A New Approach for Titration Calorimetric Data Analysis on the Binding of Magnesium Ion with Myelin Basic Protein

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Abstract The interaction of the myelin basic protein (MBP) from the bovine central nervous system with divalent magnesium ion was studied by isothermal titration calorimetry at 27 °C in aqueous solution. A simple rapid method for determination of the dissociation binding constants for Mg²⁺-MBP interaction was introduced using the isothermal titration calometric data. The binding isotherm for Mg²⁺-MBP interaction is easily obtained by carrying out a titration calorimetric experiment using only one set of concentrations of MBP. There are two identical independent intrinsic association constants equal to 0.021 μ mol·L⁻¹ in the first- and second-binding sites, respectively.

Keywords Myelin basic protein · Magnesium · Isothermal titration calorimetry · Binding parameters

1 Introduction

The energetics of biochemical reactions or molecular interactions at constant temperature are measured by Isothermal Titration Calorimetry (ITC) [1–3]. ITC gives invaluable information about biomacromolecule-ligand interactions. During the last two years we attempted to study the metal ion binding chemistry of different proteins [4–12]. They will change the conformational stability and form aggregates. The importance of metal ions such as Ca²⁺ and Mg²⁺ in determining the stability of proteins is widely reported [13–18]. There are some approximations in previous theories [18] which result in rough average values of binding parameters including a lot of errors. We have introduced a novel method which enables us to determine the binding parameters with minimal errors. This method is successfully applied to the analysis of Mg²⁺ binding MBP.

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2 Materials and Method

2.1 Materials

MBP from bovine central systems (CNS) was obtained from Sigma Chemical Co. Magnesium nitrate was purchased from Merck Co. Protein concentrations were determined from absorbance measurements at 277 nm in 1 cm quartz cuvettes. All other materials and reagents were of analytical grade, and solutions were made from double-distilled water.

2.2 Method

The isothermal titration micro-calorimetric experiments were performed with the fourchannel commercial micro calorimetric system, Thermal Activity Monitor 2277, Thermometric, Sweden. The titration vessel was made from stainless steel. The magnesium nitrate solution (500 µmol·L⁻¹) was injected by use of a Hamilton syringe into the calorimetric titration vessel, which contained 1.8 mL MBP (13.5 μ mol·L⁻¹). Thin (0.15 mm inner diameter) stainless steel hypodermic needles, permanently fixed to the syringe, reached directly into the calorimetric vessel. Injection of magnesium nitrate solution into the perfusion vessel was repeated 30 times, with 30 μ L per injection. The calorimetric signal was measured by a digital voltmeter which was part of a computerized recording system. The heat of each injection was calculated by the "Thermometric Digitam 3" software program. The enthalpy of dilution of the magnesium solution was measured as described above except MBP was excluded. The enthalpies of dilution of the magnesium solutions were subtracted from the enthalpy of Mg^{2+} -MBP interaction. The enthalpies of dilution of MBP are negligible. The micro calorimeter was frequently calibrated electrically during the course of the study. The molecular weight of MBP was taken to be 18500 Da. The enthalpies of Mg²⁺-MBP interactions were calculated in $kJ \cdot mol^{-1}$ and are shown in Fig. 1.



3 Results and Discussion

We have shown previously that the enthalpies of the solute-solvent (Mg^{2+} -MBP-water in this case) interactions in the aqueous solvent (Mg^{2+} -water in the present case) system, can be accounted for quantitatively in terms of three factors: preferential solvation by the components of a mixed solvent, weakening or strengthening of solvent-solvent bonds by the solute and the change in the enthalpy of the solute-solvent interactions [19–28]. This treatment leads to:

$$\Delta H = \Delta H_{\max} x'_{\mathrm{B}} - \delta^{\theta}_{\mathrm{A}} (x'_{\mathrm{A}} L_{\mathrm{A}} + x'_{\mathrm{B}} L_{\mathrm{B}}) - (\delta^{\theta}_{\mathrm{B}} - \delta^{\theta}_{\mathrm{A}}) (x'_{\mathrm{A}} L_{\mathrm{A}} + x'_{\mathrm{B}} L_{\mathrm{B}}) x'_{\mathrm{B}}$$
(1)

The parameters $\delta_A^{\theta} = (\alpha n + \beta N)_A^{\theta}$ and $\delta_B^{\theta} = (\alpha n + \beta N)_B^{\theta}$ are the net effects of the solute (MBP in this case) on the water-Mg²⁺ bonds in the water-rich domain and Mg²⁺-rich region, respectively, with αn resulting from the formation of a cavity wherein *n* solvent molecules become the nearest neighbors of the solute and βN reflects the enthalpy change from strengthening or weakening of solvent–solvent bonds of *N* solvent molecules ($N \ge n$) around the cavity ($\beta < 0$ indicates a net strengthening of the solven–solvent bonds). The constants α and β represent the fraction of the enthalpy of water–Mg²⁺ interaction associated with cavity formation or restructuring, respectively. The superscript θ in all cases refers to the quantities at infinite dilution of the solute. p < 1 or p > 1 indicate a preference for water or Mg²⁺, respectively; p = 1 indicates random solvation; and the x'_B are the local mole fractions of the water and surfactant in the vicinity of the lysozyme or solvation sphere, where the solvent molecules are the nearest neighbors of the solute, which can be expressed as follows:

$$x'_{\rm B} = \frac{px_{\rm B}}{x_{\rm A} + px_{\rm B}} = \frac{\nu}{g} \tag{2}$$

where x_A and x_B are bulk mole fractions and we can express them as the total Mg²⁺ concentration divided by the maximum concentration of Mg²⁺ upon saturation of all MBP as follows:

$$x_{\rm B} = \frac{[{\rm M}g^{2+}]_{\rm T}}{[{\rm M}g^{2+}]_{\rm max}}, \quad x_{\rm A} = 1 - x_{\rm B}$$
(3)

 $[Mg^{2+}]_T$ is the total concentration of surfactant and $[Mg^{2+}]_{max}$ is the maximum concentration of the Mg^{2+} upon saturation with respect to MBP. In general, there will be "g" sites for binding of Mg^{2+} per MBP molecule and ν is defined as the average moles of bound Mg^{2+} per mole of total MBP. L_A and L_B are the relative partial molar enthalpies and can be calculated from heats of dilution of Mg^{2+} in water, ΔH_{dilut} , as follows:

$$L_{\rm A} = \Delta H_{\rm dilut} + x_{\rm B} \left(\frac{\partial \Delta H_{\rm dilut}}{\partial x_{\rm B}} \right), \quad L_{\rm B} = \Delta H_{\rm dilut} + x_{\rm A} \left(\frac{\partial \Delta H_{\rm dilut}}{\partial x_{\rm B}} \right) \tag{4}$$

The enthalpies of Mg^{2+} -MBP interactions, ΔH , were fitted by Eq. 1 over the whole range of Mg^{2+} compositions. In this procedure the only adjustable parameter, p, was changed until the best agreement between the experimental and calculated data was approached over the whole range of solvent compositions (Fig. 1). Thermodynamic parameters for Mg^{2+} -MBP interactions recovered from Eq. 1 are listed in Table 1. The experimental enthalpies of Mg^{2+} -MBP interactions and calculated data from Eq. 1 are compared in Fig. 1. The agreement between the experimental enthalpies and calculated data via Eq. 1 provides good support for this equation.

Table 1 Thermodynamic parameters for Mg^{2+} -MBP interactions recovered from Eq. 1. $\Delta\Delta H_{12} < 0$ indicates that the interaction of the MBP with Mg^{2+} is stronger than with water. The precision is ± 0.005 or better

[MBP] μ mol·L ⁻¹	р	$\delta^o_{ m A}$	$\delta^o_{ m B}$	$\Delta \Delta H_{12}$
13.5	5.005	0.434	0.476	-17.774

 φ is the fraction of MBP molecule undergoing complexation with Mg²⁺ which can be expressed as follows:

$$\varphi = \frac{(\Delta H_{\rm W} - \Delta H)}{(\Delta H_{\rm W} - \Delta H_{\rm max})} \tag{5}$$

where $\Delta H_{\rm W}$ is the enthalpy of Mg²⁺-MBP interactions in the water-rich domain and $\Delta H_{\rm max} = \Delta \Delta H_{12}^{\rm o} + \delta_{\rm B}^{\rm o} \Delta H_{\rm Mg(NO_3)_2}^{\rm o} - \delta_{\rm A}^{\rm o} \Delta H_{\rm W}^{\rm o*}$ represents the heat value upon saturation of all MBP. $\Delta \Delta H_{12}^{\rm o}$ is the difference between the enthalpies of the Mg²⁺-MBP and water-MBP interactions. $\Delta \Delta H_{12} > 0$ indicates that the interaction of the MBP with Mg²⁺ is weaker than with water. $\Delta H_{\rm W}^{\rm o*}$ is the enthalpy of condensation of pure water (-44.700 kJ·mol⁻¹) and $\Delta H_{\rm Mg(NO_3)_2}^{\rm s}$ is the enthalpy of solution of magnesium nitrate in water (-3.400 kJ·mol⁻¹). The equilibrium constant values, $K_{\rm d}$, as a function of Mg²⁺ concentration can be calculated as follows:

$$K_{\rm d} = \frac{\varphi}{1 - \varphi} \tag{6}$$

The equilibrium constants, K_d , for successive replacement of the Mg²⁺ ions by water molecules are as follows:

$$K_{\rm d} = x_{\rm A}^g - \sum_{i=1}^g \beta'_i \frac{x_{\rm B}^i}{x_{\rm A}^{i-g}} \tag{7}$$

In fact the K_d values are the equilibrium constants for the equilibria:

$$MBP(H_2O)_{g-i}(Mg^{2+})_i + iH_2O \rightleftharpoons MBP(H_2O)_g + iMg^{2+},$$

$$K_d = \frac{[MBP(H_2O)_g][Mg^{2+}]^i}{[MBP(H_2O)_{g-i}(Mg^{2+})_i]}$$
(E1)

The K_d values obtained from Eq. 6, were fitted by Eq. 7 using a computer program for nonlinear least-squares fitting. Therefore, we can approximate the "g" value simply as g = 2in this work. The β'_i values are the equilibrium products for the equilibria E1. Finally, the ν values can be calculated at any concentration of Mg²⁺ via Eq. 3. The binding parameters obtained from this method are listed in Tables 2 and 3 and shown graphically in Figs. 2 and 3. The Gibbs energies as a function of the Mg²⁺ concentration can be obtained as follows:

$$\Delta G = -RT \ln K_a \tag{8}$$

where K_a is the association equilibrium constant $(1/K_d)$ as a function of Mg²⁺ concentration. Gibbs energies, ΔG , calculated from Eq. 8 are shown graphically in Fig. 4. The ΔS values were calculated using the ΔG values and are shown in Fig. 5. Therefore, for the first

Table 2 β'_i 's values recovered from the coefficients of Eq. 7, for	Number of sites	$\beta'_i/\mu \text{mol} \cdot L^{-1}$
2 individual sites of MBP	1	3.272 ± 0.026
	2	3.383 ± 0.013

Table 3	K_{d} and	v values	as a
function of	of Mg ²⁺	concent	rations
for Mg ²⁺	-MBP i	nteraction	ns. The
precisions	s are ± 0	.005 or b	etter

$[Mg^{+2}]_T/\mu mol{\cdot}L^{-1}$	$K/\mu \text{mol}\cdot L^{-1}$	ν
8.197	0	0.411
23.809	0.342	0.909
52.239	0.768	1.200
65.217	1.289	1.391
77.465	1.927	1.526
89.041	2.710	1.626
100.000	3.686	1.703
110.389	4.927	1.765
120.253	6.564	1.815
129.626	8.784	1.857
138.554	12.015	1.892
147.059	17.060	1.922
155.172	26.209	1.948
166.667	47.441	1.970



temperature.

K_d/micro molar

If the multiple binding sites on a macromolecule are identical and independent, the ligand binding sites can be reproduced by a model system of monovalent molecules $((MBP)_g \rightarrow gMBP)$ with the same set of dissociation constant, $(K_d)_{max}$, values. Setting g = i in Eq. E1, leads to:

$$MBP(Mg^{2+})_{g} + gH_{2}O \rightleftharpoons MBP(H_{2}O)_{g} + gMg^{2+},$$

$$(K_{d})_{max} = \frac{[MBP(H_{2}O)_{g}][Mg^{2+}]^{g}}{[MBP(Mg^{2+})_{g}][H_{2}O]^{g}}$$
(E2)





Equilibrium E2 is a special case of the equilibrium E1 in the Mg²⁺-rich domain where MBP is saturated by Mg²⁺ ions. In other words, the $(K_d)_{max}$ value corresponds to the point where $x'_B = 1$ (or the maximum ν value) (Fig. 3) and can be calculated easily. Binding parameters for the Mg²⁺-MBP interactions using the new model are as follows:

$$(K_{\rm d})_{\rm max} = 47.44 \,\mu{\rm mol}\cdot{\rm L}^{-1}$$
 $\Delta H = -15.10 \,\rm kJ\cdot{\rm mol}^{-1}$ $g = 2$

As we have reported previously, the free concentration of ligand, $[Mg^{2+}]_F$, can be calculated as follows:

$$[Mg^{2+}]_{\rm F} = [Mg^{2+}]_{\rm T} - \nu [MBP]_{\rm T}$$
⁽⁹⁾

The association binding constant, K_a , can be obtained from the Scatchard equation as follows:

$$\frac{\nu}{g-\nu} = K_{\rm a}[{\rm Mg}^{2+}]_{\rm F} \tag{10}$$

The association constant for the Mg²⁺-MBP interaction obtained from Eq. 10 is 0.021 μ mol·L⁻¹($K_d = 47.62 \mu$ mol·L⁻¹) [27, 28], which is in good agreement with the (K_d)_{max} value calculated from Eqs. 6 and 7. Equation 10 gives us the maximum value for K_d , but Eqs. 6 and 7 allows us to have the K_d values at every concentration of Mg²⁺ ions.

Figure 2 shows the dissociation constant as a function of the total concentration of Mg²⁺. The low K_d values in the water-rich domain reflect the higher affinity of MBP for Mg²⁺ in this domain which is in agreement with the $\Delta\Delta H_{12}$ value (Table 1). In the Mg²⁺-rich domain the slope of the curve in Fig. 2 varies sharply, indicating a lower affinity of MBP for Mg²⁺ in this region. The shallow slope of ν against K_d (Fig. 3), corresponds to the higher affinity of MBP in the water-rich domain followed by the steep slope which indicates a lower affinity of MBP in the Mg²⁺-rich domain. The negative Gibbs energies in the water-rich domain (Fig. 4) also indicate the higher affinity in this region.

We will arrive at the same conclusion using the solvation parameters recovered from Eq. 1 (Table 2). The δ_A^{θ} values reflect to the hydrophobic hydration of MBP, leading to the enhancement of the water structure. The greater the extent of this enhancement, the greater the stabilization of the MBP structure and the greater the δ_A^{θ} value and vise versa. In the Mg²⁺-rich region the δ_B^{θ} value is more positive (0.476) than that in water-rich domain, indicating that the Mg²⁺-MBP complex is more stable (lower affinity) in the Mg²⁺-rich region in comparison to the water-rich domain. The *p* value (5.005) shows the preferential tendency of Mg²⁺ for occupying the available sites on MBP.

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References

- 1. Saboury, A.A.: A review on the ligand binding studies by isothermal titration calorimetry. J. Iran. Chem. Soc. **3**, 1–21 (2006)
- Saboury, A.A.: New methods for data analysis of isothermal titration calorimetry. J. Therm. Anal. Cal. 72, 93–103 (2003)
- Saboury, A.A.: A simple method for determination of binding isotherm by isothermal titration calorimetry and its application to the interaction between Cu²⁺ and myelin basic protein. J. Therm. Anal. Cal. 77, 997–1004 (2004)
- Ataie, G., Moosavi-Movahedi, A.A., Saboury, A.A., Hakimelahi, G.H., Hwu, J.R., Tsay, S.C.: The enthalpy and enzyme activity of modified histidine residues of adenosine deaminase and diethyl pyrocarbonate complexes. Int. J. Biol. Macromol. 27, 29–33 (2000)
- Saboury, A.A., Atri, M.S., Sanati, M.H., Sadeghi, M.: Application of a simple calorimetric data analysis on the binding study of calcium ions by human growth hormone. J. Therm. Anal. Cal. 83, 175–179 (2006)
- Saboury, A.A., Atri, M.S., Sanati, M.H., Moosavi-Movahedi, A.A., Hakimelahi, G.H., Sadeghi, M.: Thermodynamic study on the interaction between magnesium ion and human growth hormone. Biopolymers 81, 120–126 (2006)
- Saboury, A.A., Atri, M.S., Sanati, M.H., Moosavi-Movahedi, A.A., Haghbeen, K.: Effects of calcium binding on the structure and stability of human growth hormone. Int. J. Biol. Macromol. 36, 305–309 (2005)
- Saboury, A.A., Kordbacheh, M., Sanati, M.H., Mizani, F., Shamsipur, M., Yakhchali, M.B., Moosavi-Movahedi, A.A.: Thermodynamics of binding copper ion by human growth hormone. Asian J. Chem. 17, 2773–2782 (2005)
- Atri, M.S., Saboury, A.A., Rezaei-Tavirani, M., Sanati, M.H., Moosavi-Movahedi, A.A., Sadeghi, M., Mansuri-Torshizi, H., Khodabandeh, N.: Binding properties and conformational change of human growth hormone upon interaction with Fe⁺³. Thermochim. Acta **438**, 178–183 (2005)
- Saboury, A.A., Ghourchaei, H., Sanati, M.H., Atri, M.S., Rezaei-Tawirani, M., Hakimelahi, G.H.: Application of a simple calorimetric data analysis on the binding study of calcium ions by human growth hormone. J. Therm. Anal. Cal. 83, 175–179 (2006)
- Hindmarsh, P.C., Brook, C.G.: Effect of growth hormone on short normal children. Br. Med. J. (Clin. Res. Ed.) 295, 573–583 (1987)
- de Voc, A.M., Ultsch, M., Kossiakoff, A.A.: Human growth hormone and extracellular domain of its receptor: Crystal structure of the complex. Science 255, 306–402 (1992)
- Filikov, A.V., Hayes, R.J., Luo, P., Stark, D.M., Chan, C., Kundu, A., Dahiyat, B.I.: Structural plasticity in a remodeled protein-protein interface. Protein Sci. 11, 1452–1461 (2002)
- Arakawa, T., Timasheff, S.N.: Mechanism of protein salting in and salting out by divalent cation salts: balance between hydration and salt binding. Biochem. 23, 5912–5923 (1984)
- MacLean, D.S., Qian, O.S., Middaugh, C.R.: Stabilization of proteins by low molecular weight multiions. J. Pharm. Sci. 91, 2220–2229 (2002)

- Nishimura, C., Uversky, V.N., Fink, A.L.: The effect of salts on the stability and folding of staphylococcal nuclease. Biochem. 40, 2113–2128 (2001)
- Curtis, R.A., Ulrich, J., Montaser, A., Blanch, H.W.: Protein-protein interactions in concentrated electrolyte solutions. Hofmeister-series effects. Biotechnol. Bioeng. 79, 367–380 (2002)
- Bordbar, A.K., Saboury, A.A., Housaindokht, M.R., Moosavi-Movahedi, A.A.: Statistical effects of the binding of ionic surfactant to protein. J. Coll. Interface Sci. 192, 415–419 (1997)
- Rezaei Behbehani, G.: Application of a new method to reproduce the enthalpies of transfer of NaI from water to aqueous methanol, ethanol and *i*PrOH solvent systems at 289.15 K. J. Bull. Korean Chem. Soc. 2, 238–240 (2005)
- Rezaei Behbehani, G.: Application of the new solvation theory to reproduce the enthalpies of transfer of LiBr, tetrabuthylammonium bromide and tetrapenthylamonium bromide from water to aqueous acetonitrile at 298 K. Acta Chim. Slovak. 52, 282–285 (2005)
- Rezaei Behbehani, G., Tazikeh, E., Saboury, A.A.: Using the new developed equation to reproduce the enthalpies of transfer of urea from water to aqueous ethanol, propan-1-ol and acetonitrile at 298 K. J. Bull. Korean Chem. Soc. 2, 208–210 (2006)
- Rezaei Behbehani, G., Ghamamy, S.: Enthalpies of transfer of formamide, N-methylformamide and N,N-dimethylformamide from water to aqueous acetonitrile mixtures at 298 K. Thermochim. Acta 444, 71–76 (2006)
- Rezaei Behbehani, G., Ghamamy, S., Waghorne, W.E.: Enthalpies of transfer of acetonitrile from water to aqueous methanol, ethanol and dimethyl sulphoxide mixtures at 298.15 K. J. Thermochim. Acta 448, 37–42 (2006)
- Rezaei Behbehani, G., Tazikeh, E., Saboury, A.A.: Using the Extension Coordination Model (ECM) to reproduce the enthalpies of transfer of tetraethylurea from water to aqueous ethanol, propan-1-ol and acetonitrile at 298 K. Acta Chim. Slovak. 53, 363–369 (2006)
- Rezaei Behbehani, G., Saboury, A.A.: Using a new solvation model for thermodynamic study on the interaction of nickel with human growth hormone. Thermochim. Acta 452, 76–79 (2007)
- Reza Bebahani, G., Dunnion, D., Falvey, P., Hickey, K., Meade, M., McCarthy, Y., Symons, M.C.R., Waghorne, W.E.: Nonelectrolyte solvation in aqueous dimethyl sulfoxide—A calorimetric and infrared spectroscopic study. J. Solution Chem. 29, 521–539 (2000)
- Rezaei Behbehani, G., Saboury, A.A.: A new method for thermodynamic study on the binding of magnesium with human growth hormone. J. Therm. Anal. Cal. 89, 852–861 (2007)
- Rezaei-Behbehani, G., Saboury, A.A., Bagheri, A.F.: A thermodynamic study on the binding of calcium ion with myelin basic protein. J. Solution Chem. 36, 1311–1320 (2007)