IR Study of Monomer–Dimer Self-association of 2,2-Dimethyl-3-ethyl-3-pentanol in *n*-Octane: Determination of the Molar Absorptivities of Monomer and Dimer Bands, and Dimerization Constants Using Novel Equations

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Abstract The monomer–dimer self-association equilibrium of 2,2-dimethyl-3-ethyl-3pentanol in *n*-octane has been studied by IR spectroscopy at four temperatures (288, 298, 308, and 318 K). The solute was chosen to restrict the self-association between solute molecules to dimerization only, owing to steric hindrance of the bulky chains in the vicinity of the OH group. Two linear equations have been derived for the treatment of the experimental data. One of these equations was used to treat the data of the concentration dependent integrated absorbance of the monomer bands for each temperature to obtain the monomer molar absorptivity, ε_m , and dimerization constant, *K*. The other equation was used to treat the data of concentration-dependent dimer bands to obtain the dimer molar absorptivity, ε_d , and *K*. Thus, the dimerization constant was determined by two methods. Since the same thermodynamic quantity *K* is obtained from either the monomer bands or the dimer bands, the difference between them at a given temperature can serve as an assessment of the quality of the experiment. The standard enthalpy and entropy of dimerization were also obtained from a van't Hoff plot.

Keywords Monomer–dimer self-association · IR spectroscopy · Hindered alcohols · Enthalpy and entropy of dimerization

1 Introduction

Hydrogen bonding is ubiquitous in the field of biological chemistry. It plays an important role in the structure of proteins [1, 2], base-pairing in DNA [3], and enzyme kinetics. Hydrogen bonding is also considered to be one of the major factors needed to describe solutions deviating from normal behavior [4], such as abnormal elevation of the boiling point and lowering of the vapor pressure. Studies of hydrogen bonding are quite extensive

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using various experimental methods as well as theoretical calculations [5–11]. Among the experimental methods, IR spectroscopy is considered to be the most convenient and a powerful one, ever since the establishment of the IR spectral characteristics of hydrogen bonding in alcohols by Errera and Mollet [12]. However, in addition to the band attributed to non hydrogen-bonded monomeric species, IR spectra may be complicated by displaying several overlapped bands corresponding to hydrogen-bonded species of various sizes and configurations due to dimerization or oligomerization. In this case, the study of self-association by IR is quite difficult in that a conjecture is needed as to what sizes and what configurations occur in the system, in order to interpret the spectra [13–19]. As a rule, conjectures are prone to inaccuracies, leading to an unreliable subsequent determination of spectral and thermodynamic parameters. Nevertheless, accurate determination of spectral and thermodynamic parameters can be still made for systems exhibiting only monomer–dimer self-association. This can be realized using hindered alcohols in dilute solution, with bulky side chains in the vicinity of the OH group. The steric hindrance due to the bulky side chains hampers alcohol–alcohol oligomerization [19–23].

In this report we choose dilute solutions of 2,2-dimethyl-3-ethyl-3-pentanol in *n*-octane as a model system to study monomer–dimer self-association [23–26]. In this system, no oligomer bands appear and monomer bands and dimer bands exhibit little overlap, which makes them amenable to spectral deconvolution. To achieve a better determination, we have derived two linear utility equations, one for the data treatment of the monomer bands (Eq. 9 below) to obtain the molar absorptivity of the monomer band, ε_m , and the dimerization constant, *K*, for each temperature; the other equation for dimer bands (Eq. 13 below) was used to obtain the molar absorptivity of the dimer band, ε_d , and the dimerization constant, *K*, for each temperature. It should be remarked that the same dimerization constant can be determined either from the monomer bands or the dimer bands.

2 Utility Equations for the Data Treatment

In a monomer–dimer self-association equilibrium, the monomer, B, and the dimer, B_2 , exhibit a dynamic equilibrium:

$$\mathbf{B} + \mathbf{B} \rightleftharpoons \mathbf{B}_2 \tag{1}$$

The concentration of the monomer, [B], and that of the dimer, [B₂], are related by

$$[B] + 2[B_2] = [B]_0 \tag{2}$$

where $[B]_0$ is the initially (prepared) concentration. In addition to this restriction, the equilibrium concentrations of [B] and $[B_2]$ are related by a temperature-dependent dimerization constant,

$$\mathbf{K} = \frac{[\mathbf{B}_2]}{[\mathbf{B}]^2} \tag{3}$$

Combining Eqs. 2 and 3 permits us to solve for [B] and [B₂] in terms of K and $[B]_0$:

$$[\mathbf{B}] = \frac{2[\mathbf{B}]_{o}}{\left(1 + 8K[\mathbf{B}]_{o}\right)^{1/2} + 1}$$
(4)

$$[\mathbf{B}_{2}] = \left(\frac{1}{2}\right) \frac{\left(1 + 8K[\mathbf{B}]_{o}\right)^{1/2} - 1}{\left(1 + 8K[\mathbf{B}]_{o}\right)^{1/2} + 1} [\mathbf{B}]_{o}$$
(5)

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Equations 4 and 5 have also been derived elsewhere for the NMR study of monomerdimer self-association [26–29].

The absorbance for monomer, $A_{\rm m}(\tilde{v})$, at a particular wavelength \tilde{v} , according to the Beer–Lambert's law, is given by:

$$A_{\rm m}(\tilde{v}) = \varepsilon_{\rm m}(\tilde{v})b[\mathbf{B}] \tag{6}$$

where $\varepsilon_{\rm m}(\tilde{v})$ is the molar absorptivity of the monomer at \tilde{v} , and b is the optical path length of the cell. The integrated absorbance of this monomer band, $A_{\rm m}$, is then obtained by integrating over the whole range of wavenumbers covered by this band, i.e.:

$$A_{\rm m} = \int \varepsilon_{\rm m}(\tilde{v}) b[{\rm B}] d\tilde{v} = \left(\int \varepsilon_{\rm m}(\tilde{v}) d\tilde{v}\right) b[{\rm B}] = \frac{2\varepsilon_{\rm m} b[{\rm B}]_{\rm o}}{\left(1 + 8K[{\rm B}]_{\rm o}\right)^{1/2} + 1}$$
(7)

where Eq. 4 has been used, and $\int \varepsilon_{\rm m}(\tilde{v})d\tilde{v}$ is denoted for brevity by $\varepsilon_{\rm m}$, the molar absorptivity of the monomer band. Equation 7 can be rearranged to obtain:

$$\frac{2\varepsilon_{\rm m}b[{\rm B}]_{\rm o}}{A_{\rm m}} = (1 + 8K[{\rm B}]_{\rm o})^{1/2} + 1 \tag{8}$$

If we subtract unity from both sides of Eq. 8, followed by taking squares, we arrive at:

$$\frac{\left[\mathbf{B}\right]_{o}}{A_{\mathrm{m}}} = \frac{1}{\varepsilon_{\mathrm{m}}b} + \left(\frac{2K}{\left(\varepsilon_{\mathrm{m}}b\right)^{2}}\right)A_{\mathrm{m}} \tag{9}$$

after some simple algebraic manipulations. Equation 9 allows us to fit the experimental data of $y = \frac{[B]_{a}}{A_{m}}$ versus $x = A_{m}$ to a straight line whose slope *p* and intercept *q* give $\varepsilon_{m} = \frac{1}{qb}$ and $K = \frac{p}{2q^{2}}$.

Following the same line of reasoning as in deriving Eq. 7, from Eq. 5 the integrated absorbance of the dimer band, A_d , is obtained to be:

$$A_{\rm d} = \varepsilon_{\rm d} b[{\bf B}_2] = \left(\frac{\varepsilon_{\rm d} b[{\bf B}]_{\rm o}}{2}\right) \frac{\left(1 + 8K[{\bf B}]_{\rm o}\right)^{1/2} - 1}{\left(1 + 8K[{\bf B}]_{\rm o}\right)^{1/2} + 1}$$
(10)

where ε_d is the molar absorptivity of the dimer band. Dividing the first and last terms in Eq. (10) by $\varepsilon_d b[B]_o/2$, and making use of the fact that if $\frac{a}{b} = \frac{c}{d}$ then $\frac{a+b}{a-b} = \frac{c+d}{c-d}$, we obtain

$$\frac{\varepsilon_{\rm d} b[\mathbf{B}]_{\rm o} + 2A_{\rm d}}{\varepsilon_{\rm d} b[\mathbf{B}]_{\rm o} - 2A_{\rm d}} = (1 + 8K[\mathbf{B}]_{\rm o})^{1/2} \tag{11}$$

Taking squares on both sides of Eq. 11 and making use of the fact that if $\frac{a}{b} = \frac{c}{d}$ then $\frac{a-b}{b} = \frac{c-d}{d}$, we obtain

$$\frac{\varepsilon_{\rm d} b A_{\rm d}}{K} = \left(\varepsilon_{\rm d} b [\mathbf{B}]_{\rm o} - 2A_{\rm d}\right)^2 \tag{12}$$

We then take square roots on both sides then divide them by $[B]_0$ on both sides to transform Eq. 12 into a linear form which is suitable for the data treatment of the integrated absorbance of the dimer band:

$$\frac{2A_{\rm d}}{[{\rm B}]_{\rm o}} = \varepsilon_{\rm d} b - \left(\frac{\varepsilon_{\rm d} b}{K}\right)^{1/2} \frac{(A_{\rm d})^{1/2}}{[{\rm B}]_{\rm o}}$$
(13)

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The experimental data of $y = \frac{2A_d}{|\mathbf{B}|_o}$ versus $x = \frac{(A_d)^{1/2}}{|\mathbf{B}|_o}$ is then fitted to Eq. 13 to obtain the best values of $\varepsilon_d = \frac{q}{b}$ and $K = \frac{q}{p^2}$ from the slope *p* and intercept *q* of the regressed line.

It is remarkable that the dimerization constant K can be independently determined either from the data treatment of the monomer band based on Eq. 9, or from the data treatment of the dimer band based on Eq. 13.

3 Experimental Section

The solute 2,2-dimethyl-3-ethyl-3-pentanol (\geq 98 %) and the solvent *n*-octane (quoted \geq 97 % in catalogue, actual assay 98.989 % by supplier) were purchased from Aldrich and Tedia, respectively, and were used without further purification. We prepared, using an analytical balance, using vials and microsyringes, the sample concentrations on the molality scale. The molalities were then converted into concentrations on the molarity scale based on the density of the solvent at different temperatures (288 K, 0.707 g·cm⁻³; 298 K, 0.701 g·cm⁻³; 308 K, 0.699 g·cm⁻³; 318 K, 0.683 g·cm⁻³), which were obtained from interpolating the published data [30, 31]. Subsequent calculations, listing and graphical plotting were then carried out in terms of molarity scale, as needed.

The spectra were taken with a FTIR spectrometer (Bio-Red Spc. 3200) with a CaF₂ cell window. The cell temperature was controlled by circulating the water from a thermostat around the cell jacket, and was measured by a thermocouple inserted into the jacket of the cell. The uncertainty of the cell temperature was estimated to be ± 1 K. A personal computer running the commercial software package PeakSolve (version 6.0, GRAMS, 2000, Galatic Industries Corp.) was connected to the spectrometer to perform the task of displaying, resolving the spectra, and calculating the integrated absorbance of the resolved spectra. Subsequent work, such as linear regression based on Eqs. 9 and 13 and plotting graphs, were carried out on another personal computer using the commercial software Mathematica.

4 Results and Discussion

Figure 1a presents a collection of the spectra exhibited by the OH fundamental stretching for different concentrations of 2,2-dimethyl-3-ethyl-3-pentanol in *n*-octane at 298 K. Two bands are observed for each concentration: the sharp one at higher wavenumber, ca. 3,630 cm⁻¹, corresponds to the OH fundamental stretching in the monomeric species, whereas the broad one at lower wavenumber, ca. 3,530 cm⁻¹, corresponds to that for the dimeric species. These two bands are not well separated. They were resolved, and their respective integrated absorbance were calculated with the commercial software PeakSolve (version 6.0, GRAMS, 2000, Galatic Industries Corp.) A typical result of this resolution is presented in Fig. 1b for the concentration 0.2798 mol·L⁻¹ at 298 K. The calculated integrated absorbance for the monomer bands, A_m , and those for the dimer bands, A_d , are listed in Table 1 for different concentrations at the four studied temperatures, 288, 298, 308, and 318 K.

From Table 1 it is seen that, at a given temperature, both $A_{\rm m}$ and $A_{\rm d}$ increase with the initial solute concentration, [B]_o. However, the observation that $A_{\rm d}/A_{\rm m}^2$ remains fairly constant over the experimental range of [B]_o at a given temperature, as shown in Fig. 2, strongly implies that monomer–dimer self-association is dominant over oligomerization. This is plausible, due to the fact that $K = \frac{[B_2]}{[B]^2} = \frac{A_d/\epsilon_d}{A_{m}^2/\epsilon_m^2}$, or $\frac{A_d}{A_{m}^2} = \frac{K\epsilon_d}{\epsilon_m^2}$, is constant for a given

Absorbance



3740 3660 3580 3500 3420 3340 Wavenumber / cm⁻¹

Fig. 1 a Collection of the spectra of the fundamental OH stretching bands for different concentrations of 2,2-dimethyl-3-ethyl-3-pentanol in n-octane at 298 K: from bottom to top: 0.2108, 0.3519, 0.4906, 0.6368, and 0.7692 mol L^{-1} , **b** curve fit for 0.2798 mol L^{-1} 2,2-dimethyl-3-ethyl-3-pentanol in *n*-octane at 298 K

temperature since the parameters on the right hand side are only functions of temperature. Once $A_{\rm m}$ and $A_{\rm d}$ are available for different concentrations at a given temperature, we are in a position to calculate $\varepsilon_{\rm m}$ and K at this temperature from the plot based on Eq. 9, and $\varepsilon_{\rm d}$ and K at this temperature from the plot based on Eq. 13.

Shown in Fig. 3 are such plots for the monomer bands based on Eq. 9 and in Fig. 4 for the dimer bands based on Eq. 13 at the four temperatures. The values of $\varepsilon_{\rm m}$, $\varepsilon_{\rm d}$, and K determined from the slope and intercept of the regressed lines at various temperatures are listed in Table 2. The values of K at different temperatures can be used to obtain the standard enthalpy (ΔH°) and entropy (ΔS°) of dimerization through a van't Hoff plot. For this system, this is done either using K from A_m or from A_d . Both plots are juxtaposed in Fig. 5 for better visual comparison. The errors associated with K, $\varepsilon_{\rm m}$ and $\varepsilon_{\rm d}$ listed in Table 2, were calculated based on a standard method [32] with the assumption of equal

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288 K			298 K			308 K			318 K		
Conc./ nol·L ⁻¹	Monomer absorbance	Dimer absorbance	Conc./ mol·L ⁻¹	Monomer absorbance	Dimer absorbance	$Conc./mol\cdot L^{-1}$	Monomer absorbance	Dimer absorbance	$Conc./mol·L^{-1}$	Monomer absorbance	Dimer absorbance
.1443	1.65	1.24	0.2108	2.14	1.68	0.1394	1.41	0.52	0.2060	1.93	0.74
0.2165	2.27	2.42	0.2798	2.74	2.78	0.2113	2.05	1.14	0.2734	2.49	1.25
).2843	2.95	3.79	0.3519	3.31	4.12	0.2781	2.61	1.87	0.3438	3.09	1.92
.3550	3.49	5.64	0.4217	3.85	5.51	0.3476	3.21	2.76	0.4121	3.59	2.68
.4252	4.01	7.28	0.4906	4.33	7.06	0.4174	3.79	3.79	0.4793	4.10	3.47
.4977	4.56	9.26	0.5651	4.88	8.83	0.4891	4.33	4.99	0.5521	4.64	4.58
).5673	5.06	11.51	0.6368	5.35	10.65	0.5547	4.75	6.16	0.6222	5.18	5.54
).6387	5.53	13.51	0.7006	5.79	12.24	0.6240	5.24	7.53	0.6846	5.62	6.33
0.7086	5.88	15.82	0.7692	6.21	14.26	0.6925	5.70	8.79	0.7516	6.07	7.55
7797.0	6.33	18.47	0.8421	6.67	16.27	0.7629	6.19	10.42	0.8228	6.55	8.80

Table 1 The integrated absorbances of OH stretching bands from the monomer A_m, and those of dimer A_d, for different initial concentrations of 2,2-dimethyl-3-ethyl-3pentanol in n-octane at different temperatures



Fig. 2 Plot for comparing the theoretical *lines* calculated from $\frac{A_4}{A_m^2} = \frac{\epsilon_4 b|B_2|}{\epsilon_m^2 b^2|B|^2} = \left(\frac{\epsilon_4}{\epsilon_m^2 b}\right) K$ versus [B]_o, with the corresponding experimental data of $\frac{A_4}{A_m^2}$ versus [B]_o of 2,2-dimethyl-3-enthyl-3-pentanol in *n*-octane at different temperatures: (*filled circle*) 288 K, (*open circle*) 298 K, (*filled square*) 308 K, and (*open square*) 318 K

variances for $A_{\rm m}$ and $A_{\rm d}$. The errors associated with ΔH° and ΔS° were also calculated based on a standard method [32] with unequal variances of *K* as listed in Table 2 at each temperature. The final value of *K* at each temperature was taken as the geometric mean of the *K* values from Eq. 9 and from Eq. 13. The results are 0.640 \pm 0.033, 0.428 \pm 0.010, 0.283 \pm 0.008, and 0.194 \pm 0.013 L·mol⁻¹, respectively, at 288, 298, 308, and 318 K.

In order to assess the goodness of the determination we also plotted, in Fig. 6a, the data points of $y = A_m$ versus $x = [B]_o$, juxtaposed with the theoretical curves calculated from Eq. 7, by inserting the determined values of *K* and ε_m for each temperature. Similar plots for $y = A_d$ versus $x = [B]_o$ and the corresponding theoretical curves calculated from Eq. 10 are presented in Fig. 6b.

As compared to the spectra taken by Huelsekopf and Ludwig [20] for the pure liquid at various temperatures, our monomer bands are basically in agreement with theirs in peak position, except that the free OH stretching of the hydrogen-bond associated complex disappears in our dilute solution case. It is a reasonable inference that the dimers in our case are mostly of the cyclic form. The bands at lower wavenumber are in disagreement, both in shape and position, due to the fact that in pure liquid state oligomerization may prevail.

It is expected that the values of K determined, either from A_m based on Eq. 9 or from A_d based on Eq. 13, should be equal, since the same physical entity is referred to. As a consequence the values of ΔH° or ΔS° that are determined either from A_m or from A_d are also expected to be equal. This fact suggests that the discrepancy between the values of K determined from A_m and from A_d at the same temperature, and the respective ΔH° or ΔS° from A_m and A_d , can serve as a consistency check.

It is obvious from Fig. 1 that the monomer bands are much sharper than the dimer bands within the experimental concentration range. Therefore, it is to be expected that the data determined from the monomer bands will be more precise than those from dimer bands.



Fig. 3 Linear plot based on Eq. 9 to treat the data of monomer bands for different concentrations of 2,2dimethyl-3-ethyl-3-pentanol in *n*-octane at different temperatures: (*filled circle*) 288 K, (*open circle*) 298 K, (*filled square*) 308 K, and (*open square*) 318 K



Fig. 4 Linear plot based on Eq. 13 to treat the data of dimer bands for different concentrations of 2,2dimethyl-3-ethyl-3-pentanol in *n*-octane at different temperatures: (*filled circle*) 288 K, (*open circle*) 298 K, (*filled square*) 308 K, and (*open square*) 318 K

However, this discrepancy should become smaller as the solute concentration increases, since the dimer bands become sharper at higher solute concentration. On account of this, it may be inappropriate to treat equilibrium constants from monomer band and from dimer band on equal footing. Nevertheless, at least the data from dimer bands can serve as a supplement to those determined from monomer bands.

ties	(ɛ _d),	and	dimerization	const

Temp/K	From monomer b	ands	From dimer band	From dimer bands	
	$K/L \cdot mol^{-1}$	$\varepsilon_{\rm m}/{\rm L}\cdot{\rm cm}^{-1}\cdot{\rm mol}^{-1}$	$K/L \cdot mol^{-1}$	$\varepsilon_d/L \cdot cm^{-1} \cdot mol^{-1}$	
288	0.629 ± 0.028	1311 ± 21	0.651 ± 0.039	12152 ± 327	
298	0.418 ± 0.008	1166 ± 7	0.438 ± 0.012	11715 ± 156	
308	0.294 ± 0.010	1082 ± 8	0.273 ± 0.009	11319 ± 190	
318	0.202 ± 0.007	1004 ± 6	0.186 ± 0.019	10851 ± 562	
$\Delta H^{\circ}/\text{kJ}\cdot\text{mol}^{-1}$	-28.61 ± 3.88		-31.81 ± 3.88		
$\Delta S^{\circ}/J \cdot mol^{-1} \cdot K^{-1}$	-103.2 ± 13.0		-113.9 ± 13.1		

Table 2 Molar monomer absorptivities (ε_m), molar dimer absorptivities (ε_d), and dimerization constants (*K*) for 2,2-dimethyl-3-ethyl-3-pentanol in *n*-octane at different temperatures, and the standard enthalpy (ΔH°) and entropy (ΔS°) of dimerization obtained from the monomer and dimer bands

Fig. 5 van't Hoff plot to obtain ΔH° and ΔS° for the dimerization of 2,2-dimethyl-3-ethyl-3-pentanol in *n*-octane from the dimerization constants obtained at different temperatures from the monomer band (*filled circle*), and from the dimer band (*open circle*)



The 2 % impurities in the solute may contain alkyl phenols, straight-chained and branched alcohols, which were introduced during the synthetic process. These sorts of alcohol or phenol impurities will contribute to the area of both monomer and dimer bands through the OH stretching of free monomers or hydrogen-bonded associates. This contribution will accordingly induce some errors in our determination.

To the best of our knowledge, most IR investigations of monomer–dimer self-association employed only the monomer bands to determine ε_m and *K*. For example, in the study of the self-association of methanol, ethanol, and *tert*-butanol in CCl₄ solution, Liddle and Becker [33] obtained *K* from the limiting slope of a plot of apparent absorptivity, $A_m/[B]_o$, against [B]_o, through the equation:

$$\lim_{[\mathbf{B}]_{0}\to 0} \left(\frac{d\left(\frac{A_{\mathbf{m}}}{[\mathbf{B}]_{0}}\right)}{d[\mathbf{B}]_{0}} \right) = -2K\varepsilon_{\mathbf{m}}$$
(14)

with $\varepsilon_{\rm m}$ obtained from another limiting slope:

$$\lim_{|\mathbf{B}|_0 \to 0} \frac{dA_{\mathrm{m}}}{d[\mathbf{B}]_{\mathrm{o}}} = \varepsilon_{\mathrm{m}} \tag{15}$$

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Fig. 6 Plot for comparison of the experimental data with theoretical curves: $\mathbf{a} A_m$ versus [B]_o with the *curves* calculated based on Eq. 9, $\mathbf{b} A_d$ versus [B]_o with the *curves* calculated based on Eq. 13, at various temperatures: (*filled circle*) 288 K, (*open circle*) 298 K, (*filled square*) 308 K, and (*open square*) 318 K

A similar method had also been adopted by Gutowsky and coworker [34] in an NMR study of the self-association of hindered phenols. The disadvantage of this method lies in the fact that a small error of the slope could induce a considerable amount of error in the determination of ε_m and *K* if the slope is steep. Prokopenko et al. [35], in an IR study of the effect of the ring size on the self-association of lactams, fitted the data of $y = A_m$ versus $x = [B]_0$ to the equation:

$$A_{\rm m} = \varepsilon_{\rm m} b \frac{\sqrt{1 + 8K[\mathbf{B}]_{\rm o}} - 1}{4K}$$
(16)

which is derivable from Eq. 7, to obtain *K* and ε_m with the help of the Levenberg–Marquardt nonlinear method. In a study of the self-association of lactams using the IR overtone, Luck [36] fitted the data of $y = A_m/[B]_o$ versus $x = A_m^2/[B]_o$ to a linear equation, which in our notation appears in the form:

$$\frac{A_{\rm m}}{[{\rm B}]_{\rm o}} = \varepsilon_{\rm m} - \frac{2K}{\varepsilon_{\rm m}} \left(\frac{A_{\rm m}^2}{[{\rm B}]_{\rm o}}\right) \tag{17}$$

This equation appears as Eq. 9 in page 163 of the above reference, it is also derivable from Eq. 8, and is used to obtain ε_m and K from the slope and intercept of the regressed line. It is remarkable that all three above-mentioned methods leave the dimer band unused. Without treating the data of the dimer bands, it is impossible to obtain the molar absorptivity of the dimer. On the contrary, in addition to the data treatment of A_m , our new approach also treats the data of A_d based on Eq. 13, allowing us to obtain both ε_d and K. Since the same K can be obtained from two independent, different sources A_m and A_d , the differences between these two determinations offers an additional opportunity to examine the consistency of the determination.

5 Conclusions

Self-association of 2,2-dimethyl-3-ethyl-3-pentanol in *n*-octane was used as a model system to illustrate a new approach to treat the data from both monomer and dimer bands. The solute compound was chosen in view of the fact that the hindrance from the neighboring bulky side chains restricts the solute molecular association to dimerization. This restriction was further examined by checking the constancy of A_d/A_m^2 over [B]_o before processing the data treatment. Two new equations, Eqs. 9 and 13, have been derived to treat the integrated absorbance of the monomer bands and those of dimer bands, A_m and A_d , respectively. The data treatment of A_m based on Eq. 9 allows us to obtain the molar absorptivity of the monomer bands, ε_m , and the dimerization constant, K, at a given temperature. The data treatment of the dimer bands allows us to obtain the molar absorptivity of the dimer bands, k_a , and the dimerization constant, K, at a given temperature. Since the same dimerization constant at a given temperature can be obtained from two independent data sources, A_m and A_d , the disparity between two determinations of K provides a vehicle to assess the precision of the determination.

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