The Phosphonium Ion Efflux System of *Escherichia coli*: Relationship to the Ethidium Efflux System and Energetic Studies

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The extent of accumulation of methyltriphenylphosphonium ion by *Escherichia coli* was shown to be dependent on the permeability of the outer membrane and the activity of an efflux system for this compound. Evidence consistent with the operation of a single efflux system for compounds such as phosphonium ions, phenanthridiniums and flavines is presented. Studies on the energy coupling mechanism for this efflux system indicated that it was driven by the transmembrane proton electrochemical gradient.

**INTRODUCTION**

Measurement of the distribution of phosphonium ions, such as methyltriphenylphosphonium ion (MTP) and tetraphenylphosphonium ion (TPP), has been used to estimate the magnitude of the membrane potential ($\Delta \psi$) in a number of biological systems (Rottenberg, 1975) including bacteria (Kashket, 1985). *Arthrobacter globiformis* and *Acinetobacter calcoaceticus* possess an energy-linked efflux system for phosphonium ions and related compounds such as tetraphenylarsonium ion (TPA) and crystal violet (Midgley et al., 1985, 1986a), and *Escherichia coli* possesses an efflux system for which phosphonium ions are a substrate (Midgley et al., 1986b). The existence in *E. coli* of an efflux system capable of extruding other complex cations has been shown from studies of the interaction of this organism with acriflavine (Nakamura, 1966, 1967; Silver et al., 1968; Kushner & Khan, 1968) and ethidium (Lambert & Le Pecq, 1984; Jones & Midgley, 1985).

The aims of the work presented here were to confirm the fluorimetric assay (Midgley et al., 1986b) for the existence of a phosphonium ion efflux system using isotopic assays, examine the relationship between the phosphonium ion efflux system and those previously detected and investigate the energy-coupling mechanism of the efflux system.

**METHODS**

*Growth and harvesting of the organism. Escherichia coli* AN 249, an uncA mutant (Cox et al., 1973), was grown and harvested as described by Midgley et al. (1986b). All experiments were done using 20 mM-HEPES/NaOH buffer (pH 7-0).

*Phosphonium ion accumulation.* This was measured using a rapid filtration technique (Midgley & Dawes, 1973). The organism was incubated aerobically at 37 °C at a cell density of 0.5 mg dry wt ml⁻¹. All assays were done in duplicate. Uptake was corrected for extracellular and filter binding using a control incubated on ice in the absence of glucose or formate. For experiments using 1 or 5 μM-MTP this value was typically 0.02 or 0.05 μmol (g dry wt)⁻¹, respectively. Unless otherwise stated the concentration of MTP used in isotopic experiments was 1 μM [specific activity 125 μCi μmol⁻¹ (4.625 MBq μmol⁻¹)]. In situ treatment with EDTA was achieved by addition of 0.3 mM-EDTA directly to the incubation mixture. Further details are given in the appropriate figure legends.

**Abbreviations:** $\Delta \psi$, (plasma) membrane potential; $\Delta \psi_p$, proton-motive force (proton electrochemical gradient in mV); $\Delta \psi_i$, electrochemical gradient; MTP, methyltriphenylphosphonium ion; TPP, tetraphenylphosphonium ion; TPA, tetraphenylarsonium ion; DMP, 2-(dimethylaminostyryl)-1-ethylpyridinium; CCCP, carbonyl cyanide m-chlorophenylhydrazone; IAA, iodoacetate.
Ethidium efflux. This was measured using a fluorimetric and an isotopic assay as described by Jones & Midgley (1985), except that 20 mM-HEPES/NaOH (pH 7.0) was used throughout. Cells were loaded with ethidium (5 μM) as previously described except that buffer was used throughout (Lambert & Le Pecq, 1984; Jones & Midgley, 1985). In the fluorimetric assay, aerobic conditions were maintained by adding H₂O₂ (0.016% vol. final concentration) along with substrate (glucose or sodium formate).

Fluorimetric measurements using DMP. These were made as described by Midgley et al. (1986b).

Energy-depleted cells. These were prepared by incubating the cells (10 mg dry wt ml⁻¹) aerobically at 37 °C in the presence of 40 μM-CCCP for 1 h. The cell suspension (10 ml) was then slowly cooled and diluted to 40 ml with buffer. The cells were sedimented, resuspended in 40 ml of buffer and again sedimented before resuspension in 2 ml of buffer.

Respiration studies. These were done at 37 °C using a Rank oxygen electrode. The cell density was 0.25 or 0.5 mg dry wt ml⁻¹.

Chemicals and radiochemicals. These were obtained from sources specified by Midgley et al. (1986a) and Jones & Midgley (1985).

RESULTS

Isotopic demonstration of MTP efflux

Washed cells of E. coli AN 249 incubated with 5 μM-MTP in the absence of an added energy source accumulated MTP (Fig. 1). This uptake was prevented by the prior addition of glucose which also supported the efflux of accumulated MTP (Fig. 1), and was inhibited by 40 μM-CCCP (data not shown). Glucose stimulated respiration approximately 14-fold to 54 μmol O₂ consumed min⁻¹ (g dry wt)⁻¹.

Factors affecting the net influx of MTP

A series of experiments was done to demonstrate the relationship of the net influx of MTP to the presence of putative substrates of the efflux system. In addition, the effect of an energy source and changes in the outer membrane permeability (produced by in situ EDTA treatment) on the relationship were investigated. Net influx of MTP was stimulated by 10 μM-TPP in the absence of an added energy source (Fig. 2), and 10 μM-TPA, 10 μM-ethylavine or 10 μM-ethidium bromide stimulated MTP uptake by 9-, 5.5- and 2.7-fold respectively, measured after 30 min. The absolute value of the control in the absence of an added energy source was 0.078 μmol MTP (g dry wt)⁻¹. The uptake of MTP in the presence of TPP was inhibited by 40 μM-CCCP (data not shown).

A stimulation of MTP uptake by TPP was not observed in the presence of glucose (Fig. 2). TPP (10 μM) was accumulated to <0.2 and 3.8 μmol TPP (g dry wt)⁻¹ after 40 min, in the presence and absence of glucose, respectively. If the cells were treated in situ with EDTA the stimulatory effect was again observed. Cells incubated with 1 μM-MTP and glucose plus EDTA or glucose, EDTA and 10 μM-TPP accumulated at the steady state 0.018 or 0.22 μmol MTP (g dry wt)⁻¹, respectively, and cells incubated with 10 μM-TPP accumulated 0.07 or 2.7 μmol TPP (g dry wt)⁻¹ in the presence of glucose or glucose plus EDTA, respectively.

The uptake of MTP by EDTA-treated cells in the presence of glucose and TPP was inhibited by 40 μM-CCCP (data not shown).

Relationship of the phosphonium ion efflux system to the ethidium efflux system

The stimulation of net MTP influx by ethidium was consistent with these two compounds sharing a common efflux system. Since it is possible to measure unidirectional efflux of ethidium from cells suitably loaded with this compound (Lambert & Le Pecq, 1984; Jones & Midgley, 1985), and since effects on unidirectional fluxes are easier to interpret, this relationship was further investigated. Using either a fluorimetric assay (Fig. 3a) or an isotopic assay (Fig. 3b) TPP, TPA and MTP were demonstrated to inhibit ethidium efflux.

The loading process involves a brief treatment with CCCP which was apparently sufficient to deplete the cells of an energy supply as ethidium efflux was substrate dependent, and could be driven by either glucose (Fig. 4a, b) or formate (Fig. 3a, b). The inhibitory effect of TPA, TPP and MTP was not attributable to inhibition of respiration; 10 mM-sodium formate supported a
Phosphonium ion efflux from *E. coli*

![Graph 1](image1)

**Fig. 1.** Substrate-induced efflux of MTP from *E. coli* AN 249. MTP uptake was assayed as described in Methods except that 5 μM-MTP was used under the following conditions: ●, no substrate added; □, plus 10 mM-glucose as indicated by the arrow; ■, plus 10 mM-glucose throughout.

**Fig. 2.** Stimulation of MTP accumulation by TPP in *E. coli* AN 249. MTP uptake was assayed as described in Methods under the following conditions: ○, no addition; ●, plus 10 μM-TPP; □, plus 10 mM-glucose or plus 10 mM-glucose and 10 μM-TPP.

![Graph 2](image2)

**Fig. 3.** Effect of MTP, TPP and TPA on the efflux of ethidium from *E. coli* AN 249. (a) Fluorimetric assay. In traces I and IV the concentration of TPP and TPA was 10 μM. For trace II 30 μM-MTP was added before 10 mM-sodium formate. Trace III is the control, with no addition other than formate. The traces are offset for the purpose of display. (b) Isotopic assay. Efflux of ethidium was assayed as described in Methods under the following conditions: ○, no addition; ●, plus 10 μM-TPA; △, plus 50 μM-MTP; □, plus 10 μM-TPP. In all cases efflux was initiated by the addition of 10 mM-sodium formate as indicated by the arrow. For clarity the initial phase in the presence of MTP and TPA is not shown.
rate of respiration of 189 \( \mu \text{mol O}_2 \text{ min}^{-1} (\text{g dry wt})^{-1} \), and this was not affected by the addition of 50 \( \mu \text{M}-\text{MTP} \) or 10 \( \mu \text{M}-\text{TPP} \) or 10 \( \mu \text{M}-\text{TPA} \). The endogenous rate of respiration was < 11 \( \mu \text{mol O}_2 \text{ min}^{-1} (\text{g dry wt})^{-1} \).

**Energetics of ethidium and phosphonium ion efflux**

Glucose-driven efflux of ethidium was inhibited by 5 mM-iodoacetate (IAA) while formate-driven efflux was insensitive to this compound (Fig. 4). IAA caused severe inhibition of glucose-supported respiration (> 90%), but not of respiration supported by formate. In the presence of glucose or formate the rate of respiration, measured using cells loaded with ethidium, was 62 or 180 \( \mu \text{mol O}_2 \text{ min}^{-1} (\text{g dry wt})^{-1} \), respectively. The endogenous rate was < 6 \( \mu \text{mol O}_2 \text{ min}^{-1} (\text{g dry wt})^{-1} \).

Since *E. coli* AN 249 is an *unc* mutant and therefore cannot produce ATP by oxidative phosphorylation, \( \Delta \mu_H^+ \) was sufficient to drive ethidium efflux. Both glucose- and formate-driven efflux were inhibited by 40 \( \mu \text{M}-\text{CCCP} \); this concentration of uncoupler did not inhibit formate oxidation (data not shown).

Evidence that the phosphonium ion efflux system was operative under the conditions where only a \( \Delta \mu_H^+ \) was generated was provided by isotopic experiments using MTP (Fig. 5) and fluorimetric experiments using DMP as a substrate (Fig. 6) (Midgley et al., 1986b). Cells of *E. coli* AN 249 were starved aerobically for 1 h in the presence of 40 \( \mu \text{M}-\text{CCCP} \), a treatment that diminished respiration to undetectable levels. Although the possibility that the cells retained some CCCP was not excluded, cells treated in this manner showed no uptake of MTP in the absence of substrate in the presence or absence of 10 \( \mu \text{M}-\text{TPP} \) (data not shown). In the presence of 10 mM-sodium formate, 10 \( \mu \text{M}-\text{TPP} \) stimulated net influx of MTP, and this was largely
Phosphonium ion efflux from *E. coli*

![Graph](image)

**Fig. 5.** Interaction of phosphonium ions with energy-depleted *E. coli* AN 249. Cells were starved and assayed for MTP uptake as described in Methods under the following conditions: ○, plus 10 mM-sodium formate or 10 mM-sodium formate plus 5 mM-IAA; △, plus 10 mM-sodium formate and 10 μM-TPP; ●, plus 10 mM-sodium formate, 10 μM-TPP and 5 mM-iodoacetate.

![Diagram](image)

**Fig. 6.** Interaction of DMP with starved *E. coli* AN 249. Cells were starved and assayed for fluorimetric interaction with DMP as described in Methods. The concentrations used were 5 μM-DMP, 10 μM-TPP, 10 mM-glucose and 10 mM-sodium formate. The incubation mixture for trace I contained 5 mM-IAA throughout. The traces are offset for the purpose of display.

unaffected by the addition of 5 mM-IAA (Fig. 5); in the presence and absence of IAA the rate of respiration was equivalent to that supported by formate in unstarved cells [180 μmol O₂ min⁻¹ (g dry wt)⁻¹].

DMP efflux could be driven in starved cells by glucose or formate; glucose-driven efflux, but not that driven by formate, was sensitive to IAA (Fig. 6). The stimulation of DMP fluorescence, previously suggested to arise from a stimulation of net influx due to an inhibition of DMP efflux (Midgley *et al.*, 1986b), was not observed in starved cells unless glucose or formate were added (Fig. 6).

**DISCUSSION**

The observations presented in this paper confirm the conclusions drawn from fluorimetric studies using DMP as a probe for the phosphonium ion efflux system (Midgley *et al.*, 1986b). The increased fluorescence of DMP on addition of phosphonium ions to *E. coli* was attributed to an inhibition of a shared efflux system, leading to an increased net influx. This interpretation is supported by the observation of a substrate-induced efflux of MTP (Fig. 1). The stimulation of
respiration accompanying the addition of glucose would predict an increase in \( \Delta \bar{\mu}_{\text{MTP}} \), due to an increase in the \( \Delta \psi \) component of the \( \Delta \bar{\mu}_{\text{H}} \) at this external pH. A clear efflux phase was observed as seen with other organisms (Midgley et al., 1985, 1986a). Thus in the presence of an oxidizable substrate such as glucose or formate, *E. coli* appears impermeable to phosphonium ions when its outer membrane is intact. Addition of putative substrates for the efflux system inhibits efflux of MTP and thus leads to a stimulation of net influx (Fig. 2). As previously discussed (Midgley et al., 1986a, b), the stimulation of net uptake of MTP by an analogue such as TPP can only be used as a positive criterion since many factors can govern the steady state intracellular concentration of such a compound. The stimulation of MTP uptake by ethidium, euflavine and TPA suggests that these compounds are substrates for this efflux system.

The rapid penetration of the organism by TPP and TPA, as shown by the rapid establishment of the inhibition of ethidium or DMP efflux (Figs 3a and 6), indicates that isotopic studies underestimate the rate of influx since a net flux rather than a unidirectional one is measured. The relative contribution of the efflux system and the outer membrane in diminishing the uptake of phosphonium ions, and related toxic cations such as euflavine, is currently unknown.

The observations of a stimulation of net MTP influx by ethidium, and the inhibition of ethidium efflux by phosphonium ions and TPA (Fig. 3a, b), are consistent with a single efflux system operating to transport these compounds, as well as DMP, out of *E. coli*. This hypothesis is further supported by the isolation of ethidium-resistant mutants of *E. coli* N43, a strain that is susceptible to ethidium because it carries the *acrA* mutation (Nakamura, 1965). These mutants are resistant to ethidium, while retaining the original mutation, due to a higher level of the efflux system. Such mutants are also resistant to MTP (M. Midgley, unpublished observations). Ethidium efflux (Fig. 4) and DMP efflux (Fig. 6), and the stimulation of net MTP influx by TPP (Fig. 5), can occur in the presence of a \( \Delta \bar{\mu}_{\text{H}} \), and ATP does not appear to have a significant role. Similar findings (I. G. Jones & M. Midgley, unpublished observations) have been made for the ethidium resistance determinant previously cloned from a staphylococcal plasmid (Jones & Midgley, 1985). A possible mechanism for these efflux systems could be an antiporter of the type \( nX^+ / \text{ethidium} \), where \( X^+ \) is a suitable cation, e.g. \( H^+ \). Further experiments, including testing the effect of adding cations other than \( Na^+ \) to the medium (as in the present work) will be of value.

The accumulation of phosphonium ions by *E. coli* corresponds to a steady state in which extrusion by an efflux system balances influx. This means that the accumulation ratio will be a function of at least the following possible variables: the outer membrane permeability, the permeability of the cytoplasmic membrane, the transmembrane \( \Delta \psi \), the transmembrane \( \Delta p \) (as a driving force for the efflux system), the surface potential at relevant membrane interfaces, the activity of the efflux system, and the concentration of the probe used. Thus many factors can be involved in determining the apparent \( \Delta \psi \) as computed from the accumulation of phosphonium ions. The majority of these factors are subject to possible variation by, for example, changing the growth conditions, the strain of *E. coli* used, or the incubation medium used. In general it is likely that the operation of such an efflux system as described here will lead to underestimates of the \( \Delta \psi \). In some cases, e.g. in vesicular systems in which the outer membrane has been removed, the operation of the efflux system may not play a significant role, but clearly much of the data derived from the application of these probes to bacteria (Kashket, 1985) will require re-evaluation in the light of these findings. The models presented to explain the anomalous interaction of phosphonium ions with *Bacillus subtilis* were derived on the assumption that no active transport system played a role in the distribution of the probes used (Zaritsky et al., 1981). If an efflux system similar to that described in this work is present in *B. subtilis* then the increase in apparent \( \Delta \psi \) recorded at increasing concentrations of the probe can be explained on the basis of an increasing contribution of the non-mediated fluxes to the steady state. This behaviour has been observed in *A. calcoaceticus*, an organism known to possess an efflux system for phosphonium ions (Midgley et al., 1986a).
Phosphonium ion efflux from E. coli

REFERENCES


