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### <sup>1</sup> Urea and guanidine salts as novel components for deep eutectic solvents

### Q1 Jozef Parnica<sup>a</sup>, Marian Antalik<sup>a,b,\*</sup>

3 <sup>a</sup> Department of Biochemistry, Faculty of Science, P.J. Safarik University, 04011 Kosice, Slovak Republic

4 <sup>b</sup> Institute of Experimental Physics, Slovak Republic

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#### 29 1. Introduction

Room temperature ionic liquids (RTILs) are low melting-temperature 30 salts (below 373 K) formed by bulky and asymmetric ions that hinder 31crystallization [1]. Properties of RTILs, such as viscosity, density, melting 32point or solubility in water can be tuned by changing either the anion 33 or cation [2]. Deep eutectic solvents (DES) were reported by Abbott and 34 co-workers [3] as extending model of ionic liquids. DES are generally 35 defined as salts with a melting point less than 100 °C exhibiting unique 36 characteristics: very low vapor pressures, non-flammability, high thermal 37 and chemical stability, possibility of recycling, good solubility for a wide 38 39 spectrum of compounds and high suitability for modifications [1,4–6]. 40 DES have large depression of the freezing point of mixtures of salts with hydrogen donors (e.g. amines, amides, alcohols and carboxylic acids) 41 [7]. The interaction of this hydrogen bond donor with the salt reduces 42 the anion-cation electrostatic force, and thus, the freezing point of the 43 44 mixture decreases. The enormously large number of salts and hydrogen bond donors that can be used to prepare DES compounds caused 45that there is no limit of combination of cations with anions. A classical 46 47 example is the combination of choline chloride (m.p. 302 °C) with urea (m.p. 132 °C), forming a DES with a melting point of 12 °C [8]. 48

Urea is a powerful protein denaturant as it disrupts the noncovalent
 bonds in the proteins. This group of chaotropic agents includes also
 ionic guanidine hydrochloride (GuHCl) and guanidine thiocyanate
 (GuSCN). These guanidine-based compounds are well known dena turing reagents and are used extensively in protein chemistry. Urea

E-mail address: antalik@saske.sk (M. Antalik).

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#### ABSTRACT

We present guanidinium salts in combination with urea, used to prepare a new combination of deep eutectic 17 solvents (DES). They have recently attracted much attention as "green" alternatives to conventional organic 18 solvents in various fields including biophysical chemistry. These were subsequently used to test the stability 19 and solubility of proteins in a water-free DES. We performed CD analyses in order to monitor global secondary 20 and tertiary structural changes of cytochrome c in the presence of Guanidine HCl-urea. Formation of the melt 21 form of DES at a room temperature was an effective way to store soluble protein in the denatured form for a 22 long time. Consequent dialysis allows one to recover proteins into their native form. 23

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and guanidine salt denaturation curves are generally used to obtain 54 knowledge of conformational stability of proteins [9–11]. The eutectic 55 system GuHCl–GuSCN (eutectic point lies at 87 °C, 60 mol% of GuSCN) 56 has been studied recently as well as several other combinations of gua-57 nidine salts such as GuFormate–GuSCN (eutecticum lies at 45 °C, 58 30 mol% of GuSCN) [12]. Eutectic points were not in a molar ratio as 59 in the case of choline chloride and urea. The paper presents the results 60 of some characteristic DES formed between urea and GuHCl as well as 61 urea and GuSCN and its effect on the structure of ferricytochrome c 62 detected by CD spectroscopy. These ionic liquids allow the high solubility 63 of cytochrome c and bovine serum albumin in these non-aqueous 64 medium and we discussed the possibility of using this type of DES for 65 long-term storage of proteins at room temperature.

#### 2. Materials and methods

Chemical denaturants, urea, GuHCl and GuSCN were raw materials of 68 deep eutectic solvents and were obtained from Sigma-Aldrich without 69 further purification. Stock solutions of GuHCl (5 M), GuSCN (5 M) and 70 urea (9 M) were prepared by weighing commercial compounds and 71 adding distilled water in the desired proportion to pH 7.0. A buffer solu-72 tion, consisting of 10 mM sodium phosphate at pH 7.0 and pH 2.0, was 73 used in some experiments and pH values of solutions were measured 74 by pH meter HI 9017 by Hanna Instruments Company by Sensorex 75 glass electrode. The eutectic mixtures were formed from the powders 76 by heating and stirring two components GuHCl-urea at 70 °C and 77 GuSCN-urea at 60 °C until homogenous liquid was formed. The molar ra-78 tios of eutectic compositions for CD measurements were found to be 1:2 79 (20 mM: 40 mM). The liquids were cooled at a rate of 1 °C/min and the 80 freezing point was taken as the temperature at which the first solid 81

<sup>\*</sup> Corresponding author at: Department of Biochemistry, Faculty of Science, P.J. Safarik University, 04011 Kosice, Slovak Republic.

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began to form. The cyt c (99%) from horse heart and BSA (98%) was
obtained from Sigma Aldrich. CD spectra were recorded with a Jasco
J-815 spectropolarimeter equipped with a Julabo Peltier temperature
control system (CDF-426S/15 model), using a cell of 0.2 mm path length
over the wavelength range from 195–250 nm, 350–500 nm. Protein
concentration was 5 mg/ml. CD signal-averaged over at least three
scans, and baseline was corrected by subtracting the reference spectrum.

#### 89 **3. Results**

We were able to prepare a new eutectic mixture with a lower eutectic 90 point by replacing one of the guanidine salts in GuHCl-GuSCN for urea. 91Isobaric phase diagram (atmospheric pressure 101.325 kPa) of these 92binary systems (Fig. 1) shows a two-dimensional space, where the 93 x-axis corresponds to the composition (in molar fractions) in accor-94 dance with the lever rule and the y-axis indicates the melting point tem-95 perature of DES (°C). Experimental determination of the phase diagram is 96 usually carried out by thermal analysis, in which the temperature of the 97studied sample during its cooling at a constant speed of 2-5 °C/min is 98 registered. When the system is cooled to the eutectic temperature it 99 will preclude the solid phase. Temperature measurements of phase 100 101 diagram were made on the basis of optical observations of crystals, using a Heidolph (Germany) heater with thermometer. 102

These values were confirmed under a light microscope (Optika 103 Microscope) with heating and cooling of samples. Guanidine salts 104mixed with urea in a 1:2 molar ratio produce colorless liquid that 105freezes at 58 °C for GuHCl-urea and 47 °C for GuSCN-urea. For GuHCl-106 urea eutecticum is formed at a composition of 67 mol% urea which is 107the same as the eutectic recently reported for the choline chloride-urea 108system [8]. The lower melting point (42 °C) relates to GuSCN-urea mix-109ture and occurs when the ratio of salt to hydrogen bond donor is 1:1.5 110 111 (60 mol% urea). The melting points of the mixtures are considerably lower than the melting points of either of the constituents (GuHCl 112 melts at 185 °C, GuSCN at 120 °C and urea at 134 °C). Concentrations of 113 the individual components at 1:2 molar ratio in the mixture GuHCl 114 (5.7 M)-urea (11.4 M) and GuSCN (5 M)-urea (10 M) are sufficient for 115 116 the complete denaturing of proteins (as in aqueous solutions).

In this report we demonstrate the ability of GuSCN-urea and GuHCl-117 urea (1:2) to dissolve higher amounts of two different proteins-bovine 118 serum albumin (BSA) and cytochrome c (cyt c). BSA is a globular water 119 120soluble protein with molecular weight ~66 kDa and isoelectric point 4.7. BSA dissolves at least 260 mg per 1 ml in both DES. Higher concentra-121 tions could not be prepared in the form of real solution because of gel 122123 forming. The second studied biomacromolecule i.e. horse hearth cyt c is a heme protein with molecular weight ~12 kDa and isoelectric point 12412510.5. Cyt c was shown to dissolve at minimum 18 mg per 1 ml in both



**Fig. 1.** Binary phase diagrams of guanidinium salts GuHCl (blue), GuSCN (red) and urea in different molar ratios. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.).

DES. Detection of higher concentrations of cytochrome c caused to observe solubility difficult because the solution was very colorful.

We performed CD analyses in order to monitor global secondary and 128 tertiary structural changes of cyt c in the presence of GuHCl-urea. 129 Although GuSCN has a stronger effect on protein denaturation as 130 GuHCl (GuSCN-urea has a lower eutectic formation temperature), the 131 disadvantage of its use in the far UV region is its stronger absorption 132 than GuHCl. Therefore we used the GuHCl-urea system in the next mea- 133 surements (the impact of DES on the CD spectrum of cytochrome c). 134 However, usage of cuvettes with path length of 0.2 mm (two separate 135 slides forming cuvette) was necessary for this DES. Cuvette constructed 136 from two parts provides an advantage when measuring viscous samples. 137 There is no problem loading the sample into the second enclosed cavity 138 portion by spreading the sample evenly throughout the entire cuvette 139 volume. Decreasing for CD signal of protein itself was offset by the high 140 concentration of protein that dissolved as well in water as in DES. Cyt c 141 allows deeper analysis of conformational changes by CD polarimetry as 142 compared to BSA because heme is the internal probe characterizing the 143 changes in the tertiary structure of the protein, Fig. 2 shows representa- 02 tive far- and near-UV CD spectra of cyt c in different conditions. 145

Fig. 2 (left) shows changes in the secondary structure of cytochrome 146 c. Native cyt c in 10 mM phosphate buffer (pH 7.0 at 25 °C) shows 147 classical dichroic minima at 208 nm and 222 nm, specific for the proteins 148 with high amount of  $\alpha$ -helicity. Water solutions of GuHCl (5 M) and urea 149 (9 M) markedly replaced these minima by a large negative minimum 150 below 210 nm suggesting an occurrence of a typical helix to random 151 coil transition. The CD spectra of the cyt c in 10 mM phosphate buffer 152 pH 2.0 at 25 °C indicate that the secondary structure is qualitatively sim- 153 ilar with spectra of the cyt c in GuHCl-urea mixture at a temperature of 154 70 °C. The spectra suggest randomly coiled conformations in both cases, 155 as well as in the individual components of a mixture. Difference in the 156 spectra may be due to expanding the unfolding of cytochrome c. Fig. 2 157 (right) shows changes around the heme of cytochrome c. Negative 158 peak at 416 nm and positive peak at 402 nm has been assigned for the 159 tertiary structure corresponding to heme-polypeptide interaction close 160 to the heme crevice and indicates native cyt c. In the presence of 161 GuHCl-urea we observed the absence of negative Cotton effect at 162 416 nm and strong positive band with a maximum at 410 nm indicating 163 a decoupling Met80 from binding to heme iron, like this observe for in 164 all individual components of DES in water as well as for acidic pH 2.0 165 buffer. 166

The guestions whether these denatured proteins in the melt form of 167 DES can renaturate again is another problem arising here. Refolding of 168 the solubilized proteins is initiated by the removal of the denaturant 169 or reducing its concentration and allowing the protein to refold into 170 the native conformation. The most often used approach here is the re- 171 moval of the solubilizing agent (DES) by dialysis (as mentioned by 172 other authors). Long-term storage of cytochrome c in GuHCl-urea DES 173 (6 months) at room temperature, and after removal of DES in 10 mM 174 phosphate buffer solution at pH 7 by means of dialysis, CD spectra 175 (Fig. 2 red lines) demonstrated folding of cytochrome c into the native 176 structure. During dialysis the concentration of the solubilizing agent 177 decreases slowly what allows to refold the protein optimally. Dialyzed 178 cyt c from GuHCl-urea solution was carried out in 10 mM phosphate 179 buffer pH 7.0 at 25 °C and subsequently measured secondary (tertia- 180 ry) structure and conformational stability. CD spectra of the refolded 181 cyt c closely resemble those of the native cyt c, showing classical di- 182 chroic minima at 208 nm and 222 nm for secondary structure and 183 negative peak at 416 nm and positive peak at 402 nm for tertiary 184 structure (Fig. 2 - red line). The protein was in DES in melt form at 185 RT for 6 month. Therefore, assuming the formation of solid melt in 186 the vicinity of the protein stabilizes the denatured state and after re- 187 moval of the denaturant lets his return to native form. These observa- 188 tions indicate that the protein could be stored at RT in the denatured 189 state and consequently renaturate anytime becoming ready for further 190 measurements. 191

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