

# A High-Resolution Technique for Multidimensional NMR Spectroscopy

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**Abstract**—In this paper, a scheme for estimating frequencies and damping factors of multidimensional nuclear magnetic resonance (NMR) data is presented. multidimensional NMR data can be modeled as the sum of several multidimensional damped sinusoids. The estimated frequencies and damping factors of multidimensional NMR data play important roles in determining protein structures. In this paper we present a high-resolution subspace method for estimating the parameters of NMR data. Unlike other methods, this algorithm makes full use of the rank-deficiency and Hankel properties of the prediction matrix composed of NMR data. Hence, it can estimate the signal parameters under low signal-to-noise ratio (SNR) by using a few data points. The effectiveness of the new algorithm is confirmed by computer simulations and it is tested by experimental data.

**Index Terms**—Damped sinusoids, high resolution, multidimensional NMR, parameter estimation.

## I. INTRODUCTION

MULTIDIMENSIONAL nuclear magnetic resonance (NMR) data can be modeled as the sum of multidimensional damped sinusoids. The frequencies and damping factors of damped sinusoids are crucial to determining protein structures using NMR spectroscopy. The frequency resolution of the fast Fourier transform (FFT)-based algorithms [1], [2] is limited by the short acquisition time of the NMR data and measurement noise. Hence, to improve the resolution, many model-based methods [3]–[27] have been proposed for the parameter estimation of one-dimensional (1-D) and two-dimensional (2-D) NMR data. The autoregressive modeling of NMR data is one of the most commonly used algorithms in the analysis of 1-D NMR data [8]. In particular, it has been shown that the methods based on the linear prediction technique can estimate parameters more accurately than the standard FFT methods do. This should also be true for 2-D NMR data.

Unfortunately, the existing model-based parameter estimation algorithms for 2-D NMR data are still sensitive to measurement noise, which limits their frequency resolution. The

subspace methods [19], [20], which provide high-resolution estimation in many signal processing applications, are good candidates for NMR spectroscopy to further improve the frequency resolution. The Multiple signal classification (MUSIC) algorithm [9] is one of the most effective and commonly used algorithms for 1-D stationary signals. The MUSIC algorithm can even achieve the *Cramér-Rao* lower bound under some mild conditions. However, the NMR signals consist of damped sinusoids, which is, therefore, nonstationary. Therefore, the original MUSIC algorithm cannot be directly applied to NMR data. In this paper, we present a novel parameter estimation method for NMR data based on the subspace techniques, which we will call M-D DMUSIC algorithm in order to reflect its capability to estimate the parameters (frequencies and damping factors) of multidimensional damped sinusoids. Since M-D MUSIC algorithm makes full use of the rank-deficiency and Hankel properties of the prediction matrix composed of NMR data, it can estimate the signal parameters under low signal-to-noise ratios (SNR's) by using only few data. The effectiveness of the new algorithm is demonstrated by computer examples and the experimental data obtained from National Institutes of Health (NIH).

The rest of the paper is organized as follows. After a brief description of the mathematical model of NMR signals in Section II, a damped MUSIC (DMUSIC) algorithm for estimating the parameters of 1-D NMR signals will be presented and analyzed in Section III. In Section IV, the DMUSIC algorithm will be generalized to multidimensional NMR signals, by first extending the results to 2-D NMR signals and then to general case of M-D NMR signals. A low-complexity peak-searching algorithm for searching the peaks of M-D DMUSIC spectrum is described in Section V. Computer examples are presented in Section VI and the application of the 2-D MUSIC algorithm to a set of experimental data from NIH is described in Section VII.

## II. MATHEMATICAL MODEL OF NMR SIGNALS

Before developing the DMUSIC algorithm for multidimensional NMR data, we briefly describe the mathematical model of M-D NMR data here. For a comprehensive review of NMR spectroscopy, interested readers should refer to [23]–[25].

Since the multidimensional NMR signal is the extension of the 2-D NMR signal, we first introduce the mathematical model of 2-D NMR signals. A 2-D NMR signal can be

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expressed by a continuous hypercomplex form  $X_h(t_1, t_2)$  as [23]–[25]

$$\begin{aligned}
 X_h(t_1, t_2) = & \sum_{k=1}^K a_k \left\{ \cos \left( \Omega_k^{(1)} t_1 + \theta_k^{(1)} \right) \right. \\
 & \cdot \cos \left( \Omega_k^{(2)} t_2 + \theta_k^{(2)} \right) \\
 & + i \sin \left( \Omega_k^{(1)} t_1 + \theta_k^{(1)} \right) \cos \left( \Omega_k^{(2)} t_2 + \theta_k^{(2)} \right) \\
 & + j \cos \left( \Omega_k^{(1)} t_1 + \theta_k^{(1)} \right) \sin \left( \Omega_k^{(2)} t_2 + \theta_k^{(2)} \right) \\
 & \left. + ij \sin \left( \Omega_k^{(1)} t_1 + \theta_k^{(1)} \right) \sin \left( \Omega_k^{(2)} t_2 + \theta_k^{(2)} \right) \right\} \\
 & \cdot e^{-t_1/T_k^{(1)} - t_2/T_k^{(2)}} \quad (1)
 \end{aligned}$$

where  $K$  is the model order,  $\Omega_k^{(1)}$  and  $\Omega_k^{(2)}$  denote the angular frequencies of the magnetization corresponding to  $t_1$  and  $t_2$ , respectively,  $T_k^{(1)}$  and  $T_k^{(2)}$  are the decay constants of the magnetization, and  $a_k$ 's are the amplitudes of damped sinusoids.

In our discussion, we employ the commonly used complex representation  $X_c(t_1, t_2)$ , that can be obtained by letting  $i = j$  and  $ij = -1$  in [2]

$$X_c(t_1, t_2) = \sum_{k=1}^K c_k e^{(-\gamma_k^{(1)} + j\Omega_k^{(1)})t_1 + (-\gamma_k^{(2)} + j\Omega_k^{(2)})t_2} \quad (2)$$

where  $c_k = a_k e^{j(\theta_k^{(1)} + \theta_k^{(2)})}$ , and  $\gamma_k^{(1)} = 1/T_k^{(1)}$ ,  $\gamma_k^{(2)} = 1/T_k^{(2)}$  which are called the decay rates.

If the continuous complex 2-D NMR signal is measured at uniform intervals,  $\Delta_1$  for  $t_1$  and  $\Delta_2$  for  $t_2$ , a set of 2-D NMR data  $\{x(n_1, n_2)\}$  will be obtained as

$$x(n_1, n_2) = \sum_{k=1}^K c_k e^{s_k^{(1)} n_1 + s_k^{(2)} n_2} \quad (3)$$

where  $s_k^{(l)} = -\alpha_k^{(l)} + j\omega_k^{(l)}$ , and  $\omega_k^{(l)} = \Omega_k^{(l)} \Delta_l$ ,  $\alpha_k^{(l)} = \gamma_k^{(l)} \Delta_l$  for  $l = 1, 2$  with  $\alpha_k^{(l)}$  being the damping factor. Without loss of generality, we assume that  $\mathbf{s}_k = (s_k^{(1)}, s_k^{(2)})$  for  $k = 1, 2, \dots, K$  are distinct. It should be pointed out that high SNR is required in order to estimate the signals with only distinct  $\alpha_k$ 's, but the same  $\omega_k$ 's, as demonstrated by Fig. 1(d)–(e). In presence of measurement error or noise  $w(n_1, n_2)$ , the measured NMR data can be expressed as

$$y(n_1, n_2) = x(n_1, n_2) + w(n_1, n_2) \quad (4)$$

for  $n_i = 0, 1, \dots, N_i - 1$  and  $i = 1, 2$ . In the above expression  $N_i$ 's are the acquisition times of each time domain. We will assume  $N_1 = N_2 = N$  in our discussion.

In (3) and (4), if we let  $N_2 = 1$ , then the mathematical model of 2-D NMR signal is degenerated into that of 1-D NMR signal, which can be rewritten as

$$y(n) = x(n) + w(n) \quad (5)$$

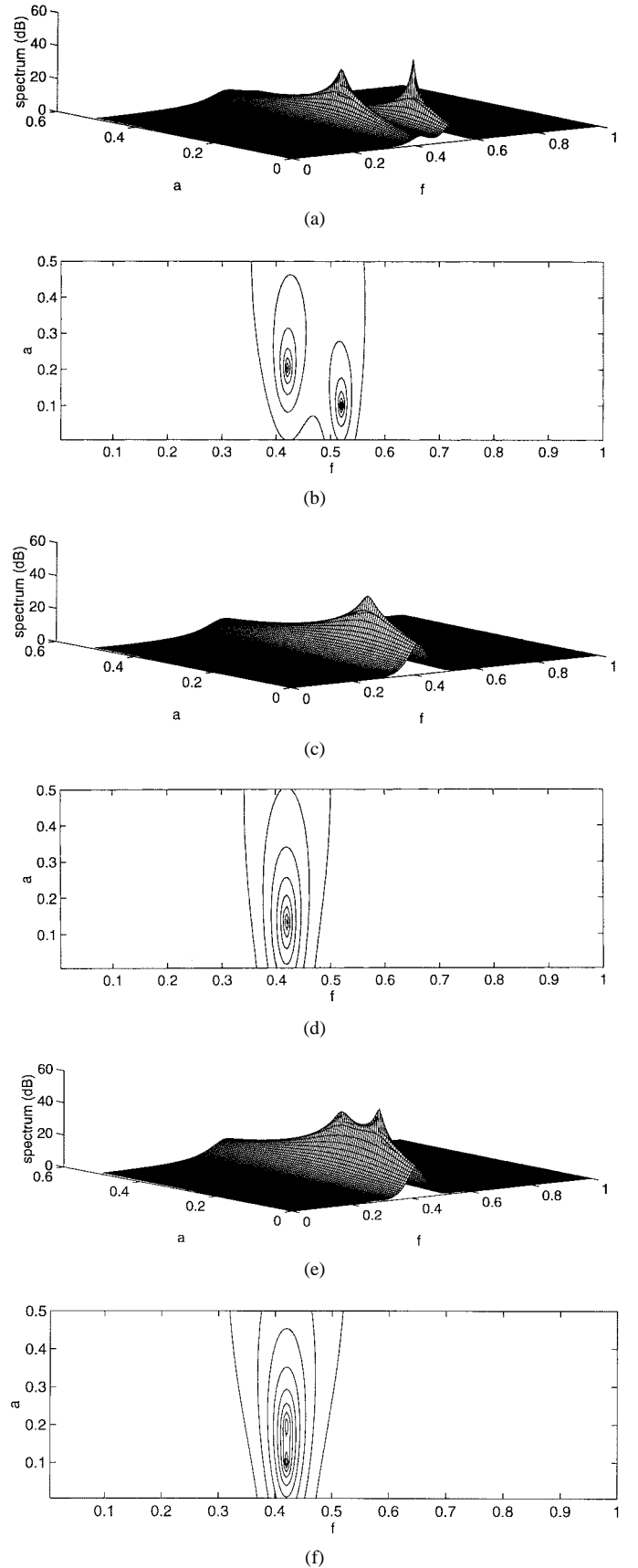


Fig. 1. Spectrum and contour of DMUSIC algorithm: (a) and (b) when  $s_1 = -0.2 + j2\pi0.42$ ,  $s_2 = -0.1 + j2\pi0.52$  and SNR = 40 dB; (c) and (d) when  $s_1 = -0.2 + j2\pi0.42$ ,  $s_2 = -0.1 + j2\pi0.42$  and SNR = 40 dB; (e) and (f) when  $s_1 = -0.2 + j2\pi0.42$ ,  $s_2 = -0.1 + j2\pi0.42$  and SNR = 60 dB.

where  $w(n)$  denotes again the measurement noise and

$$x(n) = \sum_{k=1}^K c_k e^{s_k n} \quad (6)$$

for  $n = 0, 1, \dots, N-1$ . The mathematical model of 2-D NMR signal can be easily extended to that of  $L$ -dimensional NMR signal as

$$y(\mathbf{n}) = x(\mathbf{n}) + w(\mathbf{n}) \quad (7)$$

and

$$x(\mathbf{n}) = \sum_{k=1}^K c_k e^{\mathbf{s}_k \mathbf{n}^T} \quad (8)$$

for  $\mathbf{n} \in \{0, 1, \dots, N-1\}^L$ . In the above expression,  $\mathbf{n} = [n_1, \dots, n_L]$  is a time-index vector and  $\mathbf{s}_k = [s_k^{(1)}, \dots, s_k^{(L)}]$  is a complex frequency vector. Similar to the 2-D case,  $w(\cdot)$  represents measurement noise and  $K$  denotes the model order. Normally, we have to make sure  $N > 2K$ , to be able to estimate signal parameters.

### III. 1-D DMUSIC ALGORITHM

We first present the DMUSIC algorithm for 1-D NMR signals (1-D MUSIC algorithm). To derive 1-D DMUSIC algorithm, we will set up an  $(N-J) \times J$  prediction matrix

$$\mathbf{A} = \begin{pmatrix} y(0) & y(1) & \cdots & y(J-1) \\ y(1) & y(2) & \cdots & y(J) \\ \vdots & \vdots & \vdots & \vdots \\ y(N-J-1) & y(N-J) & \cdots & y(N-1) \end{pmatrix} \quad (9)$$

where  $K \leq J \leq N-K$ . We normally choose  $J = \lceil N/2 \rceil$  to obtain the best performance [4]. The prediction matrix in the DMUSIC algorithm plays a similar role as the correlation matrix in the MUSIC algorithm [20]. From (5) and (6),  $\mathbf{A}$  can be written as (see also, [27])

$$\mathbf{A} = \sum_{k=1}^K c_k \mathbf{r}_l(s_k) \mathbf{r}_r^T(s_k) + \mathbf{W} = \mathbf{S}_l \mathbf{C} \mathbf{S}_r^T + \mathbf{W} \quad (10)$$

where  $\mathbf{r}_r(s)$  and  $\mathbf{S}_r$  are the *right signal vector* and the *right signal matrix*, respectively, defined as

$$\mathbf{r}_r(s_k) = \begin{pmatrix} 1 \\ e^{s_k} \\ \vdots \\ e^{(J-1)s_k} \end{pmatrix} \quad \mathbf{S}_r = [\mathbf{r}_r(s_1), \mathbf{r}_r(s_2), \dots, \mathbf{r}_r(s_K)] \quad (11)$$

respectively. The *left signal vector*  $\mathbf{r}_l(s)$  and the *left signal matrix*  $\mathbf{S}_l$  are similarly defined.  $\mathbf{C}$  is a  $K \times K$  diagonal matrix with  $\text{diag}(\mathbf{C}) = (c_1, c_2, \dots, c_K)$ . The *noise matrix*  $\mathbf{W}$  is given by

$$\mathbf{W} = \begin{pmatrix} w(0) & w(1) & \cdots & w(J-1) \\ w(1) & w(2) & \cdots & w(J) \\ \vdots & \vdots & \vdots & \vdots \\ w(N-J-1) & w(N-J) & \cdots & w(N-1) \end{pmatrix}. \quad (12)$$

If  $s_k$ 's are distinct, then  $\mathbf{r}_r(s_k)$  for  $k = 1, 2, \dots, K$  are linearly independent, hence  $\mathbf{S}_r$  is full column rank, and so is  $\mathbf{S}_l$ . Since the rank of  $\mathbf{C}$  is  $K$ , the rank of  $\mathbf{A}$  is also equal to  $K$  when there is no noise. Since there are many effective algorithms to estimate the model order  $K$  [28], [29], in our algorithm we will assume that  $K$  is known in advance. In [27], the authors have used a similar approach to derive their state space formulation. If there is no noise, by means of singular value decomposition,  $\mathbf{A}$  can be decomposed into the product of three matrices

$$\mathbf{A} = \mathbf{U} \mathbf{D} \mathbf{V}^H \quad (13)$$

where  $\mathbf{U}$  and  $\mathbf{V}$  are unitary matrices,  $\mathbf{D}$  is a diagonal matrix, and  $H$  denotes Hermitian, i.e., conjugate transpose, operator. Furthermore,  $\mathbf{D}$  must be of the form

$$\text{diag}(\mathbf{D}) = (\sigma_1, \sigma_2, \dots, \sigma_K, 0, \dots, 0), \quad \sigma_1 \geq \sigma_2 \geq \dots \geq \sigma_K. \quad (14)$$

From (13), we also have

$$\mathbf{A} \mathbf{V} = \mathbf{U} \mathbf{D}. \quad (15)$$

Denote  $\mathbf{v}_i$  the  $i$ -th column of  $\mathbf{V}$ ,  $\text{span}\{\mathbf{v}_1, \dots, \mathbf{v}_K\}$  is called the *signal subspace* since

$$\text{span}\{\mathbf{v}_1, \dots, \mathbf{v}_K\} = \text{span}\{\mathbf{r}_r(s_1), \dots, \mathbf{r}_r(s_K)\} \quad (16)$$

where  $\text{span}\{\cdot\}$  is referred to as the subspace that is defined by the set of all linear combinations of the vectors. From (14) and (15), we have the following *orthogonality relations*

$$\mathbf{A} \mathbf{V} \mathbf{n} = \mathbf{0}, \quad \text{or} \quad \mathbf{A} \mathbf{v}_k = \mathbf{0} \quad \text{for} \quad k = K+1, \dots, J \quad (17)$$

where  $\mathbf{V} \mathbf{n} = [\mathbf{v}_{K+1}, \dots, \mathbf{v}_L]$ . From (10), we have

$$\mathbf{S}_l \mathbf{C} \mathbf{S}_r^T \mathbf{v}_k = \mathbf{0} \quad \text{for} \quad k = K+1, \dots, J. \quad (18)$$

Since both  $\mathbf{S}_l, \mathbf{S}_r$  and  $\mathbf{C}$  are full rank,  $\mathbf{S}_r^T \mathbf{v}_k = \mathbf{0}$  for  $k = K+1, \dots, J$ , i.e.,  $\mathbf{r}_r^T(s_n) \mathbf{v}_k = 0$  for  $k = K+1, \dots, J$  and  $n = 1, 2, \dots, K$ . Hence,  $\mathbf{V}_n^T \mathbf{r}_r(s) = \mathbf{0}$  only when  $s = s_1, \dots, s_K$ . Therefore,  $s_k$  can be obtained by finding  $s$  that makes  $\|\mathbf{V}_n^T \mathbf{r}_r(s)\| = 0$ .

When noise exists, the orthogonality relations (17) no longer hold. In this case, we can search for signal vectors that are *most closely* orthogonal to the noise subspace. Hence,  $s_k$  can be obtained by finding the peak of the following DMUSIC spectrum:

$$P(s) = \frac{1}{\bar{\mathbf{r}}_r^H(s) \left( \sum_{k=K+1}^J \mathbf{v}_k^* \mathbf{v}_k^T \right) \bar{\mathbf{r}}_r(s)} \quad (19)$$

where

$$\bar{\mathbf{r}}_r = \frac{\mathbf{r}_r}{\|\mathbf{r}_r\|}. \quad (20)$$

The algorithm is summarized in Table I. The algorithm discussed above is called the damped MUSIC (DMUSIC) algorithm. Since the mathematical model of NMR data has the damped sinusoidal form, it is true that it is nonstationary in a strict mathematical sense. This is easy to prove and it

TABLE I  
DAMPED MUSIC ALGORITHM

Step 1	Forming data matrix $\mathbf{A}$ using (9)
Step 2	Finding $\mathbf{V}_n$ by making SVD to $\mathbf{A}$
Step 3	Estimating $s_k$ by finding the peaks of (19)

follows directly from standard methods for computing the autocorrelation function for a random sequence and interested readers should refer to [30] and [31]. There are two crucial differences between the DMUSIC algorithm and the MUSIC algorithm. First of all, the DMUSIC algorithm is for parameter estimation of damped sinusoidal signals which are nonstationary, as opposed to MUSIC algorithm which works for stationary signals. Since the correlation matrix is not available for nonstationary signals, in DMUSIC algorithm the prediction matrix is used instead of correlation matrix. Second, the DMUSIC algorithm searches the  $(\alpha, \omega)$  plane to estimate two parameters simultaneously.

#### IV. M-D DMUSIC ALGORITHM

The DMUSIC algorithm for 1-D NMR signals developed in the Section III can be extended to the DMUSIC algorithm for multidimensional NMR signals (M-D DMUSIC).

To obtain a DMUSIC algorithm for the 2-D NMR signals modeled in (3) and (4), we first generate an  $(N - J + 1) \times J$  matrix, (21) shown at the bottom of the page, for  $n = 1, 2, \dots, N - 1$ . Based on  $\mathbf{A}(n)$ , a  $(N - J + 1)^2 \times J^2$  matrix is formed

$$\mathbf{A} = \begin{pmatrix} \mathbf{A}(0) & \mathbf{A}(1) & \cdots & \mathbf{A}(J-1) \\ \mathbf{A}(1) & \mathbf{A}(2) & \cdots & \mathbf{A}(J) \\ \vdots & \vdots & \ddots & \vdots \\ \mathbf{A}(N-J-1) & \mathbf{A}(N-J) & \cdots & \mathbf{A}(N-1) \end{pmatrix} \quad (22)$$

where  $K \leq J \leq N - K$ . It is important to notice that the rank of matrix  $\mathbf{A}$  is  $K$ .

Using a similar procedure to the derivation of 1-D DMUSIC algorithm, we can form a 2-D DMUSIC spectrum

$$P(\mathbf{s}) = \frac{1}{\bar{\mathbf{r}}_r^H(\mathbf{s}) \left( \sum_{k=K+1}^{J^2} \mathbf{v}_k^* \mathbf{v}_k^T \right) \bar{\mathbf{r}}_r(\mathbf{s})} \quad (23)$$

where  $\mathbf{v}_k$  for  $k = K + 1, \dots, J^2$  are the right singular vectors of  $\mathbf{A}$  corresponding to the  $J^2 - K$  smallest singular values and

$$\bar{\mathbf{r}}_r(\mathbf{s}_k) = \begin{pmatrix} 1 & e^{s_k^{(2)}} & \cdots & e^{(J-1)s_k^{(2)}} & \cdots & e^{(J-1)s_k^{(1)}} \\ e^{(J-1)s_k^{(1)} + s_k^{(2)}} & \cdots & e^{(J-1)s_k^{(1)} + (J-1)s_k^{(2)}} \end{pmatrix}^T \quad (24)$$

and  $\bar{\mathbf{r}}_r = \mathbf{r}_r / \|\mathbf{r}_r\|$ . The parameter  $\mathbf{s}_k$  of 2-D NMR data can be estimated by finding  $\mathbf{s}$  to maximize the above 2-D DMUSIC spectrum.

In general, for  $L$ -dimensional NMR signal,  $(N - J + 1)^L \times J^L$  data matrix  $\mathbf{A}$  is defined as

$$\mathbf{A} = \begin{pmatrix} \mathbf{A}(0) & \mathbf{A}(1) & \cdots & \mathbf{A}(J-1) \\ \mathbf{A}(1) & \mathbf{A}(2) & \cdots & \mathbf{A}(J) \\ \vdots & \vdots & \ddots & \vdots \\ \mathbf{A}(N-J-1) & \mathbf{A}(N-J) & \cdots & \mathbf{A}(N-1) \end{pmatrix} \quad (25)$$

where indexes  $n_1, n_2, \dots, n_L$  correspond to the number of given data points in each dimension, i.e., we are using an  $n_1 \times n_2 \times \cdots \times n_L$  data matrix and (26), (27) shown at the bottom of the next page.

The *right signal vector* corresponding to the data matrix  $\mathbf{A}$  is defined as

$$\mathbf{r}(\mathbf{s}) = \begin{pmatrix} \mathbf{r}_{L-1}(\mathbf{s}) \\ e^{s_k^{(1)}} \mathbf{r}_{L-1}(\mathbf{s}) \\ \vdots \\ e^{(J-1)s_k^{(1)}} \mathbf{r}_{L-1}(\mathbf{s}) \end{pmatrix} \quad (28)$$

where

$$\mathbf{r}_l(\mathbf{s}) = \begin{pmatrix} \mathbf{r}_l(\mathbf{s}) \\ e^{s_k^{(L-l+1)}} \mathbf{r}_{l-1}(\mathbf{s}) \\ \vdots \\ e^{(J-1)s_k^{(L-l+1)}} \mathbf{r}_{l-1}(\mathbf{s}) \end{pmatrix} \quad (29)$$

for  $l = 2, 3, \dots, L - 1$ , and

$$\mathbf{r}_1(\mathbf{s}) = \begin{pmatrix} 1 \\ e^{s_k^{(L)}} \\ \vdots \\ e^{(J-1)s_k^{(L)}} \end{pmatrix}. \quad (30)$$

For  $J \geq K$ ,  $J^L \times 1$  vectors  $\mathbf{r}(\mathbf{s}_k)$ 's for  $k = 1, 2, \dots, K$  are linearly independent vectors. Therefore, similar to 1-D DMUSIC algorithm, the frequency vector  $\mathbf{s}$  can be estimated by finding the peaks of M-D DMUSIC spectrum

$$P(\mathbf{s}) = \frac{1}{\bar{\mathbf{r}}_r^H(\mathbf{s}) \left( \sum_{k=K+1}^{J^L} \mathbf{v}_k^* \mathbf{v}_k^T \right) \bar{\mathbf{r}}_r(\mathbf{s})} \quad \text{and} \quad \bar{\mathbf{r}}_r = \frac{\mathbf{r}_r}{\|\mathbf{r}_r\|} \quad (31)$$

where vectors  $\mathbf{v}_{K+1}, \dots, \mathbf{v}_{J^L}$  are  $J^L - K$  right singular vectors of  $\mathbf{A}$  corresponding to the  $J^L - K$  smallest singular values.

$$\mathbf{A}(n) = \begin{pmatrix} y(n, 0) & y(n, 1) & \cdots & y(n, J-1) \\ y(n, 1) & y(n, 2) & \cdots & y(n, J) \\ \vdots & \vdots & \ddots & \vdots \\ y(n, N-J) & y(n, N-J+1) & \cdots & y(n, N-1) \end{pmatrix} \quad (21)$$

TABLE II  
2-D PEAK-SEARCHING ALGORITHM

Step 1	Find $(\hat{\omega}_k^{(1)}, \hat{\omega}_k^{(2)})$ for $k = 1, 2, \dots, K$ maximizing $P(0, 0; \omega^{(1)}, \omega^{(2)})$
Step 2	Find $(\hat{\alpha}_1^{(1)}, \hat{\alpha}_1^{(2)})$ maximizing $P(\alpha^{(1)}, \alpha^{(2)}; \hat{\omega}_1^{(1)}, \hat{\omega}_1^{(2)})$
Step 3	Find $(\hat{\omega}_1^{(1)}, \hat{\omega}_1^{(2)})$ around $(\hat{\omega}_1^{(1)}, \hat{\omega}_1^{(2)})$ and maximizing $P(\hat{\alpha}_1^{(1)}, \hat{\alpha}_1^{(2)}, \omega^{(1)}, \omega^{(2)})$
Step 4	Repeat Step 2, 3 until the estimation of $s_1^{(1)} = -\alpha_1^{(1)} + j\omega_1^{(1)}$ $s_1^{(2)} = -\alpha_1^{(2)} + j\omega_1^{(2)}$ attains certain precision
Step 5	Repeat Step 2, 3 and 4 for $k = 2, 3, \dots, K$

## V. LOW-COMPLEXITY 2-D PEAK-SEARCHING METHOD

To estimate the parameters of L-D NMR signals, we have to find the peaks of L-D DMUSIC spectrum, which is a function of  $2L$  variables:  $\alpha^{(1)}, \dots, \alpha^{(L)}, \omega^{(1)}, \dots, \omega^{(L)}$ . To find the peak of the L-D DMUSIC spectrum, we have to calculate it in a fine lattice, which is a  $2L$ -dimensional search.

Since the damping factors of NMR signals are normally very small (usually less than 0.3), the following simplified peak-searching algorithm can be used to reduce the computation of the 2-D DMUSIC spectrum. For convenience, we rewrite  $P(s^{(1)}, s^{(2)})$  as  $P(\alpha^{(1)}, \alpha^{(2)}; \omega^{(1)}, \omega^{(2)})$ . If the damping factors of signals are small, then the maxima of  $P(\alpha^{(1)}, \alpha^{(2)}; \omega^{(1)}, \omega^{(2)})$  will be near the  $(\omega^{(1)}, \omega^{(2)})$  plane.

Because  $P(\alpha^{(1)}, \alpha^{(2)}; \omega^{(1)}, \omega^{(2)})$  is convex around its maxima,  $P(0, 0; \omega^{(1)}, \omega^{(2)})$  is also convex around the maxima. Hence,  $P(0, 0; \omega^{(1)}, \omega^{(2)})$  has maximum points  $(\hat{\omega}_k^{(1)}, \hat{\omega}_k^{(2)})$  for  $k = 1, 2, \dots, K$ , which are near the peaks of  $P(\alpha^{(1)}, \alpha^{(2)}; \omega^{(1)}, \omega^{(2)})$ . For each maximum  $(\hat{\omega}_k^{(1)}, \hat{\omega}_k^{(2)})$  of  $P(0, 0; \omega^{(1)}, \omega^{(2)})$ ,  $(\hat{\alpha}_k^{(1)}, \hat{\alpha}_k^{(2)})$  can be found to maximize  $P(\alpha^{(1)}, \alpha^{(2)}; \hat{\omega}_k^{(1)}, \hat{\omega}_k^{(2)})$  since  $P(\alpha^{(1)}, \alpha^{(2)}; \hat{\omega}_k^{(1)}, \hat{\omega}_k^{(2)})$  is convex. Then, we can find  $(\hat{\omega}_k^{(1)}, \hat{\omega}_k^{(2)})$  around  $(\hat{\omega}_k^{(1)}, \hat{\omega}_k^{(2)})$  that maximizes  $P(\hat{\alpha}_k^{(1)}, \hat{\alpha}_k^{(2)}; \omega^{(1)}, \omega^{(2)})$ . Repeating the above procedures, the peaks of  $P(\alpha^{(1)}, \alpha^{(2)}; \omega^{(1)}, \omega^{(2)})$  can be searched. The above searching procedures, which are summarized in Table II, reduces the peak search of a four-variable function to that of a two-variable function. Therefore, the computation is significantly reduced.

The parameter estimation of higher-dimensional NMR signals can be decomposed into parameter estimation of lower-dimensional NMR signals to reduce the computation. Taking three-dimensional (3-D) NMR signal  $y(n_1, n_2, n_3)$  as an example, it is a 2-D NMR signal if  $n_3$  is fixed. Hence, the

frequency pairs corresponding to time index  $n_1$  and  $n_2$  can be estimated by the 2-D MUSIC algorithm. Similarly, the frequencies corresponding to  $n_3$  can be estimated using the 1-D DMUSIC algorithm. The 3-D complex frequency vector can be found by searching the peaks of the 3-D DMUSIC spectrum near all the combinations of the frequency pairs corresponding to indices  $n_1$  and  $n_2$  and the frequencies corresponding to index  $n_3$ . In this way, parameter estimation of 3-D NMR signals is simplified into parameter estimation of 2-D NMR signals and 1-D NMR signals. In fact, the parameter estimation of higher-dimensional NMR signal can always be decomposed into the parameter estimations of corresponding 2-D NMR signals and 1-D NMR signal. It is also worth mentioning that the complexity of the  $M$ -D DMUSIC algorithm is comparable to the other singular-value decomposition (SVD)-based algorithms. Since in our algorithm high-resolution estimates can be achieved using fewer number of data points, the complexity of the SVD computation is less than that of existing methods whose performance relies on using large data matrices. But our algorithm also involves a 2L-D peak-searching process whose complexity is heavily reduced by our proposed peak-searching algorithm. Therefore, the burden of the peak-searching process is also significantly reduced in our algorithm.

## VI. SIMULATION EXAMPLES

Before presenting our estimation results for experimental data from NIH, we confirm the DMUSIC algorithms by three simulation examples in this section. In our simulation examples, the measurement noise  $w(\cdot)$  is complex white Gaussian noise with variance  $\sigma^2$  that is determined by peak

$$\mathbf{A}(n_1, \dots, n_l) = \begin{pmatrix} \mathbf{A}(n_1, \dots, n_l, 0) & \cdots & \mathbf{A}(n_1, \dots, n_l, J-1) \\ \mathbf{A}(n_1, \dots, n_l, 1) & \cdots & \mathbf{A}(n_1, \dots, n_l, J) \\ \vdots & \vdots & \vdots \\ \mathbf{A}(n_1, \dots, n_l, N-J-1) & \cdots & \mathbf{A}(n_1, \dots, n_l, N-1) \end{pmatrix} \quad (26)$$

for  $l = 1, 2, \dots, L-2$

$$\mathbf{A}(n_1, \dots, n_{L-1}) = \begin{pmatrix} y(n_1, \dots, n_{L-1}, 0) & \cdots & y(n_1, \dots, n_{L-1}, J-1) \\ y(n_1, \dots, n_{L-1}, 1) & \cdots & y(n_1, \dots, n_{L-1}, J) \\ \vdots & \vdots & \vdots \\ y(n_1, \dots, n_{L-1}, N-J) & \cdots & y(n_1, \dots, n_{L-1}, N-1) \end{pmatrix} \quad (27)$$

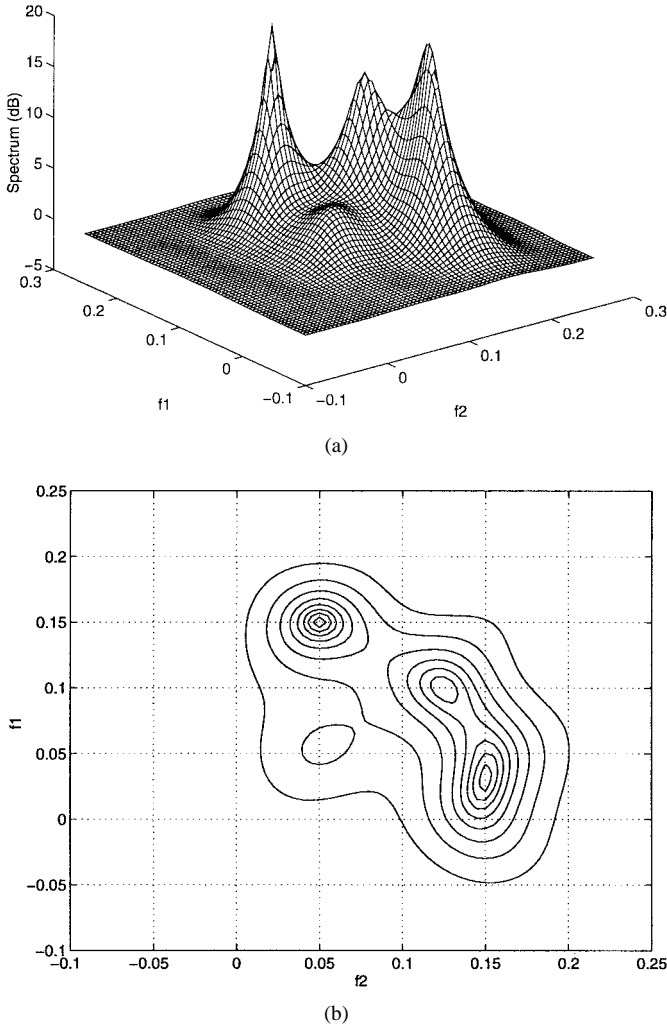


Fig. 2. The spectrum  $P(0, 0; \omega^{(1)}, \omega^{(2)})$  and its contour for synthetic 2-D NMR signal using 2-D peak-searching method when SNR = 30 dB.

SNR defined as

$$\text{SNR} = 10 \log \frac{1}{2\sigma^2}. \quad (32)$$

*Example 1:* The synthetic 1-D NMR data is generated by

$$y(n) = e^{s_1 n} + e^{s_2 n} + w(n) \quad (33)$$

where  $s_1 = -0.2 + j2\pi(0.42)$ ,  $s_2 = -0.1 + j2\pi(0.42 + \Delta)$ . The data length is  $N = 24$ , therefore, we pick  $J = 12$ .

For SNR = 40 dB,  $\Delta = 0.1$ , the 1-D DMUSIC spectrum and its contour are shown in Fig. 1(a) and (b), respectively. From these figures, the damping factors and frequencies of the signal can be estimated simultaneously by finding the peak on the spectrum. But, if SNR = 40 dB,  $\Delta = 0$ , i.e., two exponentially damped signals with the same frequency, the spectrum has just one peak, as indicated by Fig. 1(c) and (d). Hence, the damping factors of the signals cannot be estimated correctly under this condition. However, if SNR is increased to 60 dB, both the damping factors and the frequencies of the signals can be estimated again as demonstrated by Fig. 1(e) and (f).

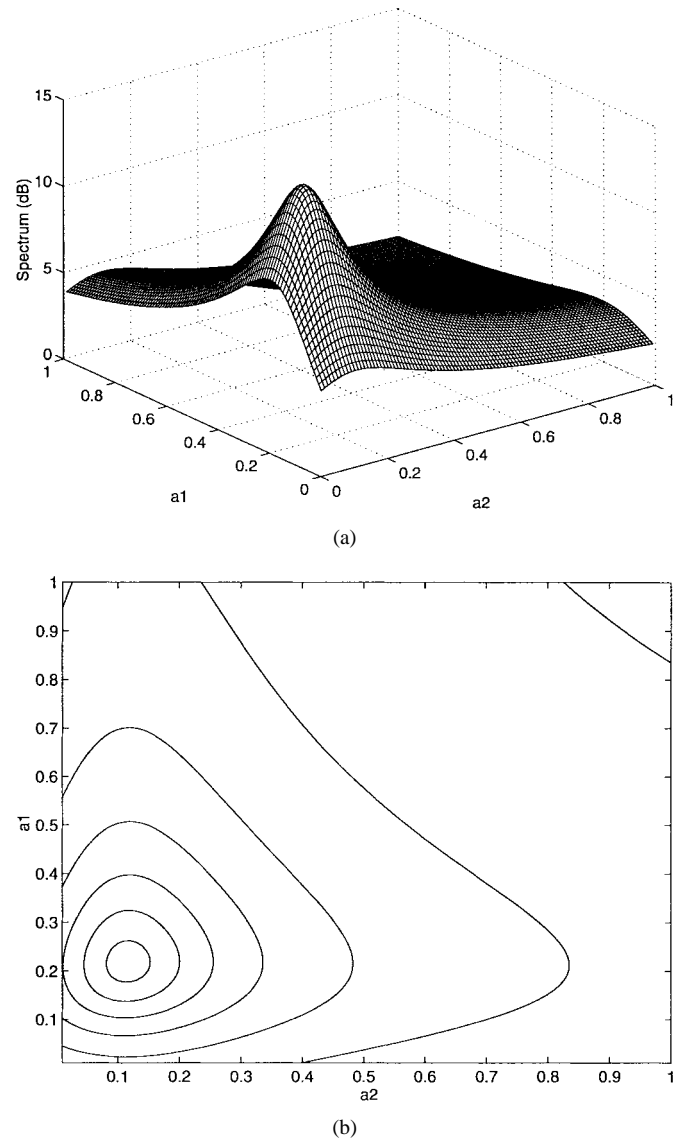


Fig. 3. The spectrum  $P(\alpha^{(1)}, \alpha^{(2)}; 0.20\pi, 0.25\pi)$  and its contour for synthetic 2-D NMR signal 2-D peak-searching method when SNR = 30 dB.

*Example 2:* In this example, a set of synthetic 2-D NMR data is first generated using (3) and (4). The model order  $K = 5$  and the frequency pairs are shown in Table III. The data length is  $N = 24$  and  $J$  is, therefore, chosen to be 12 to achieve best performance.

For SNR = 40 dB, the 2-D DMUSIC spectrum,  $P(\mathbf{s})$ , and its contour, using 2-D peak-searching algorithm are shown in Figs. 2 and 3. From the figures, we can see that  $P(\mathbf{s})$  only has four peaks on  $\omega_1 - \omega_2$  plane with  $\alpha^{(1)} = \alpha^{(2)} = 0$ , but five peaks are found by the 2-D peak-searching algorithm and the parameters can be estimated successfully as illustrated by Table III.

From the estimation results in Table III, our estimation algorithm cannot resolve the fourth and fifth frequency pairs when SNR = 20 dB, since they are too close.

*Example 3:* The 3-D synthetic NMR signal is generated by using (7) and (8). The model order is  $K = 3$ . The frequency vectors are shown in Table IV. The synthetic NMR

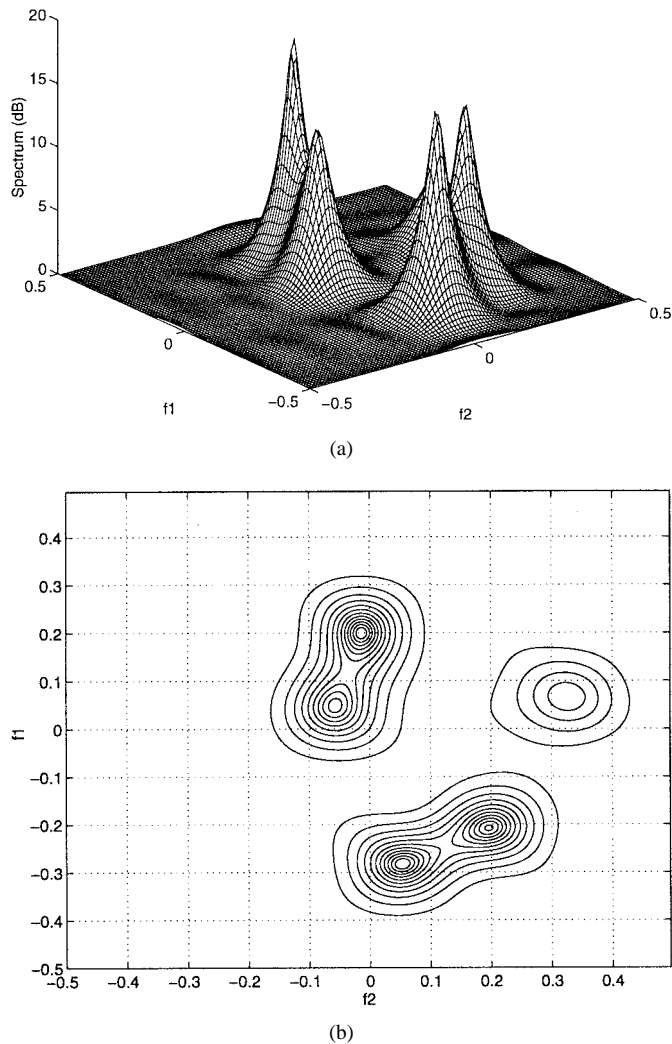


Fig. 4. The spectrum  $P(0,0;\omega^{(1)},\omega^{(2)})$  and its contour for 2-D NMR signal.

TABLE III  
ESTIMATED PARAMETERS OF A SYNTHETIC 2-D NMR SIGNAL

	k	$\alpha_k^{(1)}$	$\omega_k^{(1)}$	$\alpha^{(2)k}$	$\omega_k^{(2)}$
True values	1	-0.20	$0.10\pi$	-0.10	$0.10\pi$
	2	-0.00	$0.30\pi$	-0.00	$0.10\pi$
	3	-0.05	$0.20\pi$	-0.02	$0.25\pi$
	4	-0.02	$0.05\pi$	-0.02	$0.30\pi$
	5	-0.10	$0.06\pi$	-0.02	$0.31\pi$
Estimated values SNR=30dB	1	-0.21	$0.10\pi$	-0.11	$0.10\pi$
	2	-0.00	$0.30\pi$	-0.00	$0.10\pi$
	3	-0.05	$0.20\pi$	-0.02	$0.25\pi$
	4	-0.11	$0.05\pi$	-0.04	$0.31\pi$
	5	-0.09	$0.07\pi$	-0.01	$0.31\pi$
Estimated values SNR=20dB	1	-0.19	$0.10\pi$	-0.10	$0.10\pi$
	2	-0.00	$0.30\pi$	-0.00	$0.10\pi$
	3	-0.05	$0.20\pi$	-0.02	$0.25\pi$
	4	-1.00	$0.64\pi$	-0.00	$0.31\pi$
	5	-0.07	$0.06\pi$	-0.01	$0.31\pi$

signal is corrupted by measurement noise with SNR = 15 dB. First, we estimate the complex frequency pairs corresponding to the first two indexes using the simplified 2-D searching algorithm and the complex frequencies corresponding to the

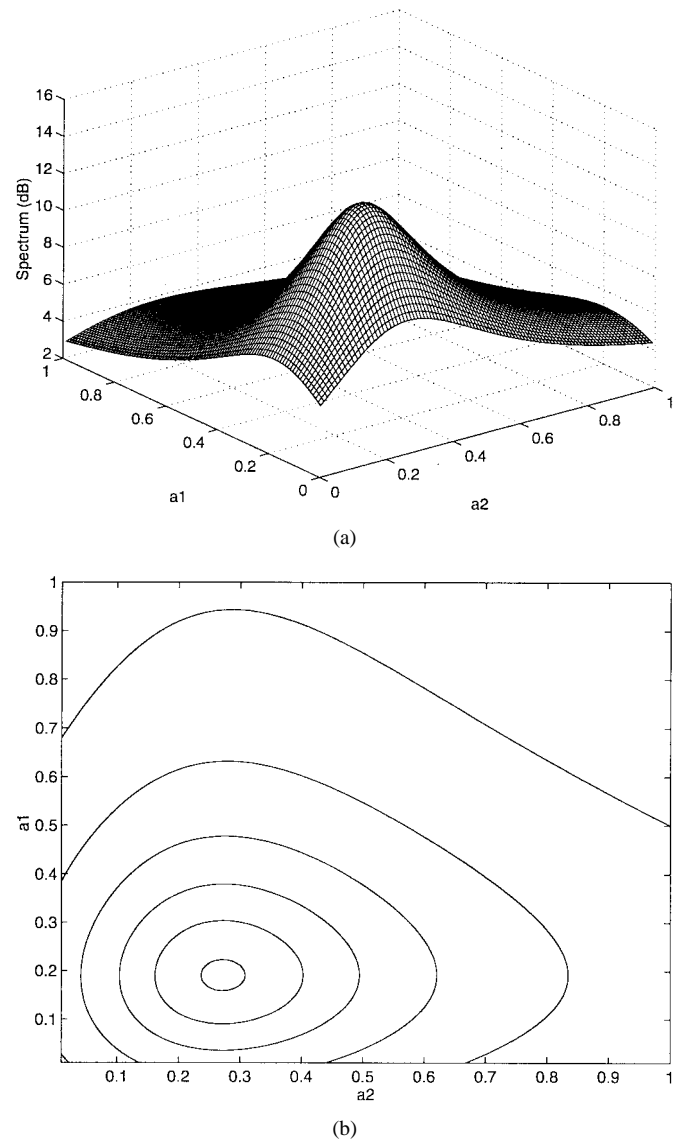


Fig. 5. The spectrum  $P(\alpha^{(1)},\alpha^{(2)};0.10\pi,-0.12\pi)$  and its contour for 2-D NMR signal.

TABLE IV  
ESTIMATED PARAMETERS OF A SYNTHETIC 3-D NMR SIGNAL

	k	$\alpha_k^{(1)}$	$\omega_k^{(1)}$	$\alpha^{(2)k}$	$\omega_k^{(2)}$	$\alpha^{(3)k}$	$\omega_k^{(3)}$
True values	1	0.040	$-0.740\pi$	0.140	$-0.820\pi$	0.100	$-0.140\pi$
	2	0.010	$0.050\pi$	0.190	$-0.190\pi$	0.170	$0.310\pi$
	3	0.140	$0.290\pi$	0.080	$-0.720\pi$	0.010	$0.150\pi$
Estimated values SNR=15dB	1	0.040	$-0.740\pi$	0.150	$-0.815\pi$	0.100	$-0.140\pi$
	2	0.010	$0.050\pi$	0.190	$-0.190\pi$	0.180	$0.310\pi$
	3	0.140	$0.295\pi$	0.080	$-0.720\pi$	0.020	$0.145\pi$

third index. Then, we find the 3-D frequency vectors by searching the maximum points at the all possible combinations of the complex frequency pairs and the complex frequencies. The estimated frequency vectors are shown in Table IV.

## VII. ESTIMATION RESULTS ON EXPERIMENTAL DATA

The 2-D MUSIC parameter estimation algorithm is used in experimental NMR data. The measured  $24 \times 24$  2-D NMR

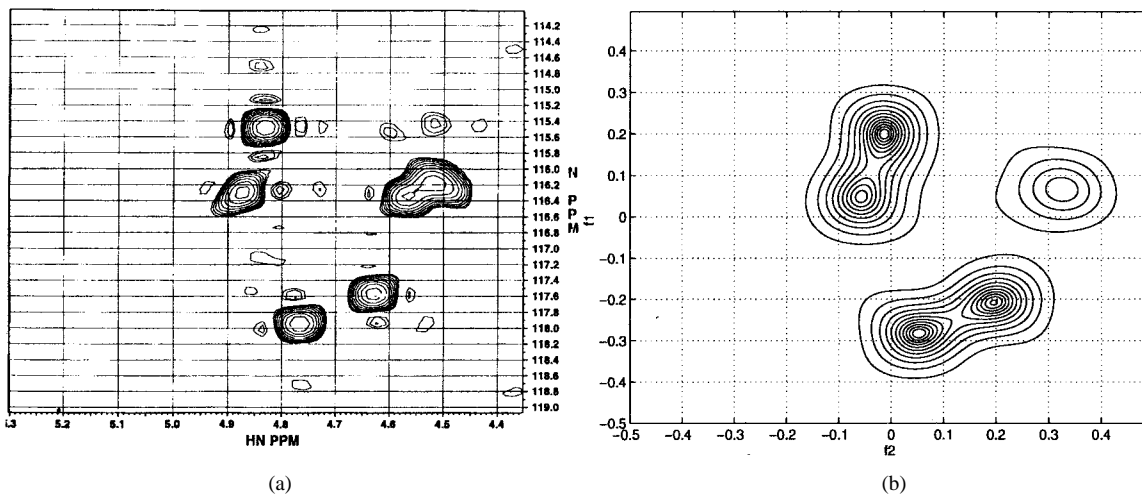


Fig. 6. Contour plot of  $P(0, 0; \omega^{(1)}, \omega^{(2)})$  for 2-D NMR signal using (a) NmrPipe and (b) our 2-D DMUSIC algorithm.

TABLE V  
ESTIMATED PARAMETERS OF A SET OF EXPERIMENTAL 2-D NMR DATA

$k$	$\alpha_k^{(1)}$	$\omega_k^{(1)}$	$\alpha_k^{(2)}$	$\omega_k^{(2)}$
1	0.06	$0.04\pi$	0.06	$-0.02\pi$
2	0.07	$-0.56\pi$	0.08	$0.10\pi$
3	0.07	$-0.42\pi$	0.09	$0.40\pi$
4	0.13	$0.10\pi$	0.09	$-0.12\pi$
5	0.21	$0.12\pi$	0.29	$0.64\pi$

data was obtained from NIH's multidimensional NMR spectroscopy. The spectrum  $P(0, 0; \omega^{(1)}, \omega^{(2)})$ , and its contour are shown in Fig. 4(a) and (b). From Fig. 4,  $P(0, 0; \omega^{(1)}, \omega^{(2)})$  has five peaks. The 2-D spectrum  $P(\alpha^{(1)}, \alpha^{(2)}; 0.10\pi, -0.12\pi)$  and its contour are shown in Fig. 5(a) and (b). By repeating Steps 2 and 3 in Table II just twice, we can successfully estimate the frequencies and the damping factors of the NMR data, which are listed in Table V.

Fig. 6 compares our algorithm with the present algorithm in the NmrPipe, which is a standard NMR processing tool in NIH used by the NMR researchers over the world. Fig. 6(a) and (b) illustrate the estimation results for the same set of experimental NMR data, respectively, by the present algorithm in the NmrPipe tool and the 2-D MUSIC algorithm. Compared with Fig. 6(b), Fig. 6(a) has many spurious signals, while our 2-D MUSIC algorithm can estimate signal parameters effectively without introducing any spurious signal. Hence, our method has been included in the NmrPipe tool as an effective algorithm for NMR data processing.

## VIII. CONCLUSION

In this paper, we have proposed a high-resolution technique for multidimensional NMR spectroscopy, which we call the *M-D MUSIC algorithm*. Since the M-D DMUSIC algorithm makes full use of the rank-deficiency and Hankel properties of the prediction matrix composed of NMR data, it can estimate the NMR signal parameters under low SNR by using very few data points. To reduce the computational complexity, we have also introduced a peak-searching method to locate the peaks of the M-D DMUSIC spectrum.

The performance of our algorithms have been tested by extensive simulation examples and experimental data. The testing results show that the M-D DMUSIC algorithm can resolve closely-spaced frequencies and damping factors, which is one of the most effective algorithms for NMR data. Hence, it has been included in the NmrPipe tool in NIH and will be used by the NMR researchers over the world.

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