Phosphorylation of elongation factor 1 in polyribosome fraction of rabbit reticulocytes

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A single protein, $M_r \sim 50000$, is shown to be phosphorylated during incubation of a mono- and polyribosome fraction of rabbit reticulocytes with [y-32P]ATP at a low ionic strength. This protein has been identified as the elongation factor 1α (EF- 1α). The phosphorylated EF- 1α , in contrast to the unmodified factor, is not detected in complexes with mono- and polyribosomes. It is suggested that the phosphorylation of EF-1α can result in its decompartmentation from polyribosomes and thus affect the rate of protein synthesis.

Elongation factor 1

Protein phosphorylation Compartmentation-decompartmentation

Polyribosome

Translation regulation

1. INTRODUCTION

It is known that reversible phosphorylation of cellular proteins can act as a physiological mechanism regulating their functions. The initiation factor 2 [1], ribosomal protein S6 [2] as well as several aminoacyl-tRNA synthetases [3,4] have been shown to be among the components of the eukaryotic translation machinery subjected to phosphorylation. Phosphorylation of eIF-2 leads to inhibition of protein synthesis [5]. The regulatory role of the phosphorylation of. ribosomal protein S6 and that of aminoacyl-tRNA synthetases has also been discussed [2-4].

have shown that intensive we phosphorylation of the elongation factor 1α (EF-1 α) is observed in vitro in the fraction of mono- and polyribosomes of rabbit reticulocytes at low ionic strength. We believe that the phosphorylation of EF-1 α can play a regulatory role in the cell.

2. MATERIALS AND METHODS

Reticulocytes were obtained from blood of rabsubcutaneous injection phenylhydrazine according to [6]. The cells were lysed with an equal volume of 5 mM MgCl₂ and the lysate was centrifuged at 12000 rpm for 15 min in a JA-14 rotor of a J-21B centrifuge (Beckman) to remove cell debris and mitochondria.

EF-1 was isolated according to a slightly modified procedure of [7].

The fraction of mono- and polyribosomes was prepared by gel filtration of the mitochondria-free extract through a column with Sephacryl S-300 (Pharmacia) equilibrated with a standard buffer solution containing 10 mM Tris-HCl (pH 7.6), 1 mM MgCl₂. The preparation was hemoglobinfree and contained about 90% of ribosomes of the initial extract.

Protein phosphorylation was performed by incubating the samples with $[\gamma^{-32}P]ATP$ 1000 Ci/ mmol (Isotope, USSR) for 5 min at 37°C.

To determine the degree of phosphorylation, the protein was precipitated with 5% trichloroacetic acid, collected on GF/C (Whatman) filters and the radioactivity counted.

Protein electrophoresis was carried out in a polyacrylamide gradient gel (10-22% acrylamide) in the presence of 0.1% SDS [8].

Two-dimensional separation of proteins was done according to a modified procedure of [9]. Due to strong phosphatase activities a phosphatase inhibitor ZnCl₂ was added to the preparations to a concentration of 10 mM. Before radioautography the gels were incubated with 1 M NaOH at 60°C for 1 h.

To obtain a radioautograph of the labeled proteins the gels were dried and exposed overnight at -70°C using an RM-V film (Tasma, USSR).

Centrifugation of the preparations was carried out in an isokinetic 15-33.5% sucrose gradient made in a standard buffer at 40000 rpm for 75 min at 4°C in an SW-41 rotor.

3. RESULTS

The fraction of mono- and polyribosomes obtained by gel-filtration of the mitochondria-free

A → 96000 → 50000 → 34000

Fig.1. Radioautograph showing phosphorylation of proteins in the mono- and polyribosome fraction at 150 mM KCl (A) and under conditions of low ionic strength (B). 100-μl samples contained 0.5 A₂₆₀ units of the polyribosome fraction and 1 μCi [γ-³²P]ATP.

extract through Sephacryl S-300 equilibrated with a standard buffer (low ionic strength) contains a considerable amount of labile associated RNA-binding proteins. Both the elongation factors 1 and 2, as well as aminoacyl-tRNA synthetases, have been found among them.

A broad range of different proteins is shown to be phosphorylated during incubation of the fraction of mono- and polyribosomes with $[\gamma^{-32}P]ATP$ in the standard buffer with 150 mM KCl (fig.1A).

Incubation of this preparation with $[\gamma^{-32}P]ATP$ in the standard buffer without KCl demonstrates that under these conditions the radioactive label is incorporated into only one polypeptide with a molecular mass of about 50 kDa (fig.1B, 2A)

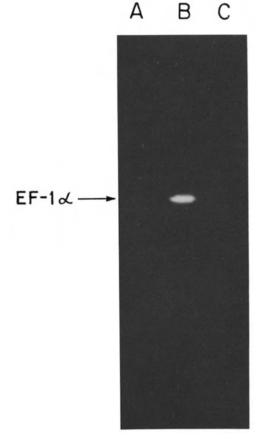


Fig.2. Radioautograph showing phosphorylation of proteins in the fraction of mono- and polyribosomes (A), in the preparation of the EF-1_H (C) and in their mixture (B) at low ionic strength. 100- μ l samples contained 0.05 A_{260} units of the polyribosome fraction (A and B), 10 μ g of EF-1_H (B and C) and 1 μ Ci [γ -³²PlATP.

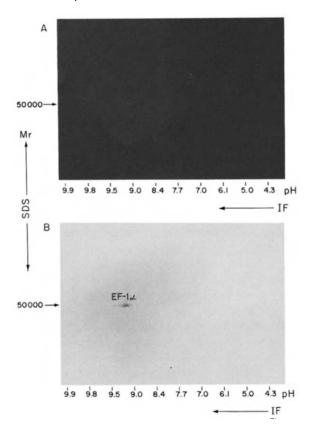


Fig.3. Two-dimensional analysis according to [7] of proteins from the mono- and polyribosome fraction incubated with $[\gamma^{-32}P]ATP$ at low ionic strength (radioautograph) (A) and of the EF-1_L (stained with Coomassie) (B).

whose position in the gel corresponds to that of the α -subunit of EF-1.

Incorporation of the radioactive label into the $M_r \sim 50000$ polypeptide band is considerably increased upon the addition of a preparation of EF-1_H to the incubation mixture (fig.2B). EF-1 α is practically not phosphorylated during control incubation of an EF-1_H preparation (95% purity) with $[\gamma^{-32}P]$ ATP (fig.2C).

Two-dimensional analysis according to [7] corroborated the identity of the approx. 50-kDa polypeptide, phosphorylated in the mono- and polyribosome fraction, to the α -subunit of EF-1 (fig.3).

Centrifugation of the mono- and polyribosome fraction in the sucrose gradient and following incubation with $[\gamma^{-32}P]$ ATP of individual fractions,

has shown that the radioactive label is incorporated into the trichloroacetic acid-insoluble product in both the poly- and monoribosomes (fig.4A). Electrophoresis and radioautography demonstrate that the label in all the individual fractions of the gradient is present in the same approx. 50-kDa polypeptide which has been identified as EF-1\alpha.

When an excess of exogenous high molecular mass RNA is added to the mono- and polyribosome fraction before centrifugation, the incorporated radioactive label is found in the postribosomal zone where exogenous RNA is distributed (fig.4B). This result correlates with the observation on the displacement of EF-1 from polyribosomes by an excess exogenous RNA [10].

Sucrose gradient centrifugation of the monoand polyribosome fraction preincubated with $[\gamma^{-32}P]$ ATP shows that all the acid-insoluble radioactive material is distributed at the top of the gradient and is absent from the zone of mono- and polyribosomes (fig. 4C).

4. DISCUSSION

Several proteins of the eukaryotic translation machinery, including the elongation factors and aminoacyl-tRNA synthetases, are known to possess a non-specific affinity for high molecular mass RNAs. This affinity provides for their labile association with mono- and polyribosomes [11].

It is proposed that the affinity for RNA serves for partial compartmentation of the proteins of the translation machinery in places of their functioning in the big volume of the eukaryotic cell [12] while a change of this affinity may serve for regulation of protein synthesis at the translation (elongation) level [11]. In particular, inhibition of protein synthesis after ADP-ribosylation of EF-2 was shown to be accompanied by a loss of its nonspecific affinity for RNA and its dissociation from loose complexes with monopolyribosomes [13]. It has also been found that phosphorylation of several RNA-binding proteins with non-identified functions changes their affinity for RNA [14].

Here, we have shown that EF-1 can be phosphorylated at the α -subunit. It is noteworthy that the kinase phosphorylating EF-1 α , just as EF-1 α itself, is found in association with mono-

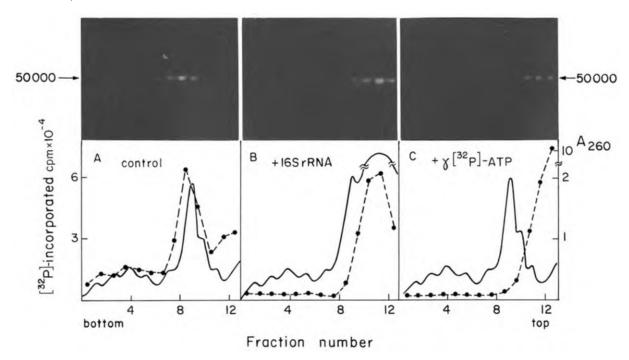


Fig. 4. Sedimentation distribution of the material of the mono- and polyribosome fraction at sucrose gradient centrifugation. A, B, $1 \mu \text{Ci} \ [\gamma^{-32}\text{P}]\text{ATP}$ was added to each gradient fraction after centrifugation; B, $20 \ A_{260}$ units of Escherichia coli 16 S ribosomal RNA were added to the polyribosome fraction before centrifugation; C, the polyribosome fraction was incubated with $13 \mu \text{Ci} \ [\gamma^{-32}\text{P}]\text{ATP}$ before centrifugation. Gel electrophoresis of the labeled proteins from the corresponding gradient fractions (radioautographs) are given at the top.

and polyribosomes. Activation of the kinase in vitro results from a decrease of the ionic strength of buffer solution and its action under these conditions is strictly selective relative to the α -subunit of EF-1. The phosphorylated form of EF-1, in contrast to the unmodified EF-1, has not been found in complexes with mono- and polyribosomes. We believe that the phosphorylation of EF-1 results in its dissociation from complexes with mono- and polyribosomes, probably due to a loss of affinity for RNA.

It is not unlikely that the phosphorylation of EF- 1α may affect the rate of polypeptide chain elongation and serve for regulation of the translation process in the cell.

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