

# Heat-stable translational inhibitor from rabbit reticulocyte lysates

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We have purified to apparent homogeneity a heat-stable (HS) factor from the postribosomal supernatant of rabbit reticulocyte lysates [(1988) FEBS Lett. 236, 479–483]. HS inhibits translation in hemin-supplemented lysates and induces phosphorylation of the  $\alpha$ -subunit of the eukaryotic initiation factor 2 as does hemin deficiency. The translational inhibition produced by addition of HS to hemin-containing reticulocyte lysates and the accompanying phosphorylation of the eIF-2 $\alpha$  subunit can be prevented or reversed by NADPH generators including glucose 6-phosphate, NADPH itself, and also by dithiols, e.g., dithiothreitol, but not by fructose 1,6-bisphosphate or by monothiols, e.g., 2-mercaptoethanol. When added to crude preparations of the proinhibitor form (proHCl) of the heme-controlled translational inhibitor (HCl), HS produces a pronounced increase of the HCl to proHCl ratio. It appeared possible that HS might be oxidized glutathione (GSSG) but this is not the case, for HS is not a substrate for highly purified glutathione reductase from rabbit erythrocytes. The spectral analysis of highly purified HS is consistent with the idea that HS could be a nucleotide derivative.

Polypeptide chain initiation; Translational inhibition; eIF-2 $\alpha$  kinase; HCl activation

## 1. INTRODUCTION

Protein synthesis in reticulocytes is regulated at the level of the eukaryotic polypeptide chain-initiation factor 2 (eIF-2) [1,2]. This factor forms a ternary complex with GTP and eukaryotic initiator methionyl tRNA (Met-tRNA<sub>i</sub>), which then binds to a 40 S ribosomal subunit. Recycling of eIF-2 is catalyzed by GEF, the GDP-GTP exchange factor, and is inhibited when the eIF-2 $\alpha$  subunit is phosphorylated by cAMP-independent protein kinases [3–5]. One of these kinases, the heme-controlled translational inhibitor (HCl), is activated in lysates incubated in the absence of added hemin or, with hemin present, upon addition of either oxidized glutathione (GSSG) [6] or Ca<sup>2+</sup>-phospholipid [7]. In reticulocyte lysates HCl is present largely in the form of an inactive precursor, proHCl. Once activated, HCl inhibits translation in hemin-containing reticulocyte lysates, but the mechanism of activation remains obscure (for reviews see [1 and 2]).

We have previously reported that activation of pro-

HCl in hemin-supplemented lysates can be elicited in a reversible fashion by Ca<sup>2+</sup> and phospholipid (PL) [7]. Like the activation of HCl in lysates by GSSG or NADPH depletion [8], the activation by Ca<sup>2+</sup> and PL could be largely prevented or reversed by NADPH-generating systems or by dithiols [9].

Recently, we have described the purification of a novel heat-stable translational inhibitor from postribosomal supernatant of rabbit reticulocyte lysates (HS) [10]. HS factor inhibits translation in hemin-supplemented lysates by promoting phosphorylation of the  $\alpha$ -subunit of eIF-2. According to the spectrum, the HS could be a nucleotide analogue and in this case, the concentration of HS required for inhibition of protein synthesis would be in the nanomolar range. Our results are consistent with the notion that HS activates the heme-stabilized proinhibitor form (proHCl) of HCl in reticulocyte lysates. Like the activation of HCl in lysates by GSSG or Ca<sup>2+</sup>-PL [6,7], the activation by HS may largely be due to oxidation of labile SH groups which are involved in the activation of HCl.

## 2. MATERIALS AND METHODS

### 2.1. Materials

G6P, FDP, GSSG and NEM were purchased from Sigma. All resins and the HPLC equipment were used earlier [10].

### 2.2. Assays

HS was assayed by its inhibitory effect on translation and its capacity for activating HCl in hemin-containing reticulocyte lysates. The effect is proportional to HS concentration up to about 70% inhibition. Translation and eIF-2 $\alpha$  phosphorylation assays were as described [7,8]. Activation of proHCl was assayed by phosphorylation of the eIF-2 $\alpha$  subunit as described in the legend to fig.3.

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*Abbreviations:* eIF-2, eukaryotic polypeptide chain-initiation factor 2; Met-tRNA<sub>i</sub>, eukaryotic initiator methionyl-tRNA; HCl, heme-controlled translational inhibitor (an eIF-2 $\alpha$  kinase); proHCl, the proinhibitor (inactive) form of HCl; HS, heat-stable translational inhibitor from rabbit reticulocyte lysates; DTT, dithiothreitol; G6P, glucose 6-phosphate; FDP, fructose 1,6-bisphosphate; NEM, *N*-ethylmaleimide; GSSG, oxidized glutathione; GEF, guanine nucleotide exchange factor; PL, phospholipid; H<sup>+</sup>-NMR, proton nuclear magnetic resonance

Glutathione reductase was assayed spectrophotometrically by following the oxidation of NADPH by GSSG at 340 nm.

### 2.3. Preparations

Reticulocyte lysates were prepared from phenylhydrazine-treated rabbits by the method of Hunt et al. [11] with slight modifications. Initiation factor eIF-2 and HCl, both from rabbit reticulocyte lysates, were purified as described [12]. Labeled ATP was prepared as described by Schendel and Wells [13]. Crude proHCl was prepared by the Gross and Rabinovitz procedure [14] as described [15]. Hemin was added to the CM-Sephadex fraction (25  $\mu$ M final concentration) and this was stored in small aliquots in liquid nitrogen. Highly purified glutathione reductase from rabbit erythrocytes was the gift of Dr C. Palomo of this laboratory.

## 3. RESULTS

### 3.1. Translational inhibition by HS

HS was purified from postribosomal supernatant of 400 ml of rabbit reticulocyte lysates as described [10]. Fig. 1A confirms our previous work [10] in that the HS activity was eluted from the last HPLC size-exclusion column of a sharp, symmetric peak, with a maximum of absorbance at 256 nm (fig. 1A, inset). The translation inhibitory activity also coincided precisely with the peak of absorbance (not shown) [10]. HS appears to be a potent inhibitor of translation in reticulocyte lysates. Inhibition is proportional to HS concentration up to about 70% (fig. 1B). The amount of highly purified HS causing 50% inhibition of hemin-dependent translation was  $0.7 \times 10^{-3}$   $A_{260}$  units (in 30  $\mu$ l assay volume). Like the translational inhibition due to heme deficiency or HCl addition to hemin-containing lysates, HS produces increased phosphorylation of the eIF-2 $\alpha$  subunit in hemin-supplemented lysates (fig. 1C). At high HS concentrations, this phosphorylation is no greater than that caused by hemin deficiency (cf. lane 1 with 3 and 4).

### 3.2. Translational inhibition by HS is prevented or reversed by NADPH generators and dithiothreitol

Like the inhibition by  $Ca^{2+}$  and phospholipid [9], the inhibitory effect of HS on translation in hemin-containing lysates is largely prevented by glucose 6-phosphate, NADPH, or dithiothreitol but not by 2-mercaptoethanol or fructose 1,6-bisphosphate that can neither be converted to G6P nor metabolized via the hexosemonophosphate shunt. However, the inhibition caused by heme deficiency is not prevented or restored by NADPH generators or by dithiols (table 1).

Fig. 2A suggests that translational inhibition produced by HS is oxidative in nature because the HS-promoted phosphorylation of the eIF-2 $\alpha$  subunit in hemin-containing lysates (cf. lanes 2 and 3) is largely prevented by dithiothreitol (lane 6), NADPH (lane 7), or glucose 6-phosphate (lane 8) but not by fructose 1,6-bisphosphate (lane 4) or 2-mercaptoethanol (lane 5). Moreover, this phosphorylation is reversed by glucose 6-phosphate and dithiothreitol (fig. 2B). Already at 5 min of incubation, the degree of phosphorylation of the eIF-2 $\alpha$  subunit is greatly increased by HS (lane 3) as compared with control lysates (lane 2). Clearly, the phosphorylation of this band is significantly diminished if the lysates are supplemented with either G6P or DTT at 5 min of incubation (cf. lane 6 with 10 and 11). The activation of HCl by heme deficiency is not prevented by G6P (lane 5). As seen before (fig. 2A) the effect of highly purified HS is largely prevented by G6P (lane 8) or DTT (lane 9).

### 3.3. Activation of HCl by HS and GSSG

A likely explanation for the inhibition of translation in hemin-containing lysates by HS is that it promotes the first step of the reaction  $\text{proHCl} \rightarrow \text{reversible HCl} \rightarrow \text{irreversible HCl}$ . To verify further this idea, we look-

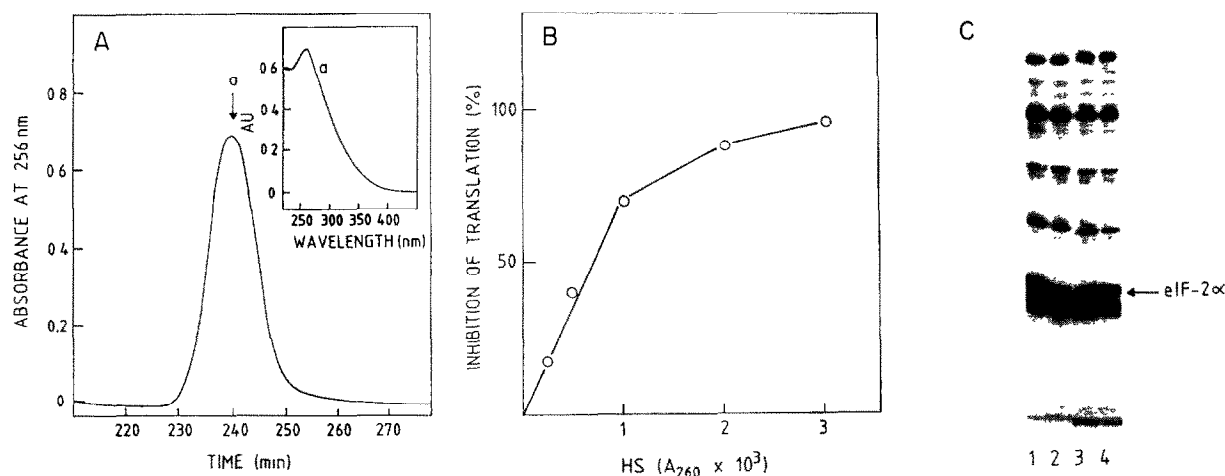


Fig. 1. (A) Rechromatography of the HS activity in the size exclusion HPLC column [10]. (Inset). Automatic overlay of spectra acquired from peak maxima (a) from 220 nm to 450 nm. (B) Increasing amounts of purified HS were assayed for translational inhibition. (C) Effect of highly purified HS on phosphorylation of the eIF-2 $\alpha$  subunit. The conditions were (lanes): (1) without hemin; (2) with 20  $\mu$ M hemin; (3) with hemin and  $2 \times 10^{-3}$   $A_{260}$  of HS; (4) with hemin and  $4 \times 10^{-3}$   $A_{260}$  of HS.

Table 1

Effect of glucose 6-phosphate (G6P), NADPH, and dithiothreitol (DTT) on translational inhibition by HS in hemin-supplemented reticulocyte lysates

Additions	[ $^{14}$ C]Leucine incorporation (cpm $\times 10^{-3}$ )		Relative hemin-dependent activity (%)
	Total	Due to hemin	
None	14.3		
G6P (1 mM)	14.8		
NADPH (1 mM)	14.5		
DTT (1 mM)	13.3		
2-mercaptoethanol (2-ME) (1 mM)	14.3		
Fructose 1,6-bisphosphate (FDP) (1 mM)	14.3		
Hemin (25 $\mu$ M)	70.1	55.8	100
+ HS <sup>a</sup> ( $6 \times 10^{-3}$ A <sub>260</sub> )	18.8	4.5	8
+ HS + G6P	60.3	45.5	81
+ HS + NADPH	51.2	36.7	66
+ HS + DTT	64.6	51.3	92
+ HS + 2-ME	17.6	3.3	6
+ HS + FDP	19.8	5.5	10

<sup>a</sup> HS from Sephadex G-15 [10] (peak I) was used

ed at the effect of HS on a crude preparation of proHCl (fig.3). Clearly, this preparation contains the inactive precursor form of HCl (lane 1) that is fully converted to HCl by NEM (lane 2) [1,2]. Incubation of crude proHCl with highly purified HS (lane 5) or with peak III of HS from Sephadex G-15 [10] (lane 3) results in HCl increase as judged by the increased phosphorylation of

the eIF-2 $\alpha$  subunit, but to produce this effect, one needs larger amounts of HS than are required with lysates. Curiously, while GSSG activates proHCl in lysates, it is less effective on partially purified proHCl (lane 4). An amount of GSSG that promotes HCl activation in hemin-supplemented lysates is not sufficient to activate crude proHCl (data not shown). This may

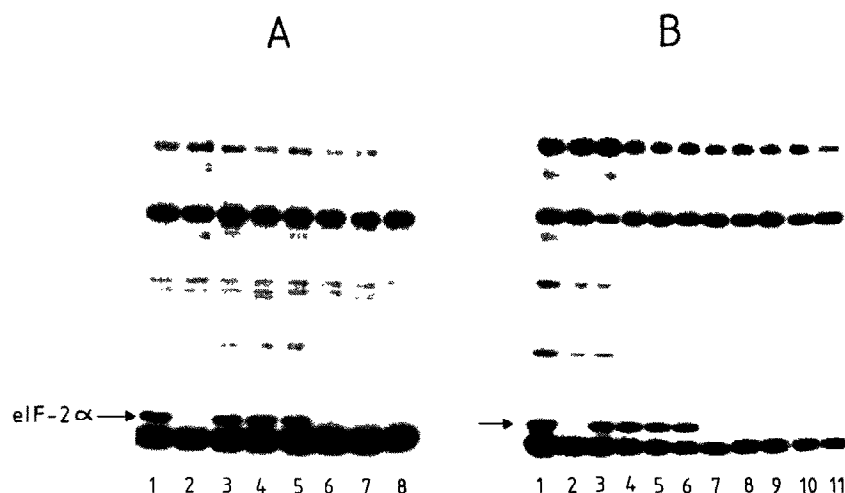


Fig.2. (A) HS-promoted phosphorylation of the 38 kDa  $\alpha$ -subunit of eIF-2 and the effect of various compounds thereon. Samples were incubated for 60 min at 30°C and aliquots were assayed for protein synthesis and phosphorylation as described. The hemin-dependent incorporation of [ $^{14}$ C]leucine (in cpm  $\times 10^{-3}$  per sample) is indicated in parentheses for each lane. Lanes: 1, without hemin; 2, with 20  $\mu$ M hemin (32.0); 3, with hemin and  $6 \times 10^{-3}$  A<sub>260</sub> of HS from Sephadex G-15 [10] (peak I) (3.4). Samples corresponding to lanes 4-8 all contained hemin and HS and, in addition, contained the following: 1 mM FDP (3.0) (lane 4); 1 mM 2-mercaptoethanol (2.6) (lane 5); 1 mM DTT (37.5) (lane 6); 1 mM NADPH (31.5) (lane 7); 1 mM G6P (30.1) (lane 8). (B) Phosphorylation of the 38 kDa  $\alpha$ -subunit of eIF-2 promoted by highly purified HS is reversed by glucose 6-phosphate or dithiothreitol. Samples were incubated for 40 min at 26°C and aliquots were assayed for protein synthesis as described. Aliquots (3  $\mu$ l) for the phosphorylation assay were taken as follows: lanes 1-3, at 5 min; lanes 4-11, at 40 min. The corresponding 40 min translation values are indicated in parentheses. Lanes: 1, without hemin; 2, with hemin; 3, with hemin and  $2 \times 10^{-3}$  A<sub>260</sub> of highly purified HS; 4, without hemin (10.0); 5, with 1 mM G6P (10.3); 6, with hemin and HS (11.1); 7, with hemin (45.3). Samples corresponding to lanes 8-11 all contained hemin and HS from the start of incubation and, in addition, contained the following: 1 mM G6P (39.3) (lane 8) or 1 mM DTT (48.0) (lane 9) from the start and 1 mM G6P (34.1) (lane 10) or 1 mM DTT (42.7) (lane 11) added at 5 min.

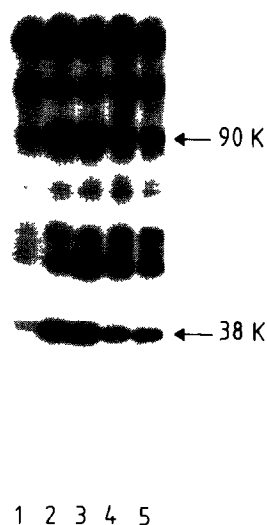


Fig.3. Effect of HS<sub>r</sub> on activation of HCl. Samples (30  $\mu$ l) containing 25 mM Tris-HCl, pH 7.6; 25 mM KCl, with proHCl (40  $\mu$ g of protein) and when present, 2 mM MEM, 2 mM GSSG;  $40 \times 10^{-3}$  A<sub>260</sub> of HS from Sephadex G-15 [10] (peak III) or  $6 \times 10^{-3}$  A<sub>260</sub> of highly purified HS, were incubated for 15 min at 34°C. The excess NEM was neutralized with 4 mM DTT. Aliquots of 6  $\mu$ l were used as a source of eIF-2 $\alpha$  kinase source in a second incubation for 9 min at 30°C in a solution (20  $\mu$ l final volume) containing 25 mM Tris-HCl, pH 7.6, 5 mM Mg(OAc)<sub>2</sub>, and 0.1 mM [ $\gamma$ -<sup>32</sup>P]ATP ( $30 \times 10^6$  cpm/sample). The 38 kDa bands of the autoradiograms were scanned at 626 nm with a Chromoscan 3 densitometer and the values are indicated in parentheses for each lane. Lanes: 1, proHCl (8  $\mu$ g of protein) (3267); 2, proHCl and NEM (11 616); 3, proHCl and HS (peak III) (12 969); 4, proHCl and GSSG (8322); 5, proHCl and highly purified HS (9085). Arrows show the position of the 90 and 38 kDa bands determined in a parallel run in which a partially purified HCl was used (not shown).

suggest that an unknown component (limiting in crude proHCl preparations) may be involved in the activation of HCl by both HS and GSSG.

#### 4. DISCUSSION

Our results provide further support for the view [10] that a homogeneous heat-stable factor (HS) from rabbit reticulocyte lysates is a potent translational inhibitor. Like GSSG, HS inhibits translation in hemin-supplemented reticulocyte lysates by activation of HCl, resulting in increased phosphorylation of the eIF-2 $\alpha$  subunit, but it appears to be much more active. Unlike GSSG, HS is not a substrate for highly purified glutathione reductase from rabbit erythrocytes (table 2). Recently, we have reported that HS appears to be neither a peptide nor an oligonucleotide since HS activity was insensitive to proteolytic or nucleolytic digestion [10]. The spectral analysis of highly purified HS ([10] and fig.1A) is consistent with the idea that HS could be a nucleotide analogue. The molar extinction coefficients of nucleotides (260 nm) are about  $1 \times 10^4$ . Therefore, the amount of the highly purified HS required for 50% inhibition of protein synthesis ( $7 \times 10^{-4}$  A<sub>260</sub>

unit)) is in the nanomolar range (70 nM). Thus, HS may be an extremely active inhibitor of protein synthesis. In attempting to identify the structure of HS, we have studied the effect of the nucleoside diphosphates (ADP, GDP, CDP and UDP) with compatible molecular masses and similar but not identical spectra (not shown). All of them inhibited protein synthesis. The concentration causing 50% inhibition in hemin-dependent translation was about 500  $\mu$ M in either case. This inhibition, however, was not accompanied by increased phosphorylation of eIF-2 $\alpha$  subunit (not shown). Furthermore, the analyses performed by the H<sup>+</sup>-NMR technique were unable to identify the structure of HS, due to the large amount required. Therefore, we have begun with the purification of HS on a preparative scale. On the other hand, using fast atom bombardment (FAB) in mass spectrometry, the molecular weight of HS appears to be 429 (not shown).

Earlier [8] we emphasized that, unlike the activation of HCl in hemin-supplemented lysates by GSSG addition or NADPH depletion, the activation caused by heme deficiency is not prevented by NADPH generators or by dithiothreitol. It was therefore argued that there are at least two fully independent mechanisms for proHCl activation: (i) heme deficiency and (ii) oxidation processes that appear to lead to the conversion of certain SH groups to disulfides [1,16]. We have also shown [9] that proHCl can be reversibly activated via oxidation by agents other than GSSG, namely polyunsaturated fatty acids and phospholipids. HCl may be activated in a number of ways [2], but the mechanisms are unknown. The activation by GSSG is of interest because of its possible physiological significance. Reversibility of the oxidative activation of HCl is a requirement for physiological significance. Although the presence of enzymes capable of phosphorylating the eIF-2 $\alpha$  subunit has been reported in non-erythroid cells [2], their mode of activation and physiological significance are unknown. Thus, although there is no doubt on the physiological relevance of the heme control of HCl activity in reticulocytes, the significance, if any, of the oxidative activation of proHCl remains a matter for speculation.

Table 2

Incubation (min)	HS is not a substrate for glutathione reductase		
	$\Delta$ OD <sub>340</sub>		
	None	GSSG	HS
1	0.006	0.139	0.004
2	0.009	0.308	0.004
3	0.013	0.437	0.003
4	0.015	0.518	0.002
6	0.015	0.575	0.002

The assay was carried out with highly purified glutathione reductase from rabbit erythrocytes. The OD<sub>340</sub> average value at 0 min was 0.602

The high activity of HS, its reversibility, and its presence in unheated lysates (data not shown) would be compatible with the notion that this factor may be involved in translational control.

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## REFERENCES

- [1] Jackson, R.J. (1982) in: Protein Biosynthesis in Eukaryotes. NATO Advanced Study Institute Series, Series A: Life Sciences, vol. 41 (Perez-Bercoff, R. ed.) pp. 363–418, Plenum, New York.
- [2] Ochoa, S. (1983) Arch. Biochem. Biophys. 233, 325–349.
- [3] Matts, R.L., Levin, D.H. and London, I.M. (1983) Proc. Natl. Acad. Sci. USA 80, 2559–2563.
- [4] Siekierka, J., Manne, V. and Ochoa, S. (1984) Proc. Natl. Acad. Sci. USA 81, 352–356.
- [5] Thomas, N.S.B., Matts, R.L., Petryshyn, R. and London, I.M. (1984) Proc. Natl. Acad. Sci. USA 81, 6998–7002.
- [6] Kosower, N.S., Vanderhoff, G.A. and Kosower, E.M. (1972) Biochim. Biophys. Acta 272, 623–637.
- [7] De Haro, C., De Herreros, A.G. and Ochoa, S. (1983) Proc. Natl. Acad. Sci. USA 80, 6843–6847.
- [8] Palomo, C., Vicente, O., Sierra, J.M. and Ochoa, S. (1985) Arch. Biochem. Biophys. 239, 497–507.
- [9] De Herreros, A.G., De Haro, C. and Ochoa, S. (1985) Proc. Natl. Acad. Sci. USA 82, 3119–3123.
- [10] Gaitero, F., Limas, G.G., Méndez, E. and De Haro, C. (1988) FEBS Lett. 236, 479–483.
- [11] Hunt, T., Vanderhoff, G.A. and London, I.M. (1972) J. Mol. Biol. 66, 471–481.
- [12] De Haro, C. and Ochoa, S. (1978) Proc. Natl. Acad. Sci. USA 75, 2713–2716.
- [13] Schendel, P.F. and Wells, R.D. (1973) J. Biol. Chem. 248, 8319–8321.
- [14] Gross, M. and Rabinovitz, M. (1972) Biochim. Biophys. Acta 287, 340–352.
- [15] Datta, A., De Haro, C., Sierra, J.M. and Ochoa, S. (1977) Proc. Natl. Acad. Sci. USA 74, 1463–1467.
- [16] Hunt, T. (1979) Miami Winter Symp. 16, 321–346.