# HQNO-sensitive NADH:DCIP Oxidoreductase of a Pathogenic *Bacillus cereus* Causing β-Hemolysis

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Membranes prepared from *Bacillus cereus* KCTC 3674, grown aerobically on a complex medium, oxidized NADH exclusively, whereas deamino-NADH was little oxidized. The respiratory chain-linked NADH oxidase system exhibited an apparent  $K_m$  value of about 65  $\mu$ M for NADH. Interestingly, the activity of NADH:DCIP oxidoreductase on NADH oxidase system was decreased remarkably by Na or K and its optimal pH was 5.5. The activity of NADH:DCIP oxidoreductase was very resistant to the respiratory chain inhibitors such as rotenone, capsaicin, and AgNO3, whereas it was inhibited by about 40% with 40  $\mu$ M 2-heptyl-4-hydroxyquinoline-N-oxide (HQNO). From the results, we suggest the possibility that the aerobic respiratory chain-linked NADH oxidase system of *B. cereus* KCTC 3674 may possess the HQNO-sensitive NADH:DCIP oxidoreductase lacking an energy coupling site.

Key words - Bacillus cereus KCTC 3674, aerobic respiratory chain, NADH:DCIP oxidoreductase

### Introduction

Although the NADH dehydrogenases (NDH) in the bacterial respiratory chains are referred to as NADH:ubiquinone oxidoreductase, many bacteria have been known to possess quinones other than ubiquinone[2]. Thus, the name NADH:quinone oxidoreductase in bacteria seems more appropriate than NADH:ubiquinone oxidoreductase.

Three types of NADH:quinone oxidoreductases in the respiratory chain of bacteria have been reported[14]. They are the H<sup>+</sup>-translocating NADH:quinone oxidoreductase (designated NDH-1), the Na<sup>†</sup>-translocating NADH:quinone oxidoreductase (designated Na<sup>+</sup>-NDH), and the NADH: quinone oxidoreductase lacking an energy coupling site (designated NDH-2). In general, NDH-1 or Na<sup>+</sup>-NDH reacts with deamino-NADH as well as with NADH, shows high affinities for NADH, and possesses an energy coupling site[4,8]. In contrast, NDH-2 reacts very little with deamino-NADH, but with NADH without any energy coupling site, and shows low affinities for NADH[4,7,8]. Escherichia coli[8], Vibrio alginolyticus[9,10], and Thermus thermophilus HB-8[13] are known to possess two different types of NADH:quinone oxidoreductases. Zymomonas mobilis is known to possess only NADH:quinone oxidoreductase lacking the energy coupling site[7].

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Respiratory chain inhibitors have proved to be a useful tool for probing the mechanism of electron transfer and proton or sodium translocation in the respiratory chain. Generally, NDH-1 is inhibited by the respiratory inhibitors rotenone and capsaicin, whereas NDH-2 is only slightly inhibited by these inhibitors[13,14]. Interestingly, Na<sup>+</sup>-NDH is known to be very resistant to rotenone and capsaicin[14], but highly sensitive to 2-heptyl-4-hydroxyquino-line-*N*-oxide[10] and AgNO<sub>3</sub>[1].

Takao Yagi showed that the respiratory inhibitor capsaicin inhibited H<sup>+</sup>-translocating NADH:quinone oxidoreductases, and did not inhibit the NADH oxidase and NADH:quinone oxidoreductase of *Bacillus subtilis*[12]. To date, little detailed study has been done on the enzymatic and energetic properties of the aerobic respiratory chain-linked NADH oxidase system in the genus *Bacillus*. *B. cereus* produces several extracellular hemolysins, which are considered potential factors of virulence of the opportunistic pathogen[11]. In this paper, we report the results for enzymatic properties of the NADH:DCIP oxidoreductase on the aerobic respiratory chain-linked NADH oxidase system of *B. cereus* KCTC 3674.

### Materials and Methods

#### Bacterial strain and conditions

The bacterial strain used in this study was *B. cereus* KCTC 3674[3,4,5]. The bacterium was grown aerobically at

37° C in liquid media containing 0.5% polypeptone and 0.5% yeast extract in 50 mM Tris-HCl buffer (pH 7.5). Preculture grown overnight was used to inoculate the main culture to give a turbidity of approximately 0.03.

# Preparation of membranes from *B. cereus* KCTC 3674

For the preparation of membranes from B. cereus KCTC 3674, the protoplast formation was carried out at 37°C. Cells harvested in logarithmic growth phase were suspended in 50 mM potassium phosphate (pH 7.5) containing 5 mM EDTA at a concentration of 1 g (wet weight) per 80 ml. Lysozyme was added at a concentration of 300 µ g/ml (freshly prepared). The mixture was incubated for 30 min, after which the protoplasts were harvested by centrifugation at 14,000×g for 30 min. Protoplasts were washed once in 50 mM potassium phosphate (pH 7.5) containing 5 mM EDTA, and centrifuged at 14,000×g for 30 min at 4°C. Washed protoplasts were resuspended in 50 mM potassium phosphate (pH 7.5) containing 5 mM EDTA to give a concentration of 20 ml/g, and protoplast suspensions were passed through a French pressure cell twice at 25,000 psi. Unbroken cells and cell debris were removed by centrifugation at 10,000×g for 10 min at 4°C, and the supernatant was centrifuged at 120,000×g for 2 h at 4°C to sediment the membrane fraction. A membrane pellet was washed in 50 mM potassium phosphate (pH 7.5) containing 5 mM EDTA. After sedimentation at 120,000×g for

2 h at 4°C, inverted membrane vesicles were rewashed in 50 mM potassium phosphate (pH 7.5) containing 10% glycerol. Membranes were resuspened in 50 mM potassium phosphate (pH 7.5) containing 10% glycerol at a protein concentration of about 25 mg/ml, and stored at -80°C.

### Measurement of enzyme activities

The activity of the NADH oxidase was measured at  $37^{\circ}\text{C}$  from a decrease in  $A_{340}$  by using varian Cary 3E spectrophotometer. The assay mixture of NADH oxidase contained  $125~\mu\text{M}$  NADH or deamino-NADH in 2 ml of 50~mM Tris-HCl (pH 8.5) containing 0.1~M KCl. The assay was started by addition of  $200~\mu\text{g}$  of membrane protein. Activity was calculated by using a millimolar extinction coefficient of 6.22. The assay mixture for NADH:DCIP oxidoreductase contained  $50~\mu\text{g}$  of membrane protein, 30~mM KCN, and  $100~\mu\text{M}$  DCIP (2.6-dichlorophenol-indophenol) in 2~ml of 50~mM MES-KOH (pH 5.5). The reaction of the NADH:DCIP oxidoreductase was started by the addition of

125  $\mu$ M NADH. The NADH:DCIP oxidoreductase activity was measured at 600 nm, and was calculated by using a millimolar extinction coefficient of 16.5.

#### Protein determination

Protein was measured by Bio-Rad protein assay, based on the method of Bradford. Bovine serum albumin was used as a standard.

#### Results

# Enzymatic properties of the membrane-bound NADH oxidase

The effects of salts and pH on the NADH oxidase activity were examined with membranes prepared from *B. cereus* KCTC 3674. As shown in Fig. 1, the membranes oxidized NADH (♠), but very little deamino-NADH (○) as a substrate. The rate of NADH oxidation was slightly activated by monovalent cations including Na<sup>+</sup> and K<sup>+</sup>, but was little activated by Li<sup>+</sup> (data not shown). The maximum activity of NADH oxidase was obtained in the presence of 0.1 M NaCl or KCl. The optimal pH for NADH oxidation in the presence of 0.1 M NaCl was 8.5 (data not shown).

The respiratory chain-linked NADH oxidase system exhibited an apparent  $K_m$  value of about 65  $\mu$ M for NADH (Fig. 2). In order to examine the enzymatic properties of NADH:quinone oxidoreductase of *B. cereus* KCTC 3674, an artificial electron acceptor 2,6-dichlorophenol-indophenol (DCIP) was introduced.

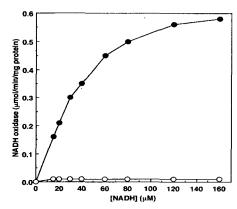
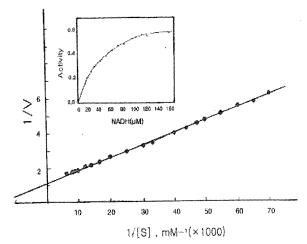


Fig. 1. Ability of NADH oxidase to oxidize NADH or deamino-NADH. The activities of NADH and deamino-NADH oxidases were measured in 2 ml of 50 mM Tris-HCl (pH 8.5) containing 0.1 M KCl and different concentrations of NADH (•) or deamino-NADH (•). The assay was started by addition of membrane suspensions containing about 200 μg of protein at 37°C.



Organisms	Km values (μM)
B. cereus KCTC 3674	65
E. coli	5.3~6.6, 48
V. alginolyticus	5~8, 25~30
Nap 1	27~35
Z. mobilis	66

Fig. 2. A Lineweaver-Berk plot of the NADH oxidase activity. The K<sub>m</sub> values of *B. cereus* KCTC 3674, *V. alginolyticus*, and Nap 1 which is a Na<sup>+</sup> pump-defective mutant of *V. alginolyticus*[9] were measured in this study.

# The effects of salts, pH, and respiratory inhibitors on the activity of the NADH:DCIP oxidoreductase

The activity of NADH:DCIP oxidoreductase was remarkably reduced by Na<sup>+</sup> and K<sup>+</sup> (Fig. 3A), and was, also, prominently reduced with an increase in the value of pH (Fig. 3B).

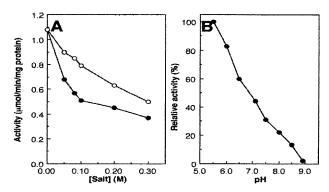


Fig. 3. Effects of salts and pH on the activity of NADH:DCIP oxidoreductase. (A) The assay mixture of NADH:DCIP oxidoreductase (2 ml) contained 50 mM MES-KOH (pH 5.5), 50 μg of membrane protein, 30 mM KCN, 100 μM DCIP, and various concentrations of NaCl (●), or KCl (○), (B) The activity of NADH:DCIP oxidoreductase was determined at various pHs. Buffers used at 50 mM was MES-KOH (pH 6 to 6.5) and Tris-HCl (pH 7.1 to 8.9). All assays were started by addition of 125 μM NADH at 37°C.

The respiratory inhibitor 2-heptyl-4-hydroxyquinoline-N-oxide(HQNO), which acts at the b cytochromes and also inhibits the Na $^{+}$ -translocating NADH:quinone oxidoreductase, inhibited the activity of NADH:DCIP oxidoreductase by about 40% at a concentration of 100  $\mu$ M (Fig. 4A). Rotenone and capsaicin, which inhibit the energy-transducing NADH: quinone oxidoreductase, inhibited the activity by about 10% at a concentration of 100  $\mu$ M and 300  $\mu$ M, respectively (Fig. 4B and 4C).

# Effect of Ag<sup>+</sup> on the enzyme activities of NADH oxidase system

AgNO<sub>3</sub> is known to inhibit Na<sup>+</sup>-translocating NADH: quinone oxidoreductase[1]. As shown in Fig. 5, the membrane-bound NADH oxidase activity was highly sensitive

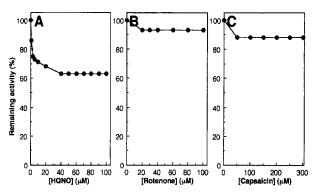


Fig. 4. Effects of respiratory chain inhibitors on the NADH: DCIP oxidoreductase. The activity of NADH:DCIP oxidoreductase was measured with various concentrations of HQNO (A), rotenone (B), and capsaicin (C). The assay mixture of NADH:DCIP oxidoreductase contained 50 μg of membrane protein, 30 mM KCN, 100 μM DCIP in 2 ml of 50 mM MES-KOH (pH 5.5). All assays were started by addition of 125 μM NADH.

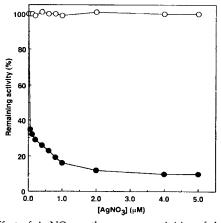


Fig. 5. Effect of AgNO₃ on the enzyme activities of the NADH oxidase and NADH:DCIP oxidoreductase. NADH oxidase (●) and NADH:DCIP oxidoreductase (○).

to  $Ag^{\dagger}$ . In contrast, the activity of NADH:DCIP oxidor-eductase was not affected by  $Ag^{\dagger}$  at all.

### Discussion

The results of this study show the possibility that the aerobic respiratory chain-linked NADH oxidase system of *B. cereus* KCTC 3674 may possess the enzymatic system lacking an energy coupling site.

Membranes prepared from B. cereus KCTC 3674 oxidized NADH, but very little deamino-NADH as a substrate, and exhibited an apparent K<sub>m</sub> value of approximately 65 µM for NADH. Generally, NADH:quinone oxidoreductase lacking an energy coupling site (NDH-2) oxidizes little deamino-NADH, and shows low affinities for NADH[7,8] (Fig. 2). The activities of NADH oxidase and NADH:ubiquinone-1 oxidoreductase of B. subtilis membranes are little affected by capsaicin[12]. In order to examine the enzymatic properties of NADH:quinone oxidoreductase of B. cereus KCTC 3674, an artificial electron acceptor 2,6-dichlorophenol-indophenol (DCIP) was introduced in this study. The activity of NADH:DCIP oxidoreductase of B. cereus KCTC 3674 was little affected by capsaicin and rotenone. AgNO3 and HQNO are known to be the potent inhibitors of Na+translocating NADH:quinone oxidoreductase[1,10]. Interestingly, the NADH:DCIP oxidoreductase of B. cereus KCTC 3674 was sensitive to HQNO, whereas was not affected by Ag+ at all. The membrane-bound NADH oxidase was highly sensitive to Ag<sup>+</sup>. Thus, it is suggested that AgNO3 does not inhibit the NADH:quinone oxidoreductase of B. cereus KCTC 3674, but inhibit the quinol oxidase.

B. cereus KCTC 3674 vigorously excretes proteases into the extracellular environment[3,6]. Thus, it is very difficult to prepare the functional inverted membrane vesicles for energy measurement experiment. Due to the lack of a functional in vitro system, little detailed study has been done on the energetic experiments in the genus Bacillus. To further examine whether the aerobic respiratory chain of B. cereus KCTC 3674 generates energy or not, we will try the preparation of inverted membrane vesicles for a functional in vitro energy measurement system.

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### References

- Asano, M., M. Hayashi, T. Unemoto and H. Tokuda. 1985. Ag<sup>+</sup>-sensitive NADH dehydrogenase in the Na<sup>+</sup>-motive respiratory chain of the marine bacterium Vibrio alginolyticus. Agric. Biol. Chem. (Biosci. Biotechnol. Biochem.) 49, 2813-2817.
- Collins, M. D. and D. Jones. 1981. Distribution of isoprenoid quinone structural types in bacteria and their taxonomic implications. *Microbiol. Rev.* 45, 316-354.
- 3. Kim, S. S., Y. J. Kim and I. Rhee. 2001. Purification and characterization of anovel extracellular protease from *Bacillus cereus* KCTC 3674. *Arch. Microbiol.* 175, 458-461.
- Kim, Y. J., S. Mizushima, and H. Tokuda. 1991. Fluorescence quenching studies on the characterization of energy generated at the NADH:quinone oxidoreductase and quinol oxidase segments of marine bacteria. J. Biochem. (Tokyo) 109, 616-621.
- Kim, S. S., Y. Park, J. Lee, J. Yoon, Y. Shin, I. Rhee and Y. J. Kim. 1998. Taxonomic studies of the beta hemolysis-causing pathogen *Bacillus cereus* isolated from sea water. J. Microbiol. Biotech. 8, 67-73.
- Kim, S. S., Y. Park, I. Rhee and Y. J. Kim. 2000. Influence of temperature, oxygen, m-chlorophenylhydrazone, cerulenin, and quinacrine on the production of extracellular proteases in *Bacillus cereus*. J. Microbiol. Biotech. 10, 103-106.
- Kim, Y. J., K. Song and S. Rhee. 1995. A novel aerobic respiratory chain-linked NADH oxidase system in Zymomonas mobilis. J. Bacteriol. 177, 5176-5178.
- 8. Matsushita K, T. Ohnishi and R. Kaback. 1987. NADH:ubiquinone oxidoreductase of the *Escherichia coli* aerobic respiratory chain. *Biochemistry* **26**, 7732-7737.
- Tokuda, H. 1983. Isolation of Vibrio alginolyticus mutants defective in the respiration-coupled Na<sup>+</sup> pump. Biochem. Biophys. Res. Commun. 114, 113-118.
- Tokuda, H and T. Unemoto. 1984. Na<sup>+</sup> is translocated at NADH:quinone oxidoreductase segment in the respiratory chain of *Vibrio alginolyticus*. J. Biol. Chem. 259, 7785-7790.
- Turnbull, P. C. B. 1986. Pharmacology of Bacterial Toxins (Dorner F and Drews J, eds). Pergamon Press, Oxford. 397-448.
- 12. Yagi T. 1990. Inhibition by capsaicin of NADH-quinone oxidoreductases is correlated with the presence of energy-coupling site 1 in various organisms. *Arch. Biochem. Biophys.* **281(2)**, 305-311.
- Yagi T, K. Hoh-nami and T. Ohnishi. 1998b. Purification and characterization of two types of NADH-quinone reductase from *Thermus thermophilus* HB-8. *Biochemistry* 27, 2008-2013.
- Yagi T, T. Yano, S. D. Bernardo, A. Matsuno-Yagi. 1998a. Procaryotic complex I (NDH-1), an overview. *Biochim. Biophys. Acta.* 1364, 125-133.

## 초록: Beta hemolysis 유발 병원균 Bacillus cereus의 HQNO-sensitive NADH: DCIP oxidoreductase

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호기적으로 자란 Bacillus cereus KCTC 3674로 부터 조제된 막은 NADH만을 산화하고, deamino-NADH는 거의 산화하지 않았다. 호흡쇄와 연계된 NADH oxidase계는 K<sub>m</sub> 값이 약 65 μM 이였다. NADH:DCIP oxidoreductase의 활성은 Na<sup>+</sup> 또는 K<sup>+</sup>에 의해 감소되었다. 그 최적 pH는 5.5 였다. NADH:DCIP oxidoreductase의 활성은 rotenone, capsaicin, AgNO<sub>3</sub>와 같은 호흡저해제에는 매우 저항적이였지만, 40 μM HQNO (2-heptyl-4-hydroxyquinoline-N-oxide) 존재하에서는 약 40% 저해되었다. 이들 결과로 부터, Bacillus cereus KCTC 3674의 호기적호흡쇄와 연계된 NADH oxidase계는 energy coupling site가 결여된 HQNO-sensitive NADH:DCIP oxidoreductase를 소유하고 있는 것으로 추정된다.