# Interaction of Cibacron Blue 3GA and Remazol Brillant Blue R with the Nucleotide Binding Site of Lactate Dehydrogenase and $(Na^+ + K^+)$ -ATPase

V. ĎURIŠOVÁ<sup>1</sup>, A. VRBANOVÁ<sup>1</sup>, A. ZIEGELHÖFFER<sup>2</sup> and A. BREIER<sup>1</sup>

 Institute of Molecular Physiology and Genetics, Slovak Academy of Sciences, Vlárska 5, 833 34 Bratislava, Czechoslovakia
Institute of Heart Research, Slovak Academy of Sciences, Dúbravská cesta 9, 842 33 Bratislava, Czechoslovakia

Abstract. The interaction of Cibacron Blue 3GA (C.I.2) and Remazol Brillant Blue R (C.I.19) with purified preparations of Lactate dehydrogenase (LDH) and  $(Na^+ + K^+)$ -ATPase was studied by means of the enzyme kinetics method. LDH was found to be inhibited by both C.I.2 and C.I.19, with the former being a stronger inhibitor. This may be explained by the fact that in contrast to C.I.19, C.I.2 resembles the whole molecule of the specific cofactor.  $(Na^+ + K^+)$ -ATPase activity was inhibited by both dyes to approximately similar degree. C.I.2 and C.I.19 resemble the ATP molecule to approximately similar extent, particularly as concerns the molecule shape and size. The results obtained confirmed the applicability of C.I.2 and C.I.19 as nucleotide-specific ligands.

Key words:  $(Na^+ + K^+)$ -ATPase — Lactate dehydrogenase —  $K^+$ -pNPPase — Cibacron Blue 3GA — Remazol Brillant Blue R

#### Introduction

Antraquinone dyes (Cibacron Blue 3GA — C.I.2 and Remazol Brillant Blue R — C.I.19) represent group-specific ligands with high affinities to nucleotidebinding sites of purine nucleotides. This specific property of C.I.2 and C.I.19 has been utilized in their applications as specific ligands in affinity chromatography of dehydrogenases such as lactate dehydrogenase (Gemeiner et. al. 1981), malate dehydrogenase (Scawen et al. 1982) or 6-phosphogluconate dehydrogenase (Kroviarski et al. 1988), and kinases like phosphofructo kinase (Kopperschläger et al. 1982), or phosphoglycerate kinase (Kulbe and Schuer 1979).

Besides the latter mentioned nucleotide-specific interactions antraquinone dyes are capable to provide also ionic- (Subramanian 1984) and hydrophobic



Fig. 1. Structures of reactive dyes C.I.2 and C.I.19 in comparison to ATP and NAD+.

interactions (Breier et al. 1987a) with proteins. The diversity of the possible interactions of C.I.2 and C.I.19 offers them almost general reactivity which may manifest itself in binding of these dyes also to different proteins with not exclusively enzymatic nature (Miribel et al. 1988). Similarities in charge displacement as well as the aromatic nature are believed to be responsible for the nucleotide-specific properties of antraquinone dyes; indeed, due to their molecule size and shape as well as to the topology of  $\pi$ -orbitals C.I.2 and C.I.19 are structurally related to purine nucleotides (Thompson et al. 1975). The high flexibility of molecules of C.I.2 and C.I.19 makes these substances suitable to mimic the shape of other molecules (Fig. 1).

Experiments with kinases have revealed that C.1.2 and C.1.19 are interacting specifically with the latter enzymes in their ATP-binding sites (Kopperschläger and Johansson 1985). Neverthelles similar data concerning the ATPases have been not available as yet. In preliminary experiments (Ďurišová et. al. 1989) both dyes were found to effectively inhibit the activity of lactate dehydrogenase (LDH) from skeletal muscle as well as that of heart sarcolemmal (Na<sup>+</sup> + K<sup>+</sup>)-ATPase. The aim of the present paper is quantification of the latter effect.

### Materials and Methods

LDH from bovine skeletal muscle with a specific activity of  $2.5 \,\mu$ mol NAD<sup>+</sup>, mg<sup>-1</sup>, min<sup>-1</sup> was prepared by affinity chromatography according to the method of Gemeiner et al. (1987). LDH activity was estimated based on the rate of NADH<sub>2</sub> formation during conversion of lactate to pyruvate. Enzyme reaction was performed in 3 ml of reaction medium containing phosphate buffer (20 mmol.1<sup>-1</sup>, pH = 7.5), L-lactate sodium salt. (1 mmol.<sup>-1</sup>), NAD<sup>+</sup> (0.125–4.000 mmol.1<sup>-1</sup>), hydrazine (1 mmol.1<sup>-1</sup>) and 100  $\mu$ g enzyme protein per ml. Reaction was started with addition of enzyme, and the rate of NADH<sub>2</sub> formation was measured spectrophotometrically at 340 nm utilizing the principle of Wartburg's optical test (Bergmeyer 1970).

Partially purified heart membrane fraction enriched on sarcolemma was isolated using the method of hypotonic shock combined with NaI treatment (Kostka et al. 1981). (Na<sup>+</sup> + K<sup>+</sup>)-ATPase activity was established as the difference in amounts of inorganic phosphate liberated during ATP-hydrolysis in the presence of NaCl. KCl, and MgCl<sub>2</sub> (100, 10 and 2 mmol.1<sup>-1</sup>) and in the presence of MgCl<sub>2</sub> only. Reaction was performed in a final volume of 0.5 ml containing imidazole-HCl buffer (25 mmol.1<sup>-1</sup>, pH 7.0). ATP (0.125–2.000 mmol.1<sup>-1</sup>), as well as the respective cations and it was started by addition of membrane fraction (final concentration 50  $\mu$ g membrane protein per ml). After 10 minutes incubation at 37 °C the reaction was terminated by addition of 1 ml of trichloroacetic acid (0.73 mol.1<sup>-1</sup>). Further details concerning the estimation of the (Na<sup>+</sup> + K<sup>+</sup>)-ATPase activity as well as those concerning the characteristics of the membrane fraction applied were published elsewhere (Ziegelhöffer et al. 1983; Breier et al. 1988). Protein concentration was determined according to Lowry et al. (1953).

K<sup>+</sup>-pNNPase activity was established as the difference in *p*-nitrophenyl (pNP) formed from *p*-nitrophenylphosphate (pNPP) in the presence and absence of KCI (5 mmol.1<sup>-1</sup>) during 10 minutes reaction at 37 °C in a medium containing MgCl<sub>2</sub> (2 mmol.1<sup>-1</sup>). Enzyme reaction was performed in a final volume of 0.5 ml imidazole-HCl buffer (25 mmol.1<sup>-1</sup>, pH 7.0) with 50  $\mu$ g of membrane protein per ml. Reaction was started with addition of pNPP (0.125–5.000 mmol.1<sup>-1</sup>) and stopped by addition of 0.5 ml NaOH (0.1 mol.1<sup>-1</sup>). Amounts of pNP formed were established spectrophotometrically at 410 nm. For more details see Breier et al. (1987b).

The activity of trypsin was established from the velocity of *p*-nitroaniline (pNA) formation from N<sup>*a*</sup>-benzoyl-DL-arginine-*p*-nitroaniline (BAPA) within 10 minutes. Reaction conditions: 2.5 ml imidazole-HCl buffer (25 mmol  $.1^{-1}$ , pH 7.0). 25 °C, and BAPA 0.025—0.25 mmol  $.1^{-1}$ . Reaction was started by adding 50 mg per ml of membrane protein and it was terminated by addition of 0.5ml acetic acid (30 per cent). The amounts of pNA liberated were determined spectrophotometrically at 410 nm. Further details have been described in a previous paper (Gemeiner and Breier 1982).

Estimation of C.I.2 binding to isolated cardiac sarcolemma: Isolated membrane fraction  $(100 \,\mu g \text{ per ml})$  was incubated together with C.I.2  $(1.25-20.00 \,\text{mmol} \,.1^{-1})$  for one hour at 37 °C in the presence and absence of ATP  $(2 \,\text{mmol} \,.1^{-1})$ . The reaction was stopped by spunning down at  $3000 \times g$ , with subsequent resuspension in imidazole-HCl buffer  $(25 \,\text{mmol} \,.1^{-1}, \text{ pH} 7.0)$  and repeated centrifugation at same g. The final pellet was dissolved in the same imidazole buffer as previously, supplemented with 5 per cent SDS. The amount of bound C.I.2 was monitored spectro-photometrically at 620 nm.

Enzyme kinetics and binding studies were evaluated using non-linear regression. The kinetic parameters obtained were characterized by 95% confidence intervals of according to Rektorys (1968).

All chemicals were of analytical grade and they were purchased from Sigma (St. Louis) and Lachema (Brno).



**Fig. 2.** Effect of NAD<sup>+</sup> on lactate dehydrogenase activity in the absence and presence of antraquinone dyes C.I.2 and C.I.19  $\odot$  — control values: • — C.I.2 (5  $\mu$ mol.1<sup>-1</sup>);  $\Box$  — C.I.19 (5  $\mu$ mol.1<sup>-1</sup>). The dyes were left to interact with the enzyme directly in a medium for estimation of enzyme activity. Results represent mean values of 6 experiments.



**Fig. 3.** Effect of ATP on  $(Na^+ + K^+)$ -ATPase activity in the absence and presence of antraquinone dyes C.I.2 and C.I.19.  $\bigcirc$  — control values;  $\bullet$  — C.I.19 (5  $\mu$ mol .1<sup>-1</sup>);  $\Box$  — C.I.2 (5  $\mu$ mol .1<sup>-1</sup>). The dyes were left to interact with the enzyme directly in a medium for the estimation of enzyme activity. Results represent mean values of 6 experiments.

#### **Results and Discusion**

C.I.2 and C.I.19,  $5 \mu \text{mol}.1^{-1}$ , had inhibitory effects on the specific activity of LDH. The kinetics of this inhibition is illustrated in Fig 2. The curves in Fig. 2. were obtained by non-linear regression and they indicated that C.I.2 had a more

Dye*	LDH		$(Na^+ + K^+)$ -ATPase	
	$\frac{K_{\rm M}}{[\mu{\rm m.l^{-1}}]}$	$V_{\text{max}}$ [ $\mu$ mol.NAD <sup>+</sup> .min <sup>-1</sup> . .mg <sup>-1</sup> ]	$\frac{K_{\rm M}}{[\mu {\rm mol}.1^{-1}]}$	$V_{\text{max}}$ [ $\mu$ mol.ATP.min <sup>-1</sup>
none	$0.55 \pm 0.03$	$2.69 \pm 0.12$	$0.33 \pm 0.02$	$0.27\pm0.01$
C.I.2	$6.24\pm0.08$	$2.69 \pm 0.09$	$1.45\pm0.10$	$0.27 \pm 0.01$
C.I.19	$0.82\pm0.02$	$2.69\pm0.14$	$1.72\pm0.12$	$0.27\pm0.01$

**Table 1.** Kinetic parameters of stimulation of LDH and  $(Na^+ + K^+)$ -ATPase by their specific nucleotide substrates in the absence and presence of antraquinone dyes C.I.2 and C.I.19.

<sup>\*</sup> The dyes  $(5 \,\mu \text{mol} \, 1^{-1})$  were applied directly into the medium for the estimation of enzyme activity.



**Fig. 4.** Effect of pNPP on K<sup>+</sup>-pNPPase activity in the absence and presence of antraquinone dye C.I.2. • — control values;  $\bigcirc$  — C.I.2 (5  $\mu$ mol.1<sup>-1</sup>). The dye was left to interact with the enzyme directly in a medium for the estimation of enzyme activity. Results represent mean values of 6 experiments.

pronounced inhibitory effect on LDH activity than did C.I.19. The respective kinetic parameters for both substances investigated are given in Table 1. and they suggest that the inhibition is of pure competitive nature. The above results are in good agreement with those reported by Mislovičová et al. (1988), with the LDH reaction kept running in direction of conversion of pyruvate to lactate.

	K <sup>+</sup> -pNPPase		Trypsin	
	$\frac{K_{\rm M}}{[{\rm mmol}.l^{-1}]}$	$V_{\text{max}}$ [ $\mu$ molpNP.h <sup>-1</sup> .mg <sup>-1</sup> ]	$K_{M}$ [mmol.1 <sup>-1</sup> ]	$V_{max}$ [ $\mu$ molpNA . h <sup>-1</sup> . mg <sup>-1</sup> ]
none	$2.58 \pm 0.12$	$10.79 \pm 0.48$	$0.19 \pm 0.01$	$11.22 \pm 0.41$
C.I.2	$2.58\pm0.17$	$6.54 \pm 0.32$	$0.19\pm0.01$	$11.22 \pm 0.37$

**Table 2.** Kinetic parameters of stimulation of  $K^+$ -pNPPase and trypsin by *p*-nitrophenylphosphate and Na<sup>*a*</sup>-benzoyl-DL-arginine-p-nitroanilide in the absence and presence of C.1.2.

The dye (5  $\mu$ mol.1<sup>-1</sup>) was applied directly into the medium for the estimation of enzyme activity. pNP = *p*-nitrophenylphosphate; pNA – *p*-nitroaniline.



**Fig. 5.** Effect of BAPA on trypsin activity in the absence and presence of antraquinone dye C.1.2.  $\odot$  — control values;  $\triangle$  — C.I.2 (5 µmol.1<sup>-</sup>). The dye was left to interact with the enzyme directly in a medium for the estimation of enzyme activity. Results represent mean values of 6 experiments.

The observation that C.I.2 is a more potent inhibitor of LDH reaction than is C.I.19 may be explained by the difference in their structure. From the latter aspect C.I.19 mimic the adenine moiety of NADH<sub>2</sub> molecule only, whereas C.I.2 may probably mimic the whole molecule of the dinucleotide.

In the presence of C.I.2 and C.I.19 the reaction of  $(Na^+ + K^+)$ -ATPase was also inhibited. The kinetics of this inhibition is shown in Fig. 3. Both substances investigated seem approximately similar potent inhibitors of the enzyme. The inhibition with both C.I.2 and C.I.19 was also found to be of nature competitive (Table 1). The minor difference observed in the  $K_m$  values of

Binding	$B_{\max}$ [nmol.mg <sup>-1</sup> ]*	$C_{0.5}$ [ $\mu$ mol.1 <sup>-1</sup> ]	Determination coefficient
Total	$47.4 \pm 1.6$	$11.57 \pm 1.09$	0.988
In the presence of ATP	37.9 ± 1.6	11.57 ± 0.99	0.992
Specific <sup>+</sup>	$9.5 \pm 0.4$	$11.66 \pm 0.77$	0.998

Table 3. Adsorption isotherm parameters of C.I.2 binding to sarcolemmal membrane.

\* expressed in nmol of C.I.2. per mg membrane protein. The parameters were counted by nonlinear regression according to:  $B = B_{\text{max}} \cdot c \cdot (c + C_{0.5})^{-1}$  (Breier et al. 1987a). Experimental data from Fig. 6 were used for the computations.



**Fig. 6.** Adsorption isotherms of C.1.2 binding to sarcolemmal membrane. Binding of C.1.2 in the absence ( $\bullet$ ) or presence ( $\bigcirc$ ) of ATP (2 mmol.1<sup>-1</sup>). Insert: Specific binding of C.1.2 expressed as the difference between the amounts of the dye bound to membrane in the absence and presence of ATP.

inhibition of  $(Na^+ + K^+)$ -ATPase between C.I.2 and C.I.19 also indicates that in this case the diversity in shape and size of the inhibitor molecules do not play a dominant role.

The inhibition of  $K^+$ -pNNPase by C.I.2 (Fig. 4, Table 2) was noncompetitive indicating that the active site of  $K^+$ -pNNPase is different from that of the (Na<sup>+</sup> + K<sup>+</sup>)-ATPase, although both enzymes are associated with a Na-pump (Vyskočil et al. 1981; Breier et al. 1987b). Ionic bounds as well as hydrophobic bounds may both participate on the interactions of C.I.2 with the K<sup>+</sup>-pNNPase. Another alternative is also plausible: C.I.2 may inhibit of K<sup>+</sup>-pNNPase activity via interaction and chemical modification of the part of the enzyme molecule responsible for  $(Na^+ + K^+)$ -ATPase activity.

The same concentrations of C.I.2 added to  $K^+$ -pNNPase, had no effect on the activity of trypsin (Fig. 5, Table 2).

C.1.2 may bind to sarcolemma membranes both specifically and non-specifically. Specific may be considered the binding which occurs in the nucleotidebinding sites, i.e. that in competitive relationship to nucleotide binding and related to inhibition of  $(Na^+ + K^+)$ -ATPase activity. An analysis of concentration-dependent binding of C.I.2 to isolated sarcolemmal membranes revealed that ATP counteracts significantly the specific interaction of C.I.2 with the membranes (Fig. 6, Table 3). However, a relatively significant amount of C.I.2 is bound to the sarcolemmal membranes in loci other than the nucleotide-binding sites. This binding considered as non-specific probably involves hydrophobic or ionic interactions and is related to high partition coefficient of C.I.2 in the membrane-water system.

From the results obtained the following conclusions may be drawn: a) the antraquinone dyes C.I.2 and C.I.19 inhibit those enzymes only which contain some nucleotide-binding sites;

b) if the nucleotide-binding site on the respective enzyme is identical with its active site, similarly as in the case of LDH or  $(Na^+ + K^+)$ -ATPase, a competitive inhibition occurs;

c) if antraquinone dyes interact with a nucleotide-binding site other than the substrate-binding site of the respective enzyme non-competitive inhibition may be observed;

d) comparing the results obtained with LDH and  $(Na^+ + K^+)$ -ATPase inhibition by C.I.2 and C.I.19 it may be concluded that both inhibitors have really a nucleotide specific character and they will act anywhere, where they can replace an adenine nucleotide and/or a nicotineamide-adenine dinucleotide;

e) a further conclusion to be drawn is that the inhibitory effect of the dyes investigated is the more expressed the more their molecule resemble in shape and size the substrate to be replaced;

f) the results obtained confirmed the suitability of the antraquinone dyes investigated for use as nucleotide-specific ligands in enzyme reactions as well as in some special cases of affinity chromatography.

## References

Bergmeyer H. V. (1970): Methoden der enzymatischen Analyse. Verlag Chemie, Mainheim

- Breier A., Gemeiner P., Ziegelhöffer A. Turi-Nagy L., Štofaniková V. (1987a): Application of a time-concentration model of adsorption for determination of the nature of adsorbed-adsorbate interaction. Coll. Polym. Sci. 265, 933-937
- Breier A., Monošíková R., Ziegelhöffer A. (1987b): Modification of primary amino groups in heart sarcolemmal by 2, 4, 6-trinitrobenzene sulfonic acid in respect to the activities of (Na<sup>+</sup> + K<sup>+</sup>)-ATPase, Na<sup>+</sup>-ATPase and pNPPase. Function of the potassium binding sites. Gen. Physiol. Biophys. 6, 103–108
- Breier A., Turi-Nagy L., Ziegelhöffer A., Monošíková R. (1988): Principles of selectivity of sodium and potassium binding sites of the Na<sup>+</sup>/K<sup>+</sup>-ATPase. A corollary hypothesis. Biochim. Biophys. Acta 946, 129–134
- Duríšová V., Ziegelhöffer A., Monošíková R., Džurba A., Breier A. (1989): Affinity labelling of ATP-binding site of heart sarcolemmal (Na<sup>+</sup> + K<sup>+</sup>)-ATPase. Physiol. Bohemslov **38**, 511
- Gemeiner P., Breier A. (1982): Aldehydic derivatives of bead cellulose relationship between the matrix structure and function in imobilization of enzymes catalyzing hydrolysis of high molecular substrates. Biotechnol. Bioeng. 24, 2573—2583
- Gemeiner P., Mislovičová D., Zemek J., Kuniak E. (1981): Antraquinone-triazine derivatives of polysaccharides: Relation between structure and affinity to lactate dehydrogenase. Collect. Czech. Chem. Commun. 46, 419–427
- Gemeiner P., Zemek J., Babušíková Ľ. (1987): Isolation of lactatedehydrogenase. AO 236136 (in Slovak)
- Kopperschläger G., Böhme H.-J., Hofmann E. (1982): Cibacron Blue F3G A and related dyes as in affinity chromatography. Adv. in Biochem. Engineering **25**, 101–137
- Kopperschläger G., Johansson G. (1985): Studies on the ATP-sensitivity of yeast phosphofructokinase by means of affinity partitioning using polymer bound Cibacron Blue F3G A. Biomed. Biochim. Acta 47, 1047–1055
- Kostka P., Ziegelhöffer A., Džurba A., Vrbjar N. (1981): A comparative study of the enzyme characteristics of different sarcolemmal preparation from the rat heart. Physiol. Bohemoslov. 3, 173
- Kroviarski Y., Cochet S., Vadon C., Truskolashi A., Boivin P., Bertrand O. (1988): Purification of human 6-phosphogluconate dehydrogenase from human haemolyzate with chromatography on an immobilized dye as the essential step and use of automation. Simultaneous purification of lactate dehydrogenase. J. Chromatogr. 449, 413–422
- Kulbe K. D., Schuer R. (1979): Large scale preparation of phosphoglycerate kinase from Saccharomyces cerevisiae using CB-Sepharose 4B pseudoaffinity chromatography. Anal. Biochem. 93, 46-51
- Lowry O. H., Rosebrough N. J., Farr A. L., Randall R. J. (1951): Protein measurement with folin-phenol reagent. J. Biol. Chem. 193, 265-275
- Miribel L., Gianazza E., Arnaud P. (1988): The use of dye-ligand affinity chromatography for the purification of non-enzymatic human plasma proteins. J. Biochem. Biophys. Meth. 16, 1–16
- Mislovičová C Gemeiner P., Breier A. (1988): Study of porous cellulose beads as an affinity adsorbent via quantitative measurements of interactions of lactate dehydrogenase with immobilized antraquinone dyes. Enzyme Microb. Technol. 10, 568–573
- Rektorys K. (1968): Handbook of Applied Mathematics. State Publishing House of Technical Literature, Prague (in Czech)

- Scawen M. D., Darbyshire J., Harvey M. J. Atkinson T. (1982): The rapid purification of 3-hydroxybutyrate dehydrogenase and malate dehydrogenase on triazine dyes affinity matrices. J. Biochem. 203, 699—705
- Subramanian S. (1984): Dye-ligand affinity chromatography: The interaction of Cibacron Blue F3G with proteins and enzymes. Crit. Rev. Biochem. 16, 169-205
- Thompson S. T., Cass K. H., Stellwagen E. (1975): Blue Dextran-Sepharose: An affinity column for the dinucleotide fold in proteins. Proc. Nat. Acad. Sci. USA 72, 669–674
- Vyskočil F., Teisinger J., Dlouha H. (1981): The disparity between effect of vanadate (V) and vanadyl (IV) ions on sodium and potassium ATPase and potassium phosphatase in skeletal muscle. Biochem. Biophys. Res. Commun. 100, 982–987
- Ziegelhöffer A., Breier A., Džurba A., Vrbjar N. (1983): Selective and reversible inhibition of heart sarcolemmal (Na<sup>+</sup> + K<sup>+</sup>)-ATPase by *p*-bromophenyl isothiocyanate. Evidence for a sulfhydryl group in the ATP-binding site of the enzyme. Gen. Physiol. Biophys. 2, 447–456

Final version accepted May 15, 1990