

The Na^+/e^- stoichiometry of the Na^+ -motive NADH : quinone oxidoreductase in *Vibrio alginolyticus*

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Abstract A method is proposed to estimate the stoichiometries of primary Na^+ -pumps in intact bacterial cells. It is based on technique when the H^+/e^- stoichiometry is measured in the presence of protonophore uncoupler and in the absence of penetrating ions other than H^+ . Under these conditions, the H^+ influx discharges membrane potential generated by the Na^+ pump so the Na^+/e^- and H^+/e^- ratios become equal. Using this approach it is shown that the Na^+/e^- ratio for the Na^+ -motive NADH : quinone oxidoreductase of *Vibrio alginolyticus* is equal to 0.71 ± 0.06 . The Na^+/e^- stoichiometry appears to be ≈ 1 , provided that the contribution of the non-coupled NADH : quinone oxidoreductase, which is resistant to low HQNO concentrations, is taken into account.

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Key words: Na^+/e^- stoichiometry; Na^+ -motive NADH : quinone oxidoreductase; *Vibrio alginolyticus*

1. Introduction

The Na^+ -motive NADH : quinone oxidoreductase is shown to operate in the initial span of the respiratory chain of the marine bacterium *Vibrio alginolyticus*. This electrogenic primary Na^+ pump generates a transmembrane difference in Na^+ electrochemical potentials ($\Delta\mu_{\text{Na}^+}$) [1,2]. The $\Delta\mu_{\text{Na}^+}$ formed is then utilised to support osmotic work, i.e. uphill transport of solutes [3], mechanic work, i.e. rotation of flagellum [4], and chemical work, i.e. oxidative phosphorylation [5–7]. Genes encoded for this enzyme were recently cloned and sequenced. It was found that the corresponding operon consists of six genes showing no substantial homology with other NADH : quinone oxidoreductases [8,9]. It was also established that the Na^+ -NADH : quinone oxidoreductase contains FAD and a FeS cluster [9,10]. The functional mechanism of this enzyme still remains obscure. In particular, such a crucial parameter as the Na^+/e^- stoichiometry is unknown. In the series of experiments described below, we established this stoichiometry to be ≈ 1 .

2. Materials and methods

2.1. Bacterial strains and growth conditions

V. alginolyticus, strain 138-2 (wild type), and a mutant strain, defi-

cient in the Na^+ -motive NADH : quinone oxidoreductase, Nap1 [2] were the generous gifts of Professor Tsutomu Unemoto. The cells were grown to the middle exponential phase at 37°C in medium containing 0.5 M NaCl, 10 mM KCl, 15 mM $(\text{NH}_4)_2\text{SO}_4$, 5 mM MgSO_4 , 0.05% yeast extract, 0.4% (v/v) glycerol, 0.5 mM Na_2HPO_4 , 50 mM Tris-HCl pH 8.6.

2.2. The stoichiometry measurement

The cells were harvested at $7000 \times g$ (10 min). Then, internal K^+ was replaced by Na^+ according to the modified method of Nakamura et al. [11]. The cells were suspended in a medium containing 0.4 M NaCl, 5 mM MgSO_4 and 50 mM diethylamine pH 8.8, incubated at room temperature for 10 min and sedimented at $7000 \times g$. This procedure was repeated twice. After this, the cells were washed twice with 0.2 M Na_2SO_4 , 5 mM MgSO_4 and 20 mM HEPES-NaOH pH 7.5. Finally, the cells were washed with the incubation medium (0.2 M Na_2SO_4 , 5 mM MgSO_4 , 10 mM glycerol, 0.1 mM HEPES-NaOH pH 7.5), re-suspended in this medium (to 1.5 mg cell protein $\times \text{ml}^{-1}$) and injected into a 1.4-ml chamber equipped with a pH electrode. After saturating the mixture with argon, the chamber was closed and the pH level was adjusted to pH 7.5 with NaOH.

To measure the H^+/e^- stoichiometry, 5 nmol O_2 was added. The pH electrode was calibrated with 10 nmol additions from the argon-saturated H_2SO_4 solution. To estimate the H^+/e^- ratio, linear extrapolation was employed according to [12].

Respiration was measured by a Clark-type oxygen electrode.

Protein concentration was estimated by means of the biuret micro-method, using bovine serum albumin (Merck) as a standard.

3. Results and discussion

In 1982, Tokuda and Unemoto [1] showed that an O_2 addition to the *V. alginolyticus* cells in the presence of a protonophore gives rise to an alkalization of the medium. This effect was interpreted as the electrophoretic, protonophore-mediated H^+ uptake supported $\Delta\Psi$ which was generated by the Na^+ -NADH : quinone oxidoreductase. Such an effect can be used to measure the Na^+/e^- stoichiometry of this enzyme. In fact, the number of H^+ ions imported by the cell in the presence of protonophore must be equal to the number of Na^+ ions exported by the Na^+ -NADH : quinone oxidoreductase, provided that H^+ is the only ion discharging the Na^+ pump-generated $\Delta\Psi$. To minimize the contribution of other ion fluxes to the $\Delta\Psi$ discharge, we excluded K^+ and Cl^- from the incubation medium and replaced intercellular K^+ with Na^+ .

As one can see in Fig. 1, the O_2 pulse in the presence of penetrating cation TPP^+ entails an H^+ efflux from the *V. alginolyticus* cells which is presumably a result of operation of the H^+ -motive terminal oxidase [13]. On the other hand, the same O_2 pulse in the presence of protonophore CCCP gives rise to an opposite effect, i.e. an H^+ influx. This could be accounted for by an electrophoretic H^+ uptake caused by a discharge of the Na^+ -NADH : quinone oxidoreductase-generated $\Delta\Psi$. No pH change occurs when both TPP^+ and CCCP

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Abbreviations: $\Delta\mu_{\text{H}^+}$ and $\Delta\mu_{\text{Na}^+}$, transmembrane differences in H^+ and Na^+ electrochemical potentials, respectively; $\Delta\Psi$, transmembrane difference in electric potentials; CCCP, m-chlorocarbonylcyanide phenylhydrazone; HQNO, 2-heptyl-4-hydroxyquinoline N-oxide; TPP^+ , tetraphenyl phosphonium; TTFB, 4,5,6,7-tetra-chloro-2-trifluoromethylbenzimidazole

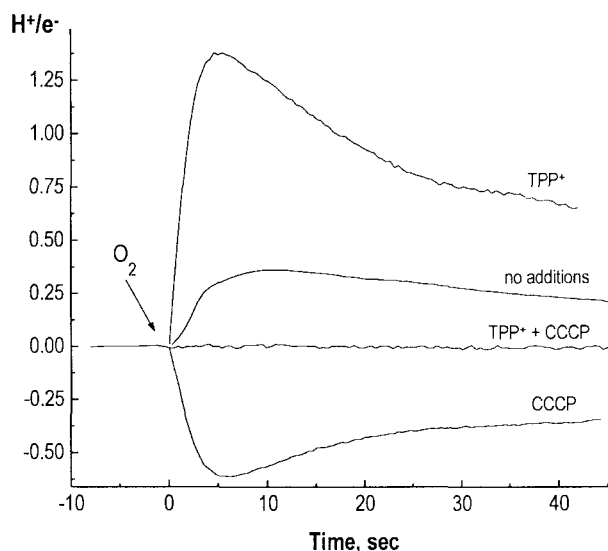


Fig. 1. The H^+/e^- stoichiometries in the *V. alginolyticus* 138-2 cells oxidizing glycerol. Additions: 5 mM TPP^+ , 15 μ M CCCP.

are added. In the absence of TPP^+ and CCCP, a very small H^+ efflux is observed.

The dependence of the H^+/e^- stoichiometry upon the CCCP concentration is shown in Fig. 2. According to these data, the maximal H^+/e^- ratio is reached at 15 μ M CCCP. This ratio provided to be equal to 0.71 ± 0.06 (mean of 8 experiments). Similar H^+/e^- stoichiometry was achieved when using TTFB as a protonophorous uncoupler (data not shown).

No pH change during O_2 pulses in the presence of CCCP is observed in a mutant strain *V. alginolyticus* lacking Na^+ -NADH:quinone oxidoreductase. On the other hand, the same mutant appears to be competent in the TPP^+ -stimulated H^+ efflux when CCCP is absent (Fig. 3A). Magnitude of the O_2 pulse-induced H^+ influx into wild-type cells in samples with CCCP is strongly decreased when lactate substitutes for glycerol as the oxidation substrate (Fig. 3B). The same figure shows that the H^+ influx is arrested by very low HQNO concentrations. Parallel measurement of the respira-

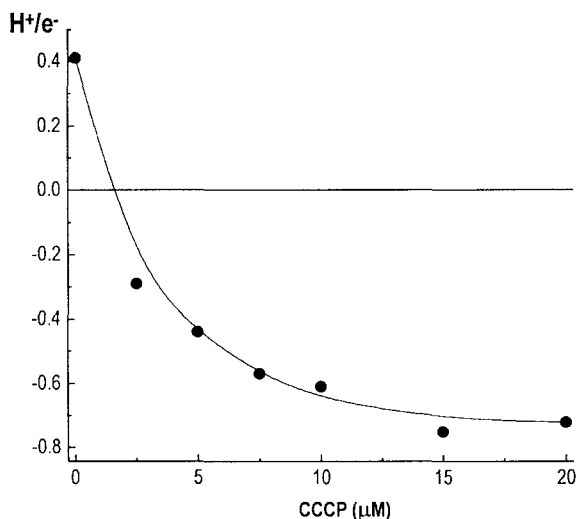


Fig. 2. The CCCP titration of measured H^+/e^- ratio in *V. alginolyticus* 138-2 cells oxidizing glycerol.

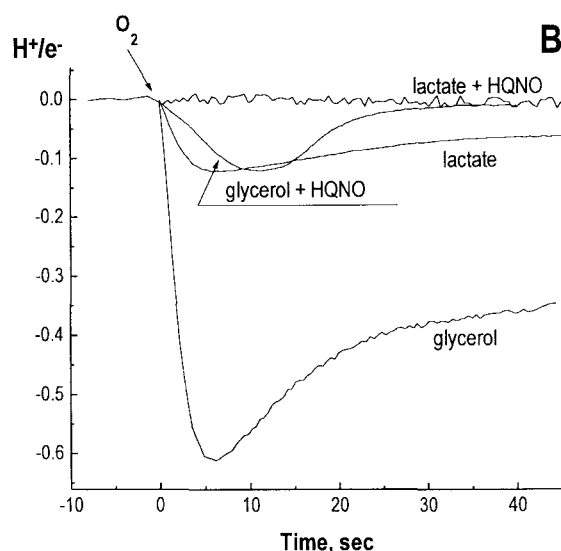
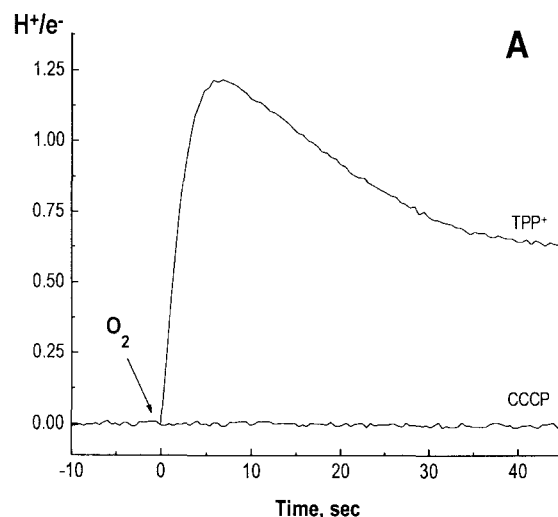


Fig. 3. (A) The H^+/e^- stoichiometries in *V. alginolyticus* mutant strain Nap1 lacking the Na^+ -NADH:quinone oxidoreductase. Substrate, glycerol. Additions: 5 mM TPP^+ , 15 μ M CCCP. (B) The H^+/e^- stoichiometries in the *V. alginolyticus* 138-2 cells oxidizing D,L-lactate or glycerol in the presence of 15 μ M CCCP. Effect of 5 μ M HQNO.

tion rate revealed that these HQNO concentrations lower the O_2 consumption by the *V. alginolyticus* cells by $\approx 70\%$ (Fig. 4). To achieve further inhibition, much higher concentrations of HQNO are required.

The above data indicate that it is the Na^+ -NADH:quinone oxidoreductase which is responsible for the CCCP-mediated H^+ influx when O_2 is added to anaerobic suspension of *V. alginolyticus* cells. This process: (i) is absent in the mutant strain lacking Na^+ -NADH:quinone oxidoreductase; (ii) is sensitive to low concentrations of HQNO specifically inhibiting this enzyme [1,14]; and (iii) cannot be effectively supported by oxidation of D-lactate (a substrate which is oxidized in this bacterium in the NAD^+ -independent fashion).

At least two facts suggest that, under the conditions employed, the bacterial membrane has a low permeability to ions other than H^+ . (1) Without TPP^+ and CCCP, the O_2 pulse produces only a small pH change. (2) The rate of decay of the

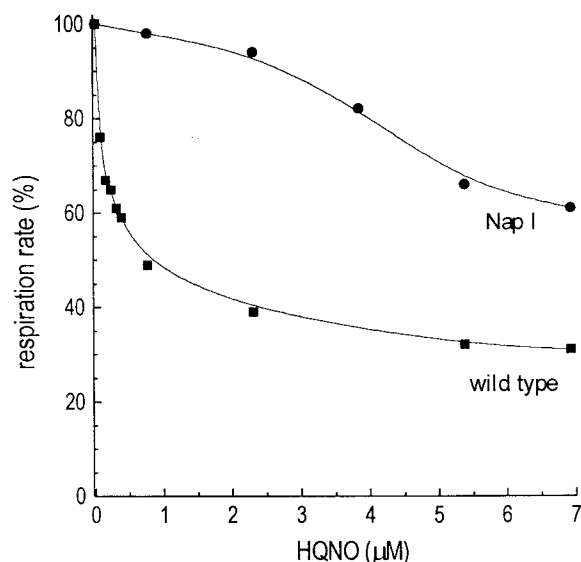


Fig. 4. The HQNO respiration inhibition in wild-type *V. alginolyticus* 138-2 cells and in the NapI mutant strain. Substrate, glycerol.

alkaline pH shift after the O_2 pulse in the presence of CCCP is much slower than the rate of formation of this pH change. Since in these experiments we used saturating CCCP concentrations, the observation in (2) indicates that penetration of ions other than H^+ are rate-limiting for the alkaline pH shift decay.

Thus, in samples with high CCCP concentrations, the membrane permeability for H^+ is much higher than for other ions so amount of the Na^+ ions extruded by the Na^+ -NADH:quinone oxidoreductase should be equal to that of the H^+ ions taken up to discharge the Na^+ efflux-generated $\Delta\Psi$. In other words, experimentally measured Na^+/e^- ratio is equal to the Na^+/e^- ratio, i.e. 0.71. In fact, this value should be lower than the true Na^+/e^- stoichiometry of the Na^+ -NADH:quinone oxidoreductase since, besides the Na^+ -NADH:quinone oxidoreductase sensitive to low [HQNO], some O_2 consumption is due to operation of the non-coupled NADH:quinone oxidoreductase which is much more resistant to HQNO [14]. Contribution of this enzyme and NAD $^+$ -

independent primary dehydrogenases, according to the HQNO titration of respiration of *V. alginolyticus* cells, is $\approx 30\%$ of the total O_2 consumption (Fig. 4). This means that the true Na^+/e^- stoichiometry of the Na^+ -NADH:quinone oxidoreductase is ≈ 1 . This value is lower than the H^+/e^- stoichiometry for the *Escherichia coli* H^+ -motive NADH:quinone oxidoreductase which we estimated in our previous work as equal to at least 1.5 [15]. Nevertheless, the measured Na^+/e^- stoichiometry is in line with Rich's schemes of the Na^+ -motive NADH:quinone oxidoreductase mechanism [9,16].

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