

The respiration-driven active sodium transport system in *E. coli* does not function with lithium

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Abstract Comparison of respiration-driven active transport of alkali cations from *E. coli* cells loaded with Na⁺ or Li⁺ showed that Li⁺ could not be expelled from the cells like Na⁺. K⁺ accumulation, which was fast in Na⁺-loaded cells, was strongly inhibited in Li⁺-loaded cells, despite high membrane potential and respiratory rate. When Li⁺-loaded cells were placed into medium containing Na⁺ instead of Li⁺, Li⁺/Na⁺ exchange took place initially, while K⁺ accumulation was delayed. Only after almost all inside Li⁺ was replaced by Na⁺ did active Na⁺ and K⁺ transport commence. These data confirm that it is a distinct active sodium transport system (AST) with Na⁺,K⁺/H⁺ antiporter activity, and not the Na⁺/H⁺ antiporters, that is responsible for active Na⁺ transport in *E. coli* [Verkhovskaya, M.L., Verkhovsky, M.I. and Wikström, M. (1996) *Biochim. Biophys. Acta* 1273, 207–216]. In contrast to the Na⁺/H⁺ antiporters, the AST system is inhibited by Li⁺.

Key words: Na⁺ transport; Na⁺/Li⁺ selectivity; *E. coli*

1. Introduction

It has been generally accepted that the respiration-dependent sodium gradient in *E. coli* is generated by the activity of two Na⁺/H⁺ antiporters, the products of the *nhaA* and *nhaB* genes (see [1] for a review). The *nhaA* and *nhaB* genes both confer resistance to Li⁺ [2,3], which is toxic for *E. coli* [4]. Thus, these gene products were considered to extrude Li⁺ as well as Na⁺ from the cells. Earlier, it was shown that Li⁺, like Na⁺, is efficient in dissipating Δ pH generated by respiration in sub-bacterial vesicles from *E. coli* [5]. Later, this was found to be due to functioning of the *nhaA* and *nhaB* gene products [3]. The *nhaB* gene product was alone responsible for dissipation of Δ pH generated by respiration in membrane vesicles from an *nhaA* gene-deleted mutant, Na⁺ and Li⁺ being equally effective dissipators [6]. Finally, the purified NhaA protein catalyses Na⁺/Li⁺ as well as Na⁺/Na⁺ exchange in reconstituted liposomes [7]. All these results prove that the two well-characterised Na⁺/H⁺ antiporters can also operate with Li⁺.

We recently suggested that the major respiration-linked Na⁺ transport in *E. coli* is not catalysed by the NhaA or NhaB antiporter as previously thought, but by an active sodium transport system (AST), which possesses Na⁺,K⁺/H⁺ antiporter activity [8]. Here we present results showing that Li⁺ cannot be actively extruded from respiring *E. coli* cells like Na⁺ which strongly corroborates our proposal.

2. Materials and methods

The *E. coli* strain GR70N was a gift from Dr. R.B. Gennis, University of Illinois. The cells were grown aerobically overnight at 37°C in synthetic salt medium, pH 7.6, containing sodium succinate, and harvested in the middle of the exponential growth phase. Then the cells were loaded with Na⁺ or Li⁺ using the diethanolamine treatment [9]. Loaded cells were concentrated to 30–40 mg protein/ml and kept on ice.

Transport of alkali cations was initiated by dilution of cell suspensions into aerated, stirred Na⁺ (or Li⁺) medium, containing 150 mM NaCl (or LiCl), 50 mM tricine-NaOH (or LiOH), pH 8.5, 5 mM MgSO₄, 10 mM KCl, 2 mM dithiotreitol (DTT) and 200 μ M ubiquinone-1 (Q-1). Then the cell suspension was filtered (Millipore, 45 μ m), the filter was washed with 5 ml 0.5 M mannitol solution, and the content of alkali cations on the filter was measured by flame photometry. Measurements and calculations of electric potential ($\Delta\psi$) in EDTA/Tris treated cells, using the tetraphenylphosphonium cation (TPP⁺) and a TPP⁺-selective electrode, were performed as described previously [8].

3. Results and discussion

Na⁺-loaded cells diluted into Na⁺ medium, containing K⁺ and respiratory substrates, showed fast respiration-driven extrusion of Na⁺ and uptake of K⁺ (Fig. 1A), as described earlier [8]. However, neither significant Li⁺ extrusion nor K⁺ uptake were observed in Li⁺-loaded cells diluted into Li⁺ medium that also contained K⁺ and respiratory substrates (Fig. 1B). To check possible toxic effects of Li⁺, we measured both the rate of respiration and the electric potential across the cell membrane. No significant difference in respiratory rates of Na⁺- and Li⁺-loaded cells was found using the Q-1/DTT substrate couple. Without added Q-1/DTT the rate of respiration was much slower in Li⁺-loaded cells indicating that they were depleted of endogenous substrates. However, after addition of Q-1/DTT the membrane potential was close to –200 mV (in the absence of K⁺ which decreases $\Delta\psi$), independently of whether the cells had been loaded with Na⁺ or Li⁺, or suspended in Na⁺- or Li⁺-containing medium.

Sodium transport could be blocked also if the Trk system, which is responsible for uphill K⁺ transport, is inhibited by Li⁺. This possibility was tested by measuring K⁺ uptake under hyperosmotic conditions when Trk is activated (see [10] for a review). The *E. coli* cells were capable of accumulating K⁺ as a response to a rapid increase of osmolarity of the medium by addition of sucrose to 0.5 M, in the presence of 100 mM Li₂SO₄ (not shown).

These data show that the sodium efflux system is incapable of extruding Li⁺. In such a case the Trk system which transports K⁺ cannot function either, despite the high $\Delta\psi$, because it is osmotically regulated [11], and since Li⁺ did not leave the cytoplasm.

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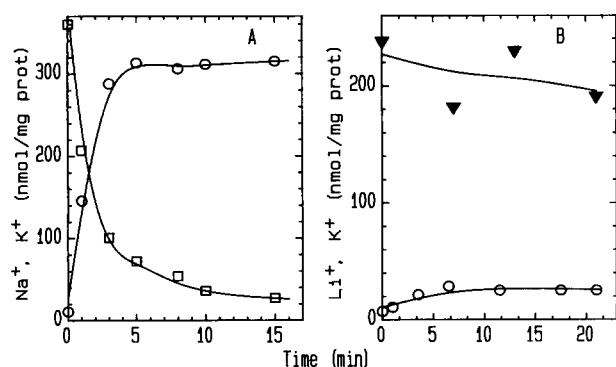


Fig. 1. Transport of alkali cations in Na^+ -loaded (A) and Li^+ -loaded (B) *E. coli* cells. At zero time the cells were diluted into medium containing 150 mM NaCl (A) or LiCl (B), 50 mM tricine NaOH (A) or LiOH (B), pH 8.5, 5 mM MgSO_4 , 10 mM KCl, 2 mM DTT and 200 μM Q-1. Squares, intracellular Na^+ ; circles, intracellular K^+ ; triangles, intracellular Li^+ .

Fast Li^+/Na^+ exchange took place initially when Li^+ -loaded cells were suspended in Na^+ -containing medium, but K^+ uptake was delayed during this process (Fig. 2). Only after nearly all inside Li^+ had been replaced by Na^+ did both uphill Na^+ efflux and K^+ influx commence (Fig. 2). This suggests that the active Na^+ efflux system is not only incapable of transporting Li^+ , but that it is actually inhibited by this cation.

We have recently shown evidence suggesting that active respiration-dependent Na^+ transport in *E. coli* is not catalysed by the electrogenic Na^+/H^+ antiporters NhaA and NhaB, but by a distinct active sodium transport system (AST) that possesses $\text{Na}^+,\text{K}^+/\text{H}^+$ antiporter activity [8]. The present data show that Li^+ is not actively transported, and that it inhibits Na^+ transport, presumably competitively. This excludes that the NhaA and NhaB antiporters are responsible for sodium

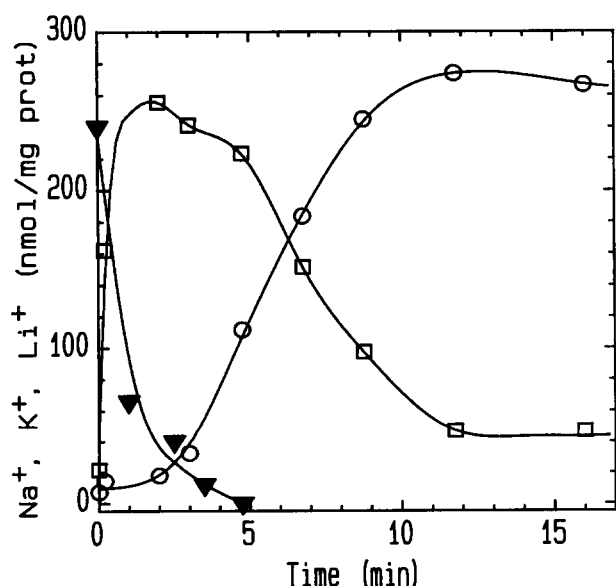


Fig. 2. Uphill transport of Na^+ and K^+ requires replacement of Li^+ with Na^+ in the cytoplasm. At zero time Li^+ -loaded cells were diluted into medium containing 150 mM NaCl, 50 mM tricine NaOH, pH 8.5, 5 mM MgSO_4 , 10 mM KCl, 2 mM DTT and 200 μM Q-1. Squares, intracellular Na^+ ; circles, intracellular K^+ ; triangles, intracellular Li^+ .

transport because, like some other Na^+ -transporting systems [12], they have been shown to operate with Li^+ (see Section 1). On the other hand, many other Na^+ transport systems are inoperative with Li^+ , such as the Na^+ -translocating NADH-quinone oxidoreductase of *Vibrio alginolyticus* [13,14]. The fast Na^+/Li^+ exchange observed here (Fig. 2) is likely due to the *nhaA* and/or *nhaB* antiporters, which then would explain why they confer resistance to Li^+ [2,3]. The difference in Li^+/Na^+ selectivity between the AST system and the Na^+/H^+ antiporters allows their distinction in vivo, and provides a good basis for future work.

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References

- [1] Padan, E., Schuldiner, S. (1994) *Biochim. Biophys. Acta* 1185, 129–151.
- [2] Padan, E., Maisler, N., Taglicht, D., Karpel, R. and Schuldiner, S. (1989) *J. Biol. Chem.* 264, 20297–20302.
- [3] Pinner, E., Kotler, Y., Padan, E. and Schuldiner, S. (1993) *J. Biol. Chem.* 268, 1729–1734.
- [4] Umeda, K., Shiota, S., Futai, M. and Tsuchiya, T. (1984) *J. Bacteriol.* 160, 812–814.
- [5] Beck, J.C. and Rosen, B.P. (1979) *Arch. Biochem. Biophys.* 194, 208–214.
- [6] Pinner, E., Padan, E. and Schuldiner, S. (1992) *J. Biol. Chem.* 267, 11064–11068.
- [7] Dibrov, P. and Taglicht, D. (1993) *FEBS Lett.* 336, 525–529.
- [8] Verkhovskaya, M.L., Verkhovsky, M.I. and Wikström, M. (1996) *Biochim. Biophys. Acta* 1273, 207–216.
- [9] Nakamura, T., Tokuda, H. and Unemoto, T. (1982) *Biochim. Biophys. Acta* 692, 389–396.
- [10] Walderhaug, M.O., Dosch, D.C. and Epstein, W. (1987) in *Ion Transport in Prokaryotes*, Rosen, B.P. and Silver, S., Eds., Academic Press, New York, 84.
- [11] Rhoads, D.B. and Epstein, W. (1978) *J. Gen. Physiol.* 72, 283.
- [12] Yazyu, H., Shiota, S., Futai, M. and Tsuchiya, T. (1985) *J. Bacteriol.* 162, 933–937.
- [13] Unemoto, T., Hayashi, M. and Hayashi, M. (1977) *J. Biochem. (Tokyo)* 82, 1389–1395.
- [14] Tokuda, H. and Unemoto, T. (1984) *J. Biol. Chem.* 259, 7785–7790.