

# Qualitative and quantitative changes in protein synthesis occur at the 8-16-cell stage of embryogenesis in the cow

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**Summary.** Cow oocytes and preimplantation embryos were cultured in medium containing radiolabelled methionine and the proteins synthesized were analysed by one-dimensional electrophoresis and fluorography. Marked changes in the pattern of synthesis were observed at the 8-16-cell stage of development. Quantitatively, a gradual decrease in the rate of protein synthesis occurred between the zygote and 8-cell stage and then the rate increased progressively to the blastocyst stage. Incorporation of radiolabelled uridine into RNA was first detected at the 16-cell stage. Taken together, these results suggest that protein synthesis is programmed by maternal mRNA up to the 8-cell stage but switches to mRNA derived from the zygote genome between the 8- and 16-cell stage.

**Keywords:** embryo; maternal mRNA; transcription; translation; cow

## Introduction

The earliest stages of embryogenesis, including those of mammalian embryos, are dependent upon stored products of the maternal genome. During oogenesis, populations of both maternal RNA and protein are accumulated within the oocyte that are sufficient for the progression of the embryo through the first cleavage division in the mouse embryo (Golbus *et al.*, 1973; Braude *et al.*, 1979), the second cleavage division of the human embryo (Braude *et al.*, 1988) and the third cleavage division of the rabbit (Manes, 1973) and the sheep embryo (Crosby *et al.*, 1988). Subsequent embryonic development is dependent upon transcriptional activation of the zygote genome. The switch from maternal to zygote genome control of development is characterized by major changes in the pattern of protein synthesis (Van Blerkom & Manes, 1974; Flach *et al.*, 1982; Norris *et al.*, 1985; Braude *et al.*, 1988; Crosby *et al.*, 1988).

In this study we examined RNA and protein synthesis to determine when the switch from maternal to zygote genome control of development occurs in the preimplantation cow embryo.

## Materials and Methods

### *Superovulation and collection of oocytes and embryos*

Heifers and cows of mixed breeds, aged 2-5 years, were obtained from ranch herds in southern Alberta and palpated rectally. Animals with normal reproductive tracts and with active ovaries were injected intramuscularly (i.m.) with 3000 i.u. PMSG (Laboratorio Elea S.A.C.I.F., Buenos Aires, Argentina) on Day 10-12 of the oestrous cycle, followed after 3 days by i.m. injection of 750 µg cloprostenol (Estrumate; Coopers, Willowdale, Canada). Induced oestrus was detected by the onset of standing heat 1.5-2 days later, and if fertilized eggs were desired, the animals were inseminated artificially 12 and 24 h after the onset of oestrus.

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Post-ovulatory oocytes and embryos to be collected on Days 1–4 after onset of oestrus were recovered surgically at Alta Genetics Incorporated (AGI) of Calgary. The ovaries from each cow were removed, and each oviduct was flushed with 15 ml PB-1 medium (Whittingham & Wales, 1969) containing 4 mg bovine serum albumin/ml (BSA; Fraction V, Sigma Chemical Co., St Louis, MO 63178, USA) and 10% heat-inactivated calf serum (Gibco, Grand Island, NY, USA). The adhering cumulus masses were removed by passing the oocytes up and down a fine-bore micropipette. Embryos to be recovered from cows on Day 5 after onset of oestrus or later were collected at AGI using a standard non-surgical procedure (Elsden *et al.*, 1976). After collection, oocytes and embryos were washed by several transfers through M2 medium (Whittingham, 1971) before use in experiments.

**Radiolabelling and electrophoresis of  $^{35}\text{S}$ -labelled material.** Labelling of embryos and electrophoresis of  $^{35}\text{S}$ -labelled proteins were carried out by the methods used by Gifford *et al.* (1987) with minor modifications. Embryos were transferred into 2–3 ml M2 medium and incubated at 37°C for 1 h. The embryos were then cultured under oil in a prewarmed 50  $\mu\text{l}$  drop of M2 medium containing 0.83  $\mu\text{M}$ -L-[ $^{35}\text{S}$ ]methionine (sp. act. 1253 Ci/mmol; Amersham, Oakville, Ontario, Canada) or 0.75  $\mu\text{M}$ -L-[ $^{35}\text{S}$ ]methionine (sp. act. 1330 Ci/mmol; Amersham) at 37°C in an humidified atmosphere of 5%  $\text{CO}_2$  for 2 h. Oocytes and embryos were then washed 3 times in 4 ml volumes of fresh, ice-cold medium free of radioactivity, with a final wash in 4 ml phosphate-buffered saline (PBS) pH 7.2 (Dulbecco & Vogt, 1954), to remove protein present in the medium. Washed individual oocytes or embryos were transferred along with a small amount of PBS into microfuge tubes and were disrupted by adding 25  $\mu\text{l}$  SDS (sodium dodecyl sulphate)-dissociation buffer (62.5 mM-Tris-HCl, pH 6.8, containing 2% (w/v) SDS, 10% (v/v) glycerol and 5% (v/v) 2-mercaptoethanol). The samples were stored at –70°C.

Measurement of trichloroacetic acid (TCA)-precipitable radioactivity was conducted on 2  $\mu\text{l}$  samples of the lysate, exactly as described by Gifford *et al.* (1987). The efficiency of counting of [ $^{35}\text{S}$ ]methionine using 'Econofluor-2' or 'Aquasol-2' was calculated to be 77.6% and 75.3%, respectively.

Samples (12  $\mu\text{l}$ ) of dissociated extracts were resolved by sodium-dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) and fluorography as described previously (Gifford *et al.*, 1987). The  $^{14}\text{C}$ -labelled molecular weight markers used were purchased from Amersham (myosin, 200 000; phosphorylase b, 92 500; bovine serum albumin, 69 000; ovalbumin, 46 000; carbonic anhydrase, 30 000; and lysozyme, 14 300).

**In-vitro labelling of embryos with [ $^3\text{H}$ ]uridine.** Embryos were transferred into a pre-warmed 50  $\mu\text{l}$  drop of M2 medium containing 5.13  $\mu\text{M}$ -[5,6- $^3\text{H}$ ]uridine (sp. act. 39 Ci/mmol; Amersham) and incubated under oil at 37°C in an humidified atmosphere of 5%  $\text{CO}_2$  air for 2 h. Measurement of acid-precipitable radioactivity in washed embryos was conducted as for protein labelling studies. Tritium counting efficiency was 36%.

## Results

A steady decrease in the rate of incorporation of L-[ $^{35}\text{S}$ ]methionine into acid-precipitable material was observed between the post-ovulatory oocyte to the 8-cell stage (Table 1). In the 8-cell embryo, the relative rate of protein synthesis was, on average, 29% of that of the post-ovulatory oocyte. In stages from the 16-cell to the blastocyst there was a progressive increase in the rate of L-[ $^{35}\text{S}$ ]methionine incorporation (Table 1). The rate in the blastocyst was observed to be 20-fold greater than at the 8-cell stage.

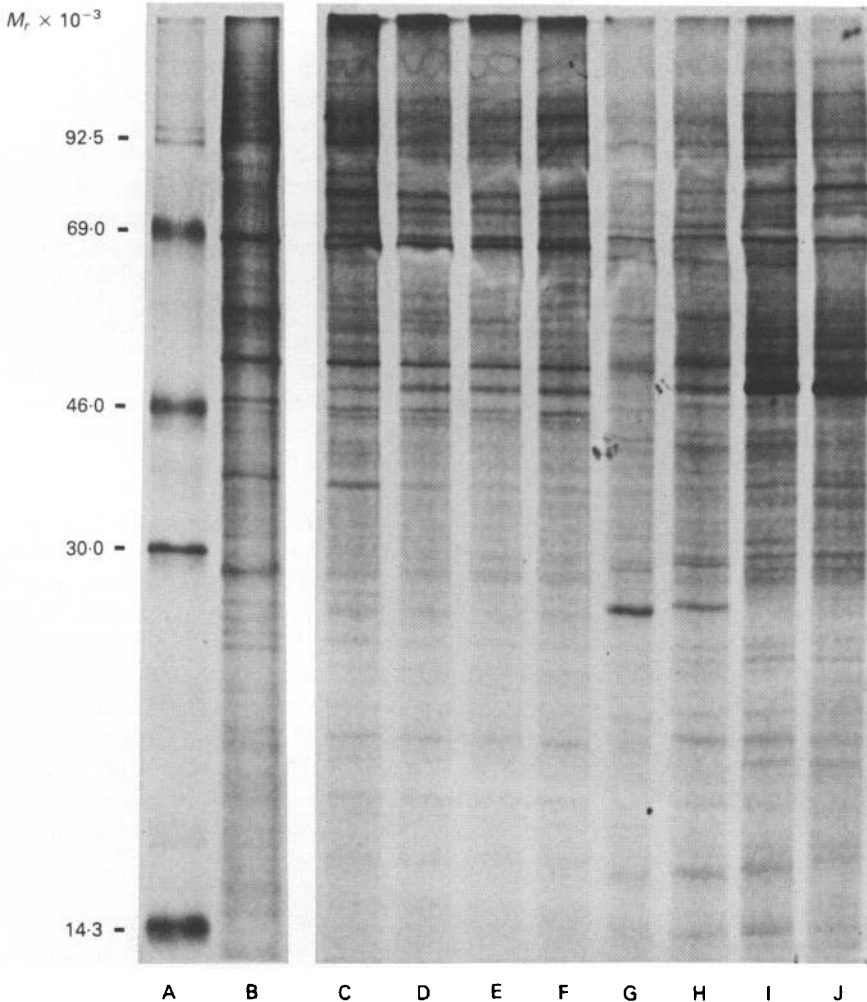
**Table 1.** Incorporation of L-[ $^{35}\text{S}$ ]methionine into TCA-precipitable material by cow oocytes and embryos

Stage	Incorporation (c.p.m./embryo/2 h*)
Postovulatory oocyte	9630 (4816–12 418, $n = 6$ )
1-cell zygote	8863 (4934–9514, $n = 9$ )
2-cell embryo	5372 (4032–6638, $n = 4$ )
4-cell embryo	4262 (4154–4372, $n = 2$ )
8-cell embryo	2819 (2650–2988, $n = 2$ )
8–16-cell stage	6520 (5122–8180, $n = 3$ )
16–32-cell stage	7900 (6256–9544, $n = 2$ )
32–48-cell stage	29 197 (18 976–39 418, $n = 2$ )
Blastocyst	56 363 (21 660–91 878, $n = 9$ )

\*Values are the average of the number of individual oocytes or embryos ( $n$ ) assayed as shown in the parentheses. The numbers in parentheses give the range of values obtained. Counting efficiency was 76%.

The qualitative patterns of proteins synthesized at different stages of development are shown in Fig. 1. While minor differences in some radiolabelled polypeptides were detected between the oocyte and 4-cell stage (Lanes B-F, Fig. 1), the patterns were quite similar at one stage and the next. Marked changes in the pattern were, however, detected in radiolabelled polypeptides derived from 8- and 16-cell stage embryos (Lanes G and H, Fig. 1). The patterns in 16- to 32-cell embryos and blastocysts were similar to each other but distinct both from those produced by embryos before the 4-cell stage and by 8-cell and 16-cell stage embryos.

To measure rates of RNA synthesis, embryos at different stages of preimplantation development were incubated *in vitro* in the presence of [<sup>3</sup>H]uridine for 2 h. We were unable to detect any



**Fig. 1.** Qualitative patterns of protein synthesis in cow oocytes and early embryos. Individual oocytes or embryos were incubated in medium containing L-[<sup>35</sup>S]methionine for 2 h. About 1800 c.p.m. of acid-insoluble radioactivity from each sample was analysed by one-dimensional electrophoresis and fluorography. The positions of <sup>14</sup>C-methylated standards are identified in Lane A with numerical values ( $\times 10^{-3}$ ). Lane B, postovulatory oocyte; Lane C, 1-cell zygote; Lane D, 2-cell embryo; Lane E, 2-cell embryo; Lane F, 4-cell embryo; Lane G, 8-cell embryo; Lane H, 8-16-cell embryo; Lane I, 16-32-cell embryo; Lane J, blastocyst (Day 7). X-ray film exposure time was 22 days.

incorporation of [ $^3\text{H}$ ]uridine into TCA-precipitable material before the 16-cell stage. The levels of radioactivity that were measured in a sample containing a pool of four 2-cell embryos and a sample containing a pool of four 8-cell embryos were indistinguishable from background. In three 16-cell embryos, an average incorporation rate of 42 c.p.m./embryo/2 h was observed. For a series of morulae to early blastocysts (>32-cells but <48 cells), the rate of [ $^3\text{H}$ ]uridine incorporation was  $760 \pm 213$  (mean  $\pm$  s.d.,  $n = 13$ ) c.p.m./embryo/2 h.

### Discussion

A major feature of the transition from maternal to zygote genome control in the mouse embryo is a marked change in the pattern of protein synthesis as the embryo proceeds through the first cleavage (Braude *et al.*, 1979; Cullen *et al.*, 1980). Some of these changes are due to differential mRNA activation, some are due to post-translational modification and some are due to differential polypeptide turnover (Howlett & Bolton, 1985). During this period there is a loss of much of the maternal messenger RNA pool but transcriptional activity from the zygote genome is also activated (Piko & Clegg, 1982; Giebelaus *et al.*, 1983; Clegg & Piko, 1983). The two-cell stage of mouse embryogenesis is, therefore, the point where developmental arrest occurs when embryos are treated with  $\alpha$ -amanitin (Golbus *et al.*, 1973). Comparable sets of studies have been conducted to suggest that the switch from maternal to zygote genome control of early development occurs at the 8-cell stage in the rabbit (Manes, 1973; Schultz *et al.*, 1973; Van Blerkom & Manes, 1974). This transition period occurs at the 4–8-cell stage of human development (Braude *et al.*, 1988) and the 8–16-cell stage of development of the sheep embryo (Crosby *et al.*, 1988).

The experiments presented in this paper indicate that, from the 1-cell to the 8-cell stage of development of the bovine zygote, there is a decrease in the level of protein synthesis, with no major changes in the qualitative patterns of protein synthesis taking place during this period. At the 8–16-cell stage, several changes in the qualitative pattern of protein synthesis are evident. A new qualitative pattern of protein synthesis is established after the 16-cell stage, with this pattern persisting up to the blastocyst stage. The rates of protein and RNA synthesis also increase dramatically between the 16-cell and the blastocyst stages of development. The biochemical data on RNA synthesis in this report are consistent with previous studies in which, through use of autoradiographic techniques, [ $^3\text{H}$ ]uridine was incorporated into the nucleoplasm and nucleoli of cow embryos only at the end of the 8-cell stage (Camous *et al.*, 1986). All of these findings are consistent with the interpretation that the maternal to zygote genome transition in the cow embryo occurs at the same time as that in the sheep.

The rate of radiolabelled methionine incorporation declines gradually between the oocyte and 8-cell stage in the cow embryo. The rate of protein synthesis also drops in the sheep embryo but, in this case, the fall is a sharp decline after the third cleavage rather than a gradual decrease in synthetic rate (Crosby *et al.*, 1988). If this decrease in translational activity is due to degradation of maternal mRNA, then the cow and sheep embryo differ in the way that the programme of maternal control of development is regulated.

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