Հայաստանի Գիտությունների Ազգային Ակադեմիա Национальная Академия Наук Армении National Academy of Sciences of Armenia



<шјшилши́р Чѣйишрши́шЦши́ <ши́рtи</li>Биологическии́ Журнал АрменииBiological Journal of Armenia

• Համարոտ հաղորդումներ • Краткие сообщения • Short communications •

Biolog. Journal of Armenia, 1 (64), 2012

# THE INTERACTION OF ZnTOEPyP4 PORPHYRIN WITH TRNA. THE INFLUENCE OF IONIC STRENGTH

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The interaction of ZnTOEPyP4 porphyrin with tRNA was studied using UV/Vis Spectrophotometry and Circular Dichroism (CD) methods. The influence of ionic strength on these interactions was investigated as well as. The preferable binding mechanism was identified.

Porphyrin s- tRNA - ionic strength

Լուսասպեկտրաչափական և շրջանային դիքրոիզմի մեթոդներով ուսումնասիրվել է ZnTOEPyP4 պորֆիրինի փոխազդեցությունը փՌՆԹ-ի հետ միջավայրի տարբեր իոնական ուժերի դեպքում։ Պարզաբանվել է փՌՆԹ-ի հետ ZnTOEPyP4 կապման մեխանիզմը։

Պորֆիրիններ - փՌՆԹ - իոնական ուժ

Методами спектрофотометрии и кругового дихроизма исследовано взаимодействие порфирина ZnTOEPyP4 с тРНК при высоких и низких ионных силах. Установлен механизм связывания ZnTOEPyP4 с тРНК.

### Порфирины - тРНК - ионная сила

In this work the influence of ionic strength on the interaction of tRNA from E.Coli with Zn(II) meso-tetra-(4N-oxyethylpyridyl) porphyrin (ZnTOEPyP4) has been studied by UV/Vis Spectrophotometry and Circular Dichroism methods. There are few works devoted to the interaction of porphyrins with tRNA [1, 2], while the interaction of ZnTOEPyP4 with tRNA is not investigated at all. It is known that conformation of tRNA strongly depends on ionic strength of medium. In case of high ionic strength ( $\mu$ =0.2 M) the molecule of tRNA has tertiary structure alike reversed letter "L" (Fig.1). At low ionic strength ( $\mu$ =0.02 M) tRNA takes a hairpin form with several short double helical regions and loops [3].



Fig.1. The structure of tRNA at high (I) and low (II) ionic strength

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*Materials and methods.* All measurements were performed in 0.1 BPSE ( $\mu$ =0.02 M) and 1 BPSE ( $\mu$ =0.2 M) buffers (1BPSE = 6 mM Na2HPO4+ 2 mM NaH2PO4 +185 mM NaCl + 1 mM Na2EDTA) pH 6.57. ZnTOEPyP4 was synthesized by Dr. R.Ghazaryan Faculty of General and Nonorganic Chemistry, Yerevan State Medical University.

**Results and Discussion.** The interaction of porphyrins has been monitored with visible absorption spectroscopy (Soret band). The solutions of constant concentration of porphyrins (10-6 M) titrated with a stock solution of tRNA. The results obtained from titration experiments were used for the calculation of the binding parameters. As it is suggested by Cantor and Schimmel the model for binding of ligands with tRNA is described via Scatchard model.

$$\frac{\mathbf{r}_{b}}{\mathbf{C}_{f}} = \frac{\mathbf{N}_{1}\mathbf{K}_{1}}{1 + \mathbf{C}_{f}\mathbf{K}_{1}} + \frac{\mathbf{N}_{2}\mathbf{K}_{2}}{1 + \mathbf{C}_{f}\mathbf{K}_{2}}$$

where  $C_f$  is the free porphyrins concentration in solution,  $r = C_b/C_{tRNA}$ , Cb is the concentration of bound porphyrins, CtRNA is the concentration of tRNA molecules. It is supposed, that there are two types of binding sites on tRNA, N<sub>1</sub> sites with binding constant K<sub>1</sub> and N<sub>2</sub> sites with binding constant K<sub>2</sub>.

The binding parameters for mentioned cases are the following:

 $\mathrm{K_{1}=8.07x109~M^{-1},~N_{1}=24.7;~K_{2}=2.09x10^{6}~M^{-1},~N_{2}=6.9~(\mu=0.02~M)}$ 

 $K_1 \!=\! 5.15 x 107 \; M^{\text{--}1}, \, N_1 \!=\! 2.2; \, K_2 \!=\! 4,\! 03 x 10^4 \; M^{\text{--}1}, \, N_2 \!=\! 38 \; (\mu \!=\! 0.2 \; M)$ 

From binding parameters it can be seen, that in both cases of ionic strength there is a preferable binding mechanism. Having one axial ligand, ZnTOEPyP4 cannot intercalate into helical sites, it binds with tRNA externally ordered. Also Zn metalloprophyirin binds with tRNA well in case of low ionic strength (corresponding binding constant is bigger).

As it is known the ICD spectra of complexes of porphyrins with nucleic acids are very informative. A rather unusual effect of ICD band modification is observed when ZnTOEPyP4 binds with tRNA in both cases of ionic strength (fig.2). From obtained data we can conclude following.



**Fig.2.** ICD-spectra of complexes ZnTOEPyP4 with tRNA in case of different ionic strength. Numbers near the curves show the ratio of porphyrins to tRNA

In case of low ionic strength it is proposed that Zn porphyrins are ordered in a stack, not only on helical sites, but also on loops of tRNA [4]. As a result of such ordered stacking, the complex resembles a short stick. These sticks can interact with each other via bound porphyrins. As a result they can form the structures that look like liquid crystals which can show the similar behavior of ICD spectra.

At high ionic strength tRNA has tertiary structure, which looks like reversed letter "L". In this condition the intensity of ICD spectra is about 2 times bigger. We assume that this kind of effect cannot be explained in terms of single molecule and it is proposed that one type of binding is with sites at the top of tertiary structure of tRNA and the other one is the intermolecular interactions via bound porphyrins on tRNA. And the growing of intensity of ICD is the result of initially stabile tertiary structure of tRNA. As a result, we suppose, that the regular structures like liquid crystals are formed.

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Received 18.01.2012