

## MECHANISM OF ALTERED CYTOSKELETON ORGANIZATION IN INFLUENZA VIRUS INFECTION

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*Summary.* — We followed the autophosphorylation of cytoskeleton (CS) isolated from control chick embryo cell membranes (CS-C) and from these membranes after influenza virus adsorption (CS-V) under conditions allowing to determine the activity of a single type protein kinase. The  $\text{Ca}^{2+}$  dependent calmodulin (CaM) kinase used different substrates from CS-V than did the c'AMP dependent protein kinase. The catalytic subunit (c-subunit) of the c'AMP dependent protein kinase added from outside phosphorylated the same polypeptides than the endogenous c'AMP dependent protein kinase, the further being more active than the latter. The purified influenza virus incorporated  $^{32}\text{P}$  in the presence of the c-subunit only. Incubation of influenza virus with the c-subunit caused morphological changes visible by electron microscopy. The pleomorphy of the particles as well as their electron transmissibility were enhanced in result of structural alterations and rarefaction of surface spikes of the haemagglutinin and neuraminidase. The contractibility of CS isolated from normal CEC and of the CS from CEC by 15 min postinfection (p.i.) was determined according to the actomyosin ATPase activity. The ATPase activity of the cytoskeleton in the presence of the  $\text{Ca}^{2+}$ /CaM and that in the presence of c'AMP were used as controls. The virus as well as the  $\text{Ca}^{2+}$ /CaM increased the ATPase activity. EGTA had no effect but did not interfere with virus stimulation, while c'AMP blocked the virus-induced enhancement of the ATPase activity.

*Key words:* influenza virus; cytoskeleton contraction; phosphorylation; actomyosin; ATPase; chick embryo cell membranes

### Introduction

The organization of actin changes due to different cell functions is regulated by different actin-binding proteins (ABP). Their activity, in turn, is also regulated at higher hierarchic level (Weber and Glenney, 1982; Rosenfeld *et al.*, 1985). Several authors assume that some ABP can bind the receptors

to cytoskeleton (CS) or that the extent of ABP phosphorylation regulates the extent of its interaction with the CS (Sato *et al.*, 1985; Zhuang *et al.*, 1984), and that phosphorylation of some ABP leads to reorganization of the CS (Kawamoto, 1984; Kometani *et al.*, 1986; Lichtfield and Ball, 1986). Phosphorylation is mediated by protein kinase, which may be activated by binding of a ligand to appropriate receptors (Helmreich and Elson, 1984) or by protein kinase C, transferred from cytosol to the membranes by the action of of 12-myristate 13-acetate (PMA) (Pontremoli *et al.*, 1987).

Modulation of the interaction between actin-binding proteins may influence the activation of myosin  $Mg^{2+}$  dependent ATPase and muscle contraction (Horiuchi *et al.*, 1986; Onii *et al.*, 1987). ABP, myosin and actin may be found in nonmuscle cells. They are all colocalized in moving or phagocytizing cells. The redistribution of F-actin as related to ABP or to myosin is important for rearrangement and orientation of forces involved in phagocytosis (Howard and Meyer, 1984). The distribution and function of CS is influenced also by the so called second messenger molecules. Contraction of CS prepared from human fibroblasts depends on  $Ca^{2+}$  and is followed by myosin light chain phosphorylation (20 kD band) by the  $Ca^{2+}$ /CaM activated myosin light chain kinase (MLC-kinase) (Masuda *et al.*, 1984). Actin polymerisation and its interaction with phosphorylated myosin and with ABP may be reversed with compounds enhancing the cytosol c'AMP concentration (Wheeler *et al.*, 1985).

We have shown that at early postinfection intervals influenza virus alters the organization and function at least of the submembraneous actin network (Križanová *et al.*, 1986). After adsorption influenza virus enhances the phosphorylation of certain CS proteins by means of an endogeneous protein kinase, but its relationship to the reorganization of CS has not been elucidated. To approach this question, we tried to identify the protein kinase involved in  $^{32}P$  incorporation to the CS and the possible relationship of the phosphorylation of individual components to its contraction.

### Materials and Methods

*Chick embryo cells (CEC)* were prepared by trypsinization of 11-day-old chick embryos. The cells were grown in Roux flasks in a modified Parker medium supplemented with 8 % inactivated bovine serum and 2 % native horse serum.

*Influenza virus A/WSN (H1N1)* purification, isolation of CEC plasma membranes, virus adsorption to the isolated membranes, autophosphorylation of CS proteins, analysis of CS components by SDS-PAGE, their autoradiography and densitometric evaluation were described previously (Križanová *et al.*, 1985; Križanová *et al.*, 1986).

*Cytoskeleton (CS)* was prepared either from isolated CEC plasma membranes or from CEC monolayer as described (Meisner and Rüegg, 1983).

*Protein content* was determined by the method of Lowry *et al.*, 1951.

*ATPase activity of the cytoskeleton (CS)*. The CS prepared from CEC monolayers was incubated for 15 min at room temperature either with virus-infected allantoic fluid (10–20 virions per cell, CS-V) or with noninfected allantoic fluid (CS-NAF). Nonadsorbed virus was removed before isolation of CS. Closely before measuring the ATPase activity the isolated CS was incubated for 20 min at 25 °C in ATPase buffer (20 mmol/l imidazol-HCl pH 7.2, 60 mmol/l KCl, 1 mmol/l  $MgCl_2$ , 0.4 mmol/l DTT, 0.2 mmol/l EGTA) which enhanced the susceptibility to  $Ca^{2+}$  (Nag and Seidel, 1983).

ATPase activity was determined in 1 mg of isolated CS in the presence of 0.3 mmol/l  $\text{CaCl}_2$ , or 0.3 mmol/l  $\text{CaCl}_2$  and 1.5  $\mu\text{mol/l}$  CaM, or in the presence of  $10^{-4}$  mol/l c'AMP or 0.5 mmol/l EGTA. The reaction started by adding of ATP in a final concentration of 2 mmol/l. After incubation for 0 or 15 min (25 °C) the proteins were precipitated with 2 vol of 15 % trichloroacetic acid (TCA) containing 1 % active carbon and centrifuged. The amount of released Pi was determined in the supernatant according to Fiske and Subbarow (1925). The ATPase activity was expressed as a difference in the amount of released Pi at 15 min and 0 min (Pi 15 min—Pi 0 min).

The effect of virus on the ATPase activity of the isolated CS was determined as follows: after incubation of CS in the imidazol buffer 1500 haemagglutination units per mg CS protein were added (CS-V). The control CS mixture (CS-C) was prepared by adding 10 mmol/l Tris-HCl and 150 mmol/l NaCl (pH 7.2). The ATPase activity was determined as described. Activity of the same amount of purified virus was subtracted from that of CS-V.

**Electron microscopy.** Virus suspension preincubated for 10 min at 30 °C in phosphorylation buffer (40 mmol/l Hepes pH 6.9, 10 mmol/l  $\text{MgCl}_2$ , 0.1 mmol/l  $\text{CaCl}_2$ , 14 mmol/l DTT) under various conditions was layered on formwar coated or carbon coated grids. After attachment (2 min), the grids were stained with 2 % phosphowolfram acid (pH 7) for 1 min and investigated in the microscope Philips EM 300 at 80 kV.

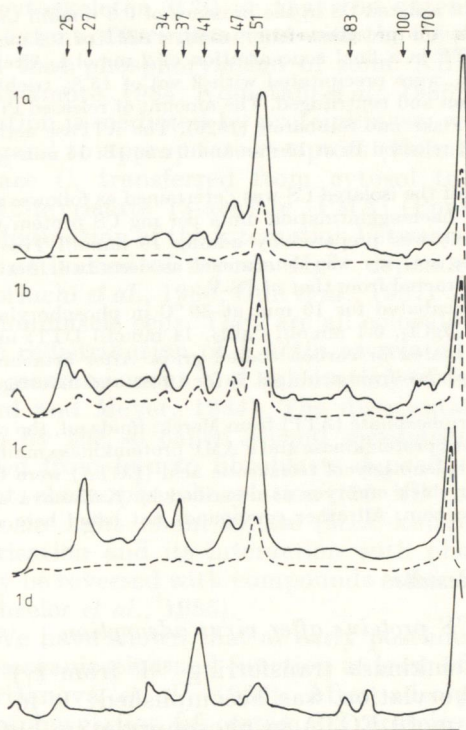
**Drugs.** Cyclic AMP (c'AMP) and adenosin triphosphate (ATP) from Merck; imidazol, the catalytic subunit (c-subunit) of the c'AMP-dependent proteinkinase and c'AMP proteinkinase inhibitor (c'AMP Prki inhibitor) were from Sigma. Ethalendisglycol tetraacetic acid (EGTA) were from Fluka. Calmodulin (CaM) was prepared from chick embryos as described by Křižanová *et al.* (1979). Dithiothreitol (DTT) was from Calbiochem. All other compounds not listed here were analytic grade.

## Results

### <sup>32</sup>P incorporation into individual CS proteins after virus adsorption

To determine the type of proteinkinase transferring <sup>32</sup>P from ( $\gamma$ )<sup>32</sup>P-ATP to individual proteins, phosphorylation was accomplished: 1. in the presence of 1  $\mu\text{mol/l}$  c'AMP and 2 mmol/l EGTA in phosphorylation buffer, 2. in the presence of 30 units of the c-subunit, 3. in the presence of 1  $\mu\text{mol/l}$  CaM, and finally 4. in the presence of 1  $\mu\text{mol/l}$  CaM and 30  $\mu\text{g}$  of c'AMP dependent kinase inhibitor. The conditions under 1 and 2 allow to determine the c'AMP dependent proteinkinase while the conditions under 3 and 4 the  $\text{Ca}^{2+}$  dependent CaM kinase. Incorporation of <sup>32</sup>P into CS observed in the presence of 2 mmol/l EGTA or 2 mmol/l EGTA and 30  $\mu\text{g}$  c'AMP kinase inhibitor corresponded to the basic activity provided by an unknown enzyme or by proteinkinase C-M, a new enzyme occurring after cleavage of proteinkinase C due to the  $\text{Ca}^{2+}$ -activated protease (Pontremoli *et al.*, 1987).

The results of CS analysis under above mentioned phosphorylation conditions are shown on Fig. 1. As visible in Fig. 1-a, in the presence of c'AMP <sup>32</sup>P has been incorporated into the following CS-V polypeptides: 170 kD, 100 kD, 85 kD, 51 kD, 47 kD, 41 kD, 37 kD, 34 kD, and 27—25 kD; in addition, one polypeptide incorporating <sup>32</sup>P moved at the front of the 10 % gel in SDS-PAGE revealing a  $M_r$  lower than 23 kD. The c-subunit of c'AMP dependent proteinkinase (Fig. 1-b) not only enhanced the phosphorylation of above mentioned polypeptides but also transferred <sup>32</sup>P, although to lower extent, from ( $\gamma$ )<sup>32</sup>P-ATP to some of above mentioned polypeptides of CS-C (polypeptides 170 kD, 100 kD, 85 and 34 kD showed minimal labelling). When CS-C was incubated under similar conditions as CS-V in the absence of the c-subunit, <sup>32</sup>P had been incorporated to the 51 kD polypeptide only.

**Fig. 1.**

Influence of phosphorylation conditions of  $^{32}\text{P}$  incorporation into CS proteins

CS proteins (45–50  $\mu\text{g}$ ) were isolated from plasma membranes of CEC, incubated for 10 min at 30  $^{\circ}\text{C}$  in the phosphorylation buffer (40 mmol/l Hepes, 10 mmol/l  $\text{MgCl}_2$ , 0.1 mmol/l  $\text{CaCl}_2$ , 0.1 mmol/l DTT, pH 6.9) to which some of substances indicated below had been added in combination with  $2 \times 10^6 - 2 \times 10^7$  cpm of  $(\gamma)^{32}\text{P}$ -ATP. The reaction was stopped by adding the solubilizing solution. After heating for 3 min (100  $^{\circ}\text{C}$ ) the CS proteins were separated in SDS-PAGE. Phosphorylated proteins were detected by autoradiography and by densitometry. Solid arrows indicate the apparent  $M_r$  in kD, interrupted arrow points at polypeptides moving with the front. Phosphorylation conditions:

1—a: in the presence of  $\mu\text{mol/l}$  c'AMP and 0.5 mmol/l EGTA (c'AMP dependent protein kinase)

1—b: 30 units of the c-subunit of c'AMP protein kinase

1—c: 1 mmol/l CaM and 30  $\mu\text{g}$  c'AMP protein kinase inhibitor ( $\text{Ca}^{2+}$ /CaM dependent protein kinase activity)

1—d: purified influenza virus in the presence c'AMP-subunit

CS-C — cytoskeleton from control membranes (— — —)

CS-V — cytoskeleton from virus treated membranes (————)

These results indicate that adsorption of the virus to the CEC membranes may enhance the activity of c'AMP dependent protein kinase in the isolated CS.

During phosphorylation of CS-V by the  $\text{Ca}^{2+}$ /CaM dependent protein kinase no incorporation of  $^{32}\text{P}$  into polypeptides of  $M_r$  170 kD, 100 kD, 85 kD, 41 kD and 25 kD was observed (Fig. 1-c). The phosphorylation of the 47 kD protein was decreased, while the phosphorylation of proteins  $M_r$  37 kD, 34 kD and 27 kD, and proteins moving at the front in SDS-PAGE was increased indicating that the  $\text{Ca}^{2+}$ /CaM dependent protein kinase acts on other substrates in CS-V than the c'AMP dependent protein kinase. When in the phosphorylation mixture,  $\text{Ca}^{2+}$  was chelated with EGTA or CaM by chlorpromazine, or the activity of c'AMP dependent protein kinase was blocked by an inhibitor of the c'AMP dependent protein kinase, the  $^{32}\text{P}$  accumulated in similar CS-V polypeptides as under action of c'AMP dependent protein kinase (results not shown).

The purified virus alone showed no protein kinase activity. Under various phosphorylation conditions in the absence of exogenous protein kinase, no

Table 1. Influence of influenza virus on the cytoskeleton ATPase activity

ATPase activity in the presence of	Expe- riment	nmol Pi released per mg CS/15 min		
		from CS-NAF	from CS-V	enhancement by the virus
0.3 mmol/l Ca <sup>2+</sup>	A	79.3*	117.7	148 %
0.3 mmol/l Ca <sup>2+</sup> and 1.5 μmol/l CaM		96.3	127.2	131 %
0.5 mmol/l EGTA		50.7	87.5	172 %
0.1 mmol/l c' AMP		54	52.4	97 %
		nmol Pi released per mg CS/15 min		
		from CS-C	from CS-V	enhancement
0.3 mmol/l Ca <sup>2+</sup> and 1.5 μmol/l CaM	B	53.2**	60.4	113.5 %
0.5 mmol/l EGTA		37	56.2	150 %
0.1 mmol/l c' AMP		32	37	115.3 %

A — CS isolated from 48-hr-old CEC monolayers to which influenza virus had been adsorbed (CS-V); NAF = normal allantoic fluid;

\* average of 4 measurements

B — Purified influenza virus (1500 HAU/mg CS) or physiological saline were added to CS (CS-V or CS-C). Viral ATPase activity was subtracted from CS-V values.

\*\* average of 3 measurements

For further details see *Mat. and Methods*.

labelled protein bands were seen in SDS-PAGE. In the presence of c-subunit of the c'AMP dependent protein kinase we observed  $^{32}\text{P}$  incorporation into several viral proteins. The 97 kD polypeptide was not labelled in CS-V and the amount of the labels in the 41 kD CS-V polypeptide was decreased; in the presence of  $\text{Ca}^{2+}/\text{CaM}$  no such CS-V band became labelled. The 34 kD band was especially prominent when CS-V was phosphorylated in the presence of  $\text{Ca}^{2+}/\text{CaM}$  and the c'AMP dependent protein kinase inhibitor.

To determine whether phosphorylation of viral components may influence the virion morphology, we investigated by electron microscopy two previously purified virion samples incubated under similar conditions with the c-subunit and with ATP. Virus samples incubated with the c-subunit only were used as controls; another controls were the virus incubated with ATP or with the buffer above. In parallel, the  $^{32}\text{P}$  incorporation from ( $\gamma$ )  $^{32}\text{P}$ -ATP was checked in the presence of c-subunit.

Incubation of influenza A/WSN in phosphorylation buffer in the presence of ATP (Fig. 2-II) did not cause morphologic changes of virions. They retained a spherical shape with a small pleomorphic variability, showed neither changes in electron density nor in the distribution of haemagglutinin and

neuraminidase spikes. Only a few particles revealed a clear-cut envelope structure. Viral particles incubated with the c-subunit (Fig. 2-III) and those incubated with the c-subunit in the presence of ATP (Fig. 2-IV) showed more evident morphologic alterations as compared with control specimens (Figs. 2-I; 2-II). Virions became more pleomorphic, more electron translucent due to rarefaction of the haemagglutinin and neuraminidase spikes. All particles incubated with the c-subunit in the presence of ATP showed more prominent envelope structure and an enlarged volume.

Fig. 3 shows (in accord with previous experiments) that viruses lacking the c-subunit did not incorporate  $^{32}\text{P}$  from  $(\gamma)$   $^{32}\text{P}$ -ATP. In contrast, several virion components were phosphorylated in the presence of the c-subunit. Whether these changes participate in the uncoating of virions remains unclear.

### *Cytoskeleton contraction*

The biochemical correlate of CS contraction are superprecipitation and activation of actinomyosin ATPase (Hidaka *et al.*, 1979; Kanamori *et al.*, 1981). The  $\text{Ca}^{2+}/\text{CaM}$  stimulates the contraction of isolated CS (Rüegg and Paul, 1982; Rüegg *et al.*, 1984; Masuda *et al.*, 1984), while c'AMP and CaM antagonists cause CS relaxation (Nagai *et al.*, 1985; Hidaka *et al.*, 1979; Rüegg and Paul, 1982; Meisner and Rüegg, 1983; Sobue *et al.*, 1985). As shown by preliminary experiments, CS prepared from 48-hr-old CEC monolayers is sensitive to  $\text{Ca}^{2+}/\text{CaM}$ . Tab. 1 shows that such CS (CS-NAF) responded to  $\text{Ca}^{2+}/\text{CaM}$  by increasing its ATPase activity. If influenza virus had been adsorbed to CEC before CS isolation, ATPase activity of CS increased also in the absence of CaM or even in the presence of EGTA. The activity detected in the presence of  $10^{-4}$  mol/l c'AMP was practically identical in CS-NAF and/or CS-V (Tab. 1). Therefore, we assume that the virus stimulated CS contraction.

Similar results were obtained when purified virus was added to already isolated CS (Tab. 1B). CS preparations which did not respond to  $\text{Ca}^{2+}/\text{CaM}$  revealed no enhancement of their ATPase activity not even after adding of the virus (results not shown).

### *Discussion*

The results of this work have shown that  $\text{Ca}^{2+}/\text{CaM}$  increased the ATPase activity of isolated CS. In the CS isolated from chick embryo fibroblasts with adsorbed virus (CS-V) the specific activity of ATPase has been increased in the absence of CaM and even in the presence of EGTA. We believe that after adsorption to corresponding cell receptors influenza virus causes reorganization of CS leading to its contraction.

The prerequisite of  $\text{Ca}^{2+}$  dependent contraction in the CS model prepared from human fibroblasts is the phosphorylation of myosin light chain (MLC) by the  $\text{Ca}^{2+}/\text{CaM}$  dependent MLC-kinase (Masuda *et al.*, 1984). The necessity of MLC phosphorylation has been described by several authors.

Electrophoresis under our conditions did not allow to determine the  $M_r$  of proteins when lower than 23 kD. In the CS isolated from CEC plasma membranes with adsorbed virus (CS-V) these polypeptides became phosphorylated also in the absence of CaM, but in the presence of  $Ca^{2+}$ /CaM their phosphorylation has more increased. The surface delineated by the phosphorylation curve in the presence of c'AMP protein kinase was 0.11, while that of outlined by the phosphorylation curve in the presence of  $Ca^{2+}$ /CaM was 0.4. In the CS isolated from noninfected plasma membranes (CS-C) the surface outlined by c'AMP kinase phosphorylation was 0.05 and that by  $Ca^{2+}$ /CaM phosphorylation 0.09. We assume that this fraction contains the phosphorylated MLC important for the  $Ca^{2+}$ -dependent contraction.

In addition to MLC phosphorylation,  $Ca^{2+}$  related contraction depends on the organization of actin filaments (Coleman and Mooseker, 1985). Several  $Ca^{2+}$ -sensitive actin binding proteins have been described which confer  $Ca^{2+}$  sensitivity for the regulation of filamentous or network CS structures. Just recently, the CaM-binding and/or CS-binding activity of two polypeptides, 34 kD and 35–36 kD has been characterized (Takahashi *et al.*, 1986; Sobue *et al.*, 1987). It should be cleared, however, whether the increased incorporation of  $^{32}P$  into polypeptides 27 kD, 35 kD and 37 kD in the presence of  $Ca^{2+}$ /CaM is associated with a similar activity.  $Ca^{2+}$  induced CS contraction may be reversed (relaxed) by c'AMP, their analogues and by substances enhancing cellular c'AMP levels (Rüegg and Paul, 1982).

Interaction actin-myosin (contraction — relaxation) may be regulated also by phosphorylation — dephosphorylation of certain actin-binding proteins. Relaxation occurs when the 150 kD protein becomes phosphorylated to bind actin and myosin (Hartzell and Glass, 1984). In turn, phosphorylation of caldesmon ( $M_r$  70 kD or 150 kD) hinders its interaction with myosin and actin (Horiuchi *et al.*, 1986). A good substrate for the c'AMP dependent protein kinase as well as for protein kinase C is the membrane skeleton-associated calmodulin-binding protein (CaM-BP 103/97) (Ling *et al.*, 1986). It binds to the spectrin-actin complex and enhances the capacity of actin filaments to interact with spectrin. The significance of its phosphorylation is unknown, however. In the presence of c'AMP and EGTA or in the presence of the c-subunit of the c'AMP dependent protein kinase we found that proteins of the infected cytoskeleton (CS-V) of apparent  $M_r$  170 kD, 100 kD, 85 kD and 41 kD were phosphorylated; their role in the reorganization of CS is still unknown. Selective phosphorylation of proteins by proteinase C is considered for a mechanism transferring environmental signals to cell interior (Pontremoli *et al.*, 1987).

Several authors found that protein kinase C is translocated from the cytosol to the plasma membranes during cell activation. The membranes release a neutral protease ( $Ca^{2+}$ -activated protease called calpain) which cleaves protein kinase C and transforms it to a new form, the C-M protein kinase. This is fully active in the absence of phospholipids and  $Ca^{2+}$  and does not bind to the plasma membranes (Pontremoli *et al.*, 1987). CS extracted with Triton X-100 is an unsuitable substrate for protein kinase C but a good substrate for protein kinase C-M, which phosphorylates more CS proteins. We found an en-

hanced phosphorylation of CS proteins in the virus-infected CS even in the presence of EGTA or of the c'AMP dependent protein kinase inhibitor.

Based on our previous observations (Križanová *et al.*, 1976; Križanová *et al.*, 1982) we believe that influenza virus at early intervals post-adsorption activates the adenylate cyclase system increasing the level of cellular c'AMP, which, in turn, activates the c'AMP dependent protein kinase. Phosphorylation by means of the latter enzyme opens the calcium channel (Hescheler *et al.*, 1987) and  $\text{Ca}^{2+}$  ions enter the cytoplasm. In addition, phosphorylation of other membrane proteins and of actin binding proteins may translocate the CaM from the membrane to cytosol (phosphorylated CaM binding proteins show a lower affinity to CaM) (Malenčík, 1982) and interrupt the association of CS with actin binding proteins in the vicinity of plasma membrane allowing rearrangement of CS subunits (or of actin microfilaments). Intracellular  $\text{Ca}^{2+}$ /CaM activates the MLC kinase which phosphorylates MLC as well as other actin binding proteins (C-M protein kinase may be phosphorylated as well). Stress (contractory) filaments are being formed allowing ATP-associated contraction important for signal transfer.

The proposed scheme may reflect early postinfection events; its validity should be investigated using specific antiglobulins.

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*Legends to the Figures (pages 203–205):*

*Fig. 2. Electron microscopy of phosphorylated virions.*

- I — Influenza virus A/WSN in the phosphorylation buffer, contrasted with 2 % PTA, magn.  $\times 99\,000$ .
- II — Influenza virus A/WSN incubated in the presence of 0.57 mmol/l ATP; magn.  $\times 99\,000$ .
- III — Influenza virus A/WSN incubated with 30 units of the c-subunit; magn.  $\times 99\,000$ .
- IV — Influenza virus A/WSN incubated in the presence of 30 units of the c-subunit and of 0.57 mmol/l ATP; magn.  $\times 99\,000$ .

*Fig. 3. Phosphorylation of influenza virus in the presence or absence of the c-subunit of c'AMP-dependent protein kinase.*

Two purified influenza A/WSN (H1N1) samples were tested

a, b — purified virions stored at 4 °C for 7 months

c, d — purified virions stored for 7 days (at 4 °C)

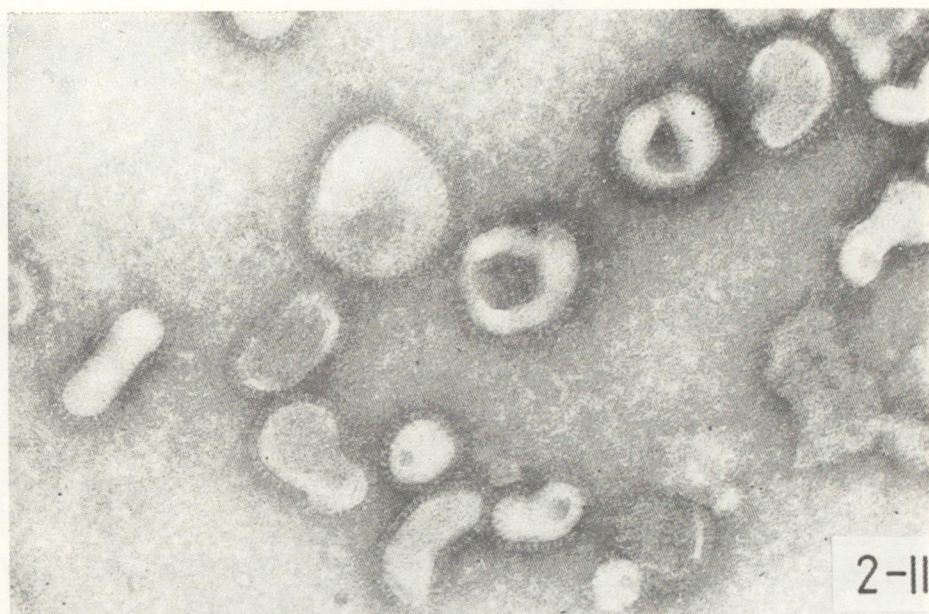
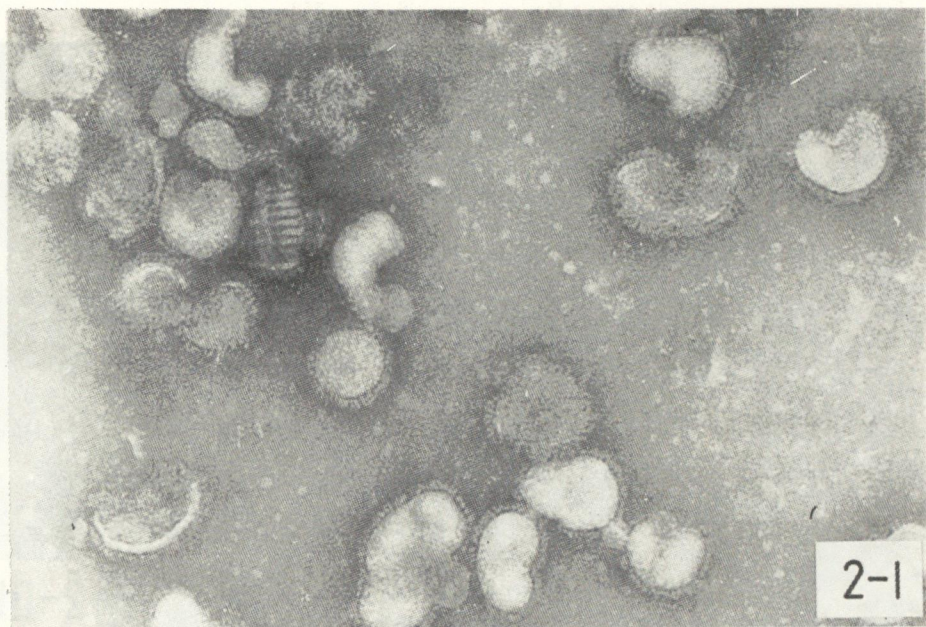
For phosphorylation conditions see Legend to Fig. 1.

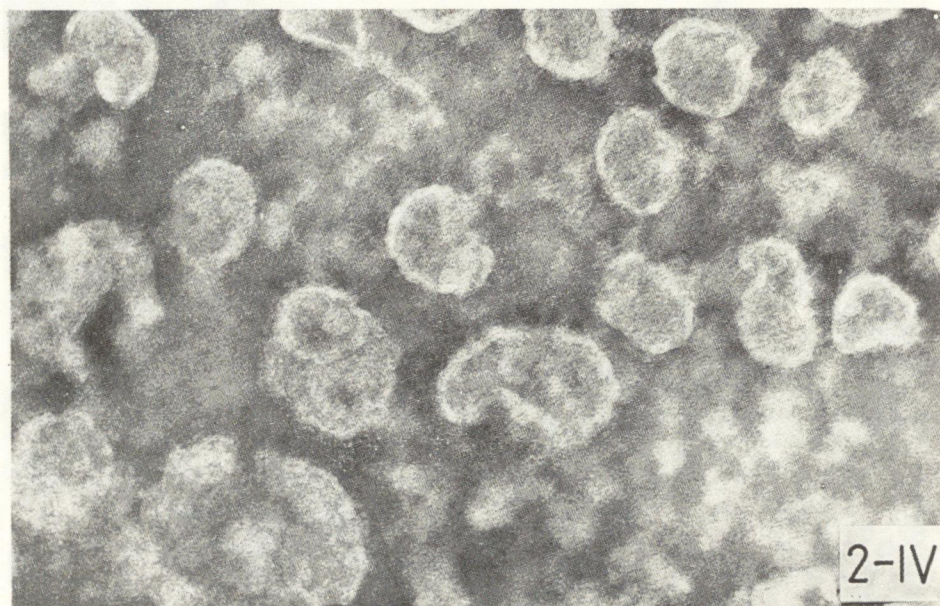
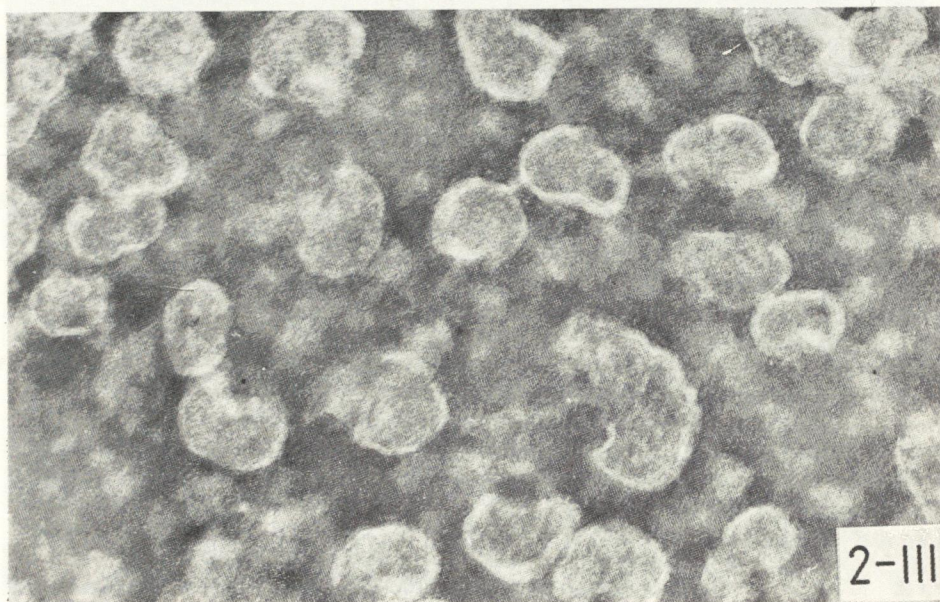
As the c-subunit had been dissolved in 100 mmol/l DDT, the same amount of DDT was added to the virus, phosphorylated in the absence of the c-subunit.

a, c — virus autophosphorylation in the absence of the c-subunit

b, d — virus phosphorylation in the presence of 30 units of the c-subunit

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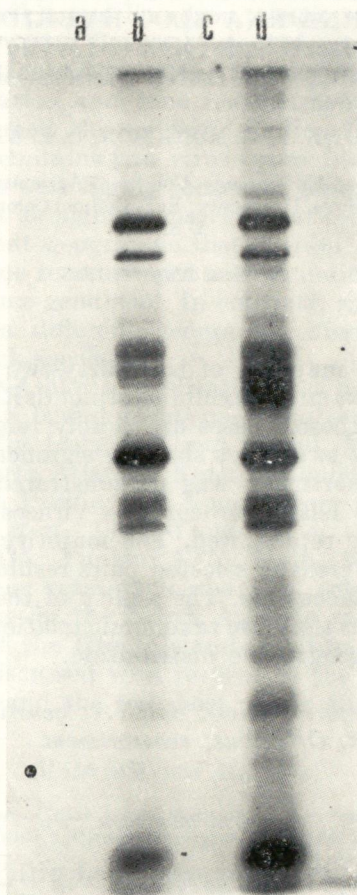


Fig. 3

For legend see page 202.