

## BBA Report

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### EVIDENCE FOR A NATURALLY OCCURRING ATPase-INHIBITOR IN *ESCHERICHIA COLI*

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#### Summary

Stimulation of the *Escherichia coli* ATPase activity by urea and trypsin shows that there is a latent ATPase activity in particles and in a crude coupling factor of *E. coli*. Moreover, crude coupling factor, completely dissociated by treatment with 7 M urea, can inhibit the ATPase activity of the crude coupling factor. It is suggested that the latency of the ATPase activity of the coupling factor is due to the presence of a protein, the ATPase-inhibitor.

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Both in mitochondria [1] and chloroplasts [2] the presence of a naturally occurring ATPase-inhibitor has been demonstrated. Ernster et al. [3] and Van de Stadt et al. [4] have presented evidence that the mitochondrial inhibitor plays a regulatory function in oxidative phosphorylation. A similar regulatory function has been postulated for the chloroplast inhibitor [5]. If such an inhibitor is an essential component of the system catalysing oxidative phosphorylation, it may be expected that it is also present in bacteria.

Several bacteria are known that possess a latent ATPase activity (ref. 6, see ref. 7 for review) which can be unmasked by trypsin. It therefore seems logical to assume that this is due to a naturally occurring bacterial ATPase inhibitor analogous to that in mitochondria and chloroplasts. The present study was undertaken to put this assumption to a test.

Subbacterial particles and a crude coupling factor from *Escherichia coli* A 428 were isolated as described previously [8]. ATPase activity was assayed by measurement of inorganic phosphate [9] liberated from ATP. Protein was determined by the method of Lowry et al. [10], with bovine serum albumin as standard.

As can be seen in Fig. 1, high concentrations of urea completely inhibit

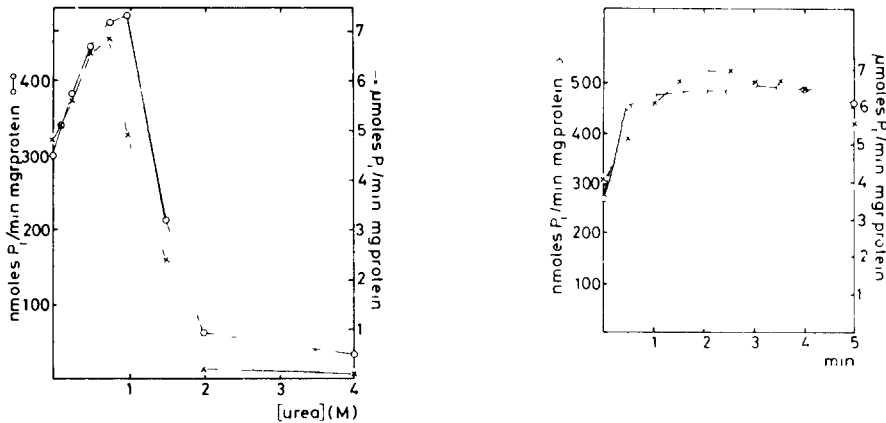


Fig 1 Effect of urea on the ATPase activity of *E. coli* particles and crude coupling factor. The crude coupling factor (16  $\mu\text{g}$ ) and particles (200  $\mu\text{g}$ ) were incubated for 5 min at  $37^\circ\text{C}$  with the indicated amounts of urea in 50 mM Tris-HCl and 2.5 mM  $\text{MgCl}_2$  (pH 7.8), in a total volume of 1 ml. After this incubation the ATPase activity was tested by addition of 5 mM ATP. X—X, crude coupling factor, O—O, particles.

Fig 2 Stimulation of the ATPase activity of *E. coli* particles and crude coupling factor by trypsin. Crude coupling factor (16  $\mu\text{g}$ ) and particles (200  $\mu\text{g}$ ) were incubated with 250  $\mu\text{g}$  TPCK-trypsin (Worthington) in 100 mM Tris-HCl and 5 mM  $\text{MgCl}_2$  (pH 7.8) in a total volume of 0.5 ml for the indicated time. Then the TPCK-trypsin was inhibited by adding 800  $\mu\text{g}$  trypsin-inhibitor (Soybean, Worthington) and the volume brought to 1 ml with water. The ATPase activity was tested in 1 ml by addition of 5 mM ATP. X—X, crude coupling factor, O—O, particles.

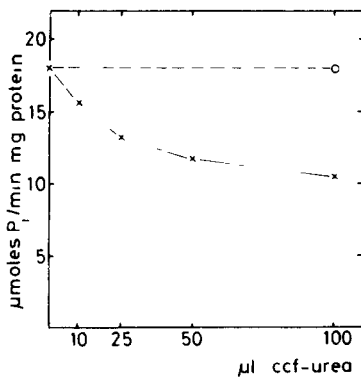


Fig 3 Inhibition of the ATPase activity of *E. coli* crude coupling factor by a completely dissociated crude coupling factor. The crude coupling factor was purified over DEAE-Sephadex A (50 to 18 units/mg protein). 'Crude coupling factor-urea' (ccf-urea). Crude coupling factor (1 mg/ml, 4–5 units/mg) was treated for 5 min with 7 M urea, then dialysed at  $4^\circ\text{C}$  for  $3 \times 1$  h against 20 mM Tris-HCl (pH 7.5) to remove most of the urea. X—X, the DEAE-purified crude coupling factor was incubated with the indicated amounts of 'ccf-urea' in 100 mM Tris-HCl, 5 mM  $\text{MgCl}_2$  (pH 7.8) in a final volume of 0.5 ml. After 10 min the volume was brought to 1 ml with  $\text{H}_2\text{O}$  and ATPase activity was tested. O—O, 100  $\mu\text{l}$  'crude coupling factor-urea' was pre-treated with TPCK-trypsin (250  $\mu\text{g}$ ) (Worthington). After 5 min the reaction was stopped by adding trypsin-inhibitor. The inhibitory effect of this 100  $\mu\text{l}$  was tested.

the ATPase activity of both membrane-bound and solubilized ATPase. This inhibition is irreversible and probably due to splitting of the ATPase into subunits. However, at lower concentrations, where the ATPase is not yet degraded, a stimulation by urea of the ATPase activity is found. A similar stimulation is illustrated in Fig 2, which shows the trypsin activation of the solubilized and membrane-bound ATPase. Apparently, in both cases there is a latent ATPase activity present. Through the proteolytic digestion of trypsin this activity is unmasked, which indicates that a protein is involved in the ATPase inhibition.

Nelson et al. [2] have shown that a 7 M urea extract of chloroplast ATPase ( $CF_1$ ) inhibits the activated  $CF_1$  and that this inhibition is due to the  $\epsilon$  subunit of the  $CF_1$ , the ATPase-inhibitor. Fig.3 shows a similar experiment with a crude solubilized ATPase from *E. coli*.

It can be seen that the ATPase activity of the coupling factor is inhibited by 40%. If we activate the crude coupling factor with trypsin, as in Fig.2, the extra ATPase activity was inhibited 80% by 100  $\mu$ l of crude coupling factor-treated with urea as described in the legend to Fig.3.

Further experiments have indicated that the interaction of the *E. coli* ATPase with the inhibitor is different from that in mitochondria. Passage of *E. coli* particles over a Sephadex G-50 column, according to Racker and Horstmann [11], which depletes submitochondrial particles of inhibitor, has no effect on the ATPase activity of *E. coli* particles. Also, sonication of *E. coli* cells in a medium containing EDTA (2 mM) at high pH (9.2) gives no enhancement of the ATPase activity.

These results indicate that the interaction of the *E. coli* ATPase with its postulated inhibitor resembles more the interaction of  $CF_1$  and its inhibitor than that of the mitochondrial ATPase. Pyridine treatment of the purified *E. coli* ATPase according to Nelson et al. [2] resulted in isolation of a fraction enriched in inhibitory activity and containing a higher proportion of a component of molecular weight 12 000, as determined by gel electrophoresis. Work is in progress to isolate a pure *E. coli* ATPase inhibitor.

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