w_0}{2\pi} = 4.44$  kHz. This corresponds to  $A_{\text{eff}} = 6.25$  kHz. The bottom panel shows the phase (in degrees) as function of time. For  $J = 140$  Hz, the time axis corresponds to  $\frac{1}{J} = 7.1$  ms.

$$c_k = 2\sqrt{c_{k+1}v_{k+1}}; \quad \bar{w}_k = v_{k+1} - c_{k+1}. \quad (34)$$

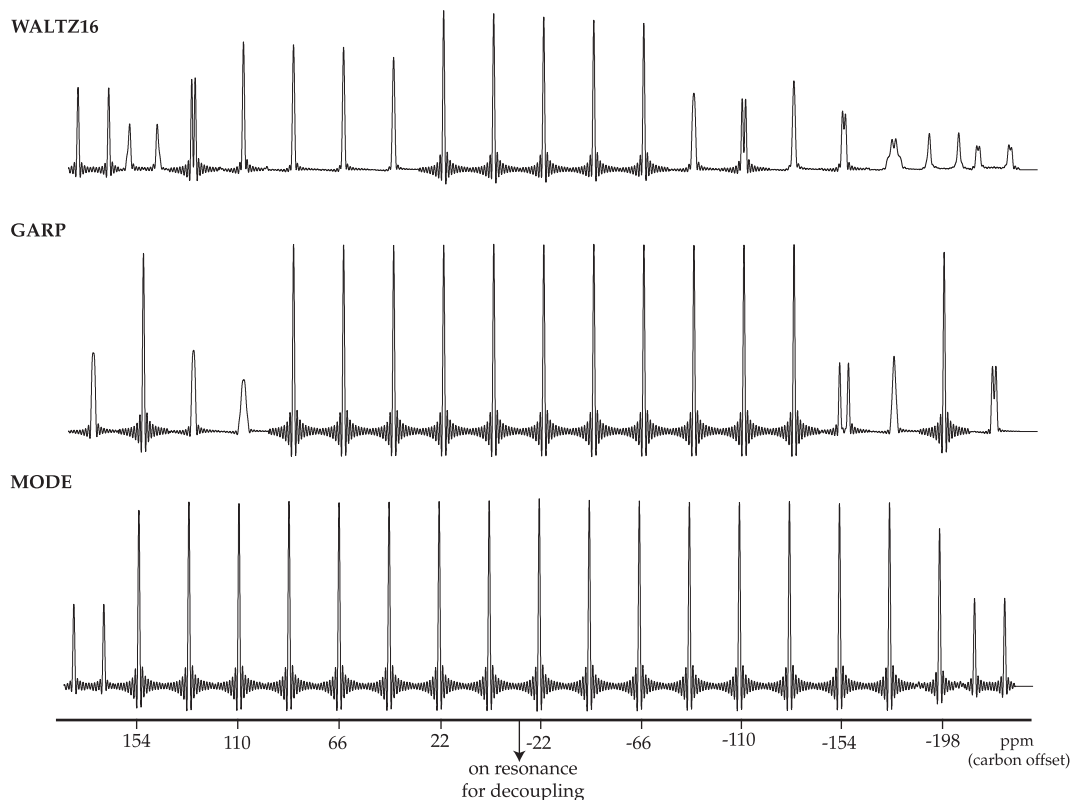
We stop, when  $\alpha_k > \alpha_0$ . All the parameters can now be renormalized to achieve desired  $A_{\text{eff}}$ . It can be shown that growth of  $\alpha_k$ , working backwards as in Eq. (34) is exponential, giving relatively fewer number of modulations for which  $\alpha_0$  is the desired value at low powers.

Adiabatic sequences are relatively broadband at low powers as seen in Fig. 6. In Fig. 7, MODE decoupling sequence is compared with an adiabatic decoupling sequence like CHIRP, in terms of the modulation sidebands.

In this paper, we introduced the multiple rotating field technique as a means to design broadband heteronuclear decoupling sequences in solution NMR. It is important to point out that the MODE sequences for the case when  $N = 1$ , simply reduce to the presence of a second oscillating field, as in [10] and the well known TPPM decoupling pulse sequence [1]. Also see [2–7]. The main contribution of this paper lies in realizing the importance of the ratio

$\alpha_k$  and showing that by adding extra modulations in the rf field and successively transforming into rotating frames, we can significantly reduce the ratio  $\alpha_k$  and improve the decoupling performance. The technique is also expected to find use in design of broadband excitation, inversion and mixing sequences, and design of broadband experiments in solid state NMR, where spread of chemical shifts can simply be removed by transforming into a suitable  $K$  frame.

The parameters of the MODE sequence for  $N = 6$ , are  $J = 140$  Hz,  $A_{\text{eff}} = 6.25$  kHz,  $w_0 = w_k = 4.44$ ,  $c_0 = 22.5$ ,  $c_1 = 9.25$ ,  $c_2 = 3.65$ ,  $c_3 = 1.35$ ,  $c_4 = .45$ ,  $c_5 = .13$ ,  $c_6 = .02$ , all in units of  $2\pi$  kHz. The modulation frequencies are  $v_1 = 13.68$ ,  $v_2 = 5.86$ ,  $v_3 = 2.46$ ,  $v_4 = 1.01$ ,  $v_5 = 0.40$  and  $v_6 = 0.1633$ , all in units of  $2\pi$  kHz. The ratio  $\frac{c_0}{w_0} = 5.07$ ,  $\frac{c_1}{w_1} = 4.17$ ,  $\frac{c_2}{w_2} = 3.29$ ,  $\frac{c_3}{w_3} = 2.44$ ,  $\frac{c_4}{w_4} = 1.63$ ,  $\frac{c_5}{w_5} = .914$ ,  $\frac{c_6}{w_6} = .355$ . The total time  $T = 85.7$  ms. The smallest frequency  $|\Delta|$  in Eq. (28) is .34 Hz, for the combination  $(v_1 - 3v_2 + 3v_4 + 3v_5 - 2v_6)$ . See Figs. 8–10 for simulation and experimental results with MODE sequence with  $N = 6$ .



**Fig. 10.** The above figure show the experimental proton spectra obtained at 500 MHz for a  $^{13}\text{C}$ -iodomethane sample, dissolved in D-chloroform (D, 99.8%, Cambridge Isotope Laboratory, Inc.). 16 scans were averaged. The top two panels show performance of WALTZ-16 [17] and GARP [19] pulse sequences as function of the offset. The rf field corresponds to the simulation parameters with  $N = 6$ . The  $A_{\text{eff}}$  for the sequences is at  $\sim 6.25$  kHz (bottom trace). The MODE sequence is implemented as small  $M = 68544$  flip angle pulses with duration  $\Delta = 1.3 \mu\text{s}$  corresponding to  $\Delta = \frac{1}{90^\circ}$ .

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