

Thermodynamics of Protonation of Amino Acid Carboxylate Groups from 50 to 125°C

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Flow calorimetry has been used to study the interaction of glycine, DL- α -alanine, DL-2-aminobutyric acid, β -alanine, 4-aminobutyric acid, and 6-aminocaproic acid with protons in aqueous solutions from 323.15 K to 398.15 K and at 1.52 MPa. LogK, ΔH° , ΔS° , and ΔC_p° for the protonation of the carboxylate groups of these amino acids have been obtained at each temperature studied. Equations are given expressing these values as functions of temperature. The protonation reactions are exothermic at lower temperatures and become endothermic as temperature increases. The logK, ΔH° , and ΔS° values are close together over the temperature range studied for the protonation of α -amino acids, i.e., glycine, DL- α -alanine, and 2-aminobutyric acid. At each temperature, the magnitudes of these thermodynamic quantities increase as the number of methylene groups between the amino group and the carboxylate group increases. The ΔC_p° value for the protonation of the carboxyl group is found to lie between those of an isocoulombic reaction and a charge reduction reaction. At 323.15 K, the protonation reactions of the carboxylate groups have larger ΔC_p° values which approach those associated with charge reduction reactions. As the temperature increases, ΔC_p° decreases and approaches those found for isocoulombic reactions. This result is explained by considering long-range and short-range solvent effects. The trend in ΔH° and ΔS° with temperature and with charge separation in the zwitterions is interpreted in terms of solvent-solute interactions and the electrostatic interaction of the two oppositely charged groups within the molecule.

KEY WORDS: Protonation; amino acids; carboxylate groups; thermodynamic quantities; aqueous solutions; solvent-solute interactions; zwitterions; elevated temperatures; and calorimetry.

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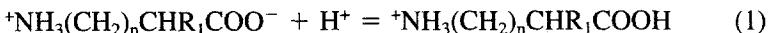
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1. INTRODUCTION

A quantitative knowledge of $\log K$ for proton ionization of proteins and their constituent amino acids in high temperature aqueous solutions is important in understanding biochemical reactions. Understanding the energetics of these reactions should help answer questions concerning the existence of life forms at high temperatures.⁽¹⁾ Recently, we have studied the protonation of the amino group for several amino acids up to 125°C.^(2,3) ΔH° and ΔS° for these protonation reactions increase while $\log K$ decreases with temperature. Separation of the zwitterion charges on the carbon chain was found to be more important than the length of the carbon chain in understanding the temperature dependence of the thermodynamic properties. These results provide information valuable for an understanding of the effect of neighboring groups in biopolymers (polypeptides, proteins, enzymes, nucleic acids, etc.) on ion association involving various functional groups.

In the present work, $\log K$, ΔH° , ΔS° , and ΔC_p° for the protonation of the carboxylate group of glycine, DL- α -alanine, DL-2-aminobutyric acid, β -alanine, 4-aminobutyric acid and 6-aminocaproic acid were determined calorimetrically from 50 to 125°C (323.15 to 398.15 K) in order to learn the effect of temperature, molecular size, and zwitterion charge separation on these quantities.

The protonation reaction of the carboxylate group of an amino acid can be expressed by



where $R_1 = H$ and $n = 0$ for glycine, $R_1 = CH_3$ and $n = 0$ for DL- α -alanine, $R_1 = CH_2CH_3$ and $n = 0$ for DL-2-aminobutyric acid; R_1 is H and $n = 1$ for β -alanine, $R_1 = H$ and $n = 2$ for 4-aminobutyric acid, and $R_1 = H$ and $n = 4$ for 6-aminocaproic acid. The effect of pressure on the protonation was not investigated in the present study, but is expected to be small at pressures from atmospheric up to the experimental pressure.^(2,4)

Values of $\log K$, ΔH° , ΔS° , and ΔC_p° below 50°C for reaction 1 for the amino acids studied have been reported.⁽⁵⁻²⁷⁾ Reviews and compilations of thermodynamic quantities for interactions of protons and metal ions with amino acids are available.⁽⁵⁻⁷⁾ Log K for the protonation of glycine have been reported up to 50°C.⁽¹⁰⁾ No ΔH° , ΔS° , and ΔC_p° values are available for reaction 1 at temperatures above 50°C for any of the amino acids studied.

2. EXPERIMENTAL

2.1. Materials

The chemicals used in this study were glycine (99+%), DL- α -alanine (99+%), β -alanine (99+%), DL-2-aminobutyric acid (99+%), 4-aminobu-

tyric acid (97%) and 6-aminocaproic acid (98%), from Aldrich, and hydrochloric acid from Fisher (standard solution). The solutions were made using distilled, deionized water and were degassed by submersion in an ultrasonic bath for 5–10 minutes. The HCl solutions were standardized by titration using a known tris[hydroxymethyl]aminomethane solution made from Sigma 99.9% pure reagent. The molar concentrations thus obtained were converted to molalities using the density data for amino acid and HCl solutions from the literature.^(28–30)

2.2. Method

The isothermal flow calorimeter used in this study has been described.⁽³¹⁾ Heat of reaction data for the titration of the amino acid solutions with HCl solutions were measured at 50, 75, 100, and 125°C and at 1.52 MPa. The concentrations of the HCl and amino acid solutions for each run were approximately the same. The flow rate for each amino acid solution was set to a constant value and that of the HCl solution was varied so that the molar ratio for proton to amino acid ranged approximately from 0.25:1 to 1.6:1. No vapor phase was present in the experiments negating the need for vaporization corrections. Details of the data reduction method are available.⁽⁴⁾

2.3. Calculations

The experimental heat of reaction data were analyzed by a computer program to derive $\log K$ and ΔH° values valid at zero ionic strength at each temperature. The Pitzer ion-interaction activity coefficient model⁽³²⁾ was used to extrapolate $\log K$ and ΔH° values to zero ionic strength and to find the $\log K$ and ΔH° which gave the best agreement between predicted and measured heats as has been described.⁽⁴⁾ The temperature dependent interaction parameters for the Pitzer activity coefficient model were found by fitting heat of dilution data for the neutral amino acids (zwitterions)⁽³⁾ and the amino acid chlorides to the cubic functions of temperature

$$\phi = A + B(T - 298.15) + C(T - 298.15)^2 + D(T - 298.15)^3 \quad (2)$$

where ϕ is $\beta^{(0)}$, $\beta^{(1)}$, C^ϕ , or λ . The value for A , the parameter at 298.15 K, was taken from the literature or estimated.^(32,33) Literature values were used for the parameters for HCl.⁽³⁴⁾ The parameters for zwitterion–zwitterion interactions were taken from our previous study.⁽³⁾ The protonation of the amino acids and the ionization of water⁽³⁵⁾ were included in the data reduction. Associated amino acid chloride and HCl(aq)^(34,36) ion pairs do not exist in significant amounts at our experimental conditions.

3. RESULTS AND DISCUSSION

Table I gives the calorimetric data at each temperature for the reaction of HCl(aq) with the amino acids studied. Heats of dilution for the amino acid chlorides are given in Table II. Each of the experimental runs was repeated three times and the heat data are the averages of the measured heats for the triplicate runs. The maximum standard deviation over the entire set of triplicate runs was 0.03 J·min⁻¹. The coefficients in Eq. (2) are listed in Table III. Table IV gives log K , ΔH° , ΔS° , and ΔC_p° for the protonation of the six amino acids studied, together with literature values for the protonation of these amino acids at zero ionic strength at lower temperatures. The uncertainties in log K , ΔH° , ΔS° , and ΔC_p° were estimated using a statistical analysis of the experimental data as described previously.⁽³⁷⁾ The Q_c values in Tables I and II were calculated using the log K and ΔH° values for reaction 1 obtained from this study.

The experimental heat data for the protonation reactions were fitted well over the temperature range of this study using

$$\log K = a + \frac{b}{T} + c \ln T + d \ln \rho_w \quad (3)$$

$$\Delta H^\circ = 2.303R \left[-b + cT + dT^2 \left(\frac{\partial \ln \rho_w}{\partial T} \right) \right] \quad (4)$$

where T is the temperature in Kelvins, ρ_w is the density of water at the experimental pressure from Haar-Gallagher-Kell equation of state for water,⁽³⁸⁾ and a , b , c , and d are fitting parameters. The fitting parameters were adjusted to allow the calculated log K and ΔH° to fit those reported in the literature at 25°C as well as the measured heat values. Temperature dependent equations for ΔS° and ΔC_p° are derived from Eqs. (3, 4) giving

$$\Delta S^\circ = 2.303R \left\{ a + c + \ln T + d \left[\ln \rho_w + T \left(\frac{\partial \ln \rho_w}{\partial T} \right) \right] \right\} \quad (5)$$

$$\Delta C_p^\circ = 2.303R \left\{ c + d \left[2T \left(\frac{\partial \ln \rho_w}{\partial T} \right) + T^2 \left(\frac{\partial^2 \ln \rho_w}{\partial^2 T} \right) \right] \right\} \quad (6)$$

Table V gives the coefficients for Eqs. (3–6) for reaction (1) as a function of temperature for each of the amino acids.

In Fig. 1, the variation with temperature of ΔH° of reaction (1) for the amino acids studied shows an increase as temperature increases. The protonation reactions for these amino acids are exothermic at the lower temperatures, and become endothermic as temperature increases. The general

Table I. Calorimetric Data for the Reaction of Amino Acids with HCl at Various Temperatures and 1.52 MPa

Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b	Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b
<i>T</i> = 323.15 K; A:0.8557 <i>m</i> HCl; B (0.4692 gH ₂ O-min ⁻¹):0.9375 <i>m</i> glycine			<i>T</i> = 373.15 K; A:0.3303 <i>m</i> HCl; B (0.4846 gH ₂ O-min ⁻¹):0.3160 <i>m</i> glycine		
0.1354	-0.3434	-0.3727	0.1371	0.1189	0.1181
0.2031	-0.5047	-0.5398	0.2057	0.1834	0.1914
0.2708	-0.6472	-0.6979	0.2742	0.2570	0.2648
0.3385	-0.7883	-0.8463	0.3428	0.3167	0.3337
0.4063	-0.9153	-0.9822	0.4114	0.3889	0.3916
0.4740	-1.0161	-1.0967	0.4799	0.4167	0.4313
0.5417	-1.0785	-1.1729	0.5485	0.4465	0.4533
0.6094	-1.1116	-1.2125	0.6170	0.4691	0.4643
0.6771	-1.1285	-1.2341			
<i>T</i> = 323.15 K; A:0.6201 <i>m</i> HCl; B (0.4759 gH ₂ O-min ⁻¹):0.6884 <i>m</i> glycine			<i>T</i> = 398.15 K; A:0.9040 <i>m</i> HCl; B (0.4686 gH ₂ O-min ⁻¹):1.0822 <i>m</i> glycine		
0.1360	-0.2294	-0.2337	0.1358	0.3602	0.3398
0.2041	-0.3376	-0.3389	0.2037	0.6294	0.5958
0.2721	-0.4349	-0.4398	0.2716	0.8836	0.8653
0.3401	-0.5322	-0.5357	0.3395	1.1476	1.1387
0.4081	-0.5989	-0.6248	0.4076	1.4094	1.4078
0.4762	-0.6550	-0.7016	0.4753	1.6457	1.6583
0.5442	-0.6913	-0.7576	0.5431	1.8326	1.8554
0.6122	-0.7185	-0.7918	0.6110	1.9375	1.9570
0.6802	-0.7414	-0.8129	0.6789	1.9706	1.9892
<i>T</i> = 348.15 K; A:1.0448 <i>m</i> HCl; B (0.4695 gH ₂ O-min ⁻¹):1.0382 <i>m</i> glycine			<i>T</i> = 398.15 K; A:0.5970 <i>m</i> HCl; B (0.4804 gH ₂ O-min ⁻¹):0.5132 <i>m</i> glycine		
0.1355	-0.0560	-0.0711	0.1365	0.3183	0.2749
0.2032	-0.0616	-0.0701	0.2047	0.5087	0.4577
0.2709	-0.0566	-0.0644	0.2730	0.6947	0.6434
0.3387	-0.0498	-0.0577	0.3412	0.8734	0.8164
0.4064	-0.0364	-0.0540	0.4095	1.0027	0.9447
0.4741	-0.0438	-0.0615	0.4777	1.0699	1.0045
0.5418	-0.0548	-0.0835	0.5460	1.0933	1.0245
0.6096	-0.0782	-0.1103	0.6142	1.1021	1.0300
0.6773	-0.0964	-0.1363	0.6825	1.1087	1.0302
<i>T</i> = 373.15 K; A:0.6713 <i>m</i> HCl; B (0.4766 gH ₂ O-min ⁻¹):0.6934 <i>m</i> glycine			<i>T</i> = 323.15 K; A:1.0629 <i>m</i> HCl; B (0.4617 gH ₂ O-min ⁻¹):1.0668 <i>m</i> DL- α -alanine		
0.1363	0.1837	0.1954	0.1354	-0.1994	-0.1458
0.2045	0.2996	0.3310	0.2031	-0.2890	-0.1898
0.2727	0.4114	0.4708	0.2708	-0.3661	-0.2401
0.3408	0.5461	0.6084	0.3385	-0.4406	-0.2997
0.4090	0.6575	0.7348	0.4063	-0.5042	-0.3670
0.4771	0.7415	0.8310	0.4740	-0.5581	-0.4332
0.5453	0.7953	0.8797	0.5417	-0.5981	-0.4888
0.6135	0.8264	0.8954	0.6094	-0.6237	-0.5372
			0.6771	-0.6407	-0.5815

Table I. Continued

Flow A ^a	Q_m^b	Q_c^b	Flow A ^a	Q_m^b	Q_c^b
$T = 323.15 \text{ K}; A: 0.5782m \text{ HCl};$ $B (0.4763 \text{ gH}_2\text{O-min}^{-1}): 0.5170m$ DL- α -alanine			$T = 398.15 \text{ K}; A: 1.0196m \text{ HCl};$ $B (0.4617 \text{ gH}_2\text{O-min}^{-1}): 1.0666m$ DL- α -alanine		
0.1365	-0.0670	-0.0546	0.1355	0.5073	0.4919
0.2048	-0.0922	-0.0621	0.2033	0.8244	0.8366
0.2731	-0.1172	-0.0665	0.2710	1.1371	1.1868
0.3414	-0.1380	-0.0698	0.3388	1.4484	1.5282
0.4096	-0.1497	-0.0732	0.4066	1.7530	1.8418
0.4779	-0.1634	-0.0773	0.4743	1.9844	2.0719
0.5462	-0.1760	-0.0819	0.5421	2.0891	2.1503
0.6144	-0.1870	-0.0866	0.6098	2.1012	2.1451
0.6827	-0.1950	-0.0913	0.6776	2.0879	2.1169
$T = 348.15 \text{ K}; A: 1.0649m \text{ HCl};$ $B (0.4624 \text{ gH}_2\text{O-min}^{-1}): 1.0410m$ DL- α -alanine			$T = 398.15 \text{ K}; A: 0.5597m \text{ HCl};$ $B (0.4763 \text{ gH}_2\text{O-min}^{-1}): 0.5172m$ DL- α -alanine		
0.1354	0.0602	0.0689	0.1365	0.3742	0.3097
0.2031	0.1239	0.1364	0.2048	0.5908	0.5133
0.2708	0.1885	0.2073	0.2730	0.8040	0.7218
0.3385	0.2554	0.2769	0.3412	0.9990	0.9226
0.4062	0.3099	0.3357	0.4095	1.1571	1.0899
0.4739	0.3405	0.3600	0.4777	1.2291	1.1867
0.5417	0.3343	0.3480	0.5460	1.2521	1.2243
0.6094	0.3185	0.3251	0.6142	1.2605	1.2377
0.6771	0.3037	0.3013	0.6825	1.2620	1.2421
$T = 373.15 \text{ K}; A: 1.0219m \text{ HCl};$ $B (0.4629 \text{ gH}_2\text{O-min}^{-1}): 1.0188m$ DL- α -alanine			$T = 323.15 \text{ K}; A: 0.7120m \text{ HCl};$ $B (0.4624 \text{ gH}_2\text{O-min}^{-1}): 0.8018m$ β -alanine		
0.1355	0.3074	0.2794	0.1354	-0.5231	-0.4856
0.2033	0.4994	0.4715	0.2031	-0.7622	-0.7007
0.2710	0.6942	0.6712	0.2708	-0.9766	-0.9085
0.3388	0.8860	0.8701	0.3385	-1.1846	-1.1108
0.4065	1.0616	1.0508	0.4062	-1.3877	-1.3082
0.4743	1.1711	1.1644	0.4739	-1.5282	-1.4987
0.5420	1.1983	1.1899	0.5416	-1.5726	-1.6358
0.6098	1.1922	1.1811	0.6094	-1.5960	-1.6830
0.6775	1.1835	1.1643	0.6771	-1.6120	-1.7165
$T = 373.15 \text{ K}; A: 0.5207m \text{ HCl};$ $B (0.4763 \text{ gH}_2\text{O-min}^{-1}): 0.5170m$ DL- α -alanine			$T = 323.15 \text{ K}; A: 0.6420m \text{ HCl};$ $B (0.4670 \text{ gH}_2\text{O-min}^{-1}): 0.6535m$ β -alanine		
0.1367	0.2181	0.2065	0.1360	-0.3639	-0.3834
0.2050	0.3417	0.3319	0.2041	-0.5272	-0.5556
0.2734	0.4711	0.4587	0.2721	-0.6750	-0.7232
0.3417	0.5925	0.5816	0.3401	-0.8207	-0.8873
0.4100	0.7102	0.6898	0.4081	-0.9563	-1.0473
0.4784	0.7880	0.7643	0.4762	-1.0891	-1.1855
0.5467	0.8339	0.7982	0.5442	-1.2000	-1.2374
0.6150	0.8558	0.8092	0.6122	-1.2389	-1.2679
0.6834	0.8676	0.8112	0.6802	-1.2577	-1.2932

Table I. Continued

Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b	Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b
<i>T</i> = 348.15 K; A:0.9310 <i>m</i> HCl; B (0.4642 gH ₂ O-min ⁻¹):0.9273 <i>m</i> β-alanine			<i>T</i> = 398.15 K; A:0.3729 <i>m</i> HCl; B (0.4753 gH ₂ O-min ⁻¹):0.5770 <i>m</i> β-alanine		
0.1355	-0.2290	-0.2449	0.1370	0.1181	0.1263
0.2032	-0.3023	-0.3169	0.2055	0.2275	0.2299
0.2709	-0.3565	-0.3772	0.2740	0.3397	0.3370
0.3387	-0.4043	-0.4292	0.3426	0.4579	0.4458
0.4064	-0.4494	-0.4750	0.4111	0.5857	0.5556
0.4741	-0.4833	-0.5166	0.4796	0.7075	0.6657
0.5418	-0.5205	-0.5546	0.5481	0.8332	0.7756
0.6096	-0.5494	-0.5875	0.6166	0.9583	0.8841
0.6773	-0.5657	-0.6161	0.6851	1.0682	0.9865
<i>T</i> = 348.15 K; A:0.5970 <i>m</i> HCl; B (0.4833 gH ₂ O-min ⁻¹):0.5004 <i>m</i> β-alanine			<i>T</i> = 398.15 K; A:0.4005 <i>m</i> HCl; B (0.4803 gH ₂ O-min ⁻¹):0.3899 <i>m</i> β-alanine		
0.1365	-0.0350	-0.0385	0.1370	0.1453	0.1807
0.2047	-0.0344	-0.0335	0.2054	0.2703	0.3019
0.2730	-0.0344	-0.0245	0.2739	0.3935	0.4240
0.3412	-0.0329	-0.0133	0.3424	0.5252	0.5455
0.4095	-0.0424	-0.0068	0.4109	0.6516	0.6635
0.4777	-0.0310	-0.0228	0.4793	0.7521	0.7481
0.5460	-0.0377	-0.0399	0.5478	0.7687	0.7572
0.6142	-0.0542	-0.0550	0.6163	0.7722	0.7543
0.6825	-0.0768	-0.0682	0.6848	0.7792	0.7503
<i>T</i> = 373.15 K; A:0.7920 <i>m</i> HCl; B (0.4670 gH ₂ O-min ⁻¹):0.9018 <i>m</i> β-alanine			<i>T</i> = 323.15 K; A:0.5122 <i>m</i> HCl; B (0.9499 gH ₂ O-min ⁻¹):0.5211 <i>m</i> DL-2-aminobutyric acid		
0.1360	0.0358	0.0908	0.1709	0.0141	-0.0166
0.2041	0.1152	0.1912	0.3417	0.0532	-0.0063
0.2721	0.1986	0.2987	0.5126	0.1001	0.0089
0.3401	0.2976	0.4100	0.6835	0.1433	0.0246
0.4081	0.3969	0.5231	0.8543	0.1855	0.0362
0.4762	0.4979	0.6352	1.0252	0.2100	0.0378
0.5442	0.5835	0.7140	1.2815	0.2193	0.0259
0.6122	0.5879	0.7083	1.5378	0.2197	0.0110
<i>T</i> = 373.15 K; A:0.6713 <i>m</i> HCl; B (0.4780 gH ₂ O-min ⁻¹):0.4743 <i>m</i> β-alanine			<i>T</i> = 323.15 K; A:0.5782 <i>m</i> HCl; B (0.7124 gH ₂ O-min ⁻¹):0.5211 <i>m</i> DL-2-aminobutyric acid		
0.1363	0.1109	0.1838	0.1707	0.0188	-0.0158
0.2045	0.2049	0.3075	0.3072	0.0487	-0.0088
0.2727	0.3000	0.4321	0.4096	0.0697	-0.0021
0.3408	0.3932	0.5348	0.5120	0.0926	0.0035
0.4090	0.4001	0.5301	0.6144	0.1144	0.0050
0.4771	0.3868	0.5135	0.7510	0.1207	-0.0023
0.5453	0.3766	0.4985	0.8875	0.1137	-0.0142
0.6135	0.3715	0.4852	1.0241	0.1040	-0.0259

Table I. Continued

Flow A ^a	Q_m^b	Q_c^b	Flow A ^a	Q_m^b	Q_c^b
<i>T = 323.15 K; A:0.5782m HCl; B (0.4726 gH₂O-min⁻¹):0.5211m</i>					
DL-2-aminobutyric acid					
0.1365	0.0087	-0.0096	0.1371	0.1811	0.1799
0.2048	0.0231	-0.0058	0.2057	0.2921	0.2828
0.2731	0.0372	-0.0013	0.2742	0.3917	0.3860
0.3414	0.0536	0.0024	0.3428	0.5019	0.4862
0.4096	0.0640	0.0033	0.4114	0.5978	0.5786
0.4779	0.0670	0.0000	0.4799	0.6798	0.6556
0.5462	0.0663	-0.0056	0.5485	0.7464	0.7097
0.6144	0.0623	-0.0117	0.6170	0.7834	0.7416
0.6827	0.0576	-0.0174			
<i>T = 348.15 K; A:1.0448m HCl; B (0.4552 gH₂O-min⁻¹):1.0550m</i>					
DL-2-aminobutyric acid					
0.1355	0.1719	0.1780	0.1355	0.5527	0.5542
0.2032	0.2865	0.3038	0.2033	0.9012	0.9033
0.2709	0.4117	0.4342	0.2710	1.2533	1.2605
0.3387	0.5320	0.5628	0.3388	1.5970	1.6149
0.4064	0.6429	0.6769	0.4066	1.9189	1.9467
0.4741	0.7093	0.7456	0.4743	2.1750	2.1960
0.5418	0.7226	0.7588	0.5421	2.2900	2.2946
0.6096	0.7132	0.7498	0.6098	2.3047	2.3102
0.6773	0.6977	0.7353	0.6776	2.2857	2.3024
<i>T = 348.15 K; A:0.5970m HCl; B (0.4705 gH₂O-min⁻¹):0.5850m</i>					
DL-2-aminobutyric acid					
0.1365	0.1646	0.1536	0.1365	0.4143	0.3867
0.2048	0.2502	0.2482	0.2048	0.6609	0.6166
0.2730	0.3527	0.3438	0.2730	0.8901	0.8485
0.3412	0.4428	0.4355	0.3412	1.1306	1.0726
0.4095	0.5237	0.5140	0.4095	1.3213	1.2679
0.4777	0.5667	0.5642	0.4777	1.4367	1.3951
0.5460	0.5946	0.5848	0.5460	1.4915	1.4478
0.6142	0.5928	0.5904	0.6142	1.5078	1.4647
0.6825	0.5986	0.5902	0.6825	1.5185	1.4686
<i>T = 373.15 K; A:0.6713m HCl; B (0.4693 gH₂O-min⁻¹):0.6186m</i>					
DL-2-aminobutyric acid					
0.1363	0.3156	0.3194	0.1709	-0.0372	-0.0638
0.2045	0.4945	0.5093	0.3417	-0.0606	-0.0838
0.2727	0.6719	0.7001	0.5126	-0.0713	-0.0956
0.3408	0.8457	0.8805	0.6835	-0.0770	-0.1024
0.4090	0.9939	1.0248	0.8543	-0.0593	-0.1059
0.4771	1.0774	1.0999	1.0252	-0.0677	-0.1201
0.5453	1.1057	1.1237	1.2815	-0.0901	-0.1666
0.6135	1.1266	1.1283	1.5378	-0.1149	-0.2032

Table I. Continued

Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b	Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b
<i>T</i> = 348.15 K; A:1.0117 <i>m</i> HCl; B (0.4558 gH ₂ O-min ⁻¹):1.0266 <i>m</i>			<i>T</i> = 398.15 K; A:0.5970 <i>m</i> HCl; B (0.4733 gH ₂ O-min ⁻¹):0.5208 <i>m</i>		
4-aminobutyric acid			4-aminobutyric acid		
0.1354	0.0147	0.0105	0.1365	0.3211	0.3278
0.2031	0.0692	0.0701	0.2048	0.5461	0.5607
0.2708	0.1348	0.1383	0.2730	0.7764	0.7993
0.3385	0.2136	0.2124	0.3412	1.0147	1.0404
0.4062	0.2918	0.2902	0.4095	1.2394	1.2669
0.4739	0.3452	0.3450	0.4777	1.2625	1.2764
0.5416	0.3092	0.3018	0.5460	1.2463	1.2602
0.6094	0.2731	0.2612	0.6142	1.2362	1.2456
0.6771	0.2475	0.2256	0.6825	1.2245	1.2326
<i>T</i> = 373.15 K; A:0.1.1256 <i>m</i> HCl; B (0.4603 gH ₂ O-min ⁻¹):1.1354 <i>m</i>			<i>T</i> = 323.15 K; A:0.4005 <i>m</i> HCl; B (0.4706 gH ₂ O-min ⁻¹):0.4201 <i>m</i>		
4-aminobutyric acid			6-aminocaproic acid		
0.1356	0.2334	0.2810	0.1370	0.0596	0.0640
0.2034	0.4221	0.5057	0.2054	0.1505	0.1193
0.2712	0.6153	0.7375	0.2739	0.2593	0.1809
0.3391	0.8254	0.9720	0.3424	0.3820	0.2471
0.4069	1.0456	1.2066	0.4109	0.5192	0.3169
0.4747	1.1818	1.3886	0.4793	0.6100	0.3872
0.5425	1.1565	1.3418	0.5478	0.5688	0.3927
0.6103	1.1141	1.2936	0.6163	0.5299	0.3798
			0.6848	0.4975	0.3683
<i>T</i> = 373.15 K; A:0.4426 <i>m</i> HCl; B (0.4733 gH ₂ O-min ⁻¹):0.4425 <i>m</i>			<i>T</i> = 348.15 K; A:1.0445 <i>m</i> HCl; B (0.4527 gH ₂ O-min ⁻¹):1.1047 <i>m</i>		
4-aminobutyric acid			6-aminocaproic acid		
0.1367	0.2047	0.2407	0.1361	0.2456	0.1903
0.2050	0.3479	0.3996	0.2041	0.4823	0.4008
0.2734	0.4848	0.5623	0.2721	0.7301	0.6411
0.3417	0.6330	0.7271	0.3401	1.0046	0.9028
0.4100	0.7831	0.8924	0.4082	1.2908	1.1799
0.4784	0.9105	1.0322	0.4762	1.5661	1.4220
0.5467	0.9172	1.0362	0.5442	1.5369	1.3808
0.6150	0.9141	1.0285	0.6122	1.4892	1.3401
0.6834	0.9157	1.0211	0.6803	1.4499	1.3044
<i>T</i> = 398.15 K; A:0.9557 <i>m</i> HCl; B (0.4611 gH ₂ O-min ⁻¹):1.0232 <i>m</i>			<i>T</i> = 348.15 K; A:0.5970 <i>m</i> HCl; B (0.4661 gH ₂ O-min ⁻¹):0.5168 <i>m</i>		
4-aminobutyric acid			6-aminocaproic acid		
0.1357	0.3169	0.3046	0.1365	0.2835	0.3386
0.2036	0.6044	0.5689	0.2048	0.4908	0.5535
0.2714	0.9112	0.8444	0.2730	0.6921	0.7761
0.3393	1.2299	1.1254	0.3412	0.8983	1.0038
0.4071	1.5535	1.4087	0.4095	1.1132	1.2059
0.4750	1.8789	1.6879	0.4777	1.1202	1.1931
0.5429	1.9115	1.7420	0.5460	1.1060	1.1760
0.6107	1.8662	1.7078	0.6142	1.0961	1.1611
0.6786	1.8387	1.6761	0.6825	1.0734	1.1481

Table I. Continued

Flow A ^a	Q_m^b	Q_c^b	Flow A ^a	Q_m^b	Q_c^b
$T = 373.15 \text{ K}; \text{A}:0.5380m \text{ HCl};$ $\text{B} (0.4668 \text{ gH}_2\text{O}\cdot\text{min}^{-1}):0.5008m$ 6-aminocaproic acid			$T = 398.15 \text{ K}; \text{A}:0.9040m \text{ HCl};$ $\text{B} (0.4527 \text{ gH}_2\text{O}\cdot\text{min}^{-1}):1.0185m$ 6-aminocaproic acid		
0.1366	0.3532	0.4381	0.1361	0.3757	0.2619
0.2050	0.6228	0.7344	0.2041	0.8420	0.6688
0.2733	0.8824	1.0416	0.2721	1.3105	1.1386
0.3416	1.1628	1.3559	0.3401	1.8256	1.6515
0.4099	1.4441	1.6729	0.4082	2.3845	2.1954
0.4782	1.5725	1.7813	0.4762	2.9281	2.7598
0.5465	1.5663	1.7697	0.5442	3.1387	2.9453
0.6149	1.5659	1.7586	0.6122	3.0950	2.9012
			0.6803	3.0602	2.8606
$T = 373.15 \text{ K}; \text{A}:0.3303m \text{ HCl};$ $\text{B} (0.4774 \text{ gH}_2\text{O}\cdot\text{min}^{-1}):0.2764m$ 6-aminocaproic acid			$T = 398.15 \text{ K}; \text{A}:0.5970m \text{ HCl};$ $\text{B} (0.4661 \text{ gH}_2\text{O}\cdot\text{min}^{-1}):0.5160m$ 6-aminocaproic acid		
0.1371	0.2730	0.3151	0.1365	0.4453	0.4258
0.2057	0.4587	0.5156	0.2048	0.8177	0.7881
0.2742	0.6325	0.7208	0.2730	1.1903	1.1762
0.3428	0.8258	0.9287	0.3412	1.5938	1.5815
0.4114	1.0001	1.0965	0.4095	1.9980	1.9477
0.4799	1.0128	1.0952	0.4777	2.0072	1.9328
0.5485	1.0118	1.0899	0.5460	1.9881	1.9123
0.6170	1.0172	1.0851	0.6142	1.9689	1.8940
			0.6825	1.9517	1.8779

^aUnits: g H₂O·min⁻¹.^bUnits: J·min⁻¹.

trend is that ΔH° increases as the number of methylene groups between the amino and the carboxylate groups increases except in the case of β -alanine. ΔC_p° for the protonation reactions decrease as temperature increases for all of the amino acids studied (Fig. 2).

Shown in Fig. 3 are the plots of $T\Delta S^\circ$ for reaction (1) as a function of temperature. The protonation reactions for the amino acids have large positive $T\Delta S^\circ$ values over the entire temperature range studied. As the temperature increases, the $T\Delta S^\circ$ values become larger and have increasing importance in determining the magnitudes of the $\log K$ values for the protonation reaction. It can also be seen in Fig. 3 that ΔS° of the α -amino acids are nearly the same at each temperature while those for the other amino acids increase as the number of methylene groups between the carboxylate and amino groups increases.

Each of the quantities K , ΔH° , ΔS° and ΔC_p° for the protonation reactions will now be discussed to show how they are affected by solvent-solute and intramolecular zwitterion interactions.

Table II. Calorimetric Data for the Dilution of the Amino Acid Chlorides at Various Temperatures and 1.52 MPa

Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b	Flow A ^a	<i>Q_m</i> ^b	<i>Q_c</i> ^b
<i>T</i> = 348.15 K; A:H ₂ O; B (0.4695 gH ₂ O-min ⁻¹): 1.0382 <i>m</i> glycine + 1.0827 <i>m</i> HCl			<i>T</i> = 373.15 K; A:H ₂ O; B (0.4629 gH ₂ O-min ⁻¹): 1.0188 <i>m</i> DL- α -alanine + 1.1660 <i>m</i> HCl		
0.1378	-0.0883	-0.0664	0.1378	-0.1937	-0.1877
0.2067	-0.0788	-0.1014	0.2067	-0.2690	-0.2665
0.2756	-0.1494	-0.1354	0.2756	-0.3311	-0.3371
0.3445	-0.1682	-0.1678	0.3445	-0.4063	-0.4006
0.4134	-0.2267	-0.1984	0.4134	-0.4695	-0.4582
0.4823	-0.1946	-0.2273	0.4823	-0.5209	-0.5107
0.5511	-0.2488	-0.2545	0.5511	-0.5636	-0.5588
0.6200	-0.2680	-0.2801	0.6200	-0.5914	-0.6032
0.6889	-0.3106	-0.3042	0.6889	-0.6516	-0.6444
<i>T</i> = 373.15 K; A:H ₂ O; B (0.4766 gH ₂ O-min ⁻¹): 0.6934 <i>m</i> glycine + 0.5661 <i>m</i> HCl			<i>T</i> = 398.15 K; A:H ₂ O; B (0.4763 gH ₂ O-min ⁻¹): 0.5172 <i>m</i> DL- α -alanine + 0.4655 <i>m</i> HCl		
0.1378	-0.0858	-0.0876	0.1378	-0.1164	-0.1169
0.2067	-0.1249	-0.1279	0.2067	-0.1645	-0.1690
0.2756	-0.1584	-0.1652	0.2756	-0.2159	-0.2166
0.3445	-0.1998	-0.1996	0.3445	-0.2491	-0.2600
0.4134	-0.2221	-0.2312	0.4134	-0.2951	-0.2998
0.4823	-0.2536	-0.2604	0.4823	-0.3324	-0.3362
0.5511	-0.2752	-0.2874	0.5511	-0.3654	-0.3697
0.6200	-0.3072	-0.3125	0.6200	-0.4004	-0.4008
0.6889	-0.3318	-0.3359	0.6889	-0.4212	-0.4295
<i>T</i> = 398.15 K; A:H ₂ O; B (0.4804 gH ₂ O-min ⁻¹): 0.5132 <i>m</i> glycine + 0.5798 <i>m</i> HCl			<i>T</i> = 348.15 K; A:H ₂ O; B (0.4642 gH ₂ O-min ⁻¹): 1.0148 <i>m</i> β -alanine + 1.0827 <i>m</i> HCl		
0.1378	-0.1450	-0.1405	0.1378	-0.0845	-0.0656
0.2067	-0.2002	-0.2013	0.2067	-0.1017	-0.0886
0.2756	-0.2580	-0.2562	0.2756	-0.0942	-0.1080
0.3445	-0.3048	-0.3061	0.3445	-0.1048	-0.1248
0.4134	-0.3444	-0.3516	0.4134	-0.1634	-0.1398
0.4823	-0.3952	-0.3932	0.4823	-0.1264	-0.1534
0.5511	-0.4336	-0.4313	0.5511	-0.1669	-0.1658
0.6200	-0.4677	-0.4666	0.6200	-0.1696	-0.1774
0.6889	-0.4990	-0.4993	0.6889	-0.2115	-0.1881
<i>T</i> = 348.15 K; A:H ₂ O; B (0.4624 gH ₂ O-min ⁻¹): 1.0410 <i>m</i> DL- α -alanine + 0.9714 <i>m</i> HCl			<i>T</i> = 373.15 K; A:H ₂ O; B (0.4780 gH ₂ O-min ⁻¹): 0.4743 <i>m</i> β -alanine + 0.6125 <i>m</i> HCl		
0.1378	-0.0955	-0.0901	0.1378	-0.0858	-0.0859
0.2067	-0.1367	-0.1305	0.2067	-0.1199	-0.1212
0.2756	-0.1738	-0.1675	0.2756	-0.1604	-0.1525
0.3445	-0.2274	-0.2015	0.3445	-0.1837	-0.1805
0.4134	-0.2377	-0.2327	0.4134	-0.2069	-0.2057
0.4823	-0.2611	-0.2615	0.4823	-0.2319	-0.2286
0.5511	-0.2766	-0.2881	0.5511	-0.2589	-0.2494
0.6200	-0.3052	-0.3128	0.6200	-0.2779	-0.2686
0.6889	-0.3270	-0.3358	0.6889	-0.3014	-0.2862

Table II. Continued

Flow A ^a	Q_m^b	Q_c^b	Flow A ^a	Q_m^b	Q_c^b
$T = 373.15 \text{ K}; \text{A:H}_2\text{O};$ $\text{B} (0.4670 \text{ gH}_2\text{O-min}^{-1})$: $0.9018m \beta\text{-alanine} + 1.1660m \text{HCl}$			$T = 348.15 \text{ K}; \text{A:H}_2\text{O};$ $\text{B} (0.4541 \text{ gH}_2\text{O-min}^{-1})$: $1.0812m \text{DL-2-aminobutyric acid}$ + $1.0827m \text{HCl}$		
0.1378	-0.1477	-0.1237	0.1378	-0.1484	-0.1278
0.2067	-0.2102	-0.1858	0.2067	-0.1645	-0.1852
0.2756	-0.2703	-0.2447	0.2756	-0.2848	-0.2376
0.3445	-0.3107	-0.2999	0.3445	-0.3140	-0.2853
0.4134	-0.3690	-0.3512	0.4134	-0.3417	-0.3289
0.4823	-0.3948	-0.3989	0.4823	-0.3587	-0.3689
0.5511	-0.4404	-0.4432	0.5511	-0.4153	-0.4056
0.6200	-0.4658	-0.4844	0.6200	-0.4710	-0.4395
0.6889	-0.4937	-0.5229	0.6889	-0.5147	-0.4710
$T = 398.15 \text{ K}; \text{A:H}_2\text{O};$ $\text{B} (0.4753 \text{ gH}_2\text{O-min}^{-1})$: $0.5770m \beta\text{-alanine} + 0.5353m \text{HCl}$			$T = 373.15 \text{ K}; \text{A:H}_2\text{O};$ $\text{B} (0.4693 \text{ gH}_2\text{O-min}^{-1})$: $0.6186m \text{DL-2-aminobutyric acid}$ + $0.5661m \text{HCl}$		
0.1378	-0.0997	-0.1061	0.1378	-0.0940	-0.1031
0.2067	-0.1520	-0.1538	0.2067	-0.1312	-0.1450
0.2756	-0.1922	-0.1975	0.2756	-0.1739	-0.1820
0.3445	-0.2262	-0.2374	0.3445	-0.1911	-0.2152
0.4134	-0.2575	-0.2738	0.4134	-0.2350	-0.2451
0.4823	-0.2939	-0.3071	0.4823	-0.2605	-0.2724
0.5511	-0.3252	-0.3377	0.5511	-0.2897	-0.2973
0.6200	-0.3550	-0.3660	0.6200	-0.3052	-0.3203
0.6889	-0.3819	-0.3921	0.6889	-0.3361	-0.3417
$T = 323.15 \text{ K}; \text{A: H}_2\text{O};$ $\text{B} (0.4726 \text{ gH}_2\text{O-min}^{-1})$: $0.5216m \text{DL-2-aminobutyric acid}$ + $0.5670m \text{HCl}$			$T = 398.15 \text{ K}; \text{A:H}_2\text{O};$ $\text{B} (0.4705 \text{ gH}_2\text{O-min}^{-1})$: $0.5850m \text{DL-2-aminobutyric acid}$ + $0.5798m \text{HCl}$		
0.1378	-0.0510	-0.0469	0.1378	-0.1287	-0.1406
0.2067	-0.0811	-0.0654	0.2067	-0.1925	-0.1979
0.2756	-0.0924	-0.0816	0.2756	-0.2341	-0.2488
0.3445	-0.0791	-0.0959	0.3445	-0.2823	-0.2943
0.4134	-0.0955	-0.1086	0.4134	-0.3305	-0.3355
0.4823	-0.1129	-0.1201	0.4823	-0.3749	-0.3729
0.5511	-0.1219	-0.1305	0.5511	-0.4151	-0.4072
0.6200	-0.1236	-0.1401	0.6200	-0.4461	-0.4388
0.6889	-0.1359	-0.1489	0.6889	-0.4794	-0.4681
$T = 348.15 \text{ K}; \text{A:H}_2\text{O};$ $\text{B} (0.4705 \text{ gH}_2\text{O-min}^{-1})$: $0.5850m \text{DL-2-aminobutyric acid}$ + $0.5798m \text{HCl}$			$T = 398.15 \text{ K}; \text{A:H}_2\text{O};$ $\text{B} (0.4544 \text{ gH}_2\text{O-min}^{-1})$: $1.0805m \text{DL-2-aminobutyric acid}$ + $1.0827m \text{HCl}$		
0.1378	-0.0747	-0.0741	0.1378	-0.2424	-0.2258
0.2067	-0.1043	-0.1041	0.2067	-0.3479	-0.3308
0.2756	-0.1276	-0.1306	0.2756	-0.4393	-0.4279
0.3445	-0.1460	-0.1543	0.3445	-0.5267	-0.5172
0.4134	-0.1789	-0.1757	0.4134	-0.5993	-0.5993
0.4823	-0.2004	-0.1952	0.4823	-0.6684	-0.6749
0.5511	-0.2223	-0.2130	0.5511	-0.7409	-0.7446
0.6200	-0.2378	-0.2294	0.6200	-0.8032	-0.8093
0.6889	-0.2433	-0.2446	0.6889	-0.8632	-0.8694

Table II. Continued

Flow A ^a	Q_m^b	Q_c^b	Flow A ^a	Q_m^b	Q_c^b
$T = 348.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4558 gH ₂ O-min ⁻¹): 1.0806 <i>m</i> 4-aminobutyric acid + 0.9714 <i>m</i> HCl			$T = 398.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4611 gH ₂ O-min ⁻¹): 0.9051 <i>m</i> 4-aminobutyric acid + 1.1660 <i>m</i> HCl		
0.1378	-0.0538	-0.0586	0.1378	-0.1871	-0.1362
0.2067	-0.0853	-0.0808	0.2067	-0.2609	-0.2091
0.2756	-0.1009	-0.1000	0.2756	-0.3206	-0.2796
0.3445	-0.1254	-0.1171	0.3445	-0.3719	-0.3466
0.4134	-0.1270	-0.1326	0.4134	-0.4244	-0.4095
0.4823	-0.1448	-0.1467	0.4823	-0.4627	-0.4684
0.5511	-0.1502	-0.1596	0.5511	-0.5003	-0.5234
0.6200	-0.1737	-0.1717	0.6200	-0.5327	-0.5749
0.6889	-0.1878	-0.1830	0.6889	-0.5617	-0.6232
$T = 373.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4733 gH ₂ O-min ⁻¹): 0.5206 <i>m</i> 4-aminobutyric acid + 0.6125 <i>m</i> HCl			$T = 348.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4527 gH ₂ O-min ⁻¹): 0.8175 <i>m</i> 6-aminocaproic acid + 0.9714 <i>m</i> HCl		
0.1378	-0.0814	-0.0617	0.1378	-0.0297	-0.0322
0.2067	-0.1162	-0.0883	0.2067	-0.0459	-0.0477
0.2756	-0.1397	-0.1124	0.2756	-0.0591	-0.0623
0.3445	-0.1690	-0.1343	0.3445	-0.0866	-0.0760
0.4134	-0.1978	-0.1543	0.4134	-0.0941	-0.0888
0.4823	-0.2218	-0.1727	0.4823	-0.1036	-0.1007
0.5511	-0.2415	-0.1895	0.5511	-0.1054	-0.1120
0.6200	-0.2516	-0.2052	0.6200	-0.1190	-0.1225
0.6889	-0.2728	-0.2197	0.6889	-0.1299	-0.1324
$T = 373.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4537 gH ₂ O-min ⁻¹): 1.0855 <i>m</i> 4-aminobutyric acid + 0.9714 <i>m</i> HCl			$T = 373.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4668 gH ₂ O-min ⁻¹): 0.5008 <i>m</i> 6-aminocaproic acid + 0.5661 <i>m</i> HCl		
0.1378	-0.1137	-0.1079	0.1378	-0.0374	-0.0476
0.2067	-0.1590	-0.1580	0.2067	-0.0825	-0.0708
0.2756	-0.1986	-0.2044	0.2756	-0.0913	-0.0926
0.3445	-0.2355	-0.2472	0.3445	-0.1113	-0.1128
0.4134	-0.2755	-0.2865	0.4134	-0.1321	-0.1316
0.4823	-0.3054	-0.3227	0.4823	-0.1408	-0.1490
0.5511	-0.3381	-0.3562	0.5511	-0.1785	-0.1651
0.6200	-0.3694	-0.3872	0.6200	-0.1743	-0.1801
0.6889	-0.3787	-0.4160	0.6889	-0.1929	-0.1940
$T = 398.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4733 gH ₂ O-min ⁻¹): 0.5208 <i>m</i> 4-aminobutyric acid + 0.5742 <i>m</i> HCl			$T = 398.15\text{ K}; \text{A:H}_2\text{O};$ B (0.4661 gH ₂ O-min ⁻¹): 0.5161 <i>m</i> 6-aminocaproic acid + 0.5742 <i>m</i> HCl		
0.1378	-0.0999	-0.0980	0.1378	-0.0724	-0.0597
0.2067	-0.1387	-0.1389	0.2067	-0.1087	-0.0891
0.2756	-0.1797	-0.1755	0.2756	-0.1382	-0.1168
0.3445	-0.2176	-0.2083	0.3445	-0.1678	-0.1426
0.4134	-0.2489	-0.2381	0.4134	-0.1899	-0.1667
0.4823	-0.2788	-0.2651	0.4823	-0.2213	-0.1890
0.5511	-0.3058	-0.2898	0.5511	-0.2451	-0.2096
0.6200	-0.3362	-0.3125	0.6200	-0.2676	-0.2289
0.6889	-0.3565	-0.3335	0.6889	-0.2889	-0.2468

Table II. Continued

Flow A ^a	Q_m^b	Q_c^b	Flow A ^a	Q_m^b	Q_c^b
<i>T</i> = 398.15 K; A:H ₂ O; B (0.4527 gH ₂ O-min ⁻¹): 0.8175 <i>m</i> 6-aminocaproic acid + 0.6125 <i>m</i> HCl					
0.1378	-0.0724	-0.0883			
0.2067	-0.1056	-0.1291			
0.2756	-0.1234	-0.1668			
0.3445	-0.1442	-0.2013			
0.4134	-0.1697	-0.2329			
0.4823	-0.1828	-0.2618			
0.5511	-0.1975	-0.2883			
0.6200	-0.2112	-0.3127			
0.6889	-0.2206	-0.3353			

^aSee footnotes in Table I.

$\log K$ for reaction (1) for the amino acids over the temperature range studied are shown in Fig. 4. As expected, the protonation reactions for the α -amino acids have $\log K$ close to each other. The same trend with chain length as those for ΔH° and $T\Delta S^\circ$ is observed for $\log K$ values among the amino acids. At each temperature, the stability of the protonated species increases as the number of methylene groups between the amino group and carboxylate group increases. This is a result of the progressively larger positive change in ΔS° as the two groups become more separated (Fig. 3), and $\log K$ are dominated by $T\Delta S^\circ$. The protonation reaction is entropy driven. Each $\log K$ vs. T plot exhibits a minimum as ΔH° changes from negative to positive. At temperatures higher than that at the minimum, $\log K$ increase with temperature.

The increase in ΔH° and $T\Delta S^\circ$ as the temperature increases and the effect of charge separation on these thermodynamic quantities can be explained in terms of solvent effects. When protonation occurs, water molecules are released from the inner sphere of hydration to the bulk phase. Considerable energy is required for the newly released water molecules to attain the same energy level as that of the bulk water. As temperature increases, water molecules in the bulk phase have higher enthalpy and entropy in part due to the decrease of the hydrogen bonding.⁽³⁹⁾ The water molecules in the hydration spheres of ions are in a more ordered state than those in bulk water due to the interaction between the water dipoles and the charged ions. The result is that progressively larger positive changes in ΔH° and ΔS° occur as temperature increases resulting also in positive ΔC_p° values.

Table III. Coefficients for Pitzer Parameters, Eq. (2)^a

Species interactions	ϕ	$10A$	10^3B	10^5C	10^7D
${}^+\text{NH}_3\text{CH}_2\text{COOH}/\text{Cl}^-$	$\beta^{(0)}$	0.80	2.4589	-3.1916	0.9235
	$\beta^{(1)}$	1.80	2.0303	-2.7057	0.9673
	C^ϕ	-0.162	-8.5442	12.526	-4.8707
${}^+\text{NH}_3\text{CH}(\text{CH}_3)\text{COOH}/\text{Cl}^-$	$\beta^{(0)}$	0.80	-3.2634	4.9276	-2.7256
	$\beta^{(1)}$	1.80	-5.7910	11.4495	-6.0393
	C^ϕ	-0.162	11.0630	-17.5215	9.2247
${}^+\text{NH}_3\text{CH}(\text{CH}_2\text{CH}_3)\text{COOH}/\text{Cl}^-$	$\beta^{(0)}$	0.80	-0.5700	0.0040	-0.1067
	$\beta^{(1)}$	1.80	1.3580	-0.1454	
	C^ϕ	-0.162	0.2684	0.2960	
${}^+\text{NH}_3(\text{CH}_2)_2\text{COOH}/\text{Cl}^-$	$\beta^{(0)}$	0.80	3.4351	-3.8680	1.1236
	$\beta^{(1)}$	1.80	0.8911	0.7861	-1.0266
	C^ϕ	-0.162	-0.04745	-1.7458	1.9886
${}^+\text{NH}_3(\text{CH}_2)_3\text{COOH}/\text{Cl}^-$	$\beta^{(0)}$	0.80	2.6048	-2.5363	0.7026
	$\beta^{(1)}$	1.80	2.3307	-2.4403	0.8442
	C^ϕ	-0.162	-3.0219	3.6656	-1.1870
${}^+\text{NH}_3(\text{CH}_2)_5\text{COOH}/\text{Cl}^-$	$\beta^{(0)}$	0.80	3.9377	-5.5425	2.1741
	$\beta^{(1)}$	1.80	6.0862	-6.1710	2.3434
	C^ϕ	-0.162	-7.5347	10.835	-4.1150
${}^+\text{NH}_3\text{CH}_2\text{COO}^-/\text{Cl}^-$	λ	0.05	-0.3274	-0.6245	0.1912
${}^+\text{NH}_3\text{CH}(\text{CH}_3)\text{COO}^-/\text{Cl}^-$	λ	0.05	-1.5564	-1.1972	0.6167
${}^+\text{NH}_3\text{CH}(\text{CH}_2\text{CH}_3)\text{COO}^-/\text{Cl}^-$	λ	0.05	-1.2584	-0.6594	0.3467
${}^+\text{NH}_3(\text{CH}_2)_2\text{COO}^-/\text{Cl}^-$	λ	0.05	-1.1271	0.3419	-0.9066
${}^+\text{NH}_3(\text{CH}_2)_3\text{COO}^-/\text{Cl}^-$	λ	0.05	-2.1163	0.3343	-0.3174
${}^+\text{NH}_3(\text{CH}_2)_5\text{COO}^-/\text{Cl}^-$	λ	0.05	0.1164	-3.5369	2.0736

^aAll parameters for ternary interactions, Ψ'_{caa} and Ψ'_{cca} ; parameters for the like charged ions, θ'_{aa} , θ'_{cc} ; $\beta^{(0)}$, $\beta^{(1)}$, and C^ϕ for $\text{OH}^-/{}^+\text{NH}_3(\text{CH}_2)_n\text{CHR}_1\text{COOH}$ pairs; and λ for zwitterions with other ionic species were taken to be zero. The maximum standard deviation of the fit, σ_s , is

$$0.031 \text{ J-min}^{-1} \text{ as calculated by } \sigma_s = \sqrt{\sum_{i=1}^n (Q_{\text{dil},ci} - Q_{\text{dil},mi})^2 / (n - m)}, \text{ where } Q_{\text{dil},ci} \text{ and } Q_{\text{dil},mi}$$

are the calculated and measured heats of dilution, respectively, in the unit of J-min^{-1} , n is the number of points measured in the data sets, and m is the number of parameters being fitted.

The magnitudes of ΔH° for the protonation of the carboxylate groups on the zwitterions are influenced by interactions between neighboring amino and carboxylate groups as well as solvation effects. Protonation of the carboxylate group results in the decrease (weakening) in the interaction between the positively charged amino group and the carboxylate group. This change in the interaction between the protonated and non-protonated groups is greatest for the α -amino acids where n is smallest and decreases as n increases. The enthalpy of the solvent system increases upon protonation since solvent molecules influenced by the charges on the proton and carboxylate groups are released to the higher energy state of the molecules in the bulk water. This change is smallest in the case of the α -amino acids since partial neutralization

Table IV. Thermodynamic Data for the Protonation of Amino Acids, Eq. (1) at Zero Ionic Strength

T	$\log K$	ΔH°	ΔS°	ΔC_p°	Method ^b	Ref
Glycine						
283.15	2.397	-5.90	24.3		C	12
298.15	2.349	-3.97	31.8		T	13
	2.35	-5.98	25.1		C	14
	2.350	-4.10	31.4		C	12
	2.351	-4.92	28.5	137.2	T	15
	2.35	-3.891	33.5		C	16
	2.347	-4.364	30.3		T	17
	2.348	-4.602	30.5		T	18
	2.358	-4.85	28.9		T	19
	2.352	-4.837	28.9		T	20
	2.352	-3.93	31.9		C	21
313.15	2.327	-1.97	38.5		C	12
323.15	2.32	-0.9	41	119	C	This Study ^c
348.15	2.32	1.9	50	103	C	This Study ^c
373.15	2.35	4.2	56	78	C	This Study ^c
398.15	2.40	5.7	60	43	C	This Study ^c
α -Alanine						
274.15	2.426	-6.318	23.4		T	22
283.15	2.392	-5.23	27.2		C	12
285.65	2.383	-5.063	27.6		T	22
293.15	2.350	-3.820	31.8		T	23
298.15	2.348	-3.14	34.3		C	12
	2.35	-2.573	33.5		C	24
	2.350	-3.368	33.9	147.7	T	15
	2.34	-3.012	34.7		T	23
	2.345	-2.43	36.8		C	21
303.15	2.332	-2.251	37.2		T	23
308.15	2.327	-1.544	39.3		T	23
310.65	2.330	-1.339	40.2		T	22
313.15	2.328	-0.96	41.4		C	12
	2.324	-0.900	41.4		T	23
318.15	2.322	-0.292	43.5		T	23
323.15	2.332	-1.046	47.7		T	22
	2.33	0.0	45	130	C	This Study ^c
348.15	2.34	3.0	53	105	C	This Study ^c
373.15	2.39	5.1	59	65	C	This Study ^c
398.15	2.44	6.1	62	11	C	This Study ^c
β -Alanine						
283.15	3.606	-6.69	45.6		C	12
298.15	3.551	-4.35	53.6		C	12
	3.55	-4.52	52.7		C	25
	3.55	-4.94	50.2		T	26
313.15	3.517	-2.59	59.0		C	12
323.15	3.51	-0.9	65	139	C	This Study ^c

Table IV. Continued

T	$\log K$	ΔH°	ΔS°	ΔC_p°	Method ^b	Ref
348.15	3.52	2.3	74	114	C	This Study ^c
373.15	3.56	4.7	81	77	C	This Study ^c
398.15	3.61	6.0	84	24	C	This Study ^c
DL-2-Aminobutyric Acid						
274.15	2.334	-4.561	28.0		T	22
283.15	2.315	-3.22	32.2		C	12
285.65	2.310	-3.138	33.1		T	22
298.15	2.286	-1.59	38.5		C	12
	2.288	-1.293	39.3	161.5	T	15
	2.286	-1.297	39.3		T	22
310.65	2.288	0.920	46.9		T	22
313.15	2.289	-0.17	43.5		C	12
323.15	2.297	3.473	54.8		T	22
	2.29	1.1	47	105	C	This Study ^c
348.15	2.31	3.6	55	95	C	This Study ^c
373.15	2.36	5.8	61	80	C	This Study ^c
398.15	2.42	7.5	65	58	C	This Study ^c
4-Aminobutyric Acid						
283.15	4.057	-3.14	66.5		C	12
298.15	4.031	-1.63	72.0		C	12
	4.03	-1.67	71.1		T	26
313.15	4.027	-0.04	77.4		C	12
323.15	4.04	2.1	84	126	C	This Study ^c
348.15	4.08	5.0	93	104	C	This Study ^c
373.15	4.14	7.2	99	69	C	This Study ^c
398.15	4.21	8.4	102	20	C	This Study ^c
6-Aminocaproic Acid						
274.15	4.420	-3.423	72.0		T	27
283.15	4.392	-0.42	82.8		C	12
285.65	4.387	-1.920	76.1		T	27
298.15	4.373	1.34	88.3		C	12
	4.373	0.033	83.7		T	27
	4.37	0.0	83.7		T	26
310.65	4.384	2.347	91.6		T	27
313.15	4.388	3.31	94.6		C	12
323.15	4.410	5.038	100.0		T	27
323.15	4.42	5.0	100	127	C	This Study ^c
348.15	4.49	8.0	109	108	C	This Study ^c
373.15	4.58	10.3	115	80	C	This Study ^c
398.15	4.68	11.9	119	40	C	This Study ^c

^aUnits: T, K; ΔH° , kJ-mol⁻¹; ΔS° , J-mol⁻¹-K⁻¹; and ΔC_p° , J-mol⁻¹-K⁻¹.

^bThe method of ΔH° determination is designated by C (calorimetric) or T (temperature variation of K).

^cThe estimated uncertainties are ± 0.03 in $\log K$, ± 0.4 in ΔH° , ± 3 in ΔS° , and ± 10 in ΔC_p° .

Table V. Coefficients for Eqs. (3–6) for Reaction (1)

Amino Acid	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>
Glycine	-86.6765	4000.204	13.27332	7.06975
DL- α -Alanine	-116.0316	5201.463	17.71994	10.97738
DL-2-Aminobutyric Acid	-63.88209	2929.324	9.891402	4.400790
β -Alanine	-115.6580	5295.963	17.80919	10.59672
4-Aminobutyric Acid	-104.9958	4709.396	16.36708	9.804667
6-Aminocaproic Acid	-92.74088	4086.123	14.64205	8.045554

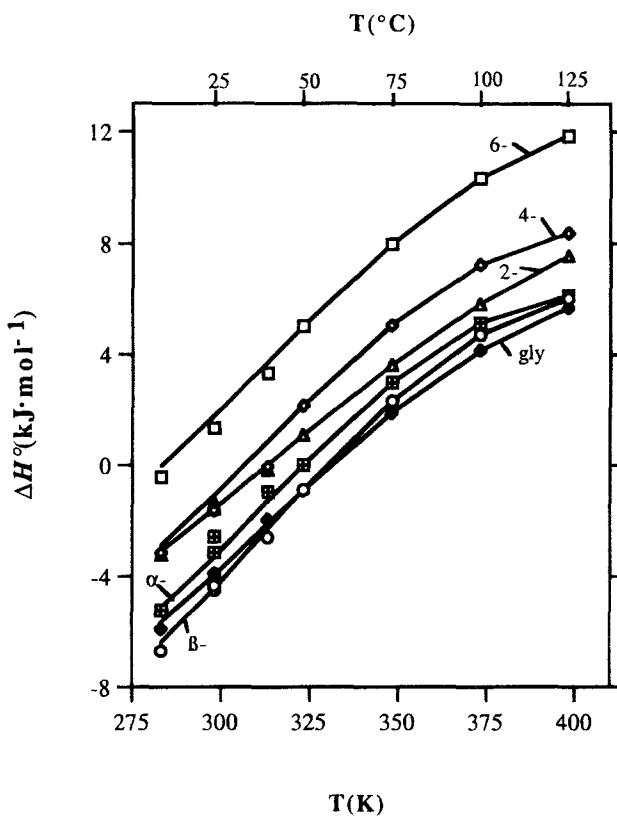


Fig. 1. Plot of ΔH° as a function of temperature for the protonation reaction: \square , 6-aminocaproic acid; \diamond , 4-aminobutyric acid; \triangle , DL-2-aminobutyric acid; \circ , β -alanine; \square , DL- α -alanine; \blacklozenge , glycine. The symbols at and above 323.15 K represent the results obtained in this study, all other points are taken from references listed in Table IV. The solid lines are based on Eq. (4).

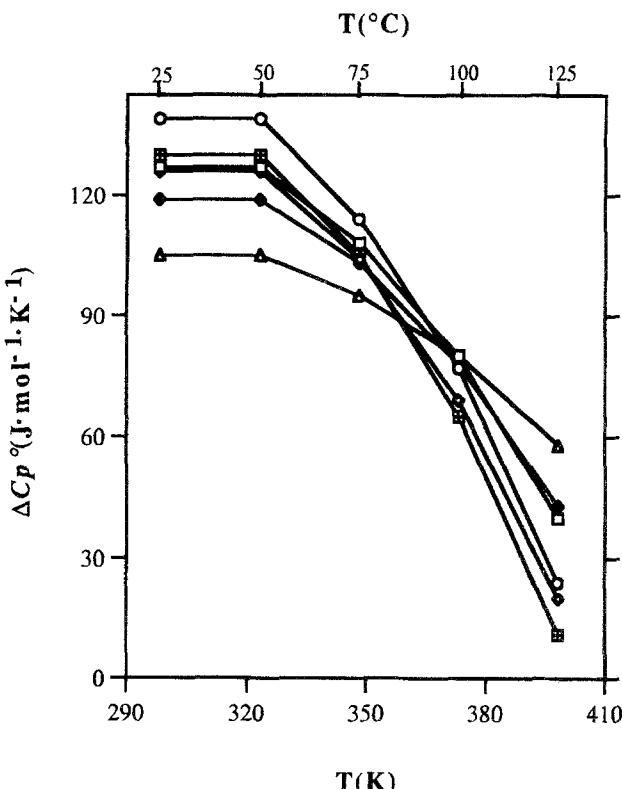


Fig. 2. Plot of ΔC_p° as a function of temperature for the protonation reaction based on Eq. (6).

of the charges on the zwitterion occurs when the two groups are close together and they have less interaction with water molecules. The net change in ΔH° as the number of carbon atoms between the groups changes is the result of these two effects. As n increases from 0 to 1 (α -amino acids to β -alanine), ΔH° at a given temperature decreases, but as n increases from 1 to 2 to 4, ΔH° increases as seen in Fig. 1. The reversal in trend with β -alanine ($n = 1$) is expected since the interaction between the groups drops off sharply resulting in less increase in ΔH° compared to that of the α -amino acids. The solute-solvent interaction increases and becomes dominant as n increases further.

The decrease in ΔC_p° with temperature for the protonation reaction can be explained in terms of long-range and short-range solvation effects. The zwitterion has no net charge, however it does contain both a positively and a negatively charged site. Therefore, the long-range effects of the zwitterion are similar to those of a neutral molecule,⁽²⁾ while the short-range effects are the same as those of two species containing single but opposite charges in close proximity. Viewed from the long-range perspective, the protonation of

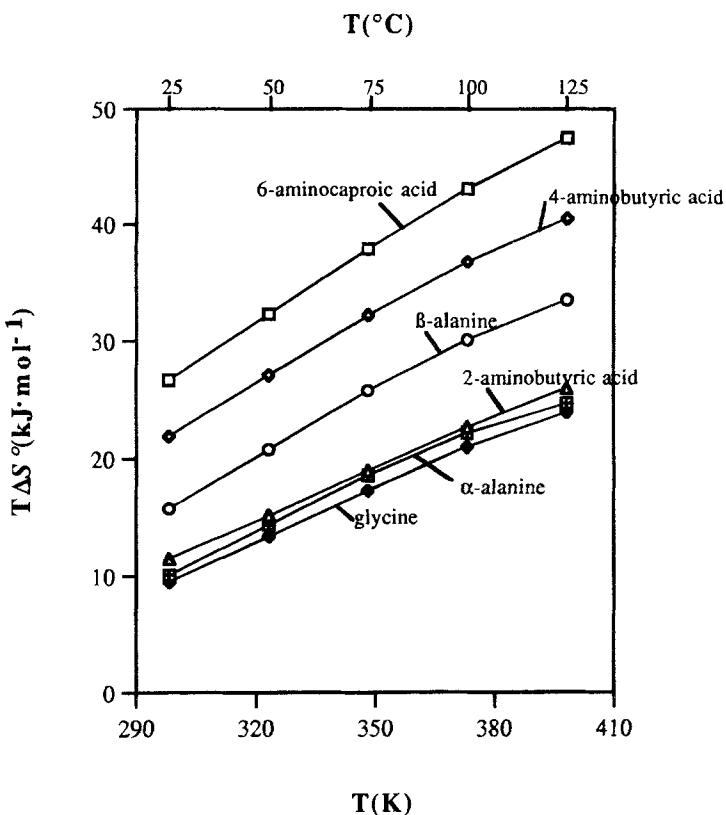


Fig. 3. Plot of $T\Delta S^\circ$ as a function of temperature for the protonation reaction: \square , 6-aminocaproic acid; \diamond , 4-aminobutyric acid; \triangle , DL-2-aminobutyric acid; \circ , β -alanine; \blacksquare , DL- α -alanine; \blacklozenge , glycine. The symbols at and above 323.15 K represent the results obtained in this study. The points at 298.15 K are extrapolated based on Eq. (5).

the carboxyl group is an isocoulombic reaction with both the reactants and products containing a single positive charge. From the short-range perspective, the reaction is the same as a neutralization reaction similar to the protonation of a simple carboxylic acid. It has been shown⁽⁴⁰⁾ that isocoulombic reactions have small ΔC_p° values resulting in relatively small, near linear changes in ΔH° with temperature. On the other hand, protonation reactions which result in charge reduction have large non-linear changes in ΔH° which accelerate as the temperature increases.^(41,42) In the relatively lower temperature range in the study, the short-range effect dominates. Therefore, larger ΔC_p° are observed due to the release of a large number of bound water molecules into the bulk phase as the carboxylate group is neutralized by a proton. As the temperature increases, the extent of hydrogen bonding decreases, causing the

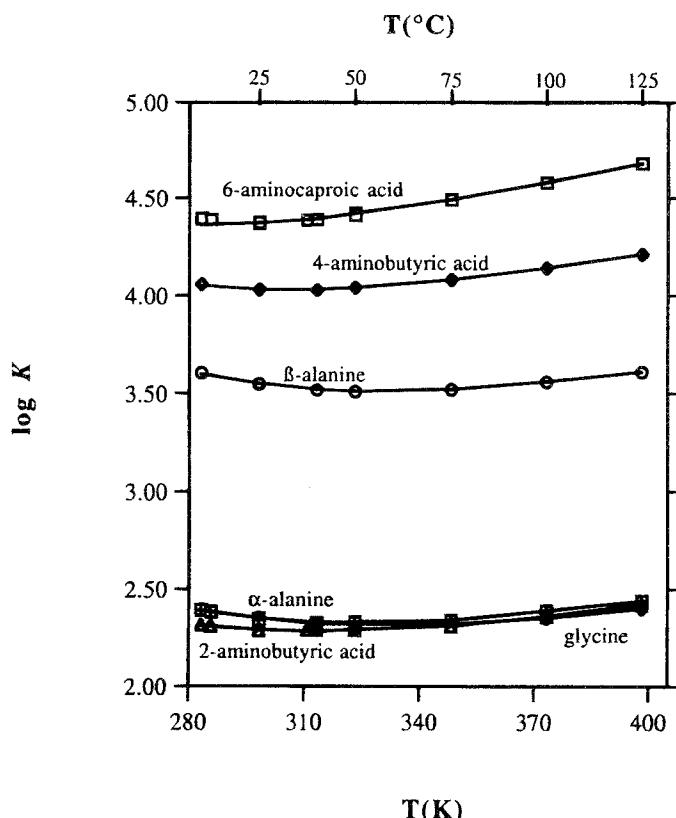


Fig. 4. Plot of $\log K$ for the protonation reaction as a function of temperature: □, 6-aminocaproic acid; ◇, 4-aminobutyric acid; △, DL-2-aminobutyric acid; ○, β-alanine; □, DL-α-alanine; ◆, glycine. The symbols at and above 323.15 K represent the results obtained in this study, all other points are taken from references listed in Table IV. The solid lines are based on Eq. (3).

dielectric constant of water to decrease and the long-range solvation effects become more important relative to short-range effects. Hence, ΔC_p° decreases as a result of the domination of the long-range solvation effects and the protonation reaction behaves more like an isocoulombic reaction. The ΔC_p° values for these reactions ranged from 105 to 139 J·mol⁻¹·K⁻¹ at lower temperatures ($T \leq 323.15$). These ΔC_p° values are approaching that for the protonation of acetate ion (~ 150 J·mol⁻¹·K⁻¹, as calculated from ΔH° from 273 to 333 K⁽⁵⁾), whereas ΔC_p° at 398.15 K ranged from 11 to 58 J·mol⁻¹·K⁻¹, which approach those of isocoulombic reactions.

The differences in the ΔS° values for the protonation reaction are primarily due to the release of solvent molecules.⁽³⁾ The smaller the value of n , the less influence the zwitterion will have on the solvent molecules. The reaction

of the amino acid zwitterion with the proton to form a positively charged species results in the smallest ΔS° when n is small (Fig. 3) because fewer water molecules are released on the formation of the positively charged species from the zwitterion which has charges close together and which appear to surrounding water molecules as a 'neutral' molecule. As n increases, the separated zwitterion charges influence an increasing number of water molecules. Thus, more water molecules are released to the bulk phase as the protonation of the carboxylate group occurs resulting in large ΔS° . Hence, the larger the n value, the larger will be ΔS° for the protonation reaction at a given temperature. This rationale is consistent with the observed trend of $T\Delta S^\circ$ with n as seen in Fig. 3. It is also seen that the difference in $T\Delta S^\circ$ among the amino acids increases as temperature increases. This "fanning out" effect was also observed in the protonation of the amino groups.⁽³⁾ The changes in the values both at a given temperature and as a function of temperature are consistent with the idea expressed earlier that the number of the released water molecules increases most with 6-aminocaproic acid and least with α -amino acids. The α -amino acids have the smallest change in $T\Delta S^\circ$ with temperature because fewer water molecules released to the bulk water of decreasing structure and dielectric constant. Therefore, as temperature increases fewer water molecules remain 'free' with a resulting increased entropy. As n increases, the charged sites have less effect on each other, more water molecules are released to the bulk solvent as the carboxylate group is protonated, and large increases in $T\Delta S^\circ$ are observed.

The increasing ΔH° and $T\Delta S^\circ$ with increasing chain length among the α -amino acids at a given temperature suggest that hydrophobic effects may be important in these ionization reactions. Details of the side chain effects on the thermodynamic quantities of the protonation of α -amino acids are discussed in a separate paper.⁽⁴³⁾

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