Interaction of Cephalexin Monohydrate and Diclofenac Sodium with Human Erythrocyte Membrane: Adsorption Study and Equilibrium Isotherm Analysis

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To elucidate the mechanism of interaction of biological surfaces with drugs, *in vitro* study of interaction of erythrocyte membrane from healthy individuals with two type of drugs was performed at 37.5 °C and at physiological pH 7.4. Cephalexin monohydrate and diclofenac sodium were chosen for this purpose. The blood samples were collected from healthy volunteers (21 males and 19 females), aged 20 to 40 years (mean age 28 ± 6 years). Hypotonic solution was used to isolate the erythrocyte membrane from red blood cells. The amount of cephalexin-H₂O or diclofenac sodium binding to erythrocyte membrane was significantly increased ($p < 0.01$) with increasing drug concentration and there was no significant difference ($p > 0.05$) in amount of drugs uptake by erythrocyte membrane of male or female. There was no significant difference ($p > 0.05$) between the amount of cephalexin-H₂O binding to erythrocyte membrane and that for diclofenac sodium. The results indicated the applicability of Langmuir and Freundlich isotherms of drugs binding to erythrocyte membrane. The models that fit the experimental data assume the presence of multiple binding sites on the surface of erythrocyte membrane. The binding data and Scatchard plot suggested the existence of non-identical independent sites, in human erythrocyte membrane, interacting with cephalexin-H₂O and having different affinities.

Keywords: Erythrocyte membrane, Interaction, Cephalexin monohydrate, Diclofenac sodium, Adsorption Isotherms.

INTRODUCTION

Erythrocyte membrane, like all other eukaryotic plasma membranes, is composed of a bilayer of several different classes of lipid along with a variety of protein molecules [1]. It contains 52 % proteins, 40 % lipids and 8 % carbohydrates that is bound to lipids (glycolipids) or proteins (glycoproteins) [2]. The lipids form a rather continuous bilayer in which the outer layer contains mainly phosphatidylcholine and sphingomyelin, and the inner one, phosphatidylethanolamine and phosphatidylserine. Lipids provide mechanical stability and a strong tendency to form closed structures [3]. Membrane proteins are classified as peripheral or integral. Peripheral proteins are molecules that are bound to membrane primarily through interactions with integral membrane proteins. The integral proteins represent more than 70 % of membrane proteins and are very firmly associated with the membrane, the two major integral proteins are band and

glycophorin [4]. The outer surface of erythrocyte membrane is negatively charged, due to presence of salicylic acid residues as well as carboxyl and phosphate groups [5]. The presence of charges on the external surface of biological membranes influences to a remarkable degree their electrical behaviour and physico-chemical properties [6]. Various materials have been shown to adsorb or interact with the erythrocyte membrane, such as adsorption of different amino acids (L-alanine, glycine, L-glutamine and L-leucine) [7], anti A or anti B-antibodies [8] and drugs [9,10]. The interaction of drugs with erythrocyte membrane may occurs through the electrostatic attraction with negatively charged lipids or with negatively charged proteins of membrane skeleton [1]. These interactions are supposed to be involved in the modulation of membrane fluidity as well as in the regulation of membrane-related processes [11].

Elferink [12] studied the binding of chlorpromazine-HCl and its quaternary analogue chlorpromazine methiodide to

open and resealed human erythrocyte ghost membranes. The results indicated that the quaternary compound is confined to the outside face of membrane. For both compounds two classes of binding sites are available. The strongest binding sites are mainly located on the inner surface of membrane. Numerous reports have demonstrated the nature of binding between certain agents and the receptor sites in red blood cell membrane. It has been suggested that binding of stilbendisulfonic acids to the protein in band 3 of red blood cell membrane occurs through the formation of covalent binding (irreversible binding) after initial rapid non-covalent binding [13]. On the other hand, it was found that the covalent bond formation between 1-fluoro-2,4-dinitrobenzene and anion permeability-controlling sites on the protein in band 3 is direct and not preceded by a step that involves non-covalent binding [14]. It has long been known that certain drugs (like salicylic acid and its derivatives) interact with the protein in red cell membrane and act as a reversible and non-competitive inhibitors of anion permeability. The inhibitory action of these drugs influence by many factors related to the structure of drug molecule, such as hydrophobicity of molecule, electron distribution within the molecule and steric effects [15].

The aim of the present work is to evaluate the interaction of cephalixin-H₂O and diclofenac sodium with erythrocyte membrane. It is planned to carry out this part of research *in vitro* aiming to get some informations regarding the mechanism of interaction of this type of biological surfaces with the drugs (cephalexin monohydrate and diclofenac sodium) under study.

EXPERIMENTAL

Subjects: Fourty blood samples have been collected from healthy Iraqi volunteers (21 males and 19 females), aged 20 to 40 years (mean age 28 ± 6 years).

Preparation of erythrocyte membrane: Erythrocyte membranes were prepared according to Reinila *et al.* [16]. Fresh blood samples (10 mL) containing heparin as an anticoagulant were centrifuged at 3000 rpm. The erythrocytes were resuspended in ice-cold isotonic saline (NaCl 0.9 %) and washed by three successive centrifugations (3000 rpm) for 5 min. Packed RBCs (3 g) were transferred into 30 mL of hypotonic medium (10 mM tris-HCl, pH 7.4, containing 0.1 mM EGTA). The tubes were centrifuged at 4 °C for 20 min at 12000 rpm. After aspiration of supernatant, the cells were resuspended in hypotonic medium and treated as above. The final pellet of RBC membranes was resuspended in tris-EGTA buffer and stored at 20 °C in deep frozen until assayed.

Adsorption on erythrocyte membrane: Solutions of each drug (cephalexin-H₂O or diclofenac sodium) of known concentrations (0.2-5 mM) in buffer solution (pH 7.4) were added to flasks containing 0.2 mL (~ 0.2 g) of erythrocyte membrane suspension. The flasks were shaken in waterbath at 37.5 °C for 45 min. After the period of incubation, the suspensions were centrifuged at 4000 rpm for 20 min. The clear supernatants were assayed for drug, after appropriate dilution, using UV-visible spectrophotometer. The amount of drug adsorbed was calculated from the following equation:

$$Q_e = \frac{V(C_o - C_e)}{m} \quad (1)$$

where Q_e is the quantity adsorbed, m is the weight of erythrocyte membrane (g), C_o is the initial concentration of drug (mg/L), C_e is the equilibrium concentration of drug (mg/L) and V is the volume of solution (L).

Protein content estimation: The protein content of erythrocyte membrane was estimated according to Lowery method [17] using bovine serum albumin as standard.

RESULTS AND DISCUSSION

The binding of cephalixin-H₂O and diclofenac sodium to erythrocyte membranes from healthy individuals at 37.5 °C and pH 7.4 was investigated. The results show a consistent and significant increase ($p < 0.01$) in the amounts of drug uptake by erythrocyte membrane with increasing drug concentration. The binding of drugs to erythrocyte membrane for both sex was studied. No significant difference ($p > 0.05$) in amount of drug uptake by erythrocyte membrane of male or female was noticed (Table-1).

TABLE-1
AMOUNT OF DRUGS UPTAKE BY THE SURFACE
OF ERYTHROCYTE MEMBRANE FOR MALE
AND FEMALE EXPRESSED (mg/g)

Cephalexin-H ₂ O		Diclofenac sodium	
Male*	Female*	Male*	Female*
0.76 ± 0.28	0.82 ± 0.19	0.54 ± 0.26	0.52 ± 0.26
1.66 ± 0.48	1.72 ± 0.30	1.04 ± 0.26	1.16 ± 0.26
3.85 ± 0.49	3.95 ± 0.26	3.21 ± 0.39	3.12 ± 0.32
5.30 ± 0.58	5.82 ± 0.98	6.76 ± 0.61	6.84 ± 0.58
9.78 ± 0.74	10.02 ± 0.77	7.88 ± 0.80	7.63 ± 0.72
10.83 ± 0.95	11.08 ± 0.96	8.24 ± 1.22	8.67 ± 1.15
11.77 ± 0.85	11.38 ± 0.98	9.31 ± 0.94	9.32 ± 1.15

*Data are presented as the mean ± S.D., N = 21 for each drug.

Fig. 1 shows the adsorption isotherms of cephalixin-H₂O and diclofenac sodium. The general shape of adsorption isotherm of cephalixin-H₂O have L₄ type isotherm, while the adsorption isotherm of diclofenac sodium is L₂ type feature of Giles classification [18].

The adsorption isotherm of cephalixin-H₂O by erythrocyte membrane (Fig. 1a) is not continuous, there is a major break (step) in which the amount of drug adsorbed appears to rise rapidly at a particular drug concentration. This result agrees with the report of Bikhazi *et al.* [19], who reported the adsorption isotherms of ouabain on hepatocytes from normal and diabetic rats are not continuous. They assumed the presence of multiple receptors on the cell surface and only when a specific fraction of total number of one receptor have effectively reacted will the other receptors initiate reaction with the ligand. They also suggested the existence of two receptors in normal and diabetic hepatocytes, interacting with ouabain and having different affinities for drug.

The experimental data were explained using the theoretical Langmuir equation and empirical Freundlich equation (eqns. 2 and 3).

$$\frac{C_e}{Q_e} = \frac{1}{Q_m K_L} + \left(\frac{1}{Q_m} \right) C_e \quad (2)$$

Fig. 1. Adsorption isotherms of (a) cephalexin-H₂O, (b) diclofenac sodium, on erythrocyte membrane at 37.5 °C

where Q_e (mg/g) is the quantity adsorbed per unit weight, C_e (mg/L) is the equilibrium concentration, Q_m (mg/g) and K_L (L/mg) are Langmuir isotherm constants.

$$\log Q_e = \log K_f + \frac{1}{n} \log C_e \quad (3)$$

where K_f and n are Freundlich constants; $1/n$ is a function of the strength of adsorption in adsorption process.

The results illustrated the applicability of Langmuir and Freundlich isotherms for both drugs as shown by the linear relationships of C_e/Q_e versus C_e (Fig. 2) and $\log Q_e$ versus $\log C_e$ (Fig. 3).

A number of binding sites are involved in the binding of adsorbate molecules (called ligand) to macromolecules, and

Fig. 2. Linear form of Langmuir isotherm of cephalexin-H₂O and diclofenac sodium on erythrocyte membrane

Fig. 3. Freundlich adsorption isotherms of cephalexin-H₂O and diclofenac sodium on erythrocyte membrane

sometimes they behave identically and act quite independently of each other. This is the simplest situation and such systems obey an equation of Langmuir form. However, in some important cases the occupation of one site affects the strength of binding at other sites; in other words, behaviour is favourable [20].

Most equations that describe the binding of adsorbate molecules to macromolecules are expressed in terms of dissociation constant (k_d) and the number of moles of adsorbate molecule bound to one mole of macromolecules (r), therefore the Langmuir equation can be written in the following form [21]:

$$r = \frac{[A]}{k_d + [A]} \quad (4)$$

where $[A]$ represent the concentration of unbound ligand at equilibrium. This equation is usually rewritten in one of two forms that allow the data to be plotted on a straight line. One form is:

$$\frac{r}{[A]} = \frac{1}{k_d} - \frac{r}{k_d} \quad (5)$$

in which a plot of $r/[A]$ versus r gives a straight line having a slope of $-1/k_d$. If all binding sites are identical and independent, the relevant equations analogous to eqn. 5 as:

$$\frac{r}{[A]} = \frac{n}{k_d} - \frac{r}{k_d} \quad (6)$$

where n is the number of binding sites and k_d is an average dissociation constant. With eqn 6 (often called Scatchard equation), a plot of $r/[A]$ versus r gives a line having a slope of $-1/k_d$ and an x-intercept of n/k_d .

If the binding sites are not both identical and independent, neither of the plots, $1/r$ versus $1/[A]$ nor $r/[A]$ versus r , will give straight lines. Eqn. 6 can be re-written in the form:

$$\frac{Q_e}{C_e} = \frac{n}{k_d} - \frac{Q_e}{k_d} \quad (7)$$

The adsorption data and eqn. 7 were used to construct a Scatchard plots for binding of cephalixin- H_2O and diclofenac sodium to erythrocyte membrane, where Q_e/C_e is plotted as a function of amount of drug uptake by membrane (Q_e) (Fig.4).

As shown in Fig. 4, the Scatchard plots of adsorption data of both drugs are different. The Scatchard plot of cephalixin- H_2O revealed the existence of two types of binding sites which are non-identical and independent. This result is in agreement with that reported by Ishida *et al.* [22] and Elferink [14].

Diclofenac sodium to erythrocyte membrane: There is an increasing evidence that erythrocyte membrane is asymmetric with regard to proteins as well as to phospholipids. In erythrocyte membrane the majority of aminophospholipids, phosphatidylserine and phosphatidylethanolamine, are located in the

inner leaflet, whereas phosphatidylcholine and sphingomyelin, are localized predominantly in the outer leaflet of lipid bilayer [23]. Most proteins of erythrocyte membrane being located in the inner half of the membrane [24]. Sheetz and Singer [25] have proposed that two halves of an asymmetric membrane can respond differently to a perturbation and thus act as bilayer couples. They attributed the different effects of amphipatic drugs on erythrocyte morphology to a differential distribution of drugs in the erythrocyte membrane as a consequence of the presence of phosphatidylserine in the inner half of membrane.

The nature of interaction of cephalixin- H_2O with erythrocyte membrane may be related to the possibility of drug adsorption on both faces of membrane. Elferink [14] showed that chlorpromazine-HCl was bound to both phases of membrane and the drug was preferentially bound to the inside face of membrane. He also found that the greater number of binding sites on the inside face is associated with the presence of phosphatidylserine in that side of the membrane and possibly with the presence of more protein binding sites on the inner half as compared to the outer half of membrane.

In erythrocytes, the inner layer of membrane is rich in phosphatidylserine – a lipid bearing net negative charge at neutral pH. Negatively charged lipids are absent in the outer layer. On the other hand, cephalixin- H_2O at neutral pH may occurs in protonated form [$pK_a = 2.7$ (4-carboxylic acid) and 6.96 (7/ α -ammonium chain)]. Hendrich *et al.* [26] found that preferential binding of thioridazine (bears positive charge at neutral pH) with the inner leaflet of membrane was due to the electrostatic attraction with negatively charge lipids as well as to its interaction with negatively charged proteins of membrane skeleton. It was suggested that phosphatidylserine in cytoplasmic half of the membrane could provide a negative field, attracting cationic drugs into cytoplasmic half [7]. The presence of charged groups at membrane surface will influence the distribution of mobile counter-ions. Surface of erythrocyte membrane normally carry net negative charge. This surface fixed charge leads to the formation of a diffuse electrical double layer in which the

Fig. 4. Scatchard plots for the binding of (a) cephalixin- H_2O , (b) diclofenac sodium to erythrocyte membrane

charge at the surface is balanced by charges of opposite sign in the medium immediately adjacent to the surface [27]. Moreover, hydrophobic properties of drug molecule may play an important role in the nature of binding to erythrocyte membrane. The amount of diclofenac sodium adsorbed was non-significantly less ($p > 0.05$) than the amount of cephalexin- H_2O uptake by erythrocyte membrane. There is no specific factor in the interpretation of this behaviour, but this may be attributed to the similarity in molecular weight of both drugs. Multiple factors can play a role in diclofenac sodium interaction with the membrane surface such as lipophilicity, stereochemistry and geometry of drug molecule, the distribution of charge on molecule and number of active functional groups in molecule bears. It is more likely the geometrical and electrostatic factors can affect diclofenac sodium bound to both sides of membrane surface.

Conclusion

The surface of human erythrocyte membrane consists of multiple binding sites with different affinities to interact with the drugs. The adsorption isotherm of cephalexin- H_2O by erythrocyte membrane is not continuous, there is a major break (step) in which the amount of drug adsorbed appears to rise rapidly at a particular drug concentration. There is no sex differences in the amounts of drugs binding to erythrocyte membrane. The amount of diclofenac sodium adsorbed was non-significantly less ($p > 0.05$) than the amount of cephalexin- H_2O uptake by erythrocyte membrane. It is more likely that geometrical and electrostatic factors can affect diclofenac sodium to both sides of membrane.

CONFLICT OF INTEREST

The authors declare that there is no conflict of interests regarding the publication of this article.

REFERENCES

- H. Li and G. Lykotrafitis, *Biophys. J.*, **107**, 642 (2014); <https://doi.org/10.1016/j.bpj.2014.06.031>.
- E.D.F. De Robertis and E.M.F. De Robertis, *Cell and Molecular Biology*, Lea and Febiger: USA, edn 8 (1987).
- M. Ikeda, A. Kihara and Y. Igarashi, *Biol. Pharm. Bull.*, **29**, 1542 (2006); <https://doi.org/10.1248/bpb.29.1542>.
- P.V. Escibá, J.M. González-Rosb, F.M. Goñi, P.K.J. Kinnunen, L. Vigh, L. Sánchez-Magraner, A.M. Fernández, X. Busquets and I. Horváth, *J. Cell. Mol. Med.*, **12**, 829 (2008); <https://doi.org/10.1111/j.1582-4934.2008.00281.x>.
- H.P. Fernandes, C.L. Cesar and M.L. Barjas-Castro, *Rev. Bras. Hematol. Hemoter.*, **33**, 297 (2011); <https://doi.org/10.5581/1516-8484.20110080>.
- P. Atkins and J. Paula, *Interações moleculares*. In: *Atkins Físico-química, Livros Técnicos e Científicos: Rio de Janeiro*, edn 7, p. 408 (2004).
- M.M. Cruz Silva, V.M.C. Madeira, L.M. Almeida and J.B.A. Custódio, *Toxicol. In Vitro*, **15**, 615 (2001); [https://doi.org/10.1016/S0887-2333\(01\)00079-0](https://doi.org/10.1016/S0887-2333(01)00079-0).
- R. Huang, M. Kiss, M. Batonick, M. Weiner and B. Kay, *Antibodies*, **5**, 11 (2016); <https://doi.org/10.3390/antib5020011>.
- M. Lucio, J.L.F.C. Lima and S. Reis, *Curr. Med. Chem.*, **17**, 1795 (2010); <https://doi.org/10.2174/092986710791111233>.
- H. Tsuchiya, *Molecules*, **20**, 18923 (2015); <https://doi.org/10.3390/molecules201018923>.
- C. Sousa, C. Nunes, M. Lúcio, H. Ferreira, J.L.F.C. Lima, J. Tavares, A. Cordeiro-da-Silva and S. Reis, *J. Pharm. Sci.*, **97**, 3195 (2008); <https://doi.org/10.1002/jps.21218>.
- J.G.R. Elferink, *Biochem. Pharmacol.*, **26**, 2411 (1977); [https://doi.org/10.1016/0006-2952\(77\)90450-6](https://doi.org/10.1016/0006-2952(77)90450-6).
- S. Ship, Y. Shami, W. Breuer and A. Rothstein, *J. Membr. Biol.*, **33**, 311 (1977); <https://doi.org/10.1007/BF01869522>.
- H. Passow, *Cell Membrane Receptors for Drugs and Hormones: A Multidisciplinary Approach*, Ravan Press: New York (1978).
- M.M. Kamel and Y.M. Syam, *Egypt. Pharm. J.*, **12**, 95 (2013); <https://doi.org/10.4103/1687-4315.124000>.
- M. Reinila, E. MacDonald, N. Salem Jr., M. Linnoila and E.G. Trams, *Anal. Biochem.*, **124**, 19 (1982); [https://doi.org/10.1016/0003-2697\(82\)90214-7](https://doi.org/10.1016/0003-2697(82)90214-7).
- O.H. Lowery, N.J. Rosebrough, A.L. Farr and R.J. Randall, *J. Biol. Chem.*, **193**, 265 (1951).
- C.H. Giles, D. Smith and A. Huitson, *J. Colloid Interface Sci.*, **47**, 755 (1974); [https://doi.org/10.1016/0021-9797\(74\)90252-5](https://doi.org/10.1016/0021-9797(74)90252-5).
- A.B. Bikhazi, K.M. Bitar, M.M. El-Kasti and E.K. Shaaban, *J. Pharm. Sci.*, **83**, 1758 (1994); <https://doi.org/10.1002/jps.2600831222>.
- K.J. Laidler, *Physical Chemistry with Biological Applications*, Benjamin Cummings Publishing Company, Inc.: California, USA (1978).
- D. Freifelder, *Physical Biochemistry: Applications to Biochemistry and Molecular Biology*, W.H. Freeman and Company: New York, edn 2 (1982).
- S. Ishida, Y. Sakiya, T. Ichikawa and Z. Taira, *Biol. Pharm. Bull.*, **16**, 293 (1993); <https://doi.org/10.1248/bpb.16.293>.
- T. Fujimoto and I. Parmyrd, *Front. Cell Dev. Biol.*, **4**, Article 155 (2017); <https://doi.org/10.3389/fcell.2016.00155>.
- A.P. Liu and D.A. Fletcher, *Biophys. J.*, **91**, 4064 (2006); <https://doi.org/10.1529/biophysj.106.090852>.
- M.P. Sheetz and S.J. Singer, *Proc Natl. Acad. Sci. USA*, **71**, 4457 (1974); <https://doi.org/10.1073/pnas.71.11.4457>.
- A.B. Hendrich, K. Lichacz, A. Burek and K. Michalak, *Cell. Mol. Biol. Lett.*, **7**, 1081 (2002).
- P.A. Janmey and P.K.J. Kinnunen, *Trends Cell Biol.*, **16**, 538 (2006); <https://doi.org/10.1016/j.tcb.2006.08.009>.