

BBA 72661

## The locus of nucleotide specificity in the reaction mechanism of $(\text{Na}^+ + \text{K}^+)$ -ATPase determined with ATP and GTP as substrates \*

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(Received January 31st, 1985)

Key words:  $(\text{Na}^+ + \text{K}^+)$ -ATPase; Nucleotide specificity; Structure transition; Reaction mechanism; Fluorescence; (Rabbit kidney)

(1) ATP and GTP have been compared as substrates for  $(\text{Na}^+ + \text{K}^+)$ -ATPase in  $\text{Na}^+$ -activated hydrolysis,  $\text{Na}^+$ -activated phosphorylation, and the  $\text{E}_2\text{K} \rightarrow \text{E}_1\text{K}$  transition. (2) Without added  $\text{K}^+$  the optimal  $\text{Na}^+$ -activated hydrolysis rates in imidazole-HCl (pH 7.2) are equal, but are reached at different  $\text{Na}^+$  concentrations: 80 mM  $\text{Na}^+$  for GTP, 300 mM  $\text{Na}^+$  for ATP. The affinities of the substrates for the enzyme are widely different:  $K_m$  for ATP 0.6  $\mu\text{M}$ , for GTP 147  $\mu\text{M}$ . The Mg-complexed nucleotides antagonize activation as well as inhibition by  $\text{Na}^+$ , depending on the affinity and concentration of the substrate. (3) The optimal 3-s phosphorylation levels in imidazole-HCl (pH 7.0) are equally high for the two substrates (3.6 nmol/mg protein). The  $K_m$  value for ATP is 0.1–0.2  $\mu\text{M}$  and for GTP it ranges from 50 to 170  $\mu\text{M}$ , depending on the  $\text{Na}^+$  concentration. The affinity of  $\text{Na}^+$  for the enzyme in phosphorylation is lower with the lower affinity substrate:  $K_m(\text{Na}^+)$  is 1.1 mM with ATP and 3.6 mM with GTP. The GTP-phosphorylated intermediate exists, like the ATP-phosphorylated intermediate, in the  $\text{E}_2\text{P}$  conformation. (4) Addition of  $\text{K}^+$  increases the optimal hydrolytic activity 30-fold for ATP (at 100 mM  $\text{Na}^+ + 10$  mM  $\text{K}^+$ ) and 2-fold for GTP (at 100 mM  $\text{Na}^+ + 0.16$  mM  $\text{K}^+$ ).  $\text{K}^+$  greatly increases the  $K_m$  values for both substrates (to 430  $\mu\text{M}$  for ATP and 320  $\mu\text{M}$  for GTP). Above 0.16 mM  $\text{K}^+$  inhibits GTP hydrolysis. (5) GTP does not reverse the quenching effect of  $\text{K}^+$  on the fluorescence of the 5-iodoacetamidofluorescein-labeled enzyme. ATP fully reverses this effect, which represents the transition from  $\text{E}_1\text{K}$  to  $\text{E}_2\text{K}$ . Hence GTP is unable to drive the  $\text{E}_2\text{K} \rightarrow \text{E}_1\text{K}$  transition.

### Introduction

Any enzyme has a certain degree of substrate specificity, some like cytochrome *c* oxidase very

narrow, others very broad.  $(\text{Na}^+ + \text{K}^+)$ -ATPase is an intermediate example, which can hydrolyse ATP, CTP, ITP, GTP, UTP and TTP with decreasing activity in that order, down to 0.5% for TTP as compared to ATP [1–3]. A relation has been sought between the increase in the dissociation constant of nucleotide binding to the native enzyme (from 0.2 to 3000  $\mu\text{M}$ ) and structural features of the substrate [3]. If this were the only cause, a sufficiently high concentration of nucleotide should give the same maximal activity, provided that the reaction steps following the binding

\* This article is No. 57 in the series Studies on  $(\text{Na}^+ + \text{K}^+)$ -activated ATPase.

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Abbreviations: 5-IAF, 5-iodoacetamidofluorescein; CDTA, *trans*-1,2-diaminocyclohexane tetraacetic acid.

step would display the same activity for all substrates. This is evidently not the case [4].

The reaction mechanism of  $(\text{Na}^+ + \text{K}^+)\text{-ATPase}$  comprises five partial reactions: (1) binding of the nucleotide to the high-affinity phosphorylating site, (2)  $(\text{Mg}^{2+} + \text{Na}^+)\text{-dependent}$  phosphorylation to an ADP-sensitive,  $\text{K}^+\text{-insensitive}$  phosphoenzyme  $\text{E}_1\text{P}$ , (3) conformational change to a  $\text{K}^+\text{-sensitive}$ , ADP-insensitive phosphoenzyme  $\text{E}_2\text{P}$ , (4)  $\text{K}^+\text{-stimulated}$  dephosphorylation of the phosphoenzyme resulting in the formation of  $\text{E}_2\text{K}$ , (5) conformational change of  $\text{E}_2\text{K}$  to  $\text{E}_1\text{K}$ , accelerated by low-affinity non-phosphorylating nucleotide binding to the enzyme, and exchange of  $\text{K}^+$  for  $\text{Na}^+$  [5].

We decided to determine which step is responsible for the substrate specificity of the enzyme and whether it involves a change in  $K_m$  only or also a change in  $V_{\max}$ . In first instance we have determined the  $(\text{Na}^+ + \text{K}^+)\text{-ATPase}$  activity, which covers all five steps, and the  $\text{Na}^+\text{-ATPase}$  activity which covers steps 1–4 (step 4 being the spontaneous or  $\text{Na}^+\text{-activated}$  dephosphorylation of  $\text{E}_2\text{P}$ ). Subsequently we have narrowed down the localization of the responsible step by more specific assays:  $\text{Na}^+\text{-activated}$  phosphorylation (step 2),  $\text{K}^+\text{-activated}$  dephosphorylation (step 4), and monitoring the change in fluorescence accompanying the nucleotide induced  $\text{E}_2\text{K} \rightarrow \text{E}_1\text{K}$  transition (step 5), using covalently bound fluorescent probes that do not affect overall activity [6,7].

In the present study we have compared GTP with ATP. GTP, which differs from ATP in having the amino group in the 2-position of the purine nucleus rather than in the 6-position, has previously been found to be hydrolyzed at 0.6–8% of the rate for ATP, depending on the enzyme source [1–3].

## Materials and Methods

**Preparation of the enzyme.**  $(\text{Na}^+ + \text{K}^+)\text{-ATPase}$  is prepared from rabbit kidney outer medulla using zonal centrifugation of the sodium dodecylsulfate treated microsomal fraction as described by Jørgensen [8]. Removal of contaminating ATP, subsequent washing and storage of the resulting preparation follows the method of Schoot et al. [9], except for the composition of the storage medium,

which contains 50 mM imidazole-HCl (pH 7.0), 0.25 M sucrose and no CDTA. The specific activity of the enzyme used in these experiments is 1000–1200  $\mu\text{mol P}_i/\text{mg protein per h}$ . Protein is determined by means of the trichloroacetic acid precipitation modification of the Lowry method against bovine serum albumin as standard [8].

**Enzyme assays.**  $\text{Na}^+\text{-stimulated ATPase}$  activity is measured at 37°C in terms of the liberation of  $^{32}\text{P}_i$  from  $[\gamma\text{-}^{32}\text{P}]\text{ATP}$ , present in micromolar or millimolar concentrations in a medium containing 0.01–500 mM NaCl, 50 mM imidazole-HCl or Tris-HCl (pH 7.2) and 0.15 mM  $\text{Mg}^{2+}$  in excess of the ATP concentration in a total volume of 0.3 ml.  $\text{Na}^+\text{-stimulated hydrolytic activity}$  with  $[\gamma\text{-}^{32}\text{P}]\text{GTP}$  (1 mM) as substrate is assayed in identical fashion. The  $\text{P}_i$  production is measured by stopping the reaction at a given time through addition of 0.3 ml 10% (w/v) trichloroacetic acid, followed by mixing with 0.4 ml of a 20% (w/v) aqueous charcoal suspension. The charcoal adsorbs adenosine phosphates from the reaction medium but not inorganic phosphate [10]. The charcoal is sedimented by centrifugation for 10 min at  $2000 \times g$  and 0°C, and a 0.2 ml aliquot of the supernatant is mixed with 4 ml liquid scintillation fluid (Aqualuma Plus). Radioactivity is measured with a liquid scintillation counter. ATPase activity is taken as the difference in the activity in the presence of NaCl and that in the absence of NaCl but with 1 mM ouabain plus 5 mM  $\text{MgCl}_2$ . The latter activity is considered to represent contaminating  $\text{Mg}^{2+}\text{-ATPase}$  activity. In order to minimize background ATPase activity, the control is incubated with ouabain and  $\text{MgCl}_2$  for 30 min at 22°C prior to assay [11].  $(\text{Na}^+ + \text{K}^+)\text{-dependent hydrolytic activity}$  is determined in identical fashion but with  $\text{K}^+$  present in the indicated concentrations. Assays are performed in duplicate and each experiment is repeated at least once. Conversion of ATP or GTP by hydrolysis is never allowed to exceed 15%.

Phosphorylation by ATP or GTP and dephosphorylation, both at 22°C, follow the methods earlier described by us [12,13]. The  $\text{E}_2\text{K} \rightarrow \text{E}_1\text{K}$  conversion is assayed by determining fluorescence changes upon addition of  $\text{K}^+$  and ATP or GTP to 5-iodoacetamidofluorescein-labeled  $(\text{Na}^+ + \text{K}^+)\text{-ATPase}$  at 22°C and pH 7.0 [7]. All kinetic data

are plotted according to the Henri-Michaelis-Menten method;  $V_{\max}$  and  $K_m$  values are determined from Scatchard plots.

**Reagents.** All chemicals are of analytical grade. Ouabain is purchased from Sigma Chemical Company (St. Louis, MO, U.S.A.). ATP and GTP are obtained from Boehringer Mannheim (F.R.G.) and are converted to the imidazole- or Tris-salt by cation exchange over a Dowex 50W X4 column in the  $H^+$ -form, followed by neutralization with imidazole or Tris.  $[\gamma\text{-}^{32}\text{P}]\text{ATP}$  and  $[\gamma\text{-}^{32}\text{P}]\text{GTP}$  are supplied by Amersham International (U.K.) and 5-iodoacetamidofluorescein by Molecular Probes, Inc. (Junction City, OR, U.S.A.). Scintillation fluid (Aqualama Plus) is from Lumac/3M, Schaesberg (The Netherlands).

## Results

### *Na<sup>+</sup>-activated substrate hydrolysis in imidazole buffer*

The  $\text{Na}^+$ -nucleosidetriphosphatase activity is measured as the  $\text{Na}^+$ -dependent hydrolysis of ATP and GTP in the absence of added  $\text{K}^+$  and at varying  $\text{Na}^+$  concentration in 50 mM imidazole-HCl buffer (pH 7.2). The concentrations for each substrate have been chosen such as to saturate the relevant binding site: 1 mM GTP (higher concentrations are inhibitory), 0.02 mM ATP (saturating the high-affinity binding site only), 3 mM ATP (saturating both high- and low-affinity binding sites). Different profiles are obtained for the two substrates (Fig. 1).

For ATP a relatively high activity is obtained in the absence of added  $\text{Na}^+$ , about half of the optimal activity at 300 mM  $\text{Na}^+$ . This represents the imidazole-activated phosphorylation, which we have recently reported [14]. Addition of 0.1–1 mM  $\text{Na}^+$  lowers the activity by 30% at low ATP concentration (0.02 mM). This inhibition corresponds to that of the  $\text{Na}^+$  pump in erythrocytes by extracellular  $\text{Na}^+$ , observed by Glynn et al. [15,16] and ascribed to inhibition of the spontaneous hydrolysis of  $\text{E}_2\text{P}$ . In line with their results, further increase of the  $\text{Na}^+$  concentration changes this inhibition into a stimulation, in which  $\text{Na}^+$  displays a  $\text{K}^+$ -like effect. The optimum is reached at 300 mM  $\text{Na}^+$ . Higher concentrations again give rise to inhibition, which may be due to either an

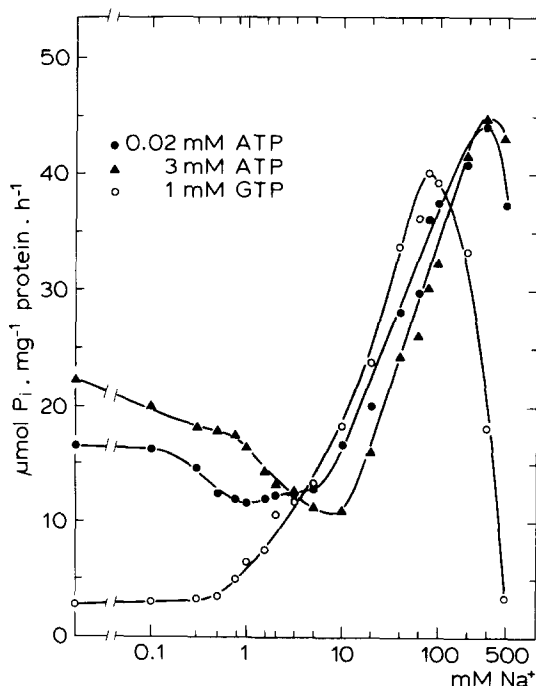


Fig. 1. Enzymatic hydrolysis of ATP or GTP in imidazole-HCl (pH 7.2), without added  $\text{K}^+$ , as a function of the  $\text{Na}^+$  concentration.  $\text{Na}^+$  and substrate concentrations as indicated, enzyme 3.3  $\mu\text{g}/\text{ml}$  for ATP and 16.7  $\mu\text{g}/\text{ml}$  for GTP, incubation 3 min (0.02 mM ATP), 120 min (3 mM ATP) and 30 min (1 mM GTP). Optimal activity for GTP is at 80 mM  $\text{Na}^+$  ( $\text{Na}_{0.5}^+ = 14.4$  mM), and for ATP at 300 mM  $\text{Na}^+$  ( $\text{Na}_{0.5}^+ = 53$  mM at 0.02 mM ATP and 100 mM at 3 mM ATP).

ionic strength effect, a slowdown of the  $\text{E}_1\text{-P} \rightarrow \text{E}_2\text{P}$  transition [17,18] or a return of the inhibition of  $\text{E}_2\text{P}$  hydrolysis. At a high ATP concentration (3 mM) the curve is shifted to the right ( $\text{Na}_{0.5}^+$  increasing from 53 to 100 mM), but the optimum remains at 300 mM and reaches the same height. An antagonism between  $\text{MgATP}$  and  $\text{Na}^+$ , as observed here, has previously been described by Robinson [19].

For GTP there is nearly no hydrolysis below 1 mM  $\text{Na}^+$  and hence no inhibition at low  $\text{Na}^+$  concentrations. The  $\text{Na}^+$ -activation curve has shifted to the left ( $\text{Na}_{0.5}^+ = 14$  mM,  $\text{Na}_{\max}^+ = 80$  mM). This indicates that there is in this case less antagonism between  $\text{Mg-nucleotide}$  and  $\text{Na}^+$ , which is probably due to a lower affinity of the enzyme for  $\text{MgGTP}$  than for  $\text{MgATP}$ . This is reflected in the increasing  $\text{Na}_{0.5}^+$  ratio of 1 : 3.4 : 6.4

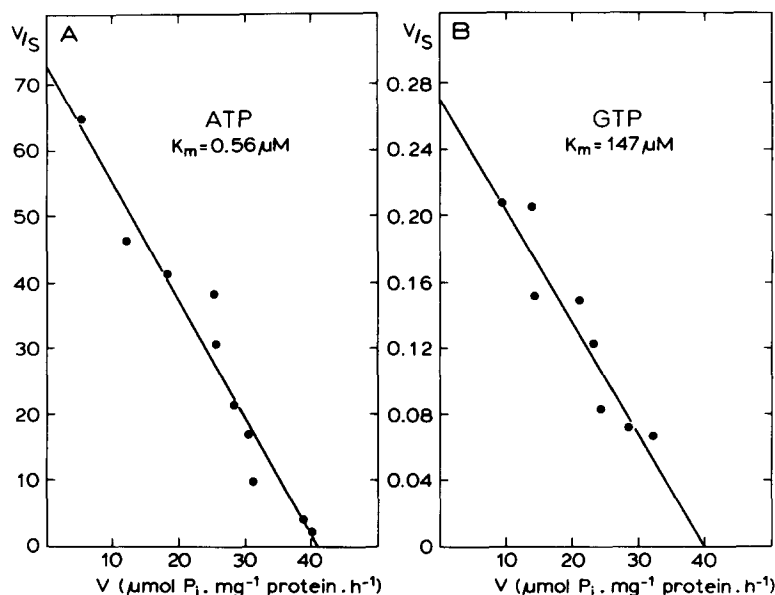


Fig. 2. Scatchard plots for hydrolysis of ATP (A) and GTP (B) in imidazole-HCl (pH 7.2) in the absence of added  $\text{K}^+$ . ATP hydrolysis is followed for 0.25–5 min in the presence of 0.1–20  $\mu\text{M}$  ATP (initial concentrations), 1.7  $\mu\text{g}/\text{ml}$  protein and 300 mM  $\text{Na}^+$ . GTP hydrolysis is followed for 15 min in the presence of 50–500  $\mu\text{M}$  GTP (initial concentrations), 5  $\mu\text{g}/\text{ml}$  enzyme and 80 mM  $\text{Na}^+$ . The substrate concentrations used for calculating  $v/S$  on the ordinate scale are averages of the initial and final concentrations. Lines have been calculated by linear regression analysis.

in the presence of 1 mM GTP, 0.02 mM ATP and 3 mM ATP, respectively.

The most remarkable observation to be made from Fig. 1 is that the peak activities for GTP and ATP are almost the same: 40 vs. 44  $\mu\text{mol}/\text{mg}$  protein per h. This is confirmed in Scatchard plots of the activity, determined at the optimal  $\text{Na}^+$  concentration for each substrate (Fig. 2A and B). While the  $V_{\text{max}}$  is nearly the same for the two substrates (approx. 40  $\mu\text{mol}/\text{mg}$  protein per h), the  $K_m$  values differ by a factor 260 (147  $\mu\text{M}$  for GTP vs. 0.56  $\mu\text{M}$  for ATP).

#### *$\text{Na}^+$ -activated substrate hydrolysis in Tris buffer*

Similar determinations of the  $\text{Na}^+$ -nucleoside triphosphatase activity have been made in Tris buffer (50 mM, pH 7.2). The results, shown in Fig. 3, have been obtained at optimal conditions for each substrate (1 mM GTP, 60 mM  $\text{Na}^+$ ; 0.02 mM ATP, 150 mM  $\text{Na}^+$ ; 3 mM ATP, 300 mM  $\text{Na}^+$ ). The triphosphatase activity for GTP (1 mM) is slightly lower than for 0.02 mM ATP (33 vs. 40  $\mu\text{mol}/\text{mg}$  protein per h), but higher than for 3 mM ATP (33 vs. 25  $\mu\text{mol}/\text{mg}$  protein per h). The

$\text{Na}_{0.5}^+$  value for GTP is, like in imidazole buffer, far lower than that for ATP: 8 mM  $\text{Na}^+$  for GTP, 25 mM for 0.02 mM ATP and 50 mM for 3 mM ATP. However, all three  $\text{Na}_{0.5}^+$  values are about

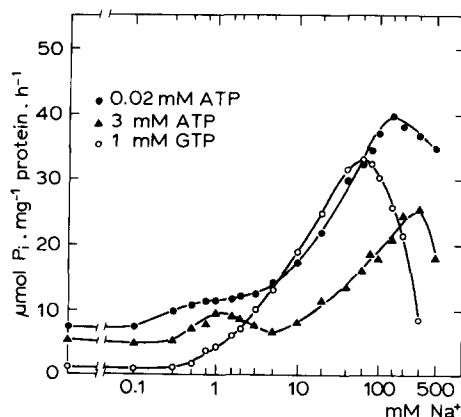


Fig. 3. Enzymatic hydrolysis of ATP and GTP in Tris-HCl (pH 7.2), without added  $\text{K}^+$  as a function of the  $\text{Na}^+$  concentration. Conditions are the same as in Fig. 1. Optimal activity for 1 mM GTP is at 60 mM  $\text{Na}^+$  ( $\text{Na}_{0.5}^+ = 7.9 \text{ mM}$ ), for 0.02 mM ATP at 150 mM  $\text{Na}^+$  ( $\text{Na}_{0.5}^+ = 25 \text{ mM}$ ) and for 3 mM ATP at 300 mM  $\text{Na}^+$  ( $\text{Na}_{0.5}^+ = 50 \text{ mM}$ ).

half of those in imidazole buffer, which is another example of an interaction between imidazole and  $\text{Na}^+$  previously noticed for the phosphorylation step [14]. The equal ratio's of the  $\text{Na}_{0.5}^+$  values in the two buffers, as well as the near equality of the  $K_m$  values for the two substrates in Tris and imidazole (100 vs. 147  $\mu\text{M}$  for GTP; 0.56 vs. 0.56  $\mu\text{M}$  for ATP, each at optimal  $\text{Na}^+$  concentration), indicate a similar antagonism between Mg-nucleotide and  $\text{Na}^+$  in these buffers.

The most striking differences between the activity profiles in imidazole and Tris are the lower buffer-stimulated ATPase activities and the strong MgATP inhibition revealed by the profile with 3 mM ATP present. The first phenomenon is due to the lower phosphorylation capability in Tris than in imidazole, which we have recently reported [14]. The second phenomenon is reminiscent of the substrate inhibition described by Robinson et al. [20] for  $\text{Na}^+$ -ATPase activity in Tris-histidine buffer, particularly at high  $\text{Na}^+$  concentration. In addition, inhibition at low  $\text{Na}^+$  concentrations is absent and is replaced by stimulation, as in previous reports on  $\text{Na}^+$ -ATPase activity in imidazole-glycylglycine [21] or Tris-HCl [16,22,23] or is shifted to higher  $\text{Na}^+$  concentrations (1–5 mM) at elevated MgATP level (3 mM), as also occurs in imidazole (Fig. 1).

#### $\text{Na}^+$ -dependent phosphorylation by GTP

Schoner et al. [4] first reported that GTP is a phosphorylating substrate for  $(\text{Na}^+ + \text{K}^+)\text{-ATPase}$ . Fig. 4 confirms this for our preparation and shows that in 3 seconds phosphorylation levels as high as 3.6 nmol/mg protein can be reached. This is the same level as obtained with ATP for the same preparation, but at a greatly increased  $K_m$  value (50–170  $\mu\text{M}$  for GTP, 0.1–0.2  $\mu\text{M}$  for ATP). The  $K_m$  value for GTP strongly depends on the  $\text{Na}^+$  concentration (ionic strength being kept constant at 140 mM by addition of choline chloride). It decreases from about 170  $\mu\text{M}$  at 1–3 mM  $\text{Na}^+$  to a minimum value of 50  $\mu\text{M}$  at 8–12 mM  $\text{Na}^+$ , and increases again to 75  $\mu\text{M}$  at 100 mM  $\text{Na}^+$ . The  $K_m$  value for ATP is fairly constant at 0.1–0.2  $\mu\text{M}$  in this range of  $\text{Na}^+$  concentrations. The  $K_m$  values for  $\text{Na}^+$  also differ: 3.6 mM for GTP vs. 1.1 mM for ATP, accompanied by a similar trend in the Hill coefficients for  $\text{Na}^+$  activation:  $n_H$  is 2.4 for GTP vs. 1.4 for ATP. The GTP-phosphorylated intermediate is predominantly in the  $\text{E}_2\text{P}$  form, since it is ADP-insensitive and  $\text{K}^+$ -sensitive. Its  $\text{K}^+$ -sensitivity appears to be equally high as that of the ATP-generated phosphoenzyme: 97%  $\text{K}^+$  (0.1 mM) induced hydrolysis within 3 s (cf. Ref. 13).

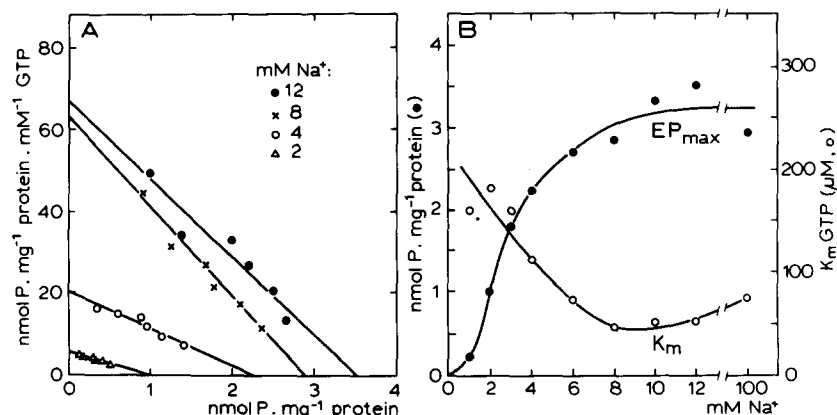


Fig. 4. Phosphorylation by GTP as a function of the  $\text{Na}^+$  concentration in 50 mM imidazole-HCl (pH 7.0). (A) Scatchard plots for 3 s phosphorylation at 22°C in the presence of 5 mM  $\text{MgCl}_2$ , 0.33 mg/ml enzyme, 20–200  $\mu\text{M}$  GTP and the indicated  $\text{Na}^+$  concentrations. Ionic strength (140 mM) is kept constant by addition of choline chloride. Lines have been calculated by linear regression analysis. (B) Michaelis-Menten plot of abscissa intercepts in (A), indicating  $\text{EP}_{\text{max}}$ , as a function of the  $\text{Na}^+$  concentration. Also shown is the  $K_m$  for GTP, determined from the slopes in A, as a function of the  $\text{Na}^+$  concentration.

*The effects of  $K^+$  on the hydrolysis of ATP and GTP*

Fig. 5 shows the effects of  $K^+$  on  $Na^+$ -activated hydrolysis of ATP and GTP at varying  $Na^+$  concentration. Again some striking differences are noticed. In the case of ATP (5 mM), in the absence of added  $K^+$  (residual  $K^+$  concentration 6  $\mu$ M) there is a peak activity at 300 mM  $Na^+$  (Fig. 5A), as previously shown in Fig. 1. Addition of  $K^+$  to the assay medium results in the appearance of a second peak at 2 mM  $Na^+$ . At 80  $\mu$ M  $K^+$  the two peaks are of the same magnitude. Further increase of the  $K^+$  concentration raises the height of both peaks and shifts them together. At 1.8 mM  $K^+$  the two peaks merge at 40 mM  $Na^+$ . The appearance of the left hand peak can be explained by  $K^+$ -stimulated hydrolysis of  $E_2P$  accompanied by a change of the rate-limiting step from the hydrolysis of  $E_2P$  to the  $ATP$ -assisted  $E_2K \rightarrow E_1Na$  transition (see Discussion). In this transition  $K^+$  and  $Na^+$  compete at the intracellular sites [24], leading to a shift of this peak to higher  $Na^+$

concentrations when  $K^+$  is increased. The  $Na^+/K^+$  concentration ratio for the left hand peak remains rather constant at 22–25.

A very different behaviour is observed in the  $Na^+$ -activated hydrolysis of GTP upon addition of  $K^+$  (Fig. 5B). The position of the peak, present at 80 mM  $Na^+$  in the absence of added  $K^+$  (6  $\mu$ M residual  $K^+$ ), is not changed. Its height is slightly increased by 0.2 mM  $K^+$ , and then decreases at higher  $K^+$  concentrations (1–5 mM). The half maximally stimulating  $Na^+$  concentration changes little. The low activity level between zero and 0.5 mM  $Na^+$  is almost doubled upon addition of 0.2–1 mM  $K^+$ , but is decreased with 5 mM  $K^+$ .

The near absence of stimulation by  $K^+$  of the GTP hydrolysis is in marked contrast to the large stimulation of the ATP hydrolysis. The absence of this effect for GTP could be due to an increased  $K_m$  value for this nucleotide at increasing  $K^+$  level. Thus we have determined the  $K_m$  and  $V_{max}$  values for GTP at constant (100 mM)  $Na^+$  concentration and variable  $K^+$  concentrations (Fig. 6).

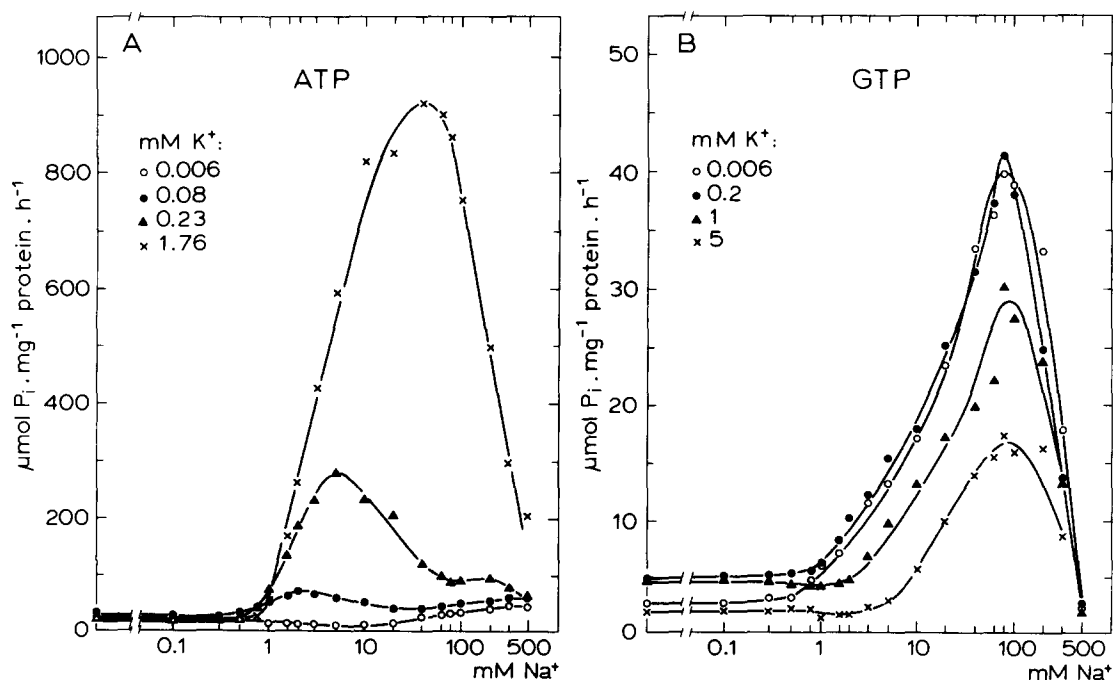


Fig. 5. Effect of  $Na^+ + K^+$  on the hydrolysis of ATP and GTP. The hydrolysis is assayed in 50 mM imidazole-HCl (pH 7.2) in the presence of the indicated  $Na^+$  and  $K^+$  concentrations. (A) Hydrolysis of ATP: ATP 5 mM, enzyme 16.7  $\mu$ g/ml, incubation for 5, 10 and 30 min at decreasing  $K^+$  concentrations. (B) Hydrolysis of GTP: GTP 1 mM, enzyme 16.7  $\mu$ g/ml and incubation for 30 min. Maximal activity at all  $K^+$  concentrations is at 80 mM  $Na^+$  ( $Na^+_{0.5} = 17$ –21 mM).

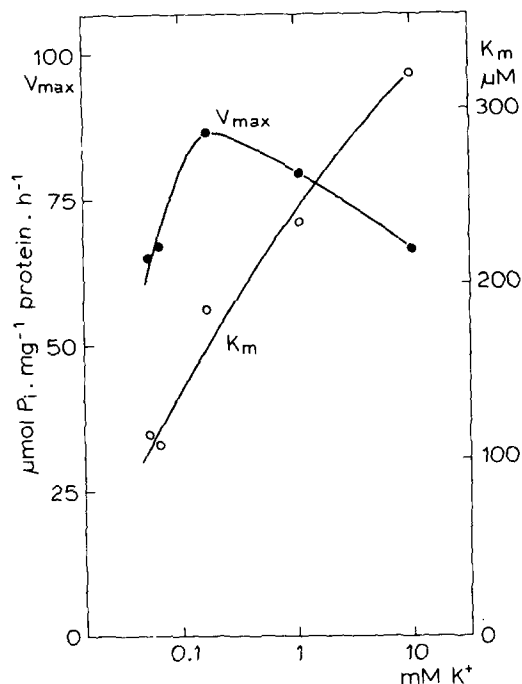


Fig. 6. Plots of  $V_{\max}$  and  $K_m$  for GTP hydrolysis as a function of the  $K^+$  concentration.  $V_{\max}$  and  $K_m$  have been determined from Scatchard plots at 0.06–1 mM GTP (inhibition above 1 mM GTP).  $Na^+$  is 100 mM,  $K^+$  as indicated, enzyme 16.7  $\mu$ g/ml, incubation 1–15 min, depending on the GTP concentration.

Up to 0.16 mM  $K^+$ , or about twice the  $K_m$  value for stimulation of  $E_2P$  hydrolysis [13], the  $V_{\max}$  value is twice that in the absence of  $K^+$  (Fig. 2B). At 10 mM  $K^+$  there is a 23% inhibition of  $V_{\max}$ , hence the stimulation is reduced to 1.6-fold. This phenomenon is accompanied by a continuous increase in the  $K_m$  value for GTP, going from 110  $\mu$ M at 0.06 mM  $K^+$  to 320  $\mu$ M at 10 mM  $K^+$ . Under these conditions the hydrolysis of ATP is optimal (1200  $\mu$ mol/mg protein per h) and 30-times higher than that in the absence of  $K^+$  (Fig. 2A). The  $K_m$  value for ATP increases even more than that for GTP: 0.6  $\mu$ M without  $K^+$  (Fig. 2A), 430  $\mu$ M at 10 mM  $K^+$  and 100 mM  $Na^+$  [25].

The GTP-phosphorylated intermediate is sensitive to  $K^+$ , like the ATP-phosphorylated intermediate. Hence, the inhibitory effect of  $K^+$  above 0.2 mM on the overall activity (Figs. 5B and 6) must occur in a step subsequent to the dephosphorylation. A likely candidate is the  $E_2K \rightarrow E_1K$

transition, which is examined in the following section.

#### The $E_2K \rightarrow E_1K$ transition

Several fluorescent probes can monitor the conformational change from  $E_2$  to  $E_1$ , but most of these are unsuitable for following the ATP-driven transition as they interfere with nucleotide binding, e.g. fluorescein isothiocyanate [26], eosin and eosin maleimide [6,27]. A covalently binding fluorescent label, which does not inhibit  $(Na^+ + K^+)$ -ATPase activity and is able to record the ATP-driven  $E_2K \rightarrow E_1K$  transition, is 5-iodoacetamidofluorescein (5-IAF, Ref. 7). Fig. 7 demonstrates the fluorescence changes upon sequential addition of  $K^+$  and ATP or GTP to the IAF-labeled enzyme. Addition of  $K^+$  up to 50  $\mu$ M quenches the fluorescence and induces the  $E_2K$  conformation. Subsequent addition of ATP (up to 0.75 mM) reverses this effect and restores the fluorescence to the original level (Fig. 7A). However, addition of GTP (up to 1 mM) is virtually without effect (Fig. 7B). Subsequent addition of ATP again gives an increase in fluorescence, but now the restoration is incomplete, probably due to inhibition exerted by

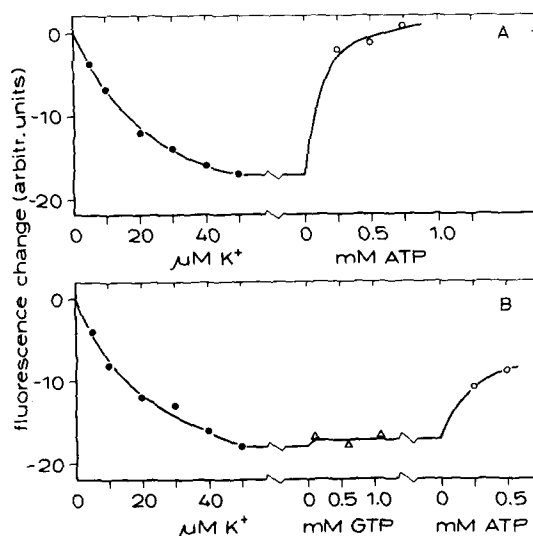


Fig. 7. Fluorescence response of 5-IAF-labeled enzyme to  $K^+$  and substrate. Fluorescence change (arbitrary units) in 50 mM imidazole-HCl (pH 7.0, 22°C), upon addition of  $K^+$  and ATP (A) or GTP (B) in concentrations as indicated. Enzyme concentration is 27  $\mu$ g/ml. Excitation at 490 nm, emission at 520 nm.

the GTP. These data indicate that GTP in millimolar concentration, unlike ATP, is unable to drive the  $E_2K \rightarrow E_1K$  transition and even inhibits the ATP-driven transition.

## Discussion

### *The $E_2K \rightarrow E_1K$ transition as rate-limiting step*

In this study we have provided evidence that in the case of GTP the  $E_2K \rightarrow E_1K$  transition is the major locus of substrate specificity in the overall reaction mechanism in the presence of  $Na^+ + K^+$ . The reduced activity with GTP as compared to ATP is not a matter of a reduced affinity for the substrate, since the  $K_m$  values for ATP (0.43 mM, Ref. 25) and GTP (0.32 mM) are not appreciably different. The  $K_d$  values for the low-affinity substrate site, which is thought to be involved in the  $E_2K \rightarrow E_1K$  transition, do not appreciably differ either [25]. Hence the reduction of the hydrolytic activity with GTP as substrate upon addition of  $K^+$  is due to a reduction of the maximal rate of the  $E_2K \rightarrow E_1K$  transition. This conclusion is supported by the 5-IAF fluorescence experiments.

Further support for this conclusion is provided by our finding that in imidazole buffer the  $Na^+$ -dependent activities in the absence of  $K^+$ , viz. the steady-state phosphorylation level and the  $Na^+$ -activated dephosphorylation, are equally high for the two substrates. This implies that steps 1–3, i.e. substrate binding, phosphorylation and  $E_1P \rightarrow E_2P$  transition, are not rate-limiting for the  $Na^+$ -activated hydrolysis of  $E_2P$  for either substrate. Neither does the  $K^+$ -stimulated dephosphorylation appear to be rate-limiting for GTP. The phosphoenzyme generated by GTP is, like that generated by ATP [13], ADP-insensitive and  $K^+$ -sensitive, indicating that the  $E_1P \rightleftharpoons E_2P$  equilibrium is poised far to the right. This explains also the two-fold increase in  $V_{max}$  of GTP hydrolysis upon addition of about twice the half-maximally stimulating concentration of  $K^+$ . This means that the large difference in GTP and ATP hydrolysis in the presence of  $K^+$  cannot be ascribed to a reduced  $K^+$ -sensitivity of the GTP-phosphorylated intermediate. Rather step 5, the  $E_2K \rightarrow E_1K$  transition following the  $K^+$ -stimulated dephosphorylation of  $E_2P$  (step 4), must be responsible. GTP is unable to drive this rate-limiting step in the overall activity.

Still another phenomenon may be explained by this insufficiency of GTP. In the hydrolysis of ATP  $K^+$  induces an additional peak of activity in the 2–40 mM  $Na^+$  concentration range ( $Na_{0.5}^+$  is 1–4 mM), which increases with increasing  $K^+$  concentration. The  $Na_{0.5}^+$  value for the  $E_2 \rightarrow E_1$  transition in the absence of added  $K^+$  (about 2 mM, Ref. 28) falls in this range of competition between  $Na^+$  and  $K^+$  ions ( $K_m Na^+ = 1.3$  mM,  $K_i K^+ = 0.23$  mM, Ref. 26). ATP apparently compensates for the greater part of the inhibition by  $K^+$ . When no such compensation by the nucleotide occurs, as in the case of GTP,  $Na^+$  must exert this action and the  $Na_{0.5}^+$  value remains high. In the absence of  $K^+$  there is also no activation but rather inhibition in the low  $Na^+$  concentration range, even though the  $E_2 \rightarrow E_1$  transition is then not hampered by  $K^+$ . However, under these conditions the dephosphorylation step is rate-limiting but not the  $E_2 \rightarrow E_1$  transition.

### *Antagonism between $Na^+$ and substrate or imidazole*

Some other interesting mechanistic effects appear from our study. In the  $Na^+$ -activated hydrolysis of GTP and ATP the activity curve strongly depends on the affinity and concentration of the Mg-complexed nucleotide. The curve for GTP, which has a 260-fold lower affinity for the enzyme in this process, is shifted to lower  $Na^+$  concentrations than the curve for ATP. The latter curve shifts even further to the right at higher ATP concentrations (3 mM). This represents an antagonism between Mg-nucleotide and  $Na^+$ . This phenomenon was previously detected by Robinson [19] for the  $(Na^+ + K^+)$ -ATPase activity, where the  $Na_{0.5}^+$  value, after extrapolation to zero  $[K^+]$ , showed an increase with the Mg-ATP concentration.

Comparison of the ratio of half-maximally activating  $Na^+$  concentrations at 1 mM GTP, 0.02 mM ATP or 3 mM ATP in either imidazole or Tris medium shows that the Mg-nucleotide- $Na^+$  antagonism is not affected by the buffer. This indicates that in the intact cell there must be an interaction between the substrate at the intracellular side and  $Na^+$  at the extracellular side of the membrane, since  $Na^+$  in that situation stimulates the hydrolysis process from the extracellular side



[15,16]. This antagonism is opposite to the synergism that appears to occur between the nucleotide and  $\text{Na}^+$  at the same (intracellular) side of the membrane. Here the ratio  $K_m$  values for intracellular  $\text{Na}^+$  in the formation of  $\text{E}_2\text{P}$  from the low-affinity substrate GTP instead of the high-affinity substrate ATP is 3.3:1, which is opposite to the 1:3.3 ratio for the stimulation of  $\text{E}_2\text{P}$  hydrolysis by extracellular  $\text{Na}^+$ .

For the  $\text{Na}^+$ -activated hydrolysis the  $\text{Na}_{0.5}^+$  values in Tris are about half of those in imidazole buffer. At high  $\text{Na}^+$  concentrations the substrate inhibition in Tris medium is stronger than in the imidazole medium. The latter phenomenon has earlier been detected by Robinson et al. [20], who found the inhibition to increase with the  $\text{Na}^+$  concentration in Tris-histidine buffer. The different effects of imidazole and Tris medium may be due to the stronger  $\text{Na}^+$ -like properties of imidazole- $\text{H}^+$  as compared to Tris- $\text{H}^+$ , as has also been observed for phosphorylation [14]. However, lacking the  $\text{K}^+$ -like properties of  $\text{Na}^+$ , which stimulates  $\text{E}_2\text{P}$  hydrolysis, imidazole buffer inhibits this  $\text{Na}^+$  effect more than Tris buffer does. Imidazole- $\text{H}^+$  increases the  $\text{Na}_{0.5}^+$  values for the hydrolysis, while Tris- $\text{H}^+$  increases the extent of the  $\text{Na}^+$ -dependent substrate inhibition. Here also, the antagonism between imidazole- $\text{H}^+$  and  $\text{Na}^+$  at the low-affinity extracellular hydrolytic sites is replaced by a synergism at the high-affinity intracellular phosphorylating sites, in which imidazole increases instead of decreases the apparent affinity for  $\text{Na}^+$  [14]. The increase in the cooperativity index  $n_H$  for  $\text{Na}^+$  in phosphorylation from 1.4 for ATP to 2.4 for GTP indicates that the  $\text{Na}^+$  sites interact more strongly when the substrate has a lower affinity [29].

#### *Substrate structure and the locus of specificity*

Nucleotides can be modified in the purine ring, the ribose group or the phosphate chain, with different effects on the location of the rate-limiting step in the reaction mechanism. Our data on the partial reactions with ATP and GTP, where only the purine ring is modified, suggest that the  $\text{E}_2\text{K} \rightarrow \text{E}_1\text{K}$  transition is the main locus of substrate specificity, since the other partial reactions are not markedly affected. Similar conclusions can be reached for other substrates with a modified pu-

rine ring, viz. CTP [30] and UTP [31]. The conclusion is supported by the low effectiveness of ITP, GTP, UTP and CTP in supporting the  $\text{K}^+ - \text{K}^+$  exchange, which involves the  $\text{E}_2\text{K} \rightleftharpoons \text{E}_1\text{K}$  transition [32]. This may be due to the relatively high  $pK$  value of the  $\text{N}_1$  nitrogen of the purine ring ( $\text{N}_3$  of the pyrimidine group) in these substances, precluding them from taking up a proton from the enzyme in this transition [33].

Modification of the phosphate chain shifts the rate-limiting step to other partial reactions. Adenylyl imidodiphosphate ( $\text{AdoPP}[\text{NH}]P$ ), modified in the  $\beta$ - $\gamma$  phosphate link, is hydrolyzed by the enzyme in the presence of  $\text{Mg}^{2+} + \text{Na}^+$ , yielding  $\text{ADP-NH}_2$  and  $\text{P}_i$  [34].  $\text{K}^+$  neither stimulates nor inhibits the activity. Our interpretation is that the phosphorylation step and not the  $\text{E}_2\text{K} \rightarrow \text{E}_1\text{K}$  transition is here the rate-limiting step. This is in agreement with the finding of Simons [35] that  $\text{AdoPP}[\text{NH}]P$  can effectively (55%) substitute for ATP in  $\text{K}^+ - \text{K}^+$  exchange transport.

When the  $\gamma$ -phosphate group is modified as in  $[\gamma\text{-S}]\text{ATP}$ , there is a general reduction in the rate constants of the partial reactions, but the  $\text{E}_1\text{P} \rightarrow \text{E}_2\text{P}$  transition (and the phosphorylation step) appears to be most retarded as reflected in the size of the ADP-sensitive phosphoenzyme fraction and its reduced  $\text{K}^+$ -sensitivity [13]. Hence, there is no constant locus of specificity for the various substrates in the reaction mechanism of  $(\text{Na}^+ + \text{K}^+) - \text{ATPase}$ . The general picture emerging is: (1) modification of the phosphate chain predominantly affects reactions involving the phosphoenzyme (formation of  $\text{E}_1\text{P}$  and its transition to  $\text{E}_2\text{P}$ ), (2) modification of the purine ring mainly affects the  $\text{E}_2\text{K} \rightarrow \text{E}_1\text{K}$  transition of the dephosphorylated enzyme, in addition to a general decrease of the substrate binding affinity [3,36].

#### **Acknowledgement**

The authors wish to acknowledge the financial support received from The Netherlands Ministry of Education and Science, which enabled the first author to work for two years in the Department of Biochemistry, University of Nijmegen, in the context of the scientific and cultural exchange program between the People's Republic of China and The Netherlands.

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