

## CONCERTED PHOSPHORYLATION OF ENDOGENOUS TRACHEAL SMOOTH MUSCLE MEMBRANE PROTEINS BY $\text{Ca}^{2+}$ · CALMODULIN-, CYCLIC GMP- AND CYCLIC AMP-DEPENDENT PROTEIN KINASES

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### 1. Introduction

The precise control of the myoplasmic free  $[\text{Ca}^{2+}]$  is an important determinant in the regulation of the contractile state of mammalian smooth muscle [1]. The control mechanism is thought to reside with membrane  $\text{Ca}^{2+}$ -transport ATPases which actively pump  $\text{Ca}^{2+}$  against a concentration gradient and maintain the intracellular  $[\text{Ca}^{2+}]$  below  $0.1 \mu\text{M}$ . The  $\text{Ca}^{2+}$ -transport ATPase has been measured in many smooth muscle types [2,3], purified [4], and its activity has been shown to be modulated by cyclic nucleotides [5,6] as well as by the  $\text{Ca}^{2+}$ -binding protein calmodulin [3].

The molecular mechanism by which calmodulin and cyclic nucleotides regulate smooth muscle  $\text{Ca}^{2+}$ -transport ATPase is unknown. The phosphorylation of the enzyme itself or the regulatory proteins may be associated with it. The latter mechanism has been developed for cardiac sarcoplasmic reticular  $\text{Ca}^{2+}$ -transport ATPase [7].

Here, we report the  $\text{Ca}^{2+}$  · calmodulin-, cyclic GMP- and the possible cyclic AMP-dependent phosphorylation of endogenous membrane proteins from tracheal smooth muscle. Two proteins, 130 000 and 80 000

$M_r$ , appear to be phosphorylated by all 3 second messengers. The functional role of the concerted phosphorylation remains unknown but may be intimately involved in the regulation of smooth muscle membrane function including  $\text{Ca}^{2+}$  transport.

### 2. Methods

$[\gamma\text{-}^{32}\text{P}]\text{ATP}$  was from New England Nuclear. Trifluoperazine was a generous gift of Smith, Kline and French.

Tracheal smooth muscle membranes were prepared as in [3] with the following modifications. The smooth muscle was homogenized in 8 vol. buffer A containing 250 mM sucrose, 10 mM Hepes (pH 7.0), 10 mM 2-mercaptoethanol, 0.5 mM PMSF, 0.05 mM TPCK, 0.5% ethanol, and 50  $\mu\text{g}/\text{ml}$  heparin. The homogenate was filtered through 3 layers of cotton gauze. The filter residue was resuspended in 4 vol. buffer A, homogenized and filtered as above. Differential centrifugation of the combined filtrate was performed as follows:  $1300 \times g$ , 10 min;  $10\,000 \times g$ , 10 min;  $132\,000 \times g$ , 60 min. The  $132\,000 \times g$  pellet used here was resuspended in buffer A lacking ethanol and heparin, placed on ice and used within 3 h.

Cyclic AMP- and cyclic GMP-dependent phosphorylation was assayed in 75  $\mu\text{l}$  with 40 mM Hepes (pH 7.0), 10 mM  $\text{KN}_3$ , 15 mM EGTA, 10 mM Mg-acetate, 1 mM 3-isobutyl-1-methylxanthine, 100  $\mu\text{g}$  protein, 5  $\mu\text{M}$  MgATP, with or without cyclic AMP or cyclic GMP.  $\text{Ca}^{2+}$  · Calmodulin-dependent phosphorylation was assayed under identical conditions minus isobutylmethylxanthine and with or with-

**Abbreviations:** Hepes, 4-(2-hydroxyethyl)-1-piperazine ethanesulfonic acid; PMSF, phenylmethylsulfonyl fluoride; TPCK, 1-tosylamide-2-phenylethyl chloromethyl ketone; EGTA, ethylene-glycol-(bis( $\beta$ -aminoethylether)- $N,N,N',N'$ -tetraacetic acid; SDS, sodium dodecyl sulfate; PKI, cyclic AMP-dependent protein kinase inhibitor protein; CaM, calmodulin; TFP, trifluoperazine

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out  $\text{Ca}^{2+}$  and/or calmodulin. The  $[\text{Ca}^{2+}]_{\text{free}}$  was determined with a  $\text{Ca}^{2+}$ -EGTA buffer system [3].

Following preincubation at  $30^\circ\text{C}$  for 4 min, reactions were begun with  $[\gamma\text{-}^{32}\text{P}]\text{ATP}$  (15 Ci/mmol), incubated for 1 min at  $30^\circ\text{C}$ , and terminated by adding 25  $\mu\text{l}$  of a solution containing 280 mM Tris (pH 6.8), 8% SDS, 32% glycerol, 4% 2-mercaptoetha-

nol and 0.008% bromophenol blue. Each sample was heated at  $100^\circ\text{C}$  for 2 min, covered, and allowed to stand overnight at room temperature.

Each sample was subjected to SDS-polyacrylamide gel electrophoresis as in [8] on 7–14% continuous polyacrylamide gradient gels. After autoradiography the appropriate protein bands were cut from the gels

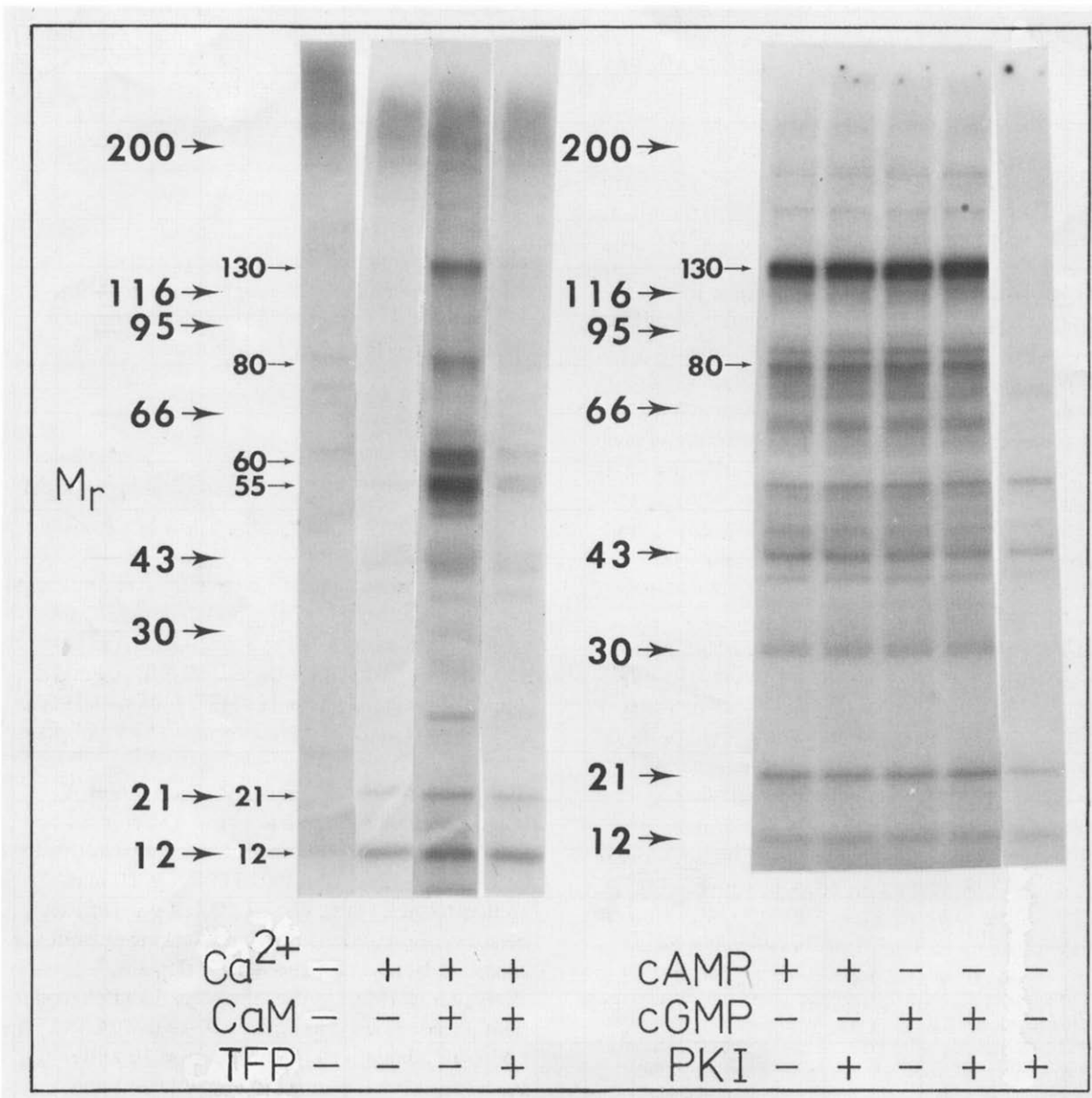


Fig.1. Autoradiograms illustrating phosphorylation of endogenous trachealis muscle membrane proteins.  $M_r$  standards ( $\times 10^{-3}$ ) were: myosin (200),  $\beta$ -galactosidase (116), phosphorylase (95), bovine serum albumin (66), ovalbumin (43), catalase (30), trypsin inhibitor (21) and cytochrome *c* (12). The  $[\text{Ca}^{2+}]_{\text{free}}$ ,  $[\text{CaM}]$ ,  $[\text{cAMP}]$  and  $[\text{cGMP}]$  were all  $10 \mu\text{M}$ . The effect of  $30 \mu\text{M}$  TFP was determined in the presence of  $10 \mu\text{M}$   $\text{Ca}^{2+}_{\text{free}}$  and  $0.3 \mu\text{M}$  CaM and the effect of  $0.4 \text{ mg PKI/ml}$  was assayed with  $10 \mu\text{M}$  cAMP or cGMP. The autoradiograms are a representative example of  $\geq 3$  detn.

and  $^{32}\text{P}$ -content assessed by scintillation spectrometry. Protein was determined as in [9].

### 3. Results and discussion

Preliminary experiments demonstrated the phosphorylation of membrane proteins to be linear for up to 90 s at  $30^\circ\text{C}$  in the presence of  $5\text{ }\mu\text{M}$  ATP. Varying the  $[\text{ATP}]$  from  $0.3\text{--}100\text{ }\mu\text{M}$  had no effect on the endogenous phosphorylation pattern.

Fig.1 illustrates the effect of various agents on membrane phosphorylation.  $\text{Ca}^{2+}$  and calmodulin together (calmodulin alone had no effect) stimulated the phosphorylation of several proteins, the most notable being 130 000, 80 000, 60 000, 55 000, 21 000 and 12 000  $M_r$ . The majority of these proteins appear to be intrinsic to the membranes as washing with various salt concentrations (60–600 mM KCl) after the phosphorylation assay had no effect on the phosphorylation pattern. The only exception was the 21 000  $M_r$  protein (unpublished). Trifluoperazine, an agent that binds to calmodulin and inhibits calmodulin-dependent processes [10], was found to specifically reverse the effect of  $\text{Ca}^{2+}$  and calmodulin.

Cyclic AMP and cyclic GMP also stimulated the incorporation of  $^{32}\text{P}$  into 2 membrane proteins (130 000 and 80 000  $M_r$ ). These results are similar to the findings in the rat vas deferens and rabbit aorta [11]. The effect of cyclic AMP on membrane phosphorylation (fig.1) was not reversed by a concentration of PKI (0.4 mg/ml) that completely inhibits cyclic AMP-dependent phosphorylation in tracheal smooth muscle cytosol preparations (unpublished). The endogenous protein substrates also appear to be intrinsic to the membranes as washing with various salt concentrations (60–600 mM KCl) had no effect on the phosphorylation pattern (unpublished).

The effect of varying the  $[\text{Ca}^{2+}]_{\text{free}}$  and  $[\text{calmodulin}]$  on  $\text{Ca}^{2+}$  · calmodulin-dependent phosphorylation was determined (fig.2). All proteins had similar  $[\text{Ca}^{2+}]_{\text{free}}$ - and  $[\text{calmodulin}]$ -dependencies with app.  $K_a$  for  $\text{Ca}^{2+}$  ( $3\text{ }\mu\text{M}$ ) and calmodulin ( $0.1\text{--}0.5\text{ }\mu\text{M}$ ) within known physiological concentration ranges [11,12].

The  $[\text{cGMP}]$ - and  $[\text{cAMP}]$ -dependencies of the 130 000  $M_r$  and 80 000  $M_r$  proteins were different with respect to their activation characteristics (fig.3). Cyclic GMP stimulated the incorporation of  $^{32}\text{P}$  into the 130 000  $M_r$  and 80 000  $M_r$  proteins in a manner

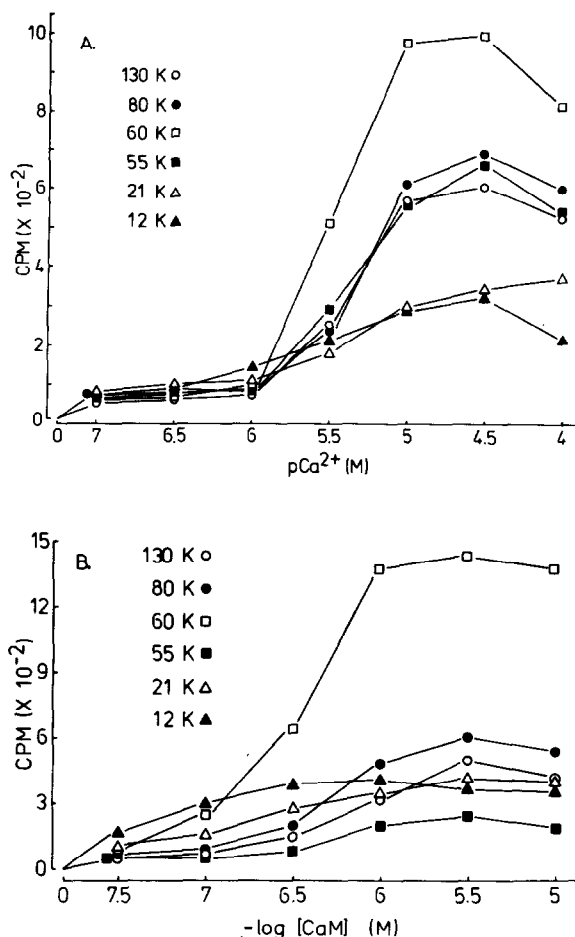


Fig.2.  $\text{Ca}^{2+}$  · CaM-dependent phosphorylation of endogenous tracheal smooth muscle membrane proteins as a function of  $[\text{Ca}^{2+}]_{\text{free}}$  (A) or  $[\text{CaM}]$  (B). The effect of the  $[\text{Ca}^{2+}]_{\text{free}}$  was determined in the presence of  $10\text{ }\mu\text{M}$  CaM. The phosphorylation measured in the presence of  $15\text{ mM}$  EGTA alone was subtracted from each value. The figures are representative examples of  $\geq 3$  detn.

similar to the activation of several physiological processes by homogenous cyclic GMP-dependent protein kinase [13]. In contrast, cyclic AMP had an effect that is similar to the activation of cyclic GMP-dependent protein kinase by cyclic AMP [13]. These results and the finding that PKI had no effect on cyclic AMP, in this membrane preparation, may be due to the stimulation of an endogenous cyclic GMP-dependent protein kinase. Additional studies must be conducted to substantiate these results.

The physiological regulation of many cellular processes by second messengers such as  $\text{Ca}^{2+}$  and cyclic

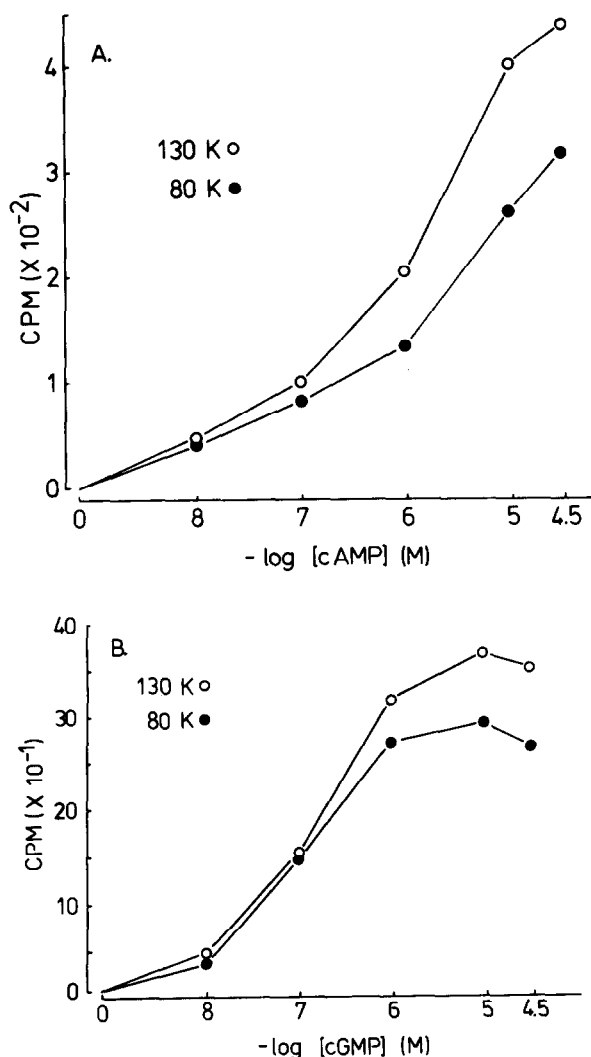


Fig.3. Phosphorylation of endogenous proteins from trachealis muscle as a function of the [cAMP] (A) or [cGMP] (B). The basal phosphorylation in the absence of the nucleotide was subtracted from each value. The figures represent an example of  $\geq 3$  detn.

nucleotides is thought to occur through reversible phosphorylation mechanisms [14]. The specificity of a particular intracellular effector for a process may reside in the cellular location of endogenous substrate proteins with a tissue [15]. In addition,  $>1$  intracellular mediator may regulate a system by a concerted phosphorylation as seen for brain protein I [16], phospholamban [17] and glycogen synthase [18].

These results suggest that  $\text{Ca}^{2+}$ ·calmodulin and cyclic GMP, but not cyclic AMP, may regulate smooth

muscle membrane function through the concerted phosphorylation of  $\geq 1$  proteins. As calmodulin [3] and cyclic GMP [6] have been shown to stimulate smooth muscle microsomal  $\text{Ca}^{2+}$  transport, the phosphorylation of the  $\text{Ca}^{2+}$ -transport ATPase by these intracellular messengers may provide a common pathway by which the myoplasmic free  $[\text{Ca}^{2+}]$  may be precisely regulated in smooth muscle.

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