

# ELECTROTOPOLOGICAL STATE STUDIES OF COPPER(II) COMPLEXES WITH $\alpha$ -AMINOACIDATES

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## SUMMARY

Descriptor combinations, consisting of the electrotopological state (E-state) indexes of some skeletal groups, the first-order  $\chi$  molecular connectivity index and the logarithm of the statistical factor, are employed to describe the second stepwise formation constants of binary and ternary copper(II) complexes with  $\alpha$ -aminoacidate ligands. Descriptor sets formed by E-state values of skeletal groups belonging to the chelate rings,  $\chi$  and the logarithm of the statistical factor lead to the best regression equations. Results suggest that differences in stability over the series of metal chelates depend mainly upon variations in the coordination tendency of the carboxylate ligand groups.

Key Words: Copper(II), aminoacidate ligand, electrotopological state, connectivity index, intramolecular hydrophobic interactions.

## INTRODUCTION

Although topological indexes derived from the chemical graph theory have been widely used for the modeling of physicochemical and biological properties of organic compounds, at present, little attention has been paid on their possible applications to coordination chemistry. Quantitative structure-stability analyses of metal complexes with organic ligands through graph theoretical indices could contribute to a better understanding of the factors governing the formation of coordination compounds. Thus, for example, it might be expected that noncovalent interactions between coordinated ligands, such as hydrophobic interactions and steric effects, which are recognized to affect the stability of metal chelates<sup>1,2</sup>, could be appropriately described by topological indexes encoding structural information about size, shape or branching. In previous reports we have successfully studied the modeling of stability constants of metal complexes of biological interest<sup>3-6</sup> by means of electrotopological state indexes computed from hydrogen-suppressed graphs of the respective free neutral ligands<sup>7,8</sup>. Moreover, for  $\alpha$ -aminoacid ligands<sup>6</sup> we have observed significant correlations between the electrotopological state indices of the potential coordinating groups and the hydrophobicity scale of aminoacid side chains<sup>9</sup>. However, in the above mentioned modeling studies somewhat large sets of independent variables were taken relative to the number of available experimental data<sup>3-6</sup>, so that the probability of incidence of chance correlations with  $r^2 \geq 0.9$  was sometimes up to 6%<sup>10</sup>. In this work the modeling of the second stepwise formation constants of binary and ternary copper(II) complexes with  $\alpha$ -aminoacidate ligands was attempted by using descriptor sets consisting of the electrotopological state indices (E-states) of some selected skeletal groups<sup>7,8</sup> and the first-order  $\chi$  molecular connectivity index<sup>11</sup>. Both types of descriptors were computed from hydrogen-suppressed graphs of the metal complexes. However, seeing that these studies deal simultaneously with both binary and ternary complexes, the logarithm of the statistical factor (sf) was also included in the descriptor sets<sup>12</sup>. Further, a fitly restricted number of independent variables was considered, so as to keep the probability of occurrence of chance correlations with  $r^2 \geq 0.9$  at levels far lesser than 1%<sup>10</sup>.

## METHOD

Logarithms of the stepwise formation constants  $K_{\text{CuAB}}^B = [\text{CuAB}]/[\text{CuA}][\text{B}]$  at 25°C were taken from the literature<sup>2</sup>. In this expression A and B denote aminoacidate ligands, and  $A = B$  or  $A \neq B$  for binary and ternary complexes, respectively. The electrotopological states of the skeletal atoms of the [CuAB] complexes were calculated from the appropriate hydrogen-suppressed graphs by means of the expression

$$S_i = I_i + \Delta I_i$$

where  $I_i$  is the intrinsic state of atom  $i$  and  $\Delta I_i$  is the perturbation of this atom due to its interactions with the remaining atoms of the molecule<sup>7,8</sup>. Though the metal ion was considered as a vertex in the hydrogen-suppressed graphs, its intrinsic-state value was always set equal to zero because the E-state descriptors have originally been defined for the s-block and p-block elements only<sup>8</sup>. For the remaining skeletal groups the  $I_i$  values were calculated through the expression<sup>7,8</sup>

$$I_i = [(2/N)^2 \delta^v + 1] / \delta$$

where  $N$  is the principal quantum number, and  $\delta^v$  and  $\delta$  are the counts of valence electrons and  $\sigma$  electrons respectively, in the skeleton of the metal complex molecule. In turn,  $\delta^v$  and  $\delta$  were computed by the equations:

$$\delta^v = Z^v - h \quad \text{and} \quad \delta = \sigma - h$$

where  $Z^v$  is the number of valence electrons,  $\sigma$  is the count of electrons in  $\sigma$  orbitals and  $h$  is the number of bonded hydrogen atoms. The nonbonded contributions were evaluated by the expression<sup>7,8</sup>

$$\Delta I_i = S_i(I_i - I_j) / (r_{ij})^2$$

where  $r_{ij}$  is the count of atoms in the shorter path between atoms  $i$  and  $j$ , including both  $i$  and  $j$  (i.e. the graph distance plus one). According to the above quoted definitions, an  $S_i$  value encodes both electronic

and topological information because the intrinsic-state  $I_i$  reflects the valence-state electronegativity of atom  $i$  whereas the perturbation term  $\Delta I_i$  embodies the influence on such atom by all the other atoms in the molecular skeleton<sup>8</sup>.  $S_i$  values for equivalent skeletal groups were added together. First-order  ${}^1\chi$  molecular connectivity indexes for the metal complexes were calculated by the expression<sup>11</sup>:

$${}^1\chi = \sum (\delta_i \delta_j)^{-0.5}$$

where  $\delta$  is the number of adjacent skeletal groups,  $i$  and  $j$  correspond to the pairs of adjacent skeletal groups and the summation is over all bonds between skeletal groups. The statistical factor<sup>12</sup> was calculated from the corresponding definition:  $sf = (r+s)!/r!s!$ , where  $r$  and  $s$  are the stoichiometric subscripts in the general formulation  $[CuA_xB_y]$ . Thus,  $\log(sf)$  takes the values 0 and  $\log 2$  for binary and ternary complexes, respectively. Different sets of descriptors, consisting of the  $\Sigma S_i$  values of two or three selected skeletal groups, the  ${}^1\chi$  index and the logarithm of the statistic factor, were alternately correlated with the logarithms of the stepwise formation constants  $K_{CuAB}^B$ . Multiple regression analyses were performed by using the software Origin 4.0<sup>13</sup> on a DTK 486 computer.

## RESULTS AND DISCUSSION

Some of the calculated topological indexes have been collected in Table I. These selected indexes were found to provide the best descriptions of  $\log K_{CuAB}^B$ . In turn, the best four- and five-descriptor combinations were  $\{\Sigma S(-O-), \Sigma S(C_1), {}^1\chi, \log(sf)\}$  and  $\{\Sigma S(-O-), \Sigma S(C_1), {}^1\chi, \Sigma S(-NH_2-), \log(sf)\}$ , respectively. The corresponding regression equations are:

$$\begin{aligned} \log K_{CuAB}^B &= 0.9254 (\pm 0.2219) \Sigma S(C_1) - 0.6451 (\pm 0.3011) \Sigma S(-O-) + \\ &0.2015 (\pm 0.0762) {}^1\chi \dots \\ &\dots + 1.4133 (\pm 0.1333) \log(sf) + 12.1403 (\pm 2.4004) \end{aligned} \quad (1)$$

$r = 0.963$ ,  $s = 0.0767$ ,  $F = 48$

and

$$\begin{aligned} \log K_{CuAB}^B &= 1.8819 (\pm 0.4432) \Sigma S(C_1) - 1.0425 (\pm 0.4346) \Sigma S(-NH_2-) - \\ &0.4268 (\pm 0.2777) \Sigma S(-O-) \dots \\ &\dots + 0.2063 (\pm 0.0665) {}^1\chi + 1.4514 (\pm 0.1173) \log(sf) + \\ &13.9085 (\pm 2.2178) \end{aligned} \quad (2)$$

$r = 0.974$ ,  $s = 0.0669$ ,  $F = 52$

In these expressions the numbers in parentheses are the standard deviations on the respective regression coefficients. From the statistics for the regression equations, it can be noticed that both correlations are significant. In Table II the values calculated through equations 1 and 2 are compared with the experimental  $\log K_{CuAB}^B$  data taken from ref. 2. As can be seen, discrepancies between the experimental and computed values, expressed as percentage, fall in the ranges 0.00 - 2.02 and 0.00 - 2.17, for equations 1 and 2, respectively. The upper limits of these ranges are smaller than or, at least, similar to the standard errors currently reported for experimental data of formation constants of Cu(II)-aminoacidate complexes<sup>14</sup>. Equations 1 and 2 reasonably suggest that some skeletal groups belonging to the chelate rings are closely related to the internal factors determining the thermodynamic stability of the metal complexes in aqueous solution<sup>12</sup>. Thus, the E-states of -O- and  $C_1$  skeletal groups can be correlated with some thermodynamic parameters which are well recognized to affect such property. Accordingly, the  $\Sigma S(-O-)$  values give a good linear correlation with the hydrophobicity scale of aminoacid side chains<sup>9</sup>. The corresponding

regression equation is

$$\begin{aligned} \Sigma \Delta f_i &= 4.0209 (\pm 0.190) \Sigma S(-O-) - 34.4877 (\pm 1.705) \\ r &= 0.981, \quad s = 0.2859, \quad F = 448 \end{aligned} \quad (3)$$

where  $\Sigma \Delta f_i$  is the sum (HA plus HB) of the group contributions to the free energy transfer of aminoacid side chains from 100% organic solvent to water at 25°C<sup>29</sup>. The plot of  $\Sigma \Delta f_i$  against  $\Sigma S(-O-)$  is shown in Figure I. These results suggest that  $\Sigma S(-O-)$  encodes some information about the contributions of the intramolecular hydrophobic interactions to the stability of the copper(II) complexes herein considered. However, if this statement were accepted, the regression coefficient for  $\Sigma S(-O-)$  would be occurring with a *wrong* sign in both equations 1 and 2. Namely, these equations indicate that  $\log K_{CuAB}^B$  should decrease as  $\Sigma S(-O-)$  increases. Instead, intramolecular hydrophobic interactions are known to enhance the stability of binary and ternary metal complexes<sup>1,2</sup>. This inconsistency can be ascribed to the fact that the descriptors here used are mutually interdependent, i.e., they are correlated to each other to some extent<sup>15,16</sup>. In fact,  $\Sigma S(-O-)$  values give a significant positive linear correlation with  ${}^1\chi$  values ( $r = 0.876$ ). Accordingly, if  $\log K_{CuAB}^B$  is correlated with the set  $\{\Sigma S(C_1), \Sigma S(-O-), \Sigma S(-NH_2-), \log(sf)\}$ , i.e., if  ${}^1\chi$  is removed from equation 2, the regression coefficient of  $\Sigma S(-O-)$  occurs with a positive sign in the resulting regression equation

$$\begin{aligned} \log K_{CuAB}^B &= 1.2935 (\pm 0.5028) \Sigma S(C_1) - 1.0020 (\pm 0.5453) \Sigma S(-NH_2-) + \\ &0.3643 (\pm 0.1384) \Sigma S(-O-) \dots \\ &\dots + 1.5606 (\pm 0.1404) \log(sf) + 7.5203 (\pm 1.0368) \end{aligned}$$

$r = 0.956$ ,  $s = 0.084$ ,  $F = 39$

In order to discern the true contribution of  $\Sigma S(-O-)$  to  $\log K_{CuAB}^B$ , the descriptors involved in equations 1 and 2 were further subjected to orthogonalization<sup>15,16</sup>. In such process,  $\Sigma S(-O-)$  was taken as the first orthogonal descriptor ( $W_1$ ). The resulting regression equations are

$$\begin{aligned} \log K_{CuAB}^B &= 0.5624 \Omega_1 + 0.2100 \Omega_2 + 0.4497 \Omega_3 + 1.4133 \Omega_4 + 4.9642 \\ \text{and} \\ \log K_{CuAB}^B &= 0.5624 \Omega_1 + 0.2100 \Omega_2 + 0.4497 \Omega_3 + 1.4133 \Omega_4 - 1.0425 \Omega_5 \\ &+ 4.9642 \end{aligned}$$

respectively.

As it can be realized from these equations,  $\log K_{CuAB}^B$  indeed increases as  $\Sigma S(-O-)$  increases, in accordance with the observed correlation between  $\Sigma S(-O-)$  and the hydrophobicity scale of aminoacid side chains.

Orthogonalization processes taking successively  $\Sigma S(C_1)$ ,  ${}^1\chi$  and  $\Sigma S(-NH_2-)$  as first orthogonal descriptors, indicate that  $\log K_{CuAB}^B$  is directly proportional to each one of these indexes.

The occurrence of a good linear correlation between  $\Sigma S(-O-)$  and  $\Sigma \Delta f_i$  is rather surprising considering that it has been shown previously that, for alkyl ethers, the E-state values of bridging oxygen S(-O-) give a good linear correlation with the oxygen partial charges computed by the STO-3G method<sup>7</sup>. Moreover, among the remaining indexes, only  ${}^1\chi$  gives a somewhat significant linear correlation with  $\Sigma \Delta f_i$ , though being rather poorer than that obtained with  $\Sigma S(-O-)$ :

$$\Sigma \Delta f_i = 0.9148 {}^1\chi - 4.7276; \quad r = 0.815, \quad s = 0.842, \quad F = 36 \quad (\text{compare with equation 3})$$

The best descriptions of  $\Sigma \Delta f_i$  by means of two-index sets were



found to be

$$\Sigma\Delta f_1 = 3.7732(\pm 0.1538)\Sigma S(-O-) + 0.9486(\pm 0.2384)\Sigma S(C_1) - 31.3276(\pm 1.4917) \quad (4)$$

$$r = 0.990, \quad s = 0.2117, \quad F = 417$$

and

$$\Sigma\Delta f_1 = 0.9456(\pm 0.0375)^1\chi + 3.5786(\pm 0.2125)\Sigma S(C_1) - 1.4035(\pm 0.3286) \quad (5)$$

$$r = 0.990, \quad s = 0.2174, \quad F = 440$$

The statistical parameters for these regression equations indicate that both correlations are highly significant. As it can be realized, these descriptions also involve E-states of skeletal groups belonging to the carboxylate ligand group. Moreover, according to equation 5,  $\Sigma S(C_1)$  and  ${}^1\chi$  should be encoding complementary structural information about  $\Sigma\Delta f_1$ .

On the other hand, the two-descriptor set containing the E-state values for -O- and  $C_1$  skeletal groups was also found to correlate significantly with the ligand basicities through  $pK_{a1}$  and the pH at the isoelectric point,  $pI = 1/2(pK_{a1} + pK_{a2})$ . Here, thermodynamic values for  $pK_{a1}$  and  $pK_{a2}$  were considered<sup>14</sup>. The respective regression equations are:

$$pK_{a1}^{av} = 1/2[pK_{a1}(H_2A^+) + pK_{a1}(HB^+)] \\ = 0.3600(\pm 0.0231)\Sigma S(C_1) - 0.1089(\pm 0.0149)\Sigma S(-O-) + 3.5755(\pm 0.1445) \quad (6)$$

$$r = 0.967, \quad s = 0.0205, \quad F = 122$$

$$pI^{av} = \frac{1}{2}[pI(HA) + pI(HB)] \\ = 0.7437(\pm 0.0427)\Sigma S(C_1) - 0.2130(\pm 0.0275)\Sigma S(-O-) + 8.5065(\pm 0.2668) \quad (7)$$

$$r = 0.973, \quad s = 0.0379, \quad F = 152$$

In both expressions the superscript *av* stands for the average property of the aminoacids HA and HB, as explicitly shown. The statistics for these regression equations indicates that both correlations are fairly significant. In Table 3 the calculated  $pK_{a1}^{av}$  and  $pI^{av}$  values are compared with the corresponding thermodynamic data. In equations 6 and 7 the parameter  $\Sigma S(C_1)$  would be reflecting partly the average electron density on the carboxylate ligand groups<sup>7</sup>. Accordingly, from orthogonalization of the descriptors involved in these equations, taking  $\Sigma S(C_1)$  as the first orthogonal descriptor, it can be realized that both  $pK_{a1}^{av}$  and  $pI^{av}$  increase as a function of  $\Sigma S(C_1)$ . This could be related to an increase in the average basicity of the carboxylate ligand groups and, hence, to an increase in  $\log K_{CuAB}^B$ <sup>17</sup>.

It should be pointed out that even using  $\Sigma S(C_1)$  as a single descriptor of  $pK_{a1}^{av}$  and  $pI^{av}$ , the resulting correlations are still somewhat significant, being characterized by correlation coefficients of 0.853 and 0.872, respectively.

The set of descriptors  $\{\Sigma S(-NH_2-), \Sigma S(-O-)\}$  was also found to give a rather significant correlation with the  $pI^{av}$  values, the corresponding statistical parameters being  $r = 0.958$ ,  $s = 0.0473$ ,  $F = 94$ . However, as can be noticed, such a correlation is rather poorer than the one obtained by the  $\{\Sigma S(C_1), \Sigma S(-O-)\}$  descriptor set (see equation 7). Good linear correlations between  $\log K_{CuAB}^B$  and  $(pK_{a1} + pK_{a2})$  have previously been reported for  $[Cu(2,2'-bipyridine)(aminoacidate)]^{+}$  ternary complexes<sup>18</sup>. Such results provide some support to the reliability of the relationships, found in this study, between the above mentioned two-descriptor sets and the thermodynamic  $pK_{a1}^{av}$  and  $pI^{av}$  data.

Since the second stepwise ionization of aminoacids involves deprotonation of an  $NH_3^+$  group, it could be expected  $\Sigma S(-NH_2-)$  to give a good correlation with  $1/2[pK_{a2}(HA) + pK_{a2}(HB)]$ . Indeed, this was not observed. Instead,  $\Sigma S(-NH_2-)$  was found to give a significant correlation with  $\Sigma S(C_1)$ , the corresponding correlation coefficient being  $r = 0.945$ .

In spite of this interdependence, description of  $\log K_{CuAB}^B$  notably impairs with respect to equation 2 on replacing  $\Sigma S(-NH_2-)$  by the E-states of other skeletal groups, such as  $\Sigma S(-OH)$ , ( $r = 0.965$ ,  $F = 38$ ), or  $\Sigma S(C_2)$ , ( $r = 0.964$ ,  $F = 37$ ).

On the other hand, when binary and ternary complexes are separately considered,  $\Sigma S(-NH_2-)$  values give moderately significant positive correlations with the logarithms of the second stepwise formation constants of both series of metal complexes. The corresponding correlation coefficients are 0.889 and 0.811, for binary and ternary complexes, respectively. As it should be expected,  $\Sigma S(C_1)$  values also give significant positive correlations with both series of experimental data, the correlation coefficients being 0.901 and 0.803, for binary and ternary complexes, respectively. Though being of moderate quality, these results allow us to realize the role of  $\log(sf)$  in equations 1 and 2. Thus, this descriptor would encode structural information which is appropriate to unite the binary and ternary complexes in the above mentioned equations. In fact, if the index  $\log(sf)$  is removed from the descriptor set leading to equation 2, the statistical parameters of the resulting regression equation turn out to be:  $r = 0.621$ ,  $s = 0.2233$ ,  $F = 2.4$ , i.e., poorer than those of equation 1. It has to be borne in mind that the latter equation also arises from a correlation between  $\log K_{CuAB}^B$  and a four-descriptor set.

It has been pointed out recently that molecular connectivity index  ${}^1\chi$  can be interpreted as the encoding of intermolecular accessibility, i.e., as the contribution of one molecule to bimolecular interactions arising from encounters of bonds among two molecules<sup>19</sup>. In the present study, as previously stated, the index  ${}^1\chi$  was found to give a moderately significant correlation with the  $\Sigma\Delta f_1$  scale. Thus, in equations 1 and 2 the molecular connectivity index seems to be merely encoding structural information about molecular size and shape, which would be relevant to characterize the contributions of intramolecular hydrophobic interactions to the stability of the copper(II) complexes herein studied.

The fact that the correlation between  ${}^1\chi$  and  $\Sigma\Delta f_1$  is only of moderate quality can be ascribed to the presence of polar substituents in the side chains of some aminoacidate ligands (-OH groups). In fact, if  ${}^1\chi$  values for metal complexes containing aminoacidate ligands with apolar side chains are separately correlated with the hydrophobicity scale, a fairly good regression equation results:  $\Sigma\Delta f_1 = 1.0003^1\chi - 4.4523$ ;  $r = 0.996$ ,  $s = 0.1415$ ,  $F = 596$ ,  $n = 7$ . Similarly, for copper(II) complexes containing a single hydroxyl substituent, for example  $A = \text{gly}$  and  $B = \text{ser}$ , a rather significant correlation between  ${}^1\chi$  and  $\Sigma\Delta f_1$  is obtained:  $r = 0.984$ ,  $s = 0.2753$ ,  $F = 207$ ,  $n = 9$ . In turn, for complexes with two hydroxyl substituents, e.g.  $A = B = \text{ser}$ , correlation of  $\Sigma\Delta f_1$  vs  ${}^1\chi$  results also to be highly significant:  $r = 0.993$ ,  $s = 0.1984$ ,  $F = 152$ ,  $n = 4$ .

Some conclusions can be drawn from the structure-property relationships found in the present study, mainly with regard to the leading role of  $\Sigma S(-O-)$  and  $\Sigma S(C_1)$  in the description of  $\log K_{CuAB}^B$ . So, according to equations 3-5, the stability enhancement arising from the intramolecular hydrophobic interaction would operate mainly through the  $Cu(II)-O(\text{carboxylate})$  bonds, i.e., through an increase in the strength

of the coordinate bonds bearing a greater degree of ionic character. This proposal would agree with a view of such interactions as solvent structure-enforced ion-pairing type contributions. In turn, the latter would account for the role of  $\chi$  in equations 1, 2 and 5. Moreover, equations 6 and 7 suggest that the contributions of the ligand basicities, as determining factors for the differences in  $\log K_{\text{CuAB}}^{\text{B}}$  over the series of binary and ternary complexes, can be connected mainly with changes in the basicities of the carboxylate groups. Thus, although the amino group should make a greater contribution to the ligand polarizability than carboxylate<sup>20</sup>, the latter group might be more sensitive to undergo changes in polarizability under the influence of differential  $\sigma$ -electronic effects arising from structural modifications of the side chain.

Finally, when binary and ternary complexes are simultaneously considered, inclusion of  $\log(\text{sf})$  in the descriptor sets appears to be essential to achieve an appropriate modeling of  $\log K_{\text{CuAB}}^{\text{B}}$ .

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## CAPTIONS TO TABLES

**Table 1:** Electrotopological state values for some selected skeletal groups<sup>a</sup> and first-order connectivity indexes for binary and ternary copper(II) complexes with  $\alpha$ -aminoacidate ligands.<sup>b</sup>

[CuAB]		$\Sigma S(\text{O}^-)$	$\Sigma S(\text{C}_1)$	$\Sigma S(\text{C}_2)$	$\Sigma S(\text{NH}_2)$	$\Sigma S(\text{OH})$	$\chi$
A	B						
1	Ser Ser	8.578	-1.368	-1.592	2.270	17.286	6.453
2	Thr Thr	8.816	-1.377	-1.674	2.320	18.363	7.198
3	Ala Ala	8.815	-0.789	-0.642	2.841	0	5.377
4	Val Val	9.358	-0.714	-0.622	3.042	0	7.198
5	Gly Gly	8.449	-0.793	0.368	2.736	0	4.536
6	Thr Ser	8.697	-1.373	-1.633	2.295	17.825	6.826
7	Leu Leu	9.500	-0.669	-0.573	3.112	0	8.165
8	Phe Ser	9.167	-1.083	-1.221	2.786	8.746	8.010
9	Ser Gly	8.514	-1.081	-0.604	2.503	8.570	5.494
10	Ala Gly	8.626	-0.791	-0.132	2.876	0	4.956
11	Ser Ala	8.697	-1.079	-1.117	2.555	8.635	5.915
12	Thr Ala	8.815	-1.083	-1.158	2.581	9.122	6.288
13	Thr Gly	8.633	-1.085	-0.642	2.528	9.056	5.867
14	Phe Thr	9.358	-1.088	-1.262	2.650	9.357	8.383
15	Thr Thr	9.239	-1.181	-1.384	2.595	18.562	8.793
16	Phe Ala	9.286	-0.793	-0.746	2.947	0	7.472
17	Tyr Ala	9.238	-0.887	-0.867	2.855	9.175	7.883
18	Tyr Gly	9.055	-0.889	-0.342	2.803	9.145	7.462
19	Phe Gly	9.102	-0.796	-0.222	2.894	0	7.051
20	Tyr Val	9.510	-0.849	-0.858	2.956	9.227	8.793

<sup>a</sup> (-O<sup>-</sup>), coordinating oxygen; (C<sub>1</sub>), carboxylate carbon; (C<sub>2</sub>),  $\alpha$ -carbon; (-NH<sub>2</sub>) coordinating amino group; (-OH), hydroxyl substituent on aminoacidate side chain.

<sup>b</sup> Gly, glycinate; Ala, alaninate; Ser, serinate; Thr, threoninate; Val, valinate; Leu, leucinate; Phe, phenylalaninate; Tyr, tyrosinate.

**Table 2:** Comparison between the experimental  $\log K_{\text{CuAB}}^{\text{B}}$  data and the values calculated with Equations 1 and 2.

[CuAB]		Equation 1				Equation 2			
A	B	$\log K_{\text{CuAB}}^{\text{B}}$ exp	$\log K_{\text{CuAB}}^{\text{B}}$ calc	residue exp - calc	error %	$\log K_{\text{CuAB}}^{\text{B}}$ calc	residue exp - calc	error %	
Ser	Ser	6.64	6.64	0.00	0.00	6.64	0.00	0.00	
Thr	Thr	6.74	6.63	0.11	1.63	6.62	0.12	1.78	
Ala	Ala	6.81	6.81	0.00	0.00	6.81	0.00	0.00	
Val	Val	6.86	6.89	-0.03	0.44	6.88	-0.02	0.29	
Gly	Gly	6.89	6.87	0.02	0.29	6.89	0.00	0.00	
Thr	Ser	6.92	7.06	-0.14	2.02	7.07	-0.15	2.17	
Leu	Leu	6.94	7.04	-0.10	1.44	7.04	-0.10	1.44	
Phe	Ser	7.14	7.26	-0.12	1.68	7.14	0.00	0.00	
Ser	Gly	7.17	7.18	-0.01	0.14	7.20	-0.03	0.42	
Ala	Gly	7.18	7.27	-0.09	1.25	7.20	-0.02	0.28	
Ser	Ala	7.19	7.15	0.04	0.56	7.16	0.03	0.42	
Thr	Ala	7.20	7.14	0.06	0.83	7.15	0.05	0.69	
Thr	Gly	7.21	7.17	0.04	0.55	7.19	0.02	0.28	
Phe	Thr	7.23	7.21	0.02	0.28	7.27	-0.04	0.55	
Tyr	Thr	7.25	7.28	-0.03	0.41	7.29	-0.04	0.55	
Phe	Ala	7.38	7.35	0.03	0.41	7.36	0.02	0.27	
Tyr	Ala	7.42	7.37	0.05	0.67	7.38	0.04	0.54	
Tyr	Gly	7.43	7.41	0.02	0.27	7.42	0.01	0.13	
Phe	Gly	7.44	7.38	0.06	0.81	7.40	0.04	0.54	
Tyr	Val	7.50	7.42	0.08	1.07	7.42	0.08	1.07	



**Table 3:** Comparison between the thermodynamic  $pK_{a1}^{av}$  and  $pI^{av}$  data and the values calculated with Equations 6 and 7, respectively.

[CuAB]		Equation 6				Equation 7			
		$pK_{a1}^{av}$		residue	error	$pI^{av}$		residue	error
A	B	obs	calc	obs-calc	%	obs	calc	obs-calc	%
Ser	Ser	2.187	2.149	0.038	1.74	5.698	5.662	0.036	0.63
Thr	Thr	2.088	2.120	-0.032	1.53	5.594	5.605	-0.011	0.20
Ala	Ala	2.347	2.332	0.015	0.64	6.108	6.042	0.066	1.08
Val	Val	2.286	2.299	-0.013	0.57	6.002	5.982	0.020	0.33
Gly	Gly	2.350	2.370	-0.020	0.85	6.064	6.117	-0.053	0.87
Thr	Ser	2.138	2.134	0.004	0.19	5.646	5.633	0.013	0.23
Leu	Leu	2.329	2.300	0.029	1.25	6.038	5.986	0.052	0.86
Phe	Ser	2.194	2.187	0.007	0.32	5.727	5.749	-0.022	0.38
Ser	Gly	2.269	2.259	0.010	0.44	5.881	5.889	-0.008	0.14
Ala	Gly	2.349	2.351	-0.002	0.09	6.086	6.081	0.005	0.08
Ser	Ala	2.268	2.240	0.028	1.23	5.903	5.852	0.051	0.86
Thr	Ala	2.218	2.226	-0.008	0.36	5.851	5.824	0.027	0.46
Thr	Gly	2.219	2.245	-0.026	1.17	5.829	5.861	-0.032	0.55
Phe	Thr	2.144	2.165	-0.021	0.98	5.675	5.704	-0.029	0.51
Tyr	Thr	2.139	2.144	-0.005	0.23	5.642	5.660	-0.018	0.32
Phe	Ala	2.274	2.279	-0.005	0.22	5.932	5.939	-0.007	0.12
Tyr	Ala	2.268	2.250	0.019	0.84	5.899	5.879	0.020	0.34
Tyr	Gly	2.270	2.269	0.001	0.04	5.877	5.917	-0.040	0.68
Phe	Gly	2.275	2.298	-0.023	1.01	5.910	5.976	-0.066	1.12
Tyr	Val	2.238	2.234	0.004	0.18	5.846	5.850	-0.004	0.07

### CAPTIONS TO FIGURES

**Fig. 1:** Correlation between  $\Sigma S(-O-)$  and the hydrophobicity scale for aminoacid side chains,  $\Sigma \Delta f_t$  (kcal mol<sup>-1</sup>). Numbers refer to compounds in Table 1.

