

EFFECTS OF K^+ DIFFUSION POTENTIALS ON DUROQUINOL-CYTOCHROME *c* REDUCTASE ACTIVITY CATALYZED BY COMPLEX III INCORPORATED INTO LIPOSOMES

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1. Introduction

It has been shown that beef heart quinol-cytochrome *c* reductase (complex III) incorporated into phospholipid vesicles can drive proton translocation [1,2]. In such vesicles, quinol-cytochrome *c* reductase activity is inhibited in the absence of uncoupling agents or valinomycin + K^+ [1–3], and it is generally assumed that inhibition is due to the presence of a membrane potential or an electrochemical gradient of protons developed between the internal and external osmotic spaces [4–6]. To date, however, little information has been published showing that quinol-cytochrome *c* reductase can be influenced in reconstituted vesicles by ionic movements between the two osmotic spaces. In the present study we show that duroquinol-cytochrome *c* reductase activity can be brought into equilibrium with a K^+ diffusion potential, and that the enzyme is inactivated as the potential becomes more negative inside the vesicles. In addition, we also show, using the cationic dye safranin, that vesicles containing complex III can drive the formation of a membrane potential.

2. Methods

Complex III was prepared from beef heart mitochondria as in [8]. Phospholipid vesicles containing complex III were prepared from soybean phospholipids (Sigma) as in [1,2] using the cholate

dialysis procedure [9]. The phospholipids were extracted with acetone [9] prior to use. Vesicles were prepared in 50 mM Tris- SO_4 , pH 7.5, containing various concentrations of KCl or NaCl as indicated in the figure legends. Safranin uptake and duroquinol-cytochrome *c* reductase activity [2] were routinely measured in 1 ml buffer containing 50 mM Tris- SO_4 , pH 7.5, 1 mM EDTA, and varying amounts of KCl or NaCl to maintain osmolarity with the vesicles. Reduction of cytochrome *c* was measured at 550–540 nm and safranin uptake was measured at 530–578 nm [10].

Membrane potentials were calculated from the initial concentrations of K^+ inside, $[K^+]_{in}$, and outside, $[K^+]_{out}$, using the Nernst equation:

$$E_m = 59 \log \frac{[K^+]_{in}}{[K^+]_{out}}$$

For these calculations, $[K^+]_{in}$ was assumed to be the same as the buffer $[K^+]$ during preparation of the vesicles. The concentration ratio of safranin, $[S^+]_{in}/[S^+]_{out}$, was calculated assuming that a decrease in absorbance at 530–578 nm is proportional to safranin uptake, $[S^+]_{in}$, and that 100% uptake was achieved with no external K^+ present.

3. Results

Figure 1 shows a typical experiment using liposomes containing complex III. As reported [11,13], a valinomycin-induced K^+ diffusion potential drives the

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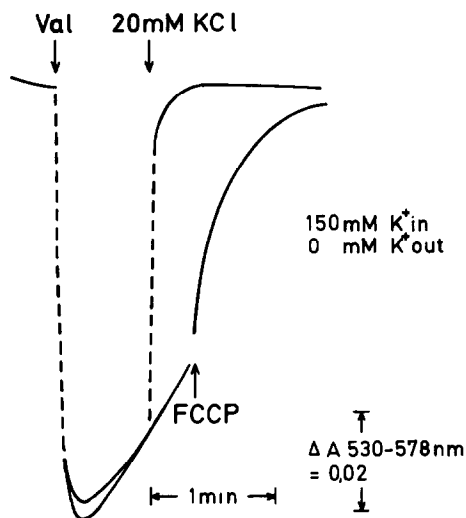


Fig.1. K^+ -driven uptake of safranin into phospholipid vesicles reconstituted with beef heart complex III. Vesicles were prepared in 50 mM Tris- SO_4 , pH 7.5, 150 mM KCl and 1 mM EDTA. The reaction media contained 50 mM Tris- SO_4 , pH 7.5, 150 mM NaCl, 1 mM EDTA, 75 μ g phospholipid and 18 μ M safranin in total vol. 1 ml. Valinomycin and FCCP were 0.5 μ g/ml and 1 μ M respectively.

uptake of safranin. Safranin uptake can be prevented or reversed by any reagent which collapses the K^+ gradient (fig.1). That safranin probes an electrical event across the liposomal membrane is shown in fig.2. As expected, the absolute external $[K^+]$ needed to prevent safranin uptake is dependent upon the $[K^+]$ inside the vesicles (fig.2A). However, when safranin uptake is plotted against the K^+ concentration gradient (fig.2B), all points fall on the same line, regardless of absolute internal $[K^+]$. A plot of the log of the accumulation ratio of safranin, $\log [S^+]_{in}/[S^+]_{out}$, against the membrane potential, gives a straight line with a slope of 1 (data not shown). It should be noted, however, that the absolute absorbance changes obtained for a particular liposomal preparation are strongly dependent upon the ratio of safranin to phospholipid, and must be taken into consideration when comparing quantitative results obtained with different liposomal preparations. Optimal absorbance changes are obtained with a ratio of 0.25 μ mol safranin/mg phospholipid.

Figure 3A shows that duroquinol-cytochrome *c* reductase activity is controlled by a K^+ diffusion

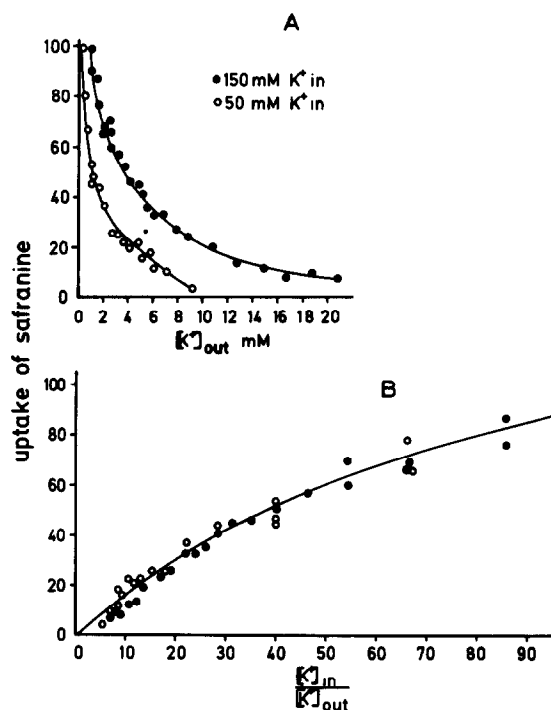


Fig.2. Effects of internal K^+ concentration on safranin uptake by complex III-phospholipid vesicles. Experimental conditions were as given in fig.1 with the exception that vesicles were made in the presence of either 50 mM or 150 mM KCl. As the external KCl concentration was raised, the external NaCl concentration was decreased proportionally to maintain osmolarity with the internal KCl.

potential. Since the membrane potential is calculated from the K^+ concentration gradient it is expressed as a positive value. It is clear, however, that as the membrane potential becomes more negative, as measured by safranin uptake, the enzyme become inactivated (fig.3A). Safranin uptake and enzyme inactivation appear to be in equilibrium with each other (fig.3B) and with the K^+ diffusion gradient (fig.3C).

Duroquinol-cytochrome *c* reductase also drives the formation of a membrane potential in liposomes, as indicated by the uptake of safranin (fig.4). Safranin uptake can be prevented or abolished by FCCP or valinomycin + K^+ , conditions which also activate duroquinol-cytochrome *c* reductase [1-3]. The slight net increase in absorbance observed in the presence of FCCP or valinomycin is due to interference from reduced cytochrome *c* at these wavelengths.

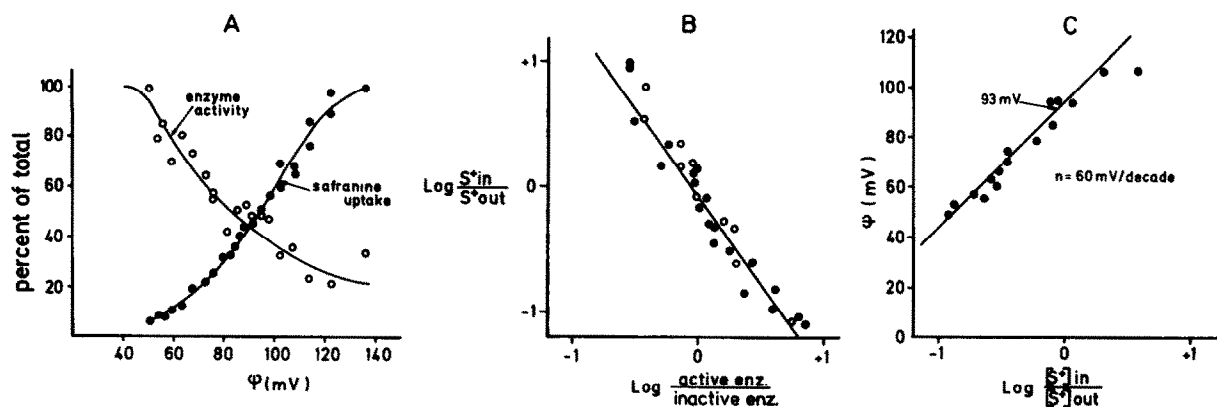


Fig.3. Control of duroquinol-cytochrome *c* reductase in complex III-phospholipid vesicles by K^+ diffusion potentials. Conditions were as in fig.2 with the exception that $60 \mu\text{M}$ duroquinol and $12 \mu\text{M}$ cytochrome *c* were present in the media during the measurements of enzyme activity. Duroquinol-cytochrome *c* reductase activity and safranin uptake were initiated by the addition of $0.5 \mu\text{g}$ valinomycin. The enzymatic rates shown in the figure are the initial rates obtained after addition of valinomycin.

4. Discussion

The present study shows that:

1. Duroquinol-cytochrome *c* reductase in reconstituted phospholipid vesicles can be brought into equilibrium with a K^+ diffusion potential, and becomes inactivated as the membrane potential becomes more negative inside the vesicle.
2. Duroquinol-cytochrome *c* reductase can drive the formation of a membrane potential in complex III vesicles, as shown by the uptake of safranin.
3. The diffusion potential-driven uptake of safranin by complex III vesicles follows the Nernst equation

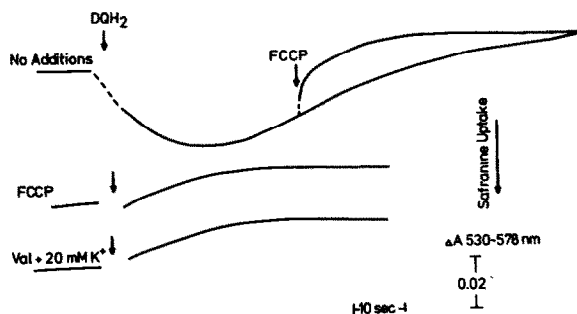


Fig.4. Formation of a membrane potential driven by duroquinol-cytochrome *c* reductase incorporated into phospholipid vesicles. The conditions of the assay were as in fig.1 and fig.3.

and is a reliable qualitative probe of transmembrane potentials.

Complex III vesicles do not catalyze the reduction of cytochrome *c* unless an uncoupling agent or valinomycin + K^+ is present [1-3]. This has been interpreted to mean that the first turnovers of the enzyme create a redox-driven electrochemical proton gradient which is sufficient to inhibit electron-transfer [1-3]. The present observation that duroquinol-cytochrome *c* reductase is able to drive the uptake of safranin shows that the enzyme can form a membrane potential, and is in keeping with the above interpretation. The present findings also clearly show that both safranin uptake and duroquinol-cytochrome *c* reductase activity can be brought into equilibrium with a K^+ diffusion gradient (fig.3), indicating that the enzyme can be controlled by ionic movements between the internal and external osmotic spaces. The effect of the ΔpH (pH gradient) on duroquinol-cytochrome *c* reductase activity has not yet been successfully determined due:

- (i) To the lack of reliable methods for measuring internal pH in liposomal systems where the proton movement is directed outwards.
- (ii) To the fact that isolated duroquinol-cytochrome *c* reductase is strongly pH dependent [3], making acid-base transition experiments [13] difficult to interpret.

It should be mentioned, however, that in several

experiments we have observed valinomycin-activated rates to be 20–40% lower than FCCP-activated rates. This difference in enzyme activity can be due to the influence of the ΔpH which is enhanced in the valinomycin-treated vesicles.

We have attempted to use the K^+ -driven uptake of safranin as a standard to quantitate [11–13] membrane potentials formed under conditions of electron transport inhibition [1–3]. The results show that the duroquinol-cytochrome *c* reductase is half-maximally activated by a K^+ diffusion potential of 90–95 mV. In contrast, duroquinol-cytochrome *c* reductase drives the formation of a potential no greater than 10 mV, as measured by safranin (data not shown), even under conditions where electron transport is controlled [1–3]. This discrepancy is most easily explained if, like cytochrome oxidase [14,15], complex III is incorporated into only a small percent of the vesicles. In this case, all vesicles would accumulate safranin in the presence of a K^+ diffusion potential, whereas only a small percentage would accumulate safranin when driven by duroquinol-cytochrome *c* reductase. Quantitation of the membrane potential based upon K^+ -driven safranin uptake [12,13] would, thus, lead to a considerable underestimation of the enzyme-driven membrane potential.

It would thus appear that the membrane potentials calculated from K^+ diffusion gradients are probably nearest to those which actually control duroquinol-cytochrome *c* reductase activity (fig.3). Even these values are open to question, however, since they are based on the assumption that the K^+ -driven safranin response is independent of the size of the liposomal vesicle. The size of the vesicle prepared by cholate dialysis can vary considerably [14]. It now appears that proteins can be inserted into vesicles of a restricted and uniform size [14]. Quantitation of membrane potentials in liposomal vesicles using safranin as a probe is, thus, associated with problems apparently not experienced with intact rat liver mitochondria [12,13].

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