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# Title: Molecular crowding has no effect on the dilution thermodynamics of the biologically relevant cation mixtures

Running title: Influence of cation mixtures on molecular crowding

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Name	Affiliations
Daria Głogocka	1. Department of Fundamental Problems of Technology, Institute of Biomedical Engineering, Wroclaw University of Technology, Wroclaw, Poland
PhD Magdalena Przybylo	1. Department of Fundamental Problems of Technology, Institute of Biomedical Engineering, Wroclaw University of Technology, Wroclaw, Poland
Prof Marek Langner	1. Department of Fundamental Problems of Technology, Institute of Biomedical Engineering, Wroclaw University of Technology, Wroclaw, Poland

Corresponding author: Daria Głogocka <daria.glogocka@pwr.edu.pl>

#### **Abstract**

The ionic composition of intracellular space is rigorously maintained in the expense of high-energy expenditure. It has been recently postulated that the cytoplasmic ionic composition is optimized so the energy cost of the fluctuations of calcium ion concentration is minimized. Specifically, thermodynamic arguments have been produced to show that the presence of potassium ions at concentrations higher than 100 mM reduce extend of the energy dissipation required for the dilution of calcium cations. No such effect has been measured when sodium ions were present in the solution or when the other divalent cation magnesium was diluted. The experimental observation has been interpreted as the indication of the formation of ionic clusters composed of calcium, chloride and potassium. In order to test the possibility that such clusters may be preserved in biological space the thermodynamics of ionic mixtures dilution in solutions containing albumins and model lipid bilayers have been measured. Obtained thermograms clearly demonstrate that the energetics of calcium/potassium mixture is qualitatively different from calcium/sodium mixture indicating that the presence of the biologically relevant quantities of proteins and membrane hydrophilic surfaces do not interfere with the properties of the intracellular aqueous phase.

Keywords: Ionic clusters in aqueous solution; Molecular crowding; Membrane crowding; Isothermal titration calorimetry

# Changelog

We modify the text so any controversial statements are avoided. Statements relating to the ionic clusters were reformulated. For example, the statement "the persistence of ionic clusters" is replaced by its modified version "the persistence of the thermodynamic properties of the dilution process is not affected by the presence of albumins or liposomes". Information about the number of repetitions was added.

1 Molecular crowding has no effect on the dilution thermodynamics of the 2 biologically relevant cation mixtures 3 Daria Głogocka<sup>1,2,\*</sup>, Magdalena Przybyło<sup>1,2</sup>, Marek Langner<sup>1,2</sup> 4 5 6 <sup>1</sup> Department of Biomedical Engineering, Faculty of Fundamental Problems of Technology, 7 Wrocław University of Technology, Wybrzeże Wyspiańskiego 27, 50-370 Wrocław, Poland, +48 71 320 4155 8 <sup>2</sup> Lipid Systems Sp. z o.o., ul. Duńska 9, 54-427 Wrocław, Poland 9 10 Influence of the molecular crowding on cation mixtures thermodynamics **HIGHLIGHTS** 11

Calorimetric data shows that biologically relevant concentration of water-soluble protein (50 mg/ml) 12

13 do not affect the dilution thermodynamics of calcium-chloride-potassium mixture.

Calorimetric data shows that neutral lipid surfaces at concentrations similar to that in the cytoplasm (7

mg/ml) do not interfere with the dilution thermodynamics of calcium-chloride-potassium mixture.

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# **KEYWORDS**

Dilution thermodynamics of ionic mixtures, Molecular crowding, Membrane crowding, Isothermal titration calorimetry, Thermodynamics of calcium intracellular signaling

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## **ABSTRACT**

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potassium ions at concentrations higher than 100 mM reduce extend of the energy dissipation 27

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ions were present in the solution or when the other divalent cation magnesium was diluted. 29

30 The experimental observation has been interpreted as the indication of the formation of ionic

clusters composed of calcium, chloride and potassium. In order to test the possibility that such 31

32 clusters may be preserved in biological space the thermodynamics of ionic mixtures dilution

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Correspondence to: Daria Głogocka, Department of Biomedical Engineering, Faculty of Fundamental Problems of Technology, Wrocław University of Technology, Wybrzeze Wyspianskiego 27, 50-370 Wroclaw, Poland E-mail address: daria.glogocka@pwr.edu.pl

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## 1. INTRODUCTION

40 Ionic homeostasis inside cell is a critical element of proper progression of all its vital functions. The ionic composition of intracellular aqueous phase affects solubility and 41 42 aggregation state of all molecules filling the cell volume (Hey, Jackson, & Yan, 2005; Huang, 43 Liu, Cheng, & Huang, 2015; Kunz, 2010). Ions also participate in many metabolic processes as necessary cofactors (Murgia, Giorgi, Pinton, & Rizzuto, 2009; J. Spitzer & Poolman, 2009). Their 44 electrochemical gradients are necessary for the energy and information transformations 45 46 (Bortner & Cidlowski, 2004; Ivannikov, Sugimori, & Llinas, 2010). The spatio-temporal distribution of ions is facilitated by an intricate system of fluxes between various 47 compartments driven by dedicated channels, pumps and chelating agents (Boldyrev, 1993; 48 Chen, Yu, & Wei, 2014; Decrock et al., 2011; Eisenberg, 2013; Lee, Davis, Roberts-Thomson, & 49 Monteith, 2011; Yu, Canzoniero, & Choi, 2001). Whereas the passive ion flux in the cell is 50 governed by electrochemical gradients the active transport is associated with energy 51 52 dissipation. Nearly 70% of adenosine triphosphate (ATP) in some cells is used to maintain the 53 ionic homeostasis (Ivannikov et al., 2010). There are four main cations present in the cytoplasm. The averaged concentrations of monovalent potassium (about 140 mM) and sodium (about 4 54 mM) as well as divalent magnesium (3 mM) are constant in time. Calcium is an unusual 55 cation, which concentration fluctuates between 10<sup>-10</sup> M and 10<sup>-4</sup> M) serving as a second 56 57 messenger in intracellular signaling cascades (Ivannikov et al., 2010; Landolfi, Curci, Debellis, Pozzan, & Hofer, 1998; Michelangeli, Ogunbayo, & Wootton, 2005). Such large concentration 58 59 changes require expenditure of large quantities of metabolic energy reaching 17% of the 60 entire energy dissipated by cells (Case et al., 2007; Eisner, Diaz, Li, O'Neill, & Trafford, 2004; Ivannikov et al., 2010). Recently, we have demonstrated that the thermodynamics of the 61 dilution of calcium ions in water is greatly affected by the concentration and type of 62 monovalent cation (Kaczynski, Borowik, Przybylo, & Langner, 2014). This finding provides the 63 thermodynamic argument for the specific ionic composition of the intracellular space. It has 64 65 been demonstrated that the thermodynamics of calcium dissolution depends on the combination of ion-water and ion-ion interactions (Arias-Moreno, Cuesta-Lopez, Millet, Sancho, 66 67 & Velazquez-Compoy, 2010; Landolfi et al., 1998; Liu, Lu, Meijer, & Wang, 2010; Yang et al., 2010). Whereas the dissolution energy of calcium is independent on the sodium chloride 68 concentration, and do not differ significantly from that in water, the effect of the potassium 69 70 chloride is dramatic. When the potassium quantity rises above the concentration of 100 mM 71 the dilution energy of calcium is dramatically reduced. No such effect is observed for 72 magnesium, the other important divalent cation present in the cytoplasm (Kaczynski et al., 2014). In order to explain the effect the formation of multi-ionic clusters have been postulated 73 74 (Kaczynski et al., 2014). Such multicomponent ionic clusters would require the sufficient level 75 of hydration, which might be not available in densely packed intracellular space (Leontidis, Christoforou, Georgiou, & Delclos, 2014; J. Spitzer & Poolman, 2009; J. J. Spitzer & Poolman, 2005). 76 77 In addition, the large quantity of, frequently charged, water-soluble proteins and membrane surfaces might affect organization, dynamics and thermodynamics of ions in the intracellular 78 79 aqueous phase (Ben-Yaakov, Andelman, Harries, & Podgornik, 2009; Collins, 1997; Collins,

Neilson, & Enderby, 2007). To evaluate the effect of two most numerous components of crowded intracellular environment, namely hydrophilic proteins and lipid surfaces, on the hydration thermodynamics of various mixtures of cations have been investigated. For that purpose the water-soluble proteins were modeled with albumins and membrane surfaces were modeled with unilamellar liposomes, both entities in biologically relevant concentrations 50 mg/ml and 7 mg/ml, respectively (Miyoshi & Sugimoto, 2008).

#### 2. MATERIALS AND METHODS

#### 2.1. Materials

Bovine serum albumin (BSA  $\geq$  98.0%) and hepes (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid) buffer were obtained from Sigma-Aldrich (Poznan, Poland), calcium chloride, propylene glycol and sodium chloride were purchased from Chempur (Piekary Slaskie, Poland) and potassium chloride was obtained from POCH (Gliwice, Poland) EggPC 90G from Lipoid GmbH (Ludwigshafen, Germany) were of analytical grade and used without further purification. 50mM hepes buffer (pH=7.4) was prepared using fresh deionized water (conductivity  $< 2\mu$ S). All solution containing mixtures of salts were tested for the presence of precipitation using the dynamic light scattering technique. In all cases no particulates larger than 2 nm were detected.

# 2.2. Liposomes preparation

EggPC at concentration 45% [w/w] was dissolved in propylene glycol and incubated for 24h at 60°C. The addition of 50mM hepes buffer was followed by further incubation for 24h at 60°C to ensure the complete lipid hydration. Next, lipid suspension was extruded 5 times through polycarbonate membrane filters (Whatman, UK). Liposomes were diluted with buffer so that the final concentration of lipid was 7 mg/ml. Size of liposomes was measured using Dynamic Light Scattering (DLS) technique (NanoSizer, Malvern GB). In order to measure the effect of the osmotic pressure difference, caused by high salt concentration, on the liposomes size distribution, the DLS experiment was performed at the end of each titration. No significant change was observed (results not shown).

## 2.3. Isothermal titration calorimetry (ITC) measurements

Sample degassing for 30 minutes preceded each titration experiment. Calorimetric measurements were performed using Thermal Activity Monitor TAM III, equipped with isothermal titration unit (TA Instruments, New Castle, USA). Measurement cell was filled with 800 µl of BSA buffered solution (50 mg/ml), liposomes buffered solution (7 mg/ml) or buffer solution alone. Solutions from syringe were added into cell in a series of 10 µl injections. Each experiment was preceded with a 1µl injection, which was neglected in the subsequent analysis. The duration of each injection was 10 s and the delay time between injections 1500 s. Experiments were carried out at 25.0°C. In the experiment buffered solutions of CaCl<sub>2</sub>, KCl, NaCl or their mixtures were titrated into the cell. Concentration of each salt in the mixture was 1.5 M. This ensures that the final concentration of salts in the cell reaches the physiological values. Concentration of BSA and lipids in measurement cell were also selected based on physiological values (Chtterjea & Shinde, 2012). Heat of the dilution was

calculated from thermograms as an area under peaks expressed in Jules. Positive or negative heat flow indicated an exothermic or endothermic process, respectively. All experiment presented in the paper were carried in triplicates using different preparations of proteins or liposomes. Control experiments were performed in duplicates.

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#### 3. RESULTS AND DISCUSSION

The ionic composition of the cell interior is tightly controlled and meticulously maintained 131 despite high-energy cost. Whereas the concentrations of most intracellular free ions are 132 invariant in time, the concentration of calcium varies widely by orders of magnitudes 133 134 facilitating the information transfer through cytoplasm (Cheng & Lederer, 2008). These fluctuations, in addition to the maintenance of trans-membrane gradients, increase the amount 135 of energy required to sustain the intracellular ionic homeostasis (Ivannikov et al., 2010). The 136 high-energy cost of information transfer facilitated by Ca<sup>2+</sup> necessitates measures to optimize 137 the physicochemical condition of Ca<sup>2+</sup> fluctuations (Kaczynski et al., 2014). The type of 138 monovalent cation present in the solution changes the dilution energetics not only 139 quantitatively but also qualitatively, specifically the dilution of calcium mixed with potassium 140 is an exothermic process but, when potassium is replaced with sodium, the dilution becomes 141 142 endothermic. Based on this observation it has been postulated that ions in aqueous phase form ionic clusters, which may dramatically change the thermodynamic properties of the solution. 143 Ionic clusters are attractive concepts, which offers straightforward explanation, albeit not 144 confirmed by other experimental evidences, of differences between dilution thermodynamics 145 of various ionic mixtures (Kaczynski et al., 2014). Data presented previously have been 146 carried out using biologically relevant concentration of ions but the experimental model has 147 not accounted for the crowding effect produced by high concentrations of proteins and 148 149 membranes. The cell volume is a maze of aqueous compartments packed with water-soluble proteins (proteins occupy about 40% of cell volume) bordered by variety of membrane 150 151 surfaces (Chandler, 2005; Chtterjea & Shinde, 2012). In such an environment the existence of intracellular "bulk" water is unlikely and the effect of various surfaces on distribution of ions 152 is likely. The possible effect of macromolecular and membrane crowding on the structure and 153 dynamics of aqueous phase leads to the model, which treats the "aqueous phase" in cell as a 154 structured mixture of water molecules, ions and small hydrophilic compounds (Leontidis et 155 al., 2014; J. Spitzer & Poolman, 2009; J. J. Spitzer & Poolman, 2005). Despite the crowded 156 conditions and low water activity the cell interior is still capable to provide conditions for the 157 propagation of intracellular signals especially these based on fluctuations of calcium 158 concentration (Swietach, Spitzer, & Vaughan-Jones, 2015; Szekely et al., 2011; Taheri-159 160 Araghi & Ha, 2010; Thurley, Skupin, Thul, & Falcke, 2012; Wakai & Fissore, 2013). Calcium is continuously released and reloaded into intracellular stores during the signaling 161 events (Joseph, Reicher, & Barda-Saad, 2014). The calcium signal can last from seconds to 162 hours therefore the intracellular aqueous compartments with elevated calcium concentration 163 should not affect the hydrophilic-hydrophobic balance, which maintain the stability of 164 concentrated intracellular suspension of proteins and membranes (Joseph et al., 2014; Thurley 165 et al., 2012). We have previously demonstrated that the dissolution thermodynamics of 166 mixtures of calcium and potassium chlorides indicate that the two cations combined with 167

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168 chloride ions may form stable clusters, which do not disintegrate upon dilution (Kaczynski et
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- al., 2014). It has also been also observed that the thermodynamics of ionic mixture dilution is
- mainly controlled by the potassium concentration regardless on the quantity of calcium
- present in the mixture what substantiate the assumption of the presence of ionic clusters.
- However, the crowded cell interior may interfere with dilution thermodynamic by altering
- local ionic composition. Surfaces of membranes and/or proteins may alter distribution of ions
- in the adjacent aqueous phases (Parsegian & Zemb, 2011; Parsons, Bostrom, Lo Nostro, &
- Ninham, 2011; Pollack, 2003; Porasso & Cascales, 2009; Przybyło, Drabik, Łukawski, &
- Langner, 2014; Zhang & Cremer, 2006). In order to determine the effect of water-soluble
- proteins and neutral membranes on the dilution thermodynamics the calorimetric experiment
- has been performed at the presence of biologically relevant concentrated protein solution and
- lipid vesicles suspension. The concentrated solution of albumin (50 mg/ml) was used as a
- model of intracellular aqueous space crowded with proteins, whereas concentrated suspension
- of lipid vesicles (7mg/ml) was used to model the effect of membrane surfaces on ions
- dissolved in the aqueous phase. Figure 1 shows thermograms of two salts mixtures
- NaCl:CaCl<sub>2</sub> (1:1) and KCl:CaCl<sub>2</sub> (1:1) titrated into buffer or concentrated albumin or
- membrane solutions.
- Panel C and F on Figure 1 show that when the buffer was titrated with mixtures of KCl:CaCl<sub>2</sub>
- 186 (1:1) and NaCl:CaCl<sub>2</sub> (1:1) qualitatively different thermograms were acquired. Whereas the
- dilution process of CaCl<sub>2</sub>:KCl mixture is endothermic the dilution of CaCl<sub>2</sub>:NaCl mixture is
- exothermic. The explanation of this difference has been offered previously (Kaczynski et al.,
- 2014). Assuming that mixtures of CaCl<sub>2</sub> and NaCl salts produce homogenous solvents their
- dilution thermograms can be approximated with the sum of dilution enthalpies of each salts
- individually. The situation is different when mixtures of CaCl<sub>2</sub> and KCl are diluted. The
- 192 quantity of energy released in this case does not differ significantly form the energy released
- when the KCl solution is diluted alone, indicating that in this case the solution is not
- homogeneous. The effect was explained by the proposition that the solution produced by
- mixtures of CaCl<sub>2</sub> and KCl salts contain dilution resistant clusters of interacting ions. Panel C
- in Figure 2 and Figure 3 demonstrate the difference between the two mixtures of salts. It has
- been proposed previously that stable ionic Ca<sup>2+</sup>:Cl<sup>-1</sup>:K<sup>+1</sup> clusters are formed in sufficiently
- high potassium concentrations preventing the change of Ca<sup>2+</sup> hydration upon dilution
- 199 (Kaczynski et al., 2014).
- 200 In order to highlight differences between titration experiments plots of cumulative enthalpy
- 201 changes are fitted with linear function and the slope is then calculated and presented in Figure
- 202 3.
- 203 Figure 3 quantitatively summarizes results obtained for each titrant, expressed as
- abovementioned slopes, derived using the regression analysis, for ions and their mixtures
- 205 titrated into buffer, protein solution or lipid suspension. Data presented in Figure 3 shows that
- energetics of CaCl<sub>2</sub>:KCl (1:1) dilution with the buffer, albumin solution or suspension of
- 207 liposomes is qualitatively similar. This indicates that postulated ionic clusters remain intact in
- 208 the aqueous phase at biologically relevant concentrations of water-soluble proteins and lipid
- 209 membranes therefore supporting the preposition that ionic clusters containing Ca<sup>2+</sup> and K<sup>+</sup>
- 210 cations may reduce the energetic cost of the intracellular signaling mediated by spatio-
- 211 temporal fluctuations of Ca<sup>2+</sup> ion concentration. No such effect is observed when mixtures of

Ca<sup>2+</sup> and Na<sup>+</sup> cations are used. Despite the lack of qualitative effects of proteins or 212 membranes, on the overall character of the dilution thermodynamics of salts and their 213 214 mixtures there are some quantitative differences. The presence of albumin and lipid 215 membrane results with the elevated enthalpies of calcium dissolutions by 11 % and 17 %, respectively. This indicates that there are some interactions between calcium and/or chlorine 216 ions with proteins and lipid surfaces. This result agrees with other studies, which have 217 218 demonstrated that polyions bind with variety of compounds and structures (Ahyayauch, Arana, Sot, Alonso, & Goni, 2009; Arias-Moreno et al., 2010; Morton, Garland, & Holden, 219 2010; Parsons et al., 2011). The effect is bigger when lipid bilayers are present in the solution. 220 Analysis of NaCl and KCl dilutions demonstrate that the effect of protein on the two ions is 221 small (8% and 5 %, respectively) as compared to the effect of lipid surfaces. The effect of 222 lipid surface on the sodium dilution (changed by 55%) is significantly stronger then on the 223 potassium dilution (changed by 34%) showing that interaction between sodium and lipid 224 surfaces is significant. The result agrees with data presented in literature (Klasczyk, Knecht, 225 226 Lipowsky, & Dimova, 2010; Przybylo et al., 2014; Vlachy et al., 2009; Yang et al., 2010). When mixtures of potassium and calcium is diluted at the presence of albumins and lipid 227 membranes the absolute value of dilution enthalpy is reduced in a meaner similar to that of 228 potassium chloride alone showing that when in the mixture calcium is not readily available 229 230 for interaction with crowding agents. When the mixture of sodium and calcium is added to 231 proteins and membranes the resulting dilution enthalpy is elevated by 46% and 66%, respectively. This result shows that the two cations interact with both proteins and membranes 232 233 with little mutual interferences.

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There are two factors, which needed to be accounted for when analyzing phenomena occurring in crowded and/or concentrated solutions. First, ions are different with the respect to their charge, surface charge density and hydration what affects their solubility and, as demonstrated previously, propensity to form ionic complexes (Collins, 1995, 1997; Collins et al., 2007). The ionic complexes may alter the Debye effect, especially at charged endomembrane surfaces (Langner & Kubica, 1999) (Vlachy et al., 2009; Yang et al., 2010). In this case the main assumptions of the Gouy-Champman theory may not be satisfied limiting dramatically applicability of the theory when intracellular processes are analyzed (Ben-Yaakov, Andelman, Podgornik, & Harries, 2011). Any local alteration of ionic composition may change hydrophobic/hydrophilic balance of surrounding macromolecules leading to their precipitation or aggregation as dramatically demonstrated by Hofmeister effect (Baldwin, 1996; Hey et al., 2005; Kunz, Lo-Nostro, & Ninham, 2004). Whereas the effect of monovalent ions on macromolecules solubility and/or propensity to aggregate is relatively weak the effect of divalent ions can be dramatic. For example, calcium may induce fusion of negatively charged membranes present in high quantities in the cell cytoplasm (Boettcher et al., 2011; Michelangeli et al., 2005; Naranjo & Mellstrom, 2012). The presence of ionic clusters containing calcium may contribute to the preservation of the electrostatic balance when the calcium concentration rises upon cell stimulation therefore ensuring the topological stability of endomembrane systems. The calcium release may induce intracellular molecular rearrangements, which may result from direct interaction of calcium ion with proteins and/or membranes triggered by the altered intracellular ionic composition (Tojyo, Morita, Nezu, & Tanimura, 2014). Data presented in the paper shows that the

thermodynamics of ionic mixtures dilution is not qualitatively altered by the presence of interfaces presented by membranes and proteins showing that underlying molecular processes are likely to be preserved in a crowded intracellular environment.

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#### 4. CONCLUSIONS

The ionic composition of the cytoplasm determines its stability (hydrophobic/hydrophilic balance) and facilitates the transmission of signals through the cell volume. Whereas the major elements of the ionic homeostasis are qualitatively and quantitatively well characterized the exact ionic composition of cell has not been convincingly justified. Recently it has been proposed that the specific composition of the cellular space is determined by the thermodynamic of the collective hydration of ions. It has been demonstrated that the thermodynamics of dilution of mixtures of ions dramatically depends on their ionic composition. Specifically, the dilution of calcium and potassium chlorides mixture is qualitatively different from all other mixtures. It has been speculated that the difference is a result of ionic clusters formation, which may ensure stable level of calcium hydration regardless on their concentration. The significance of such ionic clusters in biological space filled with tightly packed soluble proteins and membranes depend on the cluster stability at the presence of various hydrated surfaces. Results of calorimetric experiments present in the paper demonstrate that the presence biologically relevant density of hydrophilic proteins and lipid bilayers do not influence the dilution thermodynamics of ionic mixtures containing calcium cation. This result shows that the ionic composition of the cytoplasm may play critical role in the stability of crowded biological space by ensuring the solubility of various components of cell volume.

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#### 5. APPENDICES

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Figure 1. Thermograms of buffer (Panels C and F), 50 mg/mL albumin solution (Panels A and D) and 7 mg/mL lipid vesicles suspension (panel B and E) titrated with KCl:CaCl2 (1:1) mixture (A, B, C) and NaCl:CaCl2 (1:1) mixture (D, E, F).

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- Figure 2. Cumulative enthalpy changes calculated from the dilution experiments carried out for selected salt solutions as a function of the salt concentration in the reaction cell in the presence of BSA (A), liposomes (B) and buffer alone (C). CaCl2 (gray triangle). KCl dilution experiment is labeled with the reversed gray triangle, the dilution of KCl:CaCl<sub>2</sub> (1:1) mixture is labeled with gray square, NaCl labeled with black circle, and NaCl:CaCl<sub>2</sub> mixture labeled with black rhombus.
- Figure 3. The slopes of the linear dependence of the cumulative enthalpies on ions concentration as determined in the dilution experiments 50 mg/mL albumin solution and 7 mg/mL liposome suspension. The calculated values of the slopes for ionic mixtures, expressed as a sum of enthalpies obtained in the dilution experiments performed for each salt alone, are presented for comparison.

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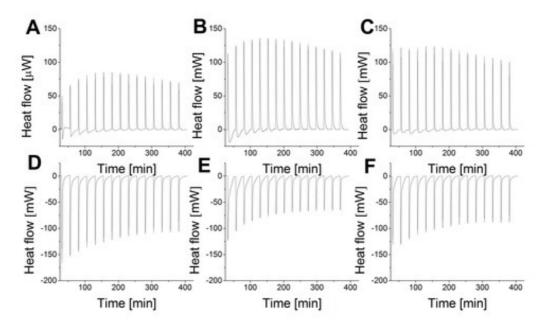


Fig. 2 <u>Download full resolution image</u>

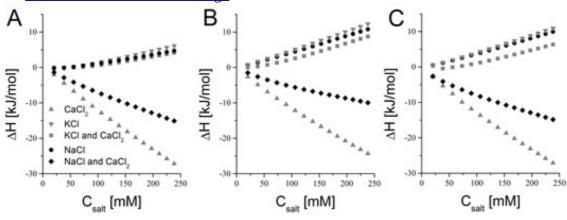


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