

## SYNTHESIS AND PROCESSING OF PRE- $\alpha_{2u}$ -GLOBULIN IN DIFFERENT TRANSLATIONAL SYSTEMS

Bea MERTENS and Guido VERHOEVEN

*Laboratorium experimentele geneeskunde, Rega Instituut, Minderbroedersstraat 10, 3000 Leuven, Belgium*

Received 26 August 1981

### 1. Introduction

Several proteins synthesized by the rat liver display some degree of sex-specificity. One of the best known examples is  $\alpha_{2u}$ -globulin, a protein synthesized by the adult male rat liver and secreted into the blood.  $\alpha_{2u}$ -Globulin is rapidly cleared by the kidney and represents a major protein in urine of male rats. Its synthesis is under complex hormonal control involving the participation of sex hormones, glucocorticoids, thyroid hormone and pituitary hormones. Here we have studied the production of sex-specific proteins in rat liver by comparing the translation products of mRNA extracts in different translation systems. Major sex-specific proteins could be demonstrated in male as well as in female liver. The predominant sex-specific protein coded by male liver mRNA could be precipitated quantitatively by  $\alpha_{2u}$ -globulin antiserum. Comparison of its size in 3 heterologous systems (wheat germ, reticulocyte lysate and *Xenopus* oocytes) demonstrates that the corresponding mRNA codes for a precursor protein that is slightly larger than the  $\alpha_{2u}$ -globulin found in rat urine. Partial processing occurs in the wheat germ system. Oocytes assure complete processing and even secrete  $\alpha_{2u}$ -globulin.

### 2. Materials and methods

#### 2.1. Preparation of mRNA

Wistar rats (90-day-old) were fasted for 24 h. The livers were removed under ether anesthesia. Polyosomes were prepared as in [1]. The resulting polyosomal pellets were dissolved in RNA buffer (30 mM Tris-HCl (pH 7.5); 2 mM Mg<sup>2+</sup>; 100 mM NaCl; 0.5%

SDS) at a final concentration of 40 A<sub>260</sub> units/ml. Polyosomes were incubated for 30 min at 37°C in the presence of self-digested proteinase K (0.5 mg/ml). After this incubation the solution was adjusted to 0.2 M NaCl and RNA was precipitated at -20°C with 2 vol. ethanol. Separation of mRNA from total polyosomal RNA was performed by affinity chromatography on oligo(dT)-cellulose (type 2, Collaborative Research) as in [2].

#### 2.2. Protein synthesis *in vitro*

The wheat germ system was prepared as outlined [3] but with omission of the preincubation step. The assay mixture contained in 100  $\mu$ l: 40  $\mu$ l wheat germ S-30 fraction; 12 mM Hepes buffer (pH 7.6); 2 mM dithiothreitol; 1 mM ATP; 20  $\mu$ M GTP, 8 mM creatine phosphate; 4  $\mu$ g creatine phosphokinase; 200  $\mu$ M of each of the 19 unlabelled amino acids; 30 mM K-acetate, 1.8 mM Mg-acetate; 80  $\mu$ M spermine; 20  $\mu$ Ci [<sup>35</sup>S]methionine; and 5  $\mu$ g mRNA. Routinely translation was performed in a 140  $\mu$ l reaction mixture during 2 h at 30°C. Incorporation of radioactivity into trichloroacetic acid-insoluble material was monitored on a 5  $\mu$ l sample using Whatman 3 MM filter papers as in [4]. Earlier experiments revealed that optimal stimulation of methionine incorporation ( $\pm$ 40-fold) was obtained using a final RNA concentration of 5  $\mu$ g/100  $\mu$ l reaction mixture. Translation in nuclease-treated reticulocyte lysate (Radiochemical Centre, Amersham) was performed as in [5]. Routinely, 7  $\mu$ g mRNA was added to a 140  $\mu$ l reaction mixture containing 56  $\mu$ Ci [<sup>35</sup>S]methionine. The reaction was allowed to proceed for 2 h at 30°C. Incorporation of radioactivity in trichloroacetic acid-insoluble products was determined on a 5  $\mu$ l sample as in [5]. Microinjection and culture of oocytes of

*Xenopus laevis* was performed as in [6]. Groups of 10 oocytes were injected either with 50 nl phosphate-buffered saline (control) or with 50 nl poly(A)-RNA. The male RNA preparation contained 0.8 mg RNA/ml. The female preparation 1 mg/ml. Oocytes were cultured in 100  $\mu$ l medium. Three hours after injection, 75  $\mu$ Ci [ $^{35}$ S]methionine was added and incubation was continued for 3 days at 19°C. After this period the medium was carefully removed and the oocytes were homogenised. The homogenate was centrifuged for 5 min at 9000  $\times$  g in an Eppendorf microfuge and the incorporation of radioactivity was monitored by trichloroacetic acid precipitation [4].

### 2.3. Product analysis

Products of synthesis *in vitro* were analysed by polyacrylamide gel electrophoresis of trichloroacetic acid precipitates or immunoprecipitates. Immunoprecipitation was performed as in [7] using a specific anti- $\alpha_{2u}$ -globulin serum raised in rabbits. To 100  $\mu$ l aliquots of the incubated reaction mixture, 5  $\mu$ g carrier,  $\alpha_{2u}$ -globulin was added and sufficient  $\alpha_{2u}$ -globulin antiserum (40  $\mu$ l) to quantitatively precipitate the added antigen.

Trichloroacetic acid precipitates and immunoprecipitates were dissolved by boiling at 100°C in 40  $\mu$ l sample buffer (10 mM Na<sub>2</sub>PO<sub>4</sub> (pH 7.2), 10% glycerol, 2% sodium dodecyl sulfate (SDS) and 2%  $\beta$ -mercaptoethanol). Samples were run on a 15% SDS-polyacrylamide slab gel for 15 h at 6 mA as in [8]. The concentration of bisacrylamide in the gel was 0.1%. Gel slabs were dried and autoradiographed on Agfa X-ray films after treatment with Enhance (New England Nuclear).

### 3. Results and discussion

A representative analysis of the [ $^{35}$ S]methionine-labelled products synthesized in a wheat germ cell-free system programmed with liver mRNA derived from male and female rats is shown in fig.1. Both mRNA preparations stimulated methionine incorporation to a similar degree but SDS-polyacrylamide gel electrophoresis reveals marked differences in the proteins responsible for this incorporation. Trichloroacetic acid precipitates from wheat germ programmed with male mRNA reveal a prominent sex specific reaction product (protein M) in the 22 000  $M_r$  region. Translation of female mRNA reveals a prominent sex-

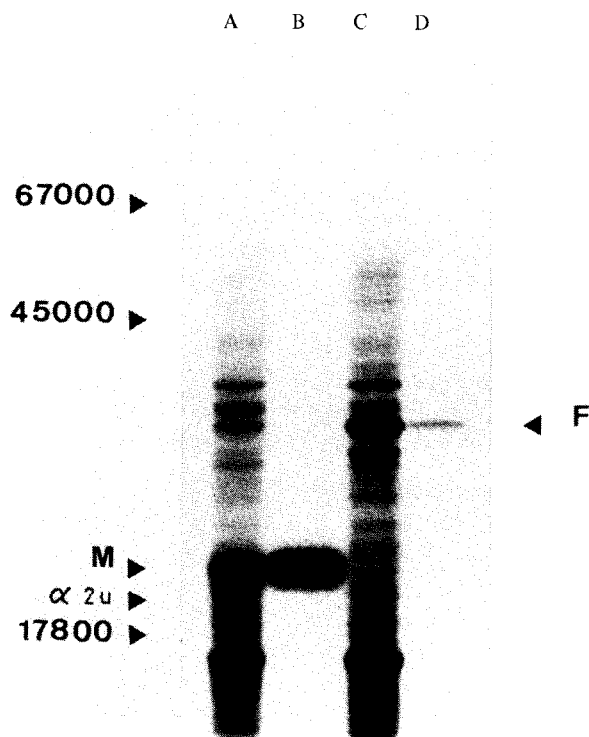


Fig.1. Translation in a wheat germ cell free system of liver mRNA derived from male and female rats, respectively: (A,B) trichloroacetic acid and immunoprecipitate of male liver mRNA translation products; (C,D) trichloroacetic acid and immunoprecipitate of female liver mRNA translation products. The marker proteins were: bovine serum albumin (6700), ovalbumin (45 000) and myoglobin (17 800).

specific protein (protein F) in the 33 000  $M_r$  region. Since it is known that one of the major proteins synthesized by male rat liver is  $\alpha_{2u}$ -globulin we investigated whether protein M could be precipitated using anti- $\alpha_{2u}$ -globulin. As shown in fig.1 this was in fact the case. Nevertheless it may be noticed that protein M is slightly larger than native  $\alpha_{2u}$ -globulin derived from rat urine. The latter protein migrates on the same gel in the 19 500  $M_r$  region. Protein F has not been identified yet. Anti- $\alpha_{2u}$ -globulin precipitates only a small fraction of the female translation products. On autoradiography the immunoprecipitate displays a weak band located in the protein F region. It cannot be excluded that this represents a non-specific interaction with the antiserum. In fact, the immunoprecipitated material represents only a minor fraction of the material migrating in the protein F region.

To further analyse the relationship between pro-

tein M and  $\alpha_{2u}$ -globulin, the translation of liver mRNA was compared in 3 systems wheat germ, reticulocyte lysate and *Xenopus* oocytes. Table 1 shows that poly(A)-RNA derived from male or female rat liver stimulates [ $^{35}$ S]methionine incorporation some 20 times in reticulocyte lysate and 30 times in wheat germ. Incorporation of amino acids in *Xenopus* oocytes was doubled after injection of male mRNA whereas female mRNA preparations consistently tended to inhibit incorporation in this system. In all 3 translational systems immunoprecipitation with anti- $\alpha_{2u}$ -globulin antiserum yields more precipitable material in reaction mixtures programmed with male mRNA. An analysis of the trichloroacetic acid-precipitable material derived from oocytes and wheat germ is shown in fig.2. In homogenates of oocytes pro-

grammed with male mRNA a major sex-specific reaction product may be noticed. This protein migrates in exactly the same region as native  $\alpha_{2u}$ -globulin. In the medium derived from the same oocytes a slight spot is observed in the same region. No specific incorporation products can be identified in oocytes programmed with female mRNA. The wheat germ system again reveals protein M when programmed with male mRNA and protein F when programmed with female mRNA. In contrast with the former experiment, however, protein M is accompanied by a slightly smaller protein that migrates in the  $\alpha_{2u}$ -globulin region. Partial processing in the wheat germ system has been described for several other secretory proteins [7,9]. Analysis of the immunoprecipitable material in fig.3 confirms that wheat germ programmed with male mRNA syn-

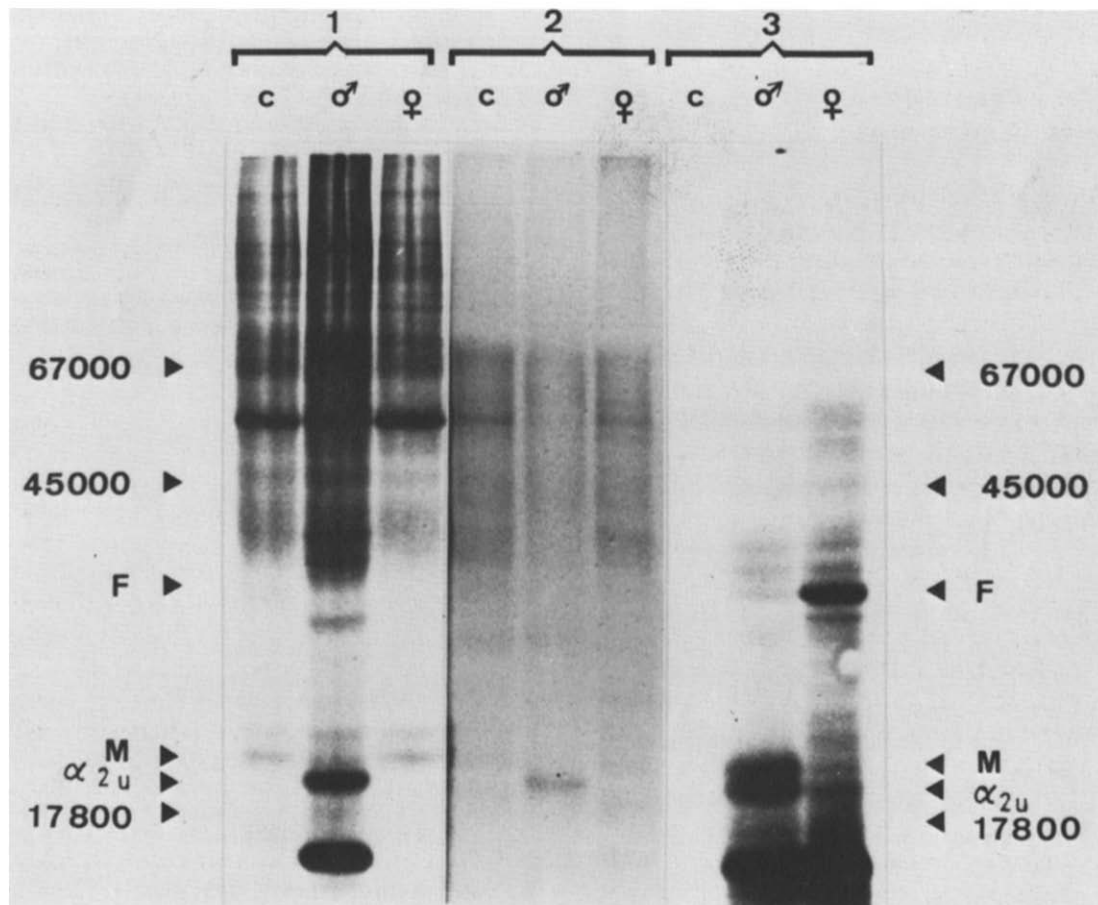


Fig.2. Comparison of the translation of liver mRNA in wheat germ and *Xenopus* oocytes: (1) oocyte homogenate; (2) oocyte medium; (3) wheat germ; (♂) male mRNA; (♀) female mRNA; (c) control (no mRNA added).

Table 1

Incorporation of [ $^{35}$ S]methionine in trichloroacetic acid-precipitable material and  $\alpha_{2u}$ -globulin antiserum immunoprecipitates in various translational systems programmed with male or female rat liver mRNA

System	mRNA prep.	cpm in total protein ( $\times 10^{-3}$ )	cpm in immunoprecipitate ( $\times 10^{-3}$ )
Reticulocyte lysate	—	74	32
	Male	1742	342
	Female	1515	135
<i>Xenopus</i> oocytes cells homogenate	—	7140	90
	Male	16 411	756
	Female	5230	50
	—	529	28
	Male	656	174
medium	Female	873	144
	—	234	12
Wheat-germ	Male	6827	748
	Female	7589	185

Male and female rat liver poly(A)-RNA was prepared from polyribosomes as in section 2. The incorporation of [ $^{35}$ S]-methionine in total trichloroacetic acid-precipitable material and in the  $\alpha_{2u}$ -immunoprecipitates was compared in 3 translational systems: wheat germ, reticulocyte lysate and *Xenopus* oocytes, respectively

thesize both forms of immunoprecipitable material but predominantly the larger  $M_r$  species. Only small amounts of immunoprecipitable material were present in the corresponding samples of systems programmed with female mRNA and those migrated in the protein F region (not shown). Taken together these data suggest that male rat liver mRNA contains a messenger that codes for pre- $\alpha_{2u}$ -globulin. The ability of *Xenopus* oocytes to convert pre- $\alpha_{2u}$ -globulin into  $\alpha_{2u}$ -globulin and to secrete this protein into the medium strongly suggests that the larger size of pre- $\alpha_{2u}$ -globulin is due to the presence of an N-terminal signal peptide related to its secretion. During the preparation of this manuscript a signal peptide containing 19 predominantly hydrophobic aminoacids has in fact been identified at the amino terminal portion of the  $\alpha_{2u}$ -globulin mRNA translation product in reticulocyte lysate [10]. The presence of such a signal sequence is compatible with studies indicating that  $\alpha_{2u}$ -globulin is synthesized on membrane bound poly-

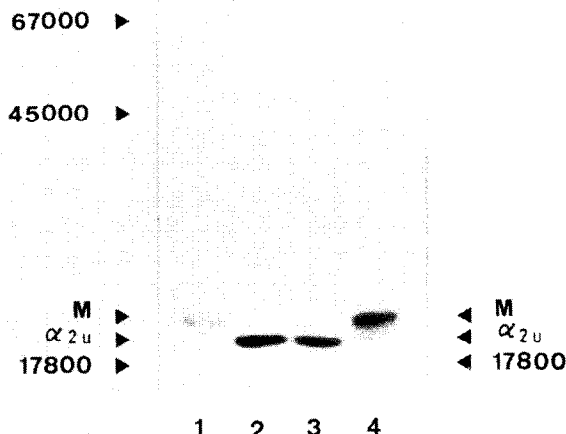


Fig.3. Incorporation of [ $^{35}$ S]methionine in immunoprecipitable translation products from reticulocyte lysate, *Xenopus* oocytes and wheat germ, respectively, programmed with male and female mRNA. Poly(A)-RNA from male and female rat liver was translated in various systems. After immunoprecipitation with  $\alpha_{2u}$ -globulin antiserum as in section 2 translation products were analysed as in fig.2: (1) reticulocyte lysate; (2) oocyte homogenate; (3) oocyte medium; (4) wheat germ.

somes and offers a further point of analogy between this protein and the major urinary proteins (MUPs) of the mouse [11].

## References

- [1] Sala-Trepat, J. M., Savage, M. J. and Bonner, J. (1978) *Biochim. Biophys. Acta* 519, 173–193.
- [2] Aviv, H. and Leder, P. (1972) *Proc. Natl. Acad. Sci. USA* 69, 1408–1412.
- [3] Roberts, B. E. and Paterson, B. M. (1973) *Proc. Natl. Acad. Sci. USA* 70, 2330–2334.
- [4] Bollum, F. J. (1968) *Methods Enzymol.* 12, 169–173.
- [5] Pelham, H. R. B. and Jackson, R. J. (1976) *Eur. J. Biochem.* 67, 247–256.
- [6] Gurdon, J. B., Lane, C. D., Woodland, H. R. and Marbaix, G. (1971) *Nature* 233, 177–182.
- [7] Peeters, B., Mous, J., Van Belleghem, H. and Rombauts, W. (1979) *Biochim. Biophys. Acta* 561, 502–516.
- [8] Laemmli, U. K. (1970) *Nature* 227, 680–685.
- [9] Shields, D., Warren, T. G., Roth, S. E. and Brenner, M. J. (1981) *Nature* 289, 511–514.
- [10] Drickamer, K., Kwoh, T. J. and Kurtz, D. T. (1981) *J. Biol. Chem.* 256, 3634–3636.
- [11] Hastie, N. J., Held, W. A. and Toole, J. J. (1979) *Cell* 17, 449–457.