Graphical Method for the Determination of the Complex NMR Shift and Equilibrium Constant for a Hetero-Association Accompanying Self-Associations

Chi-Chyuang Lin, Cheng-Yu Fang, Dah-Yu Kao, and Jenn-Shing Chen*

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We describe a novel graphical method which, in conjunction with the previously proposed graphical determination of monomer shift, dimer shift, and dimerization constant for self-association, allows us to determine the complex shift and equilibrium constant for a hetero-association, $A+B \rightleftharpoons AB$, accompanying self-associations, $A+A \rightleftharpoons A_2$, and $B+B \rightleftharpoons B_2$. The merit of the new method includes the removal of the restrictions imposed on the conventional Benesi–Hildebrand (B–H) plot: (1) that the concentration of one component must be much less than that of the other; (2) that there be no accompanying self-association. The simultaneous equilibrium of the self-association of 2-pyrrolidone (A) and that of 4-methyl- α -pyrrolidone (B) and the hetero-association between A and B in acctonitrile-d₃ at 25°C is studied. The inappropriateness of the B–H plot in dealing with this case is also pointed out.

KEY WORDS: Self-association; hetero-association; NMR shift; graphical method.

1. INTRODUCTION

Molecular association arising from hydrogen bonding or electron donorelectron acceptor charge transfer is of importance in chemistry and biochemistry, and has long been of interest to chemists. (1-4) A variety of experimental as well as theoretical means have been employed to study this phenomenon. (5-7) The traditional approach to derive spectroscopic parameters and the

equilibrium constant for a hetero-association from experimental data (UV or NMR) was pioneered by Benesi and Hildebrand (B–H).⁽⁸⁾ However, the B–H plot or its modifications, Scatchard⁽⁹⁾ and Scott⁽¹⁰⁾ plots, are subject to the limitation that the concentration of one component must be much less than that of the other. The solutions prepared under this restriction give unwanted results: (1) monitored signal is too weak to be observed, (2) an extended association such as AB₂, AB₃, etc., takes place, (3) the solution is thermodynamically nonideal so that activity, instead of concentration, should be used in the equilibrium constant expression. Another restriction is also implied, although not explicitly expressed, that the self-association of species is not allowed. This restriction, as a matter of fact, seriously limits the scope of the application of B–H method, since in many cases, self-association of the constituent species is inevitable. An outstanding example⁽¹¹⁾ is the basepairing of adenine—thymine, gutamine—cytosine, in which self-association can by no means be ignored.

In the previous study, (12) in conjunction with a graphical method for determination of monomer shift, dimer shift, and dimerization constant of self-association, (13-15) we have proposed an algorithm to deal with a system which involves simultaneous equilibria $A + B \rightleftharpoons AB$, and $A + A \rightleftharpoons A_2$, without the restrictions imposed on the B-H method. Extension of this treatment to include the self-association $B + B \rightleftharpoons B_2$ is presented here.

2. THEORY

Consider a system containing species A and B in a solvent. A and B undergo hetero-association

$$A + B \rightleftharpoons AB; \qquad K_C = \frac{[AB]}{[A][B]}$$
 (1)

and accompanying self-associations

$$A + A \rightleftharpoons A_2; K_A = \frac{[A_2]}{[A]^2}$$
 (2)

$$B + B \rightleftharpoons B_2; K_B = \frac{[B_2]}{[B]^2}$$
 (3)

Here [A], [A₂] are the equilibrium concentrations of monomer. A, and dimer A_2 , respectively; their counterparts for B are [B] and [B₂]. [AB] is the equilibrium concentration of the complex AB. K_A and K_B are respectively,

the dimerization constants of A, and B. K_C is the equilibrium constant for hetero-association. Consideration of mass balance results in

$$[A]_0 = [A] + [AB] + 2[A_2] = [A] + K_C[A][B] + 2K_A[A]^2$$
 (4)

$$[B]_0 = [B] + [AB] + 2[B_2] = [B] + K_C[A][B] + 2K_B[B]^2$$
 (5)

where [A]_o and [B]_o are, respectively, the initial concentrations of A and B. The last equalities in Eqs. (4, 5) result from the use of equilibrium constant expressions (1–3). The ratios [AB]/[A]_o and [AB]/[B]_o can be expressed as

$$\frac{[AB]}{[A]_0} = \frac{K_C[B]}{1 + K_C[B] + 2K_A[A]}$$
(6)

$$\frac{[AB]}{[B]_0} = \frac{K_C[A]}{1 + K_C[A] + 2K_B[B]} \tag{7}$$

In NMR measurements, the observed chemical shift for a nucleus in molecule A, which involves a rapid exchange between the states of monomer A, dimer A_2 , and complex AB, is the weighted average of their shifts, with the respective mole fractions as weight factors. According to Gutowsky and Saika, (16) the observed shift $\Delta \nu_{\rm obs}^{\rm A}$ is given by

$$\Delta \nu_{\text{obs}}^{A} = \frac{[A]}{[A]_{c}} \Delta \nu_{\text{m}}^{A} + \frac{2[A_{2}]}{[A]_{c}} \Delta \nu_{\text{d}}^{A} + \frac{[AB]}{[A]_{c}} \Delta \nu_{\text{c}}^{A}$$
 (8)

where $\Delta \nu_{\rm m}^{\rm A}$, $\Delta \nu_{\rm d}^{\rm A}$, and $\Delta \nu_{\rm c}^{\rm A}$ are, respectively, the chemical shifts of the monitored nucleus in molecule A, in the states of monomer, dimer, and complex. Similar result can be applied to the monitored nucleus in molecule B

$$\Delta \nu_{\text{obs}}^{\text{B}} = \frac{[\text{B}]}{[\text{B}]_{\text{o}}} \Delta \nu_{\text{m}}^{\text{B}} + \frac{2[\text{B}_{2}]}{[\text{B}]_{\text{o}}} \Delta \nu_{\text{d}}^{\text{B}} + \frac{[\text{AB}]}{[\text{B}]_{\text{o}}} \Delta \nu_{\text{c}}^{\text{B}}$$
(9)

where $\Delta \nu_m^B$, $\Delta \nu_d^B$, and $\Delta \nu_c^B$ are, respectively, the chemical shifts of the monitored nucleus in molecule B, in the states of monomer, dimer, and complex. Equations (8, 9) are valid only if there is no exchange between the monitored nuclei in A and in B.

To proceed, we define

$$\Delta \tilde{\nu}^{A} = \Delta \nu_{\text{obs}}^{A} - \left(\frac{[A]}{[A]_{\text{o}}} \Delta \nu_{\text{m}}^{A} + \frac{2[A_{2}]}{[A]_{\text{o}}} \Delta \nu_{\text{d}}^{A}\right) = \frac{[AB]}{[A]_{\text{o}}} \Delta \nu_{\text{c}}^{A}$$
(10)

$$\Delta \tilde{\nu}^{B} = \Delta \nu_{\text{obs}}^{B} - \left(\frac{[B]}{[B]_{o}} \Delta \nu_{\text{m}}^{B} + \frac{2[B_{2}]}{[B]_{o}} \Delta \nu_{\text{d}}^{B} \right) = \frac{[AB]}{[B]_{o}} \Delta \nu_{\text{c}}^{B}$$
(11)

Clearly, $\Delta \tilde{\nu}^A$ represents the portion of the contribution from the complex AB to $\Delta \nu_{obs}^A$, and $\Delta \tilde{\nu}^B$ that to $\Delta \nu_{obs}^B$. $\Delta \tilde{\nu}^A$ and $\Delta \tilde{\nu}^B$ are calculable if the self-

association parameters for A ($\Delta \nu_m^A$, $\Delta \nu_d^A$, and K_A), the self-association parameters for B ($\Delta \nu_m^B$, $\Delta \nu_d^B$, and K_B), and the hetero-association constant K_C are known.

Substitution of Eq. (6) into Eq. (10) and Eq. (7) into Eq. (11), respectively, followed by rearrangement leads to

$$\frac{1}{\Delta \tilde{\nu}^{A}} = \frac{1}{\Delta \nu_{c}^{A}} + \left(\frac{1}{[B]} + \frac{2K_{A}[A]}{[B]}\right) \frac{1}{K_{C} \Delta \nu_{c}^{A}} \equiv \frac{1}{\Delta \nu_{c}^{A}} + \frac{\Phi_{A}}{K_{c} \Delta \nu_{c}^{A}}$$
(12)

$$\frac{1}{\Delta \tilde{\nu}^{\mathrm{B}}} = \frac{1}{\Delta \nu_{\mathrm{c}}^{\mathrm{B}}} + \left(\frac{1}{[\mathrm{A}]} + \frac{2K_{\mathrm{B}}[\mathrm{B}]}{[\mathrm{A}]}\right) \frac{1}{K_{\mathrm{C}} \Delta \nu_{\mathrm{c}}^{\mathrm{B}}} \equiv \frac{1}{\Delta \nu_{\mathrm{c}}^{\mathrm{B}}} + \frac{\Phi_{\mathrm{B}}}{K_{\mathrm{c}} \Delta \nu_{\mathrm{c}}^{\mathrm{B}}}$$
(13)

A plot of $1/\Delta \tilde{v}^A vs$. Φ_A permits us to derive Δv_c^A from the intercept, and K_C from the intercept and the slope. In the same manner Δv_c^B and K_C can be obtained from a plot of $1/\Delta \tilde{v}^B vs$. Φ_B . Note that regressed values of K_C obtained from Eq. (12) and from Eq. (13) should be equal, since they represent the same hetero-association equilibrium constant. This offers an opportunity for a consistency check of the determination.

In general, the system has nine parameters to be determined: three for self-association parameters of A, their counterparts of B, and another three for hetero-association between A and B. The six self-association parameters can be determined separately by doing experiments on the self-association of A and that of B. Thus, only three parameters $\Delta \nu_c^A$, $\Delta \nu_c^B$ and K_C are left. Now we encounter a dilemma that to determine K_C from the plots based on Eq. (12) [or Eq. (13)] requires K_C to calculate $\Delta \tilde{\nu}^A$ (or $\Delta \tilde{\nu}^B$) and Φ_A and Φ_B . This difficulty can be overcome by the following strategy.

3. PROPOSED ALGORITHM

3.1. Self-Association

The algorithm for the determination of the monomer shift, dimer shift, and dimerization constant for a self-association has been published elsewhere. (13-15) We will present it only briefly here. For self-association of A, two interconvertible expressions have been derived

$$\Delta \nu_{\text{obs}}^{A} = \Delta \nu_{\text{d}}^{A} - \left(\frac{\Delta \nu_{\text{d}}^{A} - \Delta \nu_{\text{m}}^{A}}{2K_{A}}\right)^{1/2} \left(\frac{\Delta \nu_{\text{obs}}^{A} - \Delta \nu_{\text{m}}^{A}}{[A]_{0}}\right)^{1/2}$$
(14)

$$\Delta \nu_{\text{obs}}^{A} = \Delta \nu_{\text{m}}^{A} + (\Delta \nu_{\text{d}}^{A} - \Delta \nu_{\text{m}}^{A}) \frac{(1 + 8K_{\text{A}}[A]_{0})^{1/2} - 1}{(1 + 8K_{\text{A}}[A]_{0})^{1/2} + 1}$$
(15)

Our method, which combines linear and quadratic regression, is motivated by the following observation: If the value of $\Delta \nu_{\rm m}^{\rm A}$ in $X = \{(\Delta \nu_{\rm obs}^{\rm A} -$

 $\Delta \nu_{\rm m}^{\rm A}/[{\rm A}]_{\rm o}\}^{1/2}$ is correctly guessed, a linear regression on the data of $\Delta \nu_{\rm obs}^{\rm A}$ vs. X based on Eq. (14) should give a perfect linear fit to the experimental data. The true values of $\Delta \nu_{\rm d}^{\rm A}$ and $K_{\rm A}$ then can be derived from the intercept and slope of this line. If a quadratic polynomial, in the form of $\Delta \nu_{\rm obs}^{\rm A} = a + b X + C X^2$, instead of a linear one is tried and the same perfectly-fitted line is returned, the quadratic term of the adjusted equation vanishes and the linear part becomes identical to the former adjusted linear equation. On the other hand, if the value of $\Delta \nu_{\rm m}^{\rm A}$ is guessed incorrectly, the linear and quadratic regressions will be different. The adjusted linear equation should thus give values of $\Delta \nu_{\rm d}^{\rm A}$ and $K_{\rm A}$ from the intercept and slope, which are different from the values obtained with the linear part of the adjusted quadratic equation.

The new idea is to plot on the same scale regressed K_A vs. guessed $\Delta \nu_m^A$ for both linear and quadratic regressions. The point representing the true values of K_A and $\Delta \nu_m^A$ will lie upon a curve, L1, for the linear regression, or, upon Q1 for quadratic regression. We propose that the best values for K and $\Delta \nu_m^A$ should occur at the intersection, P1, of these two curves.

Now turning to Eq. (15) and following the line of the foregoing argument, we conclude that the regression of $\Delta \nu_{\text{obs}}^{\text{A}} \nu s$.

$$f_{\rm d} = \frac{(1 + 8K_{\rm A}[{\rm A}]_{\rm o})^{1/2} - 1}{(1 + 8K_{\rm A}[{\rm A}]_{\rm o})^{1/2} + 1}$$

using linear and quadratic equations will produce from the intercept the same value of $\Delta \nu_{\rm m}^{\rm A}$, if $K_{\rm A}$ is correctly guessed, or different values otherwise. Plots of guessed values of $K_{\rm A}$ vs. adjusted values of $\Delta \nu_{\rm m}^{\rm A}$, for linear and quadratic regressions, generate curves L2 and Q2, respectively. Again, the intersection, P2, of L2 and Q2 should determine the true values of $\Delta \nu_{\rm m}^{\rm A}$ and $K_{\rm A}$.

Ideally, the two intersections P1 and P2 should coincide, since they represent the same physical entities. Any departure, however, may reflect:(1) experimental errors in shift and/or solute concentration measurement, (2) extended association beyond dimerization, or (3) nonideal thermodynamic behavior of the solution. The last possibility causes the dimerization constant to depend on the solute concentration. In practical applications, the departure of P1 from P2 should provide a check on consistency as well as a method for estimating errors associated with this graphical determination. To avoid bias associated with choosing either P1 or P2, we should take both into consideration on an equal footing. Accordingly, the average value and standard deviation of $\Delta \nu_m^A$ (or K_A) calculated by using P1 and P2 together can be taken as the finally determined value for $\Delta \nu_m^A$ (or K_A) and its associated error, respectively.

The value of $\Delta \nu_{\rm m}^{\rm A}$ finally determined can then be used to compute X, and $\Delta \nu_{\rm obs}^{\rm A}$ vs. X can be fitted to Eq. (14) to obtain the true value of $\Delta \nu_{\rm d}^{\rm A}$ from

the intercept. Similarly, the value of K_A finally determined can also be used to compute f_d , and $\Delta \nu_{obs}^A$ vs. f_d can be fitted to Eq. (15) to obtain the true value of $\Delta \nu_d^A$ by evaluating $\Delta \nu_{obs}^A$ of Eq. (15) at $f_d = 1$. The average and standard deviation of $\Delta \nu_d^A$ from the two determinations can then be taken as the final value and error of $\Delta \nu_d^A$.

3.2. Hetero-Association

Equations (12, 13) are our central results on which the algorithm for obtaining the complex shifts and hetero-association constant is based. If K_A and K_B have already been determined, with a guessed value of K_C , one can solve

$$f([A]) = 2K_A[A]^2 + K_C[A][B] + [A] - [A]_0 = 0$$
 (16)

$$g([A]) = 2K_B[B]^2 + K_C[A][B] + [B] - [B]_o = 0$$
 (17)

at least numerically, for $0 < [A] < [A]_o$, and $0 < [B] < [B]_o$ from a pair of initial concentrations of $[A]_o$ and $[B]_o$. According to Descartes' rule of signs,⁽¹⁷⁾ the number of positive roots of a polynomial equation is equal to the number of variations in signs of the coefficients, If both [A] and [B] are stipulated to be positive, there is only one change in sign for f([A]) and g([A]). Hence, each equation has only one positive root. We asserted that $0 < [A] < [A]_o$, since $f([A]_o) = K_C[A]_o[B] + 2K_A[A]_o^2 > 0$, and $f(0) = -[A]_o < 0$. A similar argument leads to $0 < [B] < [B]_o$.

The solved values of [A] and [B] for a guessed value of K_C are then used to calculate $[A_2]$, and $x = \Phi_A$ in Eq. (12). Also, with the known monomer shift and dimer shift for self-association of A, one can calculate $y = \Delta \tilde{\nu}^A$. A linear plot of y vs. x based on Eq. (12) gives a regressed value of K_C . If K_C is correctly guessed, the regressed value and guessed value will be the same, otherwise they will be different. The correct, finally determined value of K_C may be obtained from the intersection of a plot for y (= regressed values of K_C) vs. x (= guessed values of K_C) and another one for x = y. The last plot is just an artifice to locate the point where K_C has the same guessed and regressed values. The regressed value of $\Delta \nu_c^A$ obtained from the intercept of the linear plot of $y = \Delta \tilde{\nu}^A$ vs. $x = \Phi_A$ by using the finally determined value of K_C is then taken as the finally determined value of $\Delta \nu_c^A$. Now turning to Eq. (13) and following the same procedure to manipulate the data of $\Delta \nu_{obs}^B$, we will obtain the corresponding results for K_C and $\Delta \nu_c^B$.

4. EXPERIMENTAL AND RESULTS

We have carried out NMR experiments at 25°C on self-association of 2-pyrrolidone (A), and of 4-methyl- α -pyrrolidone (B), and hetero-association

between them, $A + B \rightleftharpoons AB$, accompanying the foregoing two self-associations, all in acetonitrile-d₃. The 300 MHz proton spectra were taken using a Varian Unity-300 NMR spectrometer. All chemicals were highest grade from Aldrich and were used as received. Samples of different concentrations (measured in the units of molality), containing a trace of TMS as a reference, were prepared gravimetrically in small vials with the help of microsyringes. Samples of known concentration were then transferred into 5 mm O. D. NMR tubes, and were sealed without further degassing.

We monitored the concentration dependence of the shift of the NH peak in A and/or B. These data were then manipulated and the results plotted by programs written in the *Mathematica*⁽¹⁸⁾ language and executed on a Digital DECstation 5000/25.

Figure 1 represents a graphic determination of monomer shift and dimerization constant for the self-association of 2-pyrrolidone (A) in acetonitriled₃ at 25°C from the dilution shift data listed in Table I. The curves L1 and Q1 represent the plots for regressed values of K_A vs. guessed values of $\Delta \nu_m^A$, respectively, for linear and quadratic regressions based on Eq. (14).

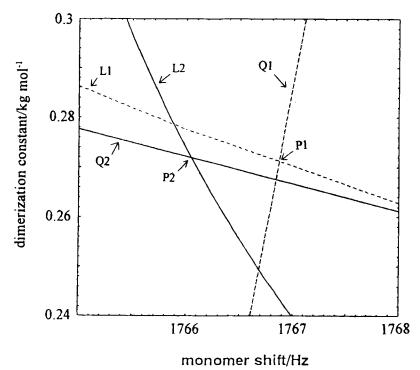


Fig. 1. Graphical determination of monomer shift and dimerization constant for self-association of 2-pyrrolidone in acetonitrile-d₃ at 25°C using data listed in Table I.

Table I. Dilution Shifts of the NH Proton in 2-Pyrrolidone, $\Delta \nu_{\text{obs}}^{A}$, at Various Initial Concentrations in Acetonitrile-d₃ at 25°C^a

[A] _o (mol-kg ⁻¹)	$\Delta v_{\rm obs}^{\rm A}$ (Hz)	[A] _o (mol-kg ⁻¹)	$\Delta v_{\rm obs}^{\rm A}$ (Hz)
0.101	1804.3	1.429	2013.6
0.211	1832.5	1.508	2017.1
0.297	1852.7	1.611	2025.9
0.430	1886.0	1.737	2037.9
0.508	1902.1	1.864	2043.5
0.617	1916.0	1.913	2047.7
0.738	1935.5	2.097	2059.6
0.831	1951.2	2.123	2061.1
0.938	1965.4	2.275	2069.0
1.034	1975.2	2.393	2075.3
1.116	1983.8	2.564	2085.2
1.216	1991.1	2.611	2088.5
1.303	1999.9		

^a Finally determined values: $\Delta \nu_{\rm m}^{\rm A} = 1766.2 \pm 0.3$ Hz; $\Delta \nu_{\rm d}^{\rm A} = 2492.4 \pm 0.6$ Hz; $K_{\rm A} = 0.27 \pm 0.01$ kg-mol⁻¹.

The curves L2 and Q2 represent the plots of guessed values of $K_A vs$. regressed values of $\Delta \nu_m^A$, respectively, for linear and quadratic regressions based on Eq. (15). The intersection P1 of L1 and Q1 then determines the values of $\Delta \nu_m^A$ and K_A based on Eq. (14), while P2 from L2 and Q2 determines those based on Eq. (15). With P1 and P2, the finally determined values of self-association parameters are then read from their midpoint to be $\Delta \nu_m^A = 1766.2 \pm 0.3$ Hz, $\Delta \nu_d^A = 2492.4 \pm 0.6$ Hz, $K_A = 0.27 \pm 0.01$ kg/mol⁻¹. For the self-association of 4-methyl- α -pyrrolidone (B) under the same conditions, the parameters determined by the same procedure are: $\Delta \nu_m^B = 3011.7 \pm 1.1$ Hz, $\Delta \nu_d^B = 4564.7 \pm 1.8$ Hz, $K_B = 0.98 \pm 0.14$ kg-mol⁻¹, respectively, from the dilution shift data listed in Table II. As shown in Fig. 2, the goodness of determination for self-association of A is appraised by a comparison of the theoretical curve obtained by putting the determined parameters into Eq. (15) with the experimental points in the plot of $\Delta \nu_{obs}^A vs$. [A]_o. A similar plot of $\Delta \nu_{obs}^B vs$. [B]_o for self-association of B is shown in Fig. 3.

Once $\Delta \nu_{\rm m}^{\rm A}$, $\Delta \nu_{\rm d}^{\rm A}$, $K_{\rm A}$ for self-association of A, and $\Delta \nu_{\rm m}^{\rm B}$, $\Delta \nu_{\rm d}^{\rm B}$, and $K_{\rm B}$ for self-association of B have been determined, we are in a position to determine $\Delta \nu_{\rm c}^{\rm A}$, $\Delta \nu_{\rm c}^{\rm B}$, and $K_{\rm C}$ from the experimental data of $\Delta \nu_{\rm obs}^{\rm A}$, and $\Delta \nu_{\rm obs}^{\rm B}$ for a pair of initial concentrations [A]_o and [B]_o as listed in Table III. With an additional guessed value of $K_{\rm C}$, and following the procedure stated in Section 3, we are able to make a plot from the data of $\Delta \nu_{\rm obs}^{\rm A}$ for various combinations of [A]_o and [B]_o, as shown in Fig. 4, to determine the hetero-association constant as 1.05 kg-mol⁻¹. Since the $K_{\rm C}$ determination is solely from the data of

Table II. Dilution Shifts of the NH Proton in 4-Methyl- α -Pyrrolidone, $\Delta \nu_{\rm obs}^{\rm B}$, at Various Initial Concentrations in Acetonitrile-d₃ at 25°C^a

[B] _o (mol-kg ⁻¹)	$\Delta \nu_{\rm obs}^{\rm B}$ (Hz)	[B] _o (mol-kg ⁻¹)	$\Delta u^{ m B}_{ m obs~(Hz)}$
0.081	3201.9	0.239	3413.9
0.099	3231.9	0.264	3434.6
0.133	3293.7	0.282	3452.6
0.146	3308.0	0.308	3480.1
0.161	3327.8	0.320	3496.5
0.184	3350.2	0.334	3498.0
0.204	3372.8	0.363	3514.6
0.218	3391.9		

^a Finally determined values: $\Delta \nu_{\rm m}^{\rm B} = 3011.7 \pm 1.1 \ {\rm Hz}; \Delta \nu_{\rm d}^{\rm B} = 4564.7 \pm 1.8 \ {\rm Hz}; K_{\rm B} = 0.98 \pm 0.14 \ {\rm kg-mol}.$

 $\Delta \nu_{\rm obs}^{\rm A}$, we rename it $K_{\rm C}^{\rm A}$. Determination of the hetero-association constant from the data of $\Delta \nu_{\rm obs}^{\rm B}$ in Table III is shown in Fig. 5 to have $K_{\rm C}^{\rm B}=1.39$ kg-mol⁻¹. By using these values just determined and following the procedure outlined in Section 3, $\Delta \nu_{\rm C}^{\rm A}=2784.6$, and $\Delta \nu_{\rm c}^{\rm B}=3632.5$ Hz are then obtained

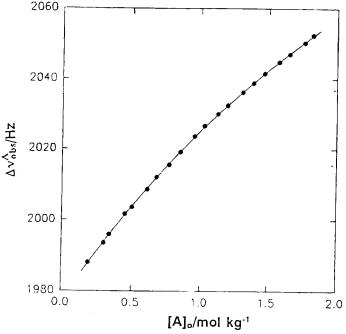


Fig. 2. Plot of observed shift vs. initial concentration for assessment of the determinations of monomer shift, dimer shift, and dimerization constant for self-association of 2-pyrrolidone (A) in acetonitrile-d₃ at 25°C.

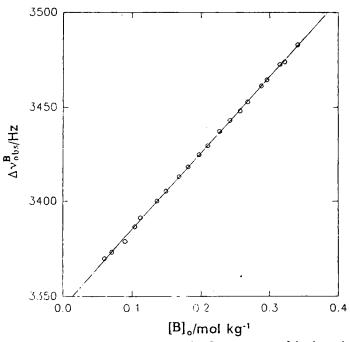


Fig. 3. Plot of observed shift vs. initial concentration for assessment of the determinations of monomer shift, dimer shift, and dimerization constant for self-association of 4-methyl- α -pyrrolidone (B) in acetonitrile-d₃ at 25°C.

from the intercepts of the linear plots based on Eqs. (12, 13), respectively, as shown in Figs. 6 and 7. We have also collected in Table IV the determined values of monomer, dimer, and complex shifts of A and B, the dimerization constants of A and B, and the hetero-association constants determine from the A side and from the B side.

The goodness of determination of the six parameters K_A , K_B , K_C^A , $\Delta \nu_m^A$, $\Delta \nu_d^A$, and $\Delta \nu_c^A$ can then be appraised by comparison with the theoretical values (juxtaposed in parentheses) with the observed values, $\Delta \nu_{\rm obs}^A$ in Table III. The former are calculated from Eq. (8). A similar comparison is also made for the B side. It appears that the determination of K_C from the A side is better and more reliable.

5. DISCUSSION

In the previous publication, (12) dealing with the system of simultaneous equilibria, $A + A \rightleftharpoons A_2$, and $A + B \rightleftharpoons AB$, we derived an equation which

Table III. Chemical Shifts of the NH Proton in 2-Pyrrolidone (A) and in 4-Methyl-α-Pyrrolidone (B) for Various Combinations of Initial Concentrations of $[A]_{o}$ and $[B]_{o}$ in Acetonitrile-d₃ at 25°C^a

[A] _o (mol-kg ⁻¹)	[B] _o (mol-kg ⁻¹)	$\Delta \nu_{ m obs}^{ m A}$ (Hz)	$\Delta \nu_{\rm obs}^{\rm B}$ (Hz)	
0.192	0.341	1988.1(1987.2) ^b	3482.9(3487.2) ^b	
0.303	0.332	1993.6(1992.3)	3473.9(3471.4)	
0.341	0.315	1995.9(1993.9)	3472.5(3466.4)	
0.452	0.296	2001.7(1998.9)	3464.4(3454.4)	
0.502	0.288	2003.7(2001.4)	3461.3(3450.1)	
0.612	0.268	2008.7(2005.9)	3452.9(3440.7)	
0.679	0.257	2012.1(2008.9)	3448.1(3436.5)	
0.770	0.242	2015.6(2013.0)	3443.1(3431.6)	
0.852	0.227	2019.2(2016.2)	3437.4(3427.1)	
0.955	0.210	2023.6(2020.7)	3429.8(3423.3)	
1.028	0.197	2026.6(2023.5)	3425.0(3420.7)	
1.126	0.181	2030.0(2027.6)	3418.5(3418.3)	
1.197	0.168	2032.4(2030.1)	3413.2(3416.3)	
1.308	0.149	2036.2(2034.4)	3405.5(3414.2)	
1.388	0.136	2038.9(2037.5)	3400.3(3413.3)	
1.471	0.112	2041.6(2038.6)	3391.2(3408.8)	
1.575	0.104	2044.9(2044.2)	3386.6(3411.3)	
1.652	0.090	2047.1(2046.7)	3378.8(3410.5)	
1.762	0.071	2050.3(2050.5)	3373.2(3410.0)	
1.826	0.060	2052.5(2052.6)	3369.9(3409.8)	

^a Finally determined values: $\Delta \nu_c^A = 2784.6 \text{ Hz}$; $\Delta \nu_c^B = 3632.6 \text{ Hz}$; $K_C^A = 1.05 \text{ kg-mol}^{-1}$; $K_C^B = 1.39 \text{ kg-mol}^{-1}$. ^bTheoretical calculated values in parentheses.

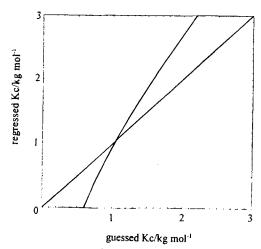


Fig. 4. Graphical determination of K_C^A based on Eq. (12) from data of Δv_{obs}^A for various combination of the initial concentrations of [A]_o and [B]_o listed in Table III.

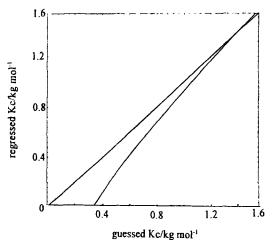


Fig. 5. Graphical determination of K_C^B based on Eq. (13) from the data of $\Delta \nu_{\rm obs}^B$ for various combinations of initial concentrations of [A]_o and [B]_o listed in Table III.

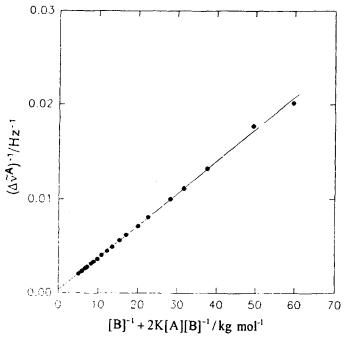


Fig. 6. Linear plot of $\Delta \bar{\nu}^A$ vs. Φ_A based on Eq. (12) using the determined $K_C^A = 1.05 \text{ kg-mol}^{-1}$ to determine $\Delta \nu_c^A$.

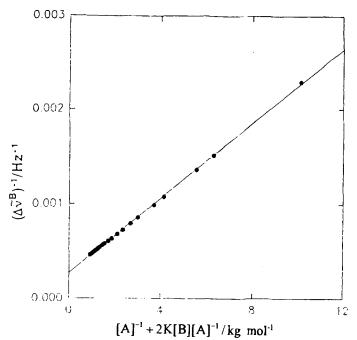


Fig. 7. Linear plot of $\Delta \bar{v}^B$ vs. Φ_B based on Eq. (13) using the determined $K_C^B = 1.39 \text{ kg-mol}^{-1}$ to determine Δv_c^B .

governs the chemical shifts as a function of concentrations of A and B

$$\frac{[B]_o}{\Delta \tilde{\nu}^A[A]_o} = \frac{1}{\Delta \nu_c^A} + \frac{1}{K_C \Delta \nu_c^A} \frac{1}{[A]}$$
 (18)

where $\Delta \tilde{\nu}^A$ is defined as in Eq. (10). This system is a special case of our general system excluding the self-association of B. Therefore, it is expected

Table IV. Collected Data of the Final Determination of Monomer, Dimer, and Complex Shifts and the Self- and Hetero-Association Constants

Self-association					
A + A = A2	$B + B \rightleftharpoons B2$				
$\delta_{\rm m}^{\rm A} = 1766.2 \pm 0.3 \text{ Hz}$	$\delta_{m}^{B} = 3011.7 \pm 1.1 \text{ Hz}$				
$\delta_{\rm d}^{\rm A} = 2492.4 \pm 0.6 \text{ Hz}$	$\delta_{d}^{B} = 4564.7 \pm 1.8 \text{ Hz}$				
$K_{\rm d}^{\rm A} = 0.27 \pm 0.01 \text{ kg-mol}^{-1}$	$K_{d}^{B} = 0.98 \pm 0.14 \text{ kg-mol}^{-1}$				
Hetero-association $A + B \rightleftharpoons AB$					
From signal of A	From signal of B				
$\Delta v_c^A = 2784.6 \text{ Hz}$	$\Delta v_c^B = 3632.6 \text{ Hz}$				
$K_C^A = 1.05 \text{ kg-mol}^{-1}$	$K_C^B = 1.39 \text{ kg-mol}^{-1}$				

that Eq. (12) can be reduced to Eq. (18) with $K_B = 0$. In order to prove this, Eq. (12) is rewritten

$$\frac{1}{\Delta \tilde{\nu}^{A}} = \frac{1}{\Delta \nu_{c}^{A}} + (1 + 2K_{A}[A] + K_{C}[B] - K_{C}[B]) \frac{1}{[B]} \frac{1}{K_{C} \Delta \nu_{c}^{A}}$$
(19)

With Eq. (4) this becomes

$$\frac{1}{\Delta \tilde{\nu}^{A}} = \frac{1}{\Delta \nu_{c}^{A}} + \left(\frac{[A]_{o}}{[A]} - K_{c}[B]\right) \frac{1}{[B]} \frac{1}{K_{c} \Delta \nu_{c}^{A}}$$

$$= \frac{[A]_{o}}{[A]} \frac{1}{[B]} \frac{1}{K_{c} \Delta \nu_{c}^{A}} \tag{20}$$

If Eq. (5) is used with $K_B = 0$, i.e., $[B_2] = 0$, then $1/[B] = (1 + K_c[A])/[B]_o$. Substituting this into Eq. (20), restores Eq. (18).

Moreover, it is also expected that, if the monitored nucleus is in B, Eq. (13), or its alternative Eq. (11), can be reduced to the B-H equation if without any self-association, and $[A]_o >> [B]_o$. In such a case, $[B_2] = 0$, and Eq. (11) is then reduced to

$$\Delta \tilde{\nu}^{B} = \Delta \nu_{\text{obs}}^{B} - \frac{[B]}{[B]_{o}} \Delta \nu_{m}^{B} = \frac{[AB]}{[B]_{o}} \Delta \nu_{c}^{B}$$
 (21)

subtracting ([AB]/[A]_o) $\Delta \nu_m^A$ from both sides of the last equality, Eq. (21) becomes

$$\Delta \nu_{\text{obs}}^{\text{B}} - \Delta \nu_{\text{m}}^{\text{B}} = \frac{[\text{AB}]}{[\text{B}]_{\circ}} (\Delta \nu_{\text{c}}^{\text{B}} - \Delta \nu_{\text{m}}^{\text{B}})$$
 (22)

which can be rearranged to be in the form of the B-H equation

$$\frac{1}{\Delta \nu_{\rm obs}^{\rm B} - \Delta \nu_{\rm m}^{\rm B}} = \frac{1}{\Delta \nu_{\rm c}^{\rm B} - \Delta \nu_{\rm m}^{\rm B}} + \frac{1}{(\Delta \nu_{\rm c}^{\rm B} - \Delta \nu_{\rm m}^{\rm B}) K_{\rm C}} \frac{1}{[{\rm A}]_{\rm o}}$$
(23)

after the use of Eq. (7) and the approximation of $[A] \cong [A]_o$ have been made. The approximation is justified by the conditions $K_A = 0$, and $[A]_o >> [B]_o$.

In order to appreciate the superiority of our method over B–H method, we also treated the data using B–H method pretending that equilibria (2) and (3) do not occur. To fulfil the requirements of the B–H method, we only sort from the data of $\Delta \nu_{\rm obs}^{\rm B}$ vs. [A]_o the portion with [A]_o ≥ 1.388 mol-kg⁻¹, where [B]_o/[A]_o < 0.1. The result of such a B–H plot based on Eq. (23) with $\Delta \nu_{\rm m}^{\rm B} = 3011.7$ Hz is presented in Fig. 8, where a fairly good fitt to a straight line is found. From the plot we obtain a negative value of the hetero-association constant. Thus, ignoring the self-association of the components

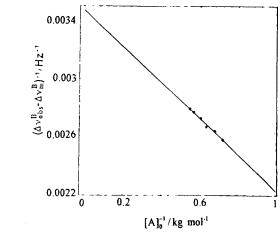


Fig. 8. B-H plot using $\Delta \nu_m^B = 3011.7$ Hz for the determination of $\Delta \nu_c^B$ and K_C from the portion of the data of $\Delta \nu_{obs}^B vs.$ [A]_o with [A]_o ≥ 1.388 mol-kg⁻¹ in Table III. Note that the heteroassociation constant is negative from this determination.

in a system where the hetero-association coexisting with self-association in using the B-H plot may lead to an absurd determination of the hetero-association constant.

6. CONCLUSION

In conjunction with the graphical method for the determination of monomer shift, dimer shift, and dimerization constant from the dilution shift data of a self-association system, we have proposed a new algorithm which enables us to determine the values of complex shift and equilibrium constant for a hetero-association $A + B \rightleftharpoons AB$, accompanying self-associations $A + A \rightleftharpoons A_2$, $B + B \rightleftharpoons B_2$. Our method is superior to the conventional B-H method because the limitation of the assumption that the concentration of one constituent must be much less than that of the other is removed, and the possibility of self-associations of A and B is taken into account. With a practical example of the coexisting self-association and hetero-association of 2-pyrrolidone (A) and that of 4-methyl- α -pyrrolidone (B) in acetonitrile-d₃ at 25°C, we have demonstrated that the new algorithm gives better results than the B-H method. We believe that our new algorithm will lend itself to reliable thermodynamic studies of association systems by NMR or other spectroscopic means.

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