



# КЛАСТЕР КОНФЕРЕНЦИЙ 2021

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ДОКЛАДОВ

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XIV МЕЖДУНАРОДНАЯ НАУЧНАЯ КОНФЕРЕНЦИЯ  
«ПРОБЛЕМЫ СОЛЬВАТАЦИИ И  
КОМПЛЕКСООБРАЗОВАНИЯ В РАСТВОРАХ»

XI МЕЖДУНАРОДНАЯ НАУЧНАЯ КОНФЕРЕНЦИЯ  
«КИНЕТИКА И МЕХАНИЗМ КРИСТАЛЛИЗАЦИИ.  
КРИСТАЛЛИЗАЦИЯ И МАТЕРИАЛЫ НОВОГО  
ПОКОЛЕНИЯ»

VI МЕЖДУНАРОДНАЯ НАУЧНАЯ КОНФЕРЕНЦИЯ  
ПО ХИМИИ И ХИМИЧЕСКОЙ ТЕХНОЛОГИИ

XIII ВСЕРОССИЙСКАЯ ШКОЛА-КОНФЕРЕНЦИЯ  
МОЛОДЫХ УЧЕНЫХ «ТЕОРЕТИЧЕСКАЯ И  
ЭКСПЕРИМЕНТАЛЬНАЯ ХИМИЯ ЖИДКОФАЗНЫХ  
СИСТЕМ» (КРЕСТОВСКИЕ ЧТЕНИЯ)



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## PARTITIONING OF L-TRYPTOPHAN IN AQUEOUS BIPHASIC SYSTEMS CONTAINING DIALKYLMIDAZOLIUM IONIC LIQUID WITH DIFFERENT ANIONS AND KOSMOTROPIC SALT

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Separation and purification of biomolecules (e.g. amino acids) are the crucial tasks in biotechnology [1]. Some ionic liquids (ILs) can serve as a safe alternative to organic solvents, many of which are toxic or flammable. Aqueous biphasic systems (ABSs) formed by IL, water and inorganic salt are of particular interest for bioextraction. An ability to form two liquid phases with the distribution of solute between these phases makes these systems applicable for the required partitioning of biomolecules. One of the important advantages of ILs is a possibility to modify their chemical structure in order to obtain the desired properties such as an extraction capacity. Among the modified ILs, the amino acid ILs (AAILs) attract much attention due to their low toxicity in comparison with conventional ILs [2]. However, extraction ability of ABSs with ILs or especially with AAIALs is poorly studied.

The main aim of this work was to study the distribution of a solute between the liquid phases effect in ABS containing 1-alkyl-3-methylimidazolium ILs with halide and amino acid anions [ $C_n\text{mim}$ ]X (where n - number of

carbon atoms in the alkyl chain of AAiL = 4, 6, 8; X = Cl – chloride, Br – bromide, [Leu] – L-Leucinate, [Val] – L-Valinate, [Lys] – L-Lysinate) and K<sub>2</sub>HPO<sub>4</sub> and K<sub>3</sub>PO<sub>4</sub> as inorganic salts. L-tryptophan was chosen as a model solute due to its high hydrophobicity.

As a result, new data on partition coefficients of L-tryptophan in the selected ABSs were obtained. The results showed that in ABSs with some AAiLs, higher distribution coefficients are observed than in the systems with halide ILs. Moreover, a correlation was established between the partition and phase behavior (the results were obtained by us earlier [3]) of ABSs. It was found that, in general, partition coefficients of L-tryptophan are larger for ABSs with a wider heterogeneous region.

It should be concluded that ABSs with AAiLs are promising for the extraction of small biomolecules. The results can be useful for understanding the driving force of phase separation and distribution of the solutes.

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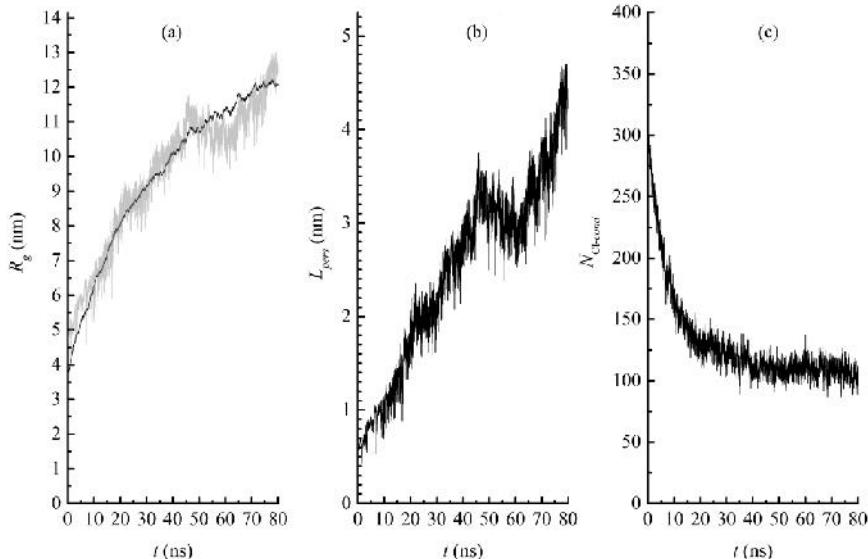
## POLYMERIC IONIC LIQUIDS IN DILUTE AQUEOUS SOLUTION: MD SIMULATION OF POLY(DIMETHYL DIALLYL AMMONIUM) SALTS

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Polymeric ionic liquids (PILs) containing amino acid anions is an important new class of PILs that attract an increasing attention. The effect of the anion chemistry on the conformational behavior of the PIL chain is one of the key issues.

In this work we report the results of molecular dynamics simulations for dilute aqueous solutions of poly(dimethyl diallyl ammonium) chloride (PDADMAC) and poly(dimethyl diallyl ammonium) lysinate (PDADMALys). Evolution of the polycation chain from a compact globule into an expanded coil is examined and discussed in terms of theoretical models. The effect of presence of ionic pairs on the free energy of the chain is estimated from the theory of dipolar chains [1, 2]. Conformational and diffusive characteristics of PDADMAC and PDADMALys are compared, revealing the effects of size and shape of the counterion.



**Figure 1.** The gyration radius (a), the persistent length (b), and the number of Cl<sup>-</sup> anions condensed on the chain (c) for a single poly(dimethyl diallyl ammonium) chain (400 monomers) in PDADMAC/water during its evolution from a globule to an expanded coil. Black, MD results; gray, the Benoit – Doty approximation for  $R_g$  using the persistent length from (b).

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