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Effect of potassium phosphate buffer on volumetric behavior of glycine in aqueous solutions at different temperatures

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article info abstract

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The apparent molar volume (V_{\emptyset}) of glycine in water and in 100, 500 and 1000 mM aqueous potassium phosphate buffer (KPB) solutions have been obtained from density measurements at different temperatures (288.15 to 328.15 K, including physiological temperature) and at different pH values (1.00, 7.40 and 14.00) by using vibrating-tube digital density meter. The estimated partial molar volumes at infinite dilution (V_2^0) have been used to obtain the corresponding transfer volume ($\Delta_{tr}V_2^o$) from water to aqueous potassium phosphate buffer solutions. Interaction parameters have been calculated from $\Delta_{tr}V_2^o$ data. The partial molar expansibilities $(\partial V_2^o|\partial T)$ _P at infinite dilution and $(\partial^2V_2^o|\partial T^2)$ _P values have also been determined from the V_2^o data at different temperatures. The hydration numbers (n_H) of glycine at pH 7.4 have been determined from the V_2^o data in the presence of KPB at different temperatures. The results have been discussed in terms of various interactions operating in these systems.

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1. Introduction

The study of interactions between salts and proteins is useful to understand the physiological systems [\[1](#page--1-0)–5]. However, proteins are complex molecules and their behavior in solutions is governed by a variety of interactions. Therefore the direct study of solute–solvent interactions of these systems is difficult [\[6\]](#page--1-0). One useful approach that reduces the degree of complexity in the study of these interactions is to study the interactions in systems containing smaller biomolecules such as amino acids. Amino acids are the building blocks of the proteins; their study provides important information which can be related to the behavior of larger biomolecules such as proteins [\[7\]](#page--1-0).

The physical and chemical characteristics of virtually all biologically relevant molecules depend strongly upon the pH of their aqueous environment. In order to maintain the biological function and catalytic activity, it is desirous that proteins maintain their native structure under physiological conditions. However, most proteins are sensitive to slightest change in cellular and environmental conditions like temperature, pH, pressure and the presence of salts. Moderate change in solution pH and/ or strength often results in drastic change in the conformational state of the protein and/or dissociation of protein complex.

Buffer solutions contain the salts that undergo reversible protonation and thus, aid in maintaining the pH of solution. This is particularly

Corresponding author. E-mail address: gdeepchem@yahoo.co.in (G. Singh). important in biological reactions that are often sensitive to small change in pH [\[8\].](#page--1-0) The intracellular and extracellular fluids of all organisms tend to have a characteristic and constant pH, which is regulated by various biological activities. Moreover, the first line of defense of living organisms against change in their internal pH is provided by buffer systems [\[9\]](#page--1-0). The phosphate buffer system, important in intracellular fluid, consists of conjugate acid–base pair $H_2PO_4^-$, as proton donor, and HPO_4^{2-} , as proton acceptor tends to resist change in pH in the range of (6.1 to 7.7) and would therefore be effective in providing buffering power in intracellular fluid whose pH is in the range of (6.9 to 7.4). Extensive work has been done on thermodynamic properties of amino acids and peptides in aqueous and mixed aqueous solutions of various additives which provide information about solute-solvent and solute-solute interactions operating in these systems [\[10](#page--1-0)–24]. However, these studies were not focused on physiological conditions or biological environment, i.e. at controlled pH and physiological temperature which is a prerequisite to understand the complex biological phenomena like protein folding/unfolding, protein dynamics, energetics, and stability. Literature survey reveals that no study on thermodynamic properties of amino acids in the presence of buffer solutions at controlled pH is available. Our aim in the present study is to determine the interactions of glycine in the presence of buffer solutions. So, we are the first to report, the partial molar volume of glycine in the presence of potassium phosphate buffer (KPB) solutions at different temperatures (288.15 to 328.15 K, including physiological temperature) and at specific pH values i.e. 7.40, 1.00 and 14.00, where amino acids exist predominantly in zwitter ionic, protonated and deprotonated form, respectively.

2. Experimental

Glycine, potassium di-hydrogen orthophosphate ($KH₂PO₄$) and dipotassium hydrogen orthophosphate (K₂HPO₄) of highest purity ≥99% were obtained from S.D. Fine Chem. Ltd. These were used as such without further purification. However, these were dried in vacuum oven and kept in a desiccator at room temperature before use. Double-distilled, deionized water was used to prepare the solutions. All the solutions were prepared afresh on weight basis in by using electronic balance (CITIZEN CY 204) having precision $\pm 1 \times 10^{-4}$ g. The required pH for water was attained by using 1 M KOH and 1 M Phosphoric acid (H_3PO_4) . Potassium phosphate buffer was prepared by adding KH_2PO_4 and $K₂HPO₄$. The desired pH value 14.00 and 1.00 was maintained by using KOH and H_3PO_4 . We were not able to attain pH 14.00 in KPB, so the present work has been carried out at pH 13.40. The solution densities were measured using vibrating-tube digital density meter (Anton Paar, DMA 4500 M). Temperature around the cell was controlled to \pm 1 × 10⁻² K by built in a solid state thermostat. The accuracy and precision in density measurements was found to be $\pm 5 \times 10^{-5}$ g \cdot cm⁻³ and \pm 1 × 10⁻⁵ g⋅cm⁻³, respectively. The working of density meter was checked by measuring the densities of aqueous sodium chloride solutions, which agreed well with the literature values [\[25\].](#page--1-0)

Table 1

 $^{\text{a}}$ m is the molality of glycine in water; standard uncertainties u in molality, u(m) = $1.3 \cdot 10^{-4}$ mol·kg⁻¹ and apparent molar volume, u(V_∅) = 0.08 to 0.001 cm³·mol⁻¹.

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