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# Matrix-assisted diffusion-ordered NMR spectroscopy with an invisible matrix: a vanishing surfactant†

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The addition of co-solutes to aid the separation of signals from molecules of similar size in DOSY experiments – matrix-assisted DOSY – is a potentially powerful technique for mixture analysis. The additional signals introduced by a co-solute can however greatly complicate analysis. By suitable choice of sample conditions, the NMR peaks of a surfactant matrix can be suppressed, allowing the clear resolution of molecular species according to chemistry and structure, without extraneous interference.

## 1 Introduction

DOSY (diffusion-ordered spectroscopy) is an effective way of analysing mixtures by NMR. Separation of the signals of mixture components is typically achieved when the components are significantly different in size, and therefore diffuse at measurably different rates. However, many mixtures of interest contain molecules that are very similar in size, *e.g.* isomers. The possibility of deliberately adding a co-solute to a mixture, changing the matrix within which the different species diffuse, in order to perturb diffusion differentially was quickly recognised as a useful tool in DOSY.<sup>1–4</sup> A wide range of methods, under names such as matrix-assisted DOSY (MAD), micelle-assisted DOSY, and chromatographic DOSY, have been used to aid mixture resolution by DOSY. These include adding chromatographic stationary phases,<sup>5</sup> polymers,<sup>6</sup> cyclodextrins,<sup>7</sup> lanthanide shift reagents<sup>8</sup> and, most commonly, surfactants.<sup>9–15</sup>

The role of the matrix in these experiments is analogous to that of the stationary phase in chromatography. There is the same potential for using different matrices, with different functionalities and properties, to manipulate systematically the diffusion coefficients of species in a sample. The choice of matrix is limited by some key considerations: it must have a suitable solubility; if it is a surfactant, it must have a Krafft temperature and critical micelle concentration (cmc) that allow a suitable concentration of micelles to form under the conditions required; and its proton spectrum should not overlap with signals of interest in the mixture. This last condition arises from

the difficulty of separating individual contributions to the diffusional decays of overlapping signals,<sup>16,17</sup> and the fact that the co-solute will often have much stronger signals than the analytes. A matrix with only a few signals, preferably far from the spectral region of interest, is a good choice where possible, but most useful matrices have a significant <sup>1</sup>H footprint. One exception is polydimethylsiloxane, with a single peak close to 0 ppm, which has been shown to separate signals in a number of mixtures, including some based on the Suzuki reaction.<sup>18</sup> Polyethylene glycol, with a single peak at around 3.5 ppm, has been shown to resolve mixtures of compounds including natural products such as  $\beta$ -estradiol and testosterone.<sup>19</sup> Deuterated<sup>9,20</sup> and fluorinated<sup>14</sup> surfactants are <sup>1</sup>H-NMR invisible; both are commercially available and have been used successfully in MAD experiments.

We show here that the same advantage of invisibility can be conferred on a common cationic surfactant, hexadecyltrimethylammonium bromide (also known as cetyltrimethylammonium bromide, CTAB), by using solution conditions that cause it to form very large structures with fast  $T_2$  relaxation. Using a DOSY pulse sequence that incorporates significant transverse relaxation weighting then allows the matrix signals to be filtered out. CTAB has long been identified as a potential matrix for MAD experiments<sup>3</sup> and its use has recently been revisited.<sup>21</sup> In addition to spherical micelles, CTAB can form a range of structures in solution, from rod shapes<sup>22</sup> to extended networks of large worm-like micelles.<sup>23</sup> Various experimental conditions are known to promote the formation of these extended structures; for example, suitable choice of counter-ions can facilitate the formation of larger species, by screening the head group charge.<sup>24</sup> Both hydrotropic species, such as salicylate ions<sup>25</sup> and catechol,<sup>26</sup> and high ionic strength solutions<sup>22,27</sup> have been shown to induce a transition from spherical to rod-like and worm-like micelles in cationic surfactants. Like surfactants, hydrotropes contain hydrophilic

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and hydrophobic parts, but do not spontaneously form micelles on their own accord. They facilitate the formation of micelles by other surfactants and, in the case of CTAB, the hydroxyl groups screen the cationic head-groups.

The formation of very large species often causes problems in NMR experiments, because the resultant slow molecular motion results in rapid  $T_2$  relaxation and hence broad spectral lines, but here it is used to simplify the acquired spectra. A new pulse sequence, PROJECT-Oneshot, is used to generate diffusion-weighted spectra in which the surfactant signals have been rendered invisible by  $T_2$  weighting. The 'invisible matrix' approach demonstrated here produces DOSY spectra in which the CTAB co-solute disperses the analyte signals without itself contributing any signals to the spectrum. This greatly expands the range of mixture analysis problems amenable to micellar matrix-assisted DOSY.

## 2 Experimental

### Materials and reagents

All chemicals used were commercially available and used without further purification. The TSP reference solution was contained in a capillary insert to avoid interactions with the micelles. All experiments were carried out with temperature regulation at 28 °C, to ensure that the samples were above the normal Krafft temperature of CTAB.

### NMR experiments

NMR measurements for the results presented in the paper were carried out, non-spinning, on a 400 MHz Varian INOVA spectrometer using a 5 mm indirect detection probe equipped with a z gradient coil producing a nominal maximum gradient of 30 G cm<sup>-1</sup>. Additional NMR experiments, reported in the ESI,<sup>†</sup> were carried out on a 300 MHz Bruker AVANCE spectrometer using a 5 mm indirect detection probe equipped with a z gradient coil producing a nominal maximum gradient of 53 G cm<sup>-1</sup>.

DOSY data were acquired using the Oneshot method<sup>28</sup> and the new PROJECT-Oneshot pulse sequence of Fig. 1, in which a PROJECT sequence<sup>29</sup> is used to add  $T_2$  weighting to Oneshot while avoiding J modulation. Both sequences used a total diffusion-encoding pulse duration,  $\delta$ , of 2.5 ms, a diffusion delay,  $\Delta$ , of 0.1 s, and 12 nominal gradient amplitudes ranging from 3.0 to 27.6 G cm<sup>-1</sup> in equal steps in gradient squared. The echo time,  $4\tau$ , in the PROJECT sequence was 4 ms, and 25 cycles were used to give 100 ms of  $T_2$  weighting, sufficient to remove all surfactant signals.

DOSY spectra were constructed in the DOSY Toolbox<sup>30</sup> by fitting to a modified Stejskal–Tanner equation, parametrized to take into account the effects of pulsed field gradient non-uniformity.<sup>2</sup>

## 3 Results

Fig. 2 compares Oneshot (Fig. 2a–c) and PROJECT-Oneshot (Fig. 2d) DOSY spectra of a 10 mM mixture of 2-methylpropan-1-ol (isobutanol) and butan-2-ol (*sec*-butanol) in (Fig. 2a) D<sub>2</sub>O, (Fig. 2b) a 150 mM solution of CTAB in D<sub>2</sub>O, and (Fig. 2c and d) a 150 mM solution of CTAB in D<sub>2</sub>O saturated with NaCl. The NaCl causes the CTAB to form long worm-like micelles that produce an entangled, highly viscous gel that restricts the CTAB motion sufficiently to reduce the  $T_2$ s of the CTAB protons to less than 10 ms. The formation of a gel complicates sample handling, but is very effective at suppressing convection.<sup>31,32</sup> It is possible to make short  $T_2$  CTAB gels with lower salt concentrations if a hydrotrope such as catechol is included, but some hydrotrope signal survives the  $T_2$  filtration so such mixtures are less useful. The effectiveness of the filtration and the use of matrices containing hydrotropes are illustrated in more detail in the ESI.<sup>†</sup>

In simple D<sub>2</sub>O solution the butanol signals are, as expected, not resolved in the diffusion dimension of a standard DOSY spectrum (Fig. 2a). In a micellar solution of CTAB in D<sub>2</sub>O (Fig. 2b), almost all the butanol signals are swamped by the much stronger CTAB signals. The addition of NaCl (Fig. 2c) greatly broadens the CTAB peaks, making it easier to distinguish some of the butanol signals but still not allowing clean diffusion resolution. However, when  $T_2$  filtration is added to the Oneshot experiment, to give the pulse sequence of Fig. 1, the resultant experimental data contain no significant CTAB signal and the clean, well-resolved DOSY spectrum of Fig. 2d results. The peak at 3.4 ppm in Fig. 2d is most probably a low MW CTAB impurity.

The separation can be interpreted on the basis of the hydrophobicities of the species in the sample. 2-Methylpropan-1-ol ( $\log P = 0.8$ ) interacts with the core of the micelles more strongly than butan-2-ol ( $\log P = 0.68$ ) and the effective diffusion coefficients observed in Fig. 2d reflect this.

A further demonstration of the technique is shown in Fig. 3, using a sample of 4 mM nicotinic and 3 mM 4-aminobenzoic acid. The 'invisible matrix' solution contained 150 mM CTAB in a saturated solution of NaCl in 98% D<sub>2</sub>O/2% DMSO, with the DMSO added to improve the solubility of the analytes.<sup>33</sup> While the two species exhibit reduced diffusion coefficients, both as a result of interaction with the surfactant and due to obstruction effects, the small difference in diffusion coefficient observed in aqueous solution is enhanced almost 5 times by the addition of the 'invisible matrix'. This behaviour is a reflection of the relative hydrophobicities of the two species; 4-aminobenzoic acid ( $\log P = 0.68$ ) is the more hydrophobic ( $\log P$  of nicotinic acid = -2.43). In this example the use of hydrotropic species such as catechol would have introduced overlapping signals that would only have been partially suppressed by the  $T_2$ -filter.

### PROJECT-Oneshot

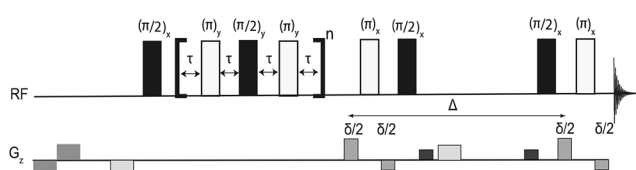


Fig. 1 PROJECT-Oneshot pulse sequence for measuring  $T_2$ -filtered spectra with diffusion weighting.



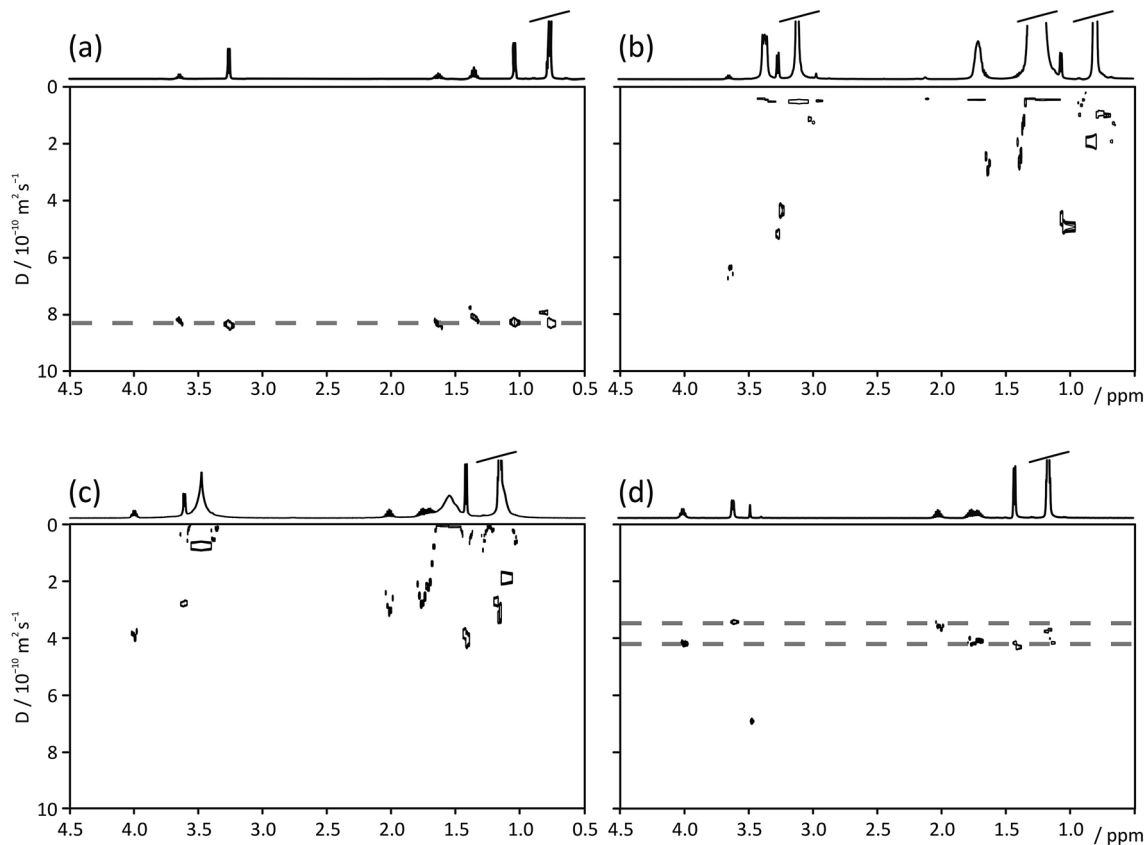


Fig. 2 DOSY spectra of a solution of 10 mM each of 2-methylpropan-1-ol (isobutanol) and butan-2-ol (sec-butanol) in (a)  $D_2O$ , (b)  $D_2O$  and 150 mM CTAB, (c) the 'invisible matrix' solution, containing 150 mM CTAB in a saturated solution of NaCl in  $D_2O$ , all acquired with Oneshot; and (d) DOSY spectrum of 2-methylpropan-1-ol and butan-2-ol in the 'invisible matrix' solution, acquired with the PROJECT-Oneshot sequence of Fig. 1. Dashed lines in (a) and (d) indicate diffusion coefficients of butanol isomers.

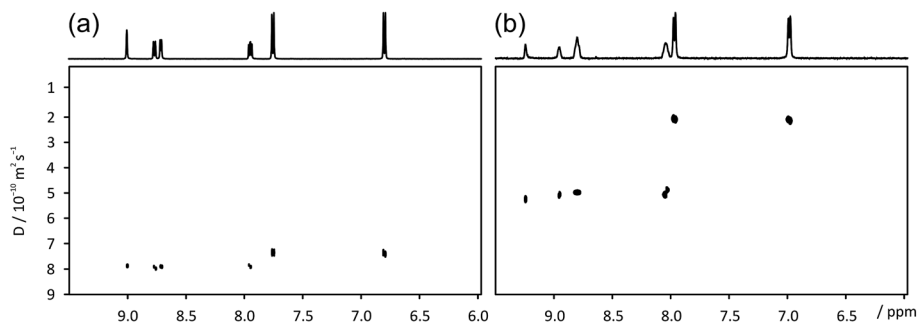


Fig. 3 DOSY spectra of a solution of 4 mM nicotinic and 3 mM 4-aminobenzoic acid in (a) the absence and (b) the presence of an invisible matrix solution consisting of 150 mM CTAB in a saturated solution of NaCl in 98%  $D_2O$ /2% DMSO.

## 4 Conclusions

This work shows the use of 'invisible matrices' – using the surfactant chemistry of CTAB to make large, worm-like micelles and removing their NMR signals by  $T_2$  weighting – as an alternative to purchasing expensive or specialist NMR-silent chemicals in matrix-assisted DOSY experiments. Very good suppression of surfactant signals can be obtained by appropriate choice of sample conditions and NMR experimental parameters, as shown in Fig. SI 3C of the ESI.† The 'invisible

matrix' solution separates both positional and structural isomers, as well as the pair of structurally similar substituted benzoic acids of Fig. 3. This approach will allow a much wider range of mixtures to be analysed, as sample signals across the entire range of proton chemical shifts can be observed.

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