

EXPRESSION OF FIBRONECTIN AND INTERSTITIAL COLLAGEN GENES IN SMOOTH MUSCLE CELLS: MODULATION BY LOW MOLECULAR WEIGHT HEPARIN FRAGMENTS AND SERUM*

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Abstract—The effect of low *M*, heparin fragments (CY222) and fetal calf serum (FCS) on the level of fibronectin and fibrillar collagen mRNAs was investigated in smooth muscle cells (SMC) in culture. In the absence of FCS, addition of CY222 (100 µg/10⁶ cells) to postconfluent early passage SMC resulted in a decrease in mRNA level of type III collagen. In contrast, mRNA levels coding for type I collagen, fibronectin and GAPDH (used as control of cellular activity) were not modified. Addition of 5% FCS (without CY222) to the culture medium did not affect mRNA levels of type I and type III collagens nor that of GAPDH. The level of fibronectin mRNA, however, increased in the presence of 5% FCS. In the presence of both 5% FCS and CY222, we observed a decrease in type III collagen mRNA and fibronectin mRNA levels (this level remained, however, above the control value without FCS and the level with CY222 alone). Our results demonstrate that low *M*, heparin fragments can modulate the steady-state levels of type III collagen and fibronectin mRNAs.

Key words: fibronectin; collagens; biosynthesis; gene expression; messenger level; low molecular weight heparin fragments; modulation of phenotype; smooth muscle cells

Fibronectin, type I collagen, and type III collagen genes code for the three major proteins of the ECM§ [1, 2]. The regulation of the genes coding for ECM is an important part of many cellular functions such as adhesion, proliferation, migration and differentiation. Quantitative and qualitative modifications in ECM gene expression vary as a function of cell type, age and related pathological processes [3, 4]. SMC, the main cell type of the media of arteries, play an important role in the atheroarteriosclerotic process. They change from a contractile to a migratory and synthetic phenotype [5] and produce large quantities of ECM components, among them collagens and fibronectin, and decreased quantities of contractile proteins [6, 7]. Heparin and related compounds can inhibit cellular proliferation [8–12] by maintaining cells in the G1 phase [13]. Heparin can also modulate the biosynthetic

phenotype of SMC, particularly the biosynthesis of fibrous collagens [14]. We were able to show that low *M*, heparin fragments (CY222) [15], administered *in vivo* to diabetic mice modulated skin type III collagen and fibronectin biosyntheses [16]. Recently, we demonstrated that the biosynthetic phenotype of SMC could be modulated *in vitro* by CY222 [17]. These results illustrate the complex interactions between cells and ECM in the control of many cellular functions. In this study, we report that gene expression is modified differently in the presence of low *M*, heparin fragments and/or in the presence of serum. Furthermore, serum components could interfere with heparin and related compounds such as CY222 [18]. We therefore studied the effect of CY222 with and without FCS in the culture medium. The results of these studies are described below.

MATERIALS AND METHODS

Cell culture. Pig aorta SMC were grown from media explants in Dulbecco's modified Eagle's medium supplemented with 20% FCS, 100 IU/mL penicillin, 100 µg/mL streptomycin, and 2 mM glutamine. At confluency, cells were dissociated with 0.5% trypsin in PBS. Cells were serially passaged with 10% FCS up to the third subculture, where they were used at overconfluency for RNA extraction. Cells were incubated at 37° in a 95% air, 5% CO₂ atmosphere. The day before the experiments, the culture medium was discarded, and the cells were incubated in the presence of

* This paper is dedicated to the memory of Dr Jean Choay who initiated the study of heparin fragments prepared in his laboratory.

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§ Abbreviations: DMEM, Dulbecco's modified Eagle's medium; SMC, smooth muscle cells; ECM, extracellular matrix; FCS, fetal calf serum; GAPDH, glyceraldehyde-3-phosphate-dehydrogenase; DEPC, diethylpyrocarbonate; MOPS, 3-(*N*-morpholino)-propanesulfonic acid.

Table 1. Comparison of GAPDH, pro α 1(I) collagen, pro α 1(III) and fibronectin mRNA levels by SMC

Probes	Control	CY222	Control + 5% FCS	CY222 + 5% FCS
GAPDH	13,216 \pm 5221	11,169 \pm 3007	15,474 \pm 4124	16,417 \pm 5064
Coll. III/GAPDH	1.31 \pm 0.12	0.84† \pm 0.09	1.29 \pm 0.03	0.62‡ \pm 0.02
Coll. I/GAPDH	1.83 \pm 0.14	2.02 \pm 0.16	1.89 \pm 0.08	2.10 \pm 0.07
FN/GAPDH	0.57 \pm 0.03	0.49 \pm 0.06	0.92* \pm 0.04	0.73§ \pm 0.08

Control, with 5% FCS, or with 100 mg/mL of CY222, with or without 5% FCS. Relative amounts of the specific mRNA were quantified by densitometric analysis of slot-blots. The values represent the mean \pm SD of eight independent determinations.

* Control 5% FCS vs control (DMEM alone), $P < 0.001$.

† CY222 vs control without CY222, $P < 0.001$.

‡ CY222 with 5% FCS vs control with 5% FCS, $P < 0.001$.

§ CY222 with 5% FCS vs CY222 alone, $P < 0.001$.

100 μ g/10⁶ cells heparin fragments CY222 (Institut Choay/Sanofi Recherches, Paris, France) with or without 5% FCS for 24 hr. These CY222 fragments (1500–8000 *M_r*) were obtained by chemical depolymerization of porcine mucosal heparin with HNO₂ and gel filtration as described by Barzu *et al.* [15]. They correspond to a mixture of hexamers and octamers.

RNA analysis. Total ribonucleic acids were extracted from confluent cultures washed free of media with 4 M guanidium isothiocyanate, according to the method of Chomczynsky and Sacchi [19]. Briefly, SMC were rapidly homogenized in a 4 M guanidium isothiocyanate, 25 mM sodium citrate, pH 7, 0.5% Sarcosyl, 0.1 M 2-mercaptoethanol buffer. Sequentially, 2 M sodium acetate, pH 4, phenol (water-saturated) and chloroform-isoamyl alcohol mixture (49:1) were added. RNAs were precipitated from the aqueous phase by isopropanol, and left overnight at -20° . After collection by centrifugation, pellets were rinsed with 1 mL of 75% ethanol and dissolved in 5 mM EDTA, pH 7.0, treated with 0.1% DEPC. Approximately 25 μ g of total RNAs was extracted from 10⁶ cells by this procedure.

For quantitative assessment, aliquots of 1, 1.5 and 2 μ g of total RNA were heat-denatured at 65° for 15 min in a 50% formamide, 6% formaldehyde, 0.2 M MOPS, 0.05 M sodium acetate, pH 7.0, and 0.01 M EDTA buffer. Total RNA was then transferred to Nylon filters (Hybond N⁺, Amersham, France), using a commercial "slot-blot" apparatus (Minifold II, Schleicher & Schull, Dassel, Germany). RNA was fixed to the filter by alkali treatment (5 min with 0.05 M NaOH, pH 5) and then washed with 2 \times SSC (1 \times : 0.15 M NaCl, 0.015 M sodium citrate, pH 7.0).

DNA probes. The following cDNA probes were used: pHCAL1U specific for human pro α 1(I) collagen and pHFS3 specific for human pro α 1(III) collagen, provided by Vuorio and Mäkelä [20, 21]; a clone of rat fibronectin from the exon encoding the first extra type III region, given by R. O. Hynes and described in Paul *et al.* [22]; and the GAPDH (glyceraldehyde-3-phosphate-dehydrogenase) probe described by Fort *et al.* [23].

The specificity of the probes was tested by

Northern blot analysis [20, 23]. Briefly, total RNA was separated by electrophoresis in a 1% agarose-formaldehyde gel. Resulting Northern blots were probed for α 1(I) and α 1(III) collagens and fibronectin with cDNA probes labelled as described below.

Slot-blot analysis. Prehybridization was carried out at 42° for 24 hr in a solution (A) containing 50% (v/v) formamide, 5 \times SSPE (1 \times SSPE: 180 mM NaCl, 10 mM Na₂HPO₄, 1 mM EDTA, pH 7.4), 0.5% (w/v) SDS, 0.1% each of Ficoll, polyvinylpyrrolidone, and BSA and 200 μ g/mL of heat-denatured sonicated salmon DNA. Hybridization was carried out in fresh buffer A, containing 3–5 \times 10⁸ cpm/mg of the appropriate ³²P-labelled probe for 24 hr at 42° . Labelling was performed using a nick translation kit (Boehringer Mannheim, Germany) with [α -³²P]dCTP (111 TBq/mmol, Amersham, France). Labelled probes were purified by chromatography through an NACS Prepac column (Bethesda Research Labs, Bethesda, France). The filters were washed at 42° with 2 \times SSPE, 1 \times SSPE, 0.1 \times SSPE, sequentially with 0.1% SDS. The dried filters were exposed to X-ray films at -80° . The autoradiograms were quantified by densitometric scanning with a laser densitometer (Ultrascan 2202 LKB, Integrator 2220 LKB).

Statistical analyses were performed using the Student's *t*-test.

RESULTS

Specificity of the probes was demonstrated by Northern blot analysis (Fig. 1). For α 1(I) mRNA and α 1(III) mRNAs, the two spots arise from usage of alternative polyadenylation signals [20, 21].

Linearity of densitometric analysis from slot-blots

When specific mRNAs were quantified by densitometric analysis we obtain a linear relationship between the amounts of mRNA applied to the filter for hybridization. To illustrate this, Fig. 2 presents the calibration curve based upon the integrated intensity from autoradiograms of GAPDH mRNAs with a correlation coefficient $r = 0.97$, $P < 0.01$.

Effect of heparin fragment CY222 on extracellular matrix and GAPDH mRNAs levels

The aortic smooth muscle cells at postconfluence

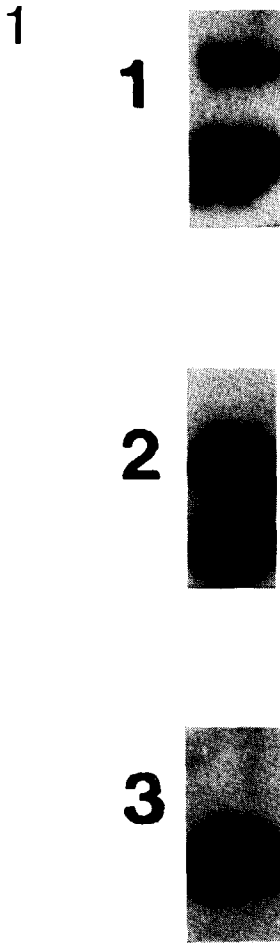


Fig. 1. Northern blot analysis of collagen and fibronectin mRNAs expressed in smooth muscle cells in culture. Total RNA was isolated from cultures of SMC and samples ($10 \mu\text{g}/\text{lane}$) were separated by electrophoresis in 1% agarose-formaldehyde gels. Resulting Northern blots were probed for $\alpha 1(\text{I})$ collagen mRNA (1), $\alpha 1(\text{III})$ collagen mRNA (2) and fibronectin mRNA (3). The two $\alpha 1(\text{I})$ mRNA species as well as the two $\alpha 1(\text{III})$ mRNA species arise from usage of alternative polyadenylation signals [20, 21].

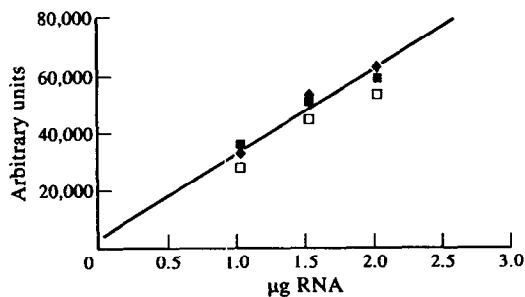


Fig. 2. Calibration curve based upon the integrated intensity from the autoradiogram of GAPDH RNA. Abscissa: amounts of total RNA applied to filters for hybridization with the GAPDH probe; ordinates: integrated autoradiography intensity in arbitrary units. Correlation coefficient $r = 0.97$.

incubated with $100 \mu\text{g}$ CY222/ 10^6 cells showed markedly reduced levels of $\text{pro}\alpha 1(\text{III})$ mRNA, compared with cultures incubated with DMEM alone. In contrast, cultures incubated without heparin fragments (Fig. 3A) and those treated with CY222 alone (Fig. 3B) showed comparable levels of $\text{pro}\alpha 1(\text{I})$ mRNAs. Likewise, the level of GAPDH mRNA remained relatively constant (Table 1).

These results are in agreement with the biosynthetic measurements previously obtained [16] which show that the addition of CY222 leads to inhibition of collagen III biosynthesis without change in collagen type I and total protein biosyntheses.

Effect of DMEM supplemented with 5% FCS or with a combination of CY222 and 5% FCS on extracellular matrix and GAPDH mRNAs levels

In these experiments, SMC were incubated in DMEM supplemented with 5% FCS. The levels of fibronectin mRNAs appeared to be increased by the presence of FCS, when compared with the levels seen with DMEM alone. Nevertheless, the addition of 5% FCS did not significantly affect GAPDH, $\text{pro}\alpha 1(\text{I})$ collagen or $\text{pro}\alpha 1(\text{III})$ collagen mRNA levels (Fig. 4).

The combination of 5% FCS and heparin fragments CY222 significantly down-regulated the mRNA levels of fibronectin and $\text{pro}\alpha 1(\text{III})$ collagen without affecting GAPDH and $\text{pro}\alpha 1(\text{I})$ collagen mRNA levels (Table 1), as compared with SMC incubated with DMEM supplemented with 5% FCS. The action of CY222 appeared to be greater in the presence of serum, since steady-state levels of collagen type III mRNA were decreased by 36% in the presence of CY222 alone. In the presence of CY222 and 5% FCS this decrease reached 52% ($P < 0.001$). Likewise, CY222 alone produced a slight decrease in fibronectin mRNA (14%, not significant). When SMC were incubated with a combination of CY222 and FCS, the mRNA fibronectin level was depressed by approximately 21% ($P < 0.001$).

DISCUSSION

The effect of polyanions on the phenotype of vascular cells has been studied by several authors [10, 14].

The proliferation of vascular smooth muscle cells (SMC) after endothelial injury and secretion of the extracellular matrix (ECM) components are postulated to be important early events in the pathogenesis of atherosclerosis [5–7].

Our results demonstrate that low M_r heparin fragments (CY222) can modulate ECM gene expression. Our results confirm and extend results from our and other laboratories on the action of heparin and derivatives on arterial SMCs [16] and diabetic skin explants of KK mice treated with CY222 [17]. SMC proliferation and biosynthesis *in vitro* were shown to be strongly influenced by heparin and related compounds [8–12, 14, 16, 17, 24–28]. Postconfluent SMCs and vascular endothelial cells both produce heparin-like substances exhibiting antiproliferative properties for smooth muscle cells [9, 28]. Binding and internalization of heparin by vascular SMCs and endothelial cells via specific sites

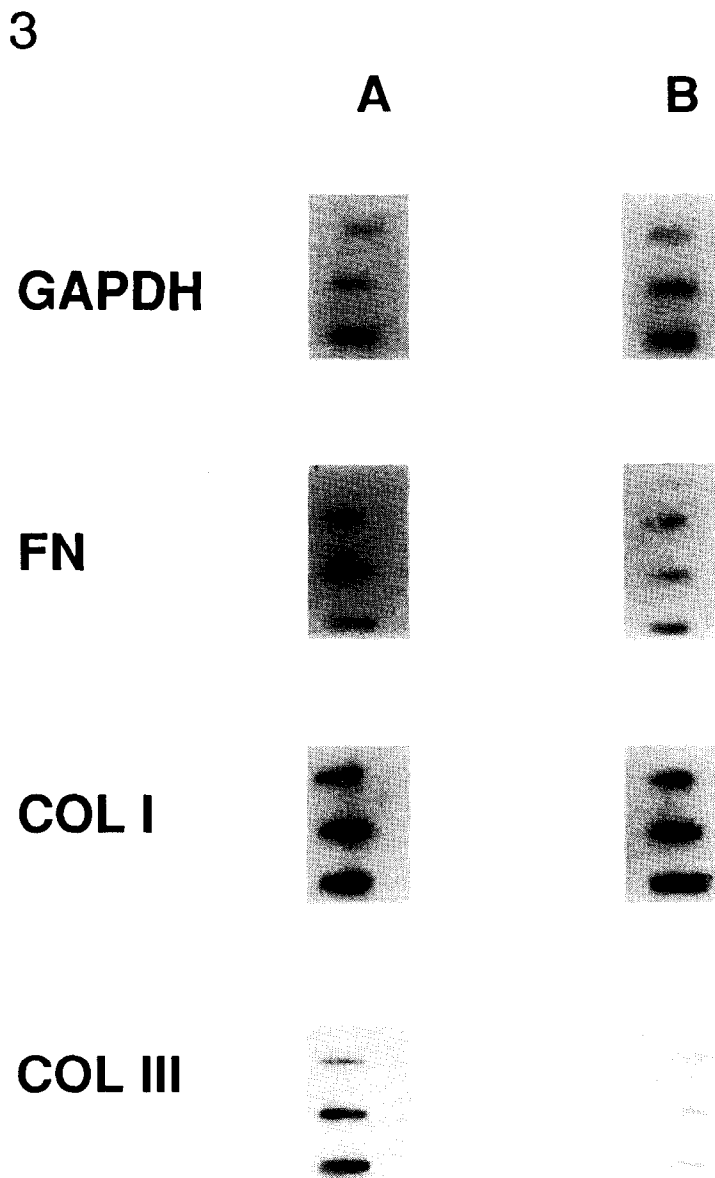


Fig. 3. Slot-blot analysis of GAPDH, fibronectin III A (FN), pro α 1(I) (Col I), and pro α 1(III) (Col III) collagen mRNA levels in pig aorta SMC cultures. Total RNA from SMC at the third passage was extracted with 4 M guanidium thiocyanate as described in Materials and Methods. Increasing quantities of total RNA (1, 1.5, and 2 mg) were applied to the filters. (A) Culture medium alone; (B) addition of 100 mg CY222/ 10^6 cells.

have been demonstrated [15, 25]. Moreover, the presence of heparan sulfate and other heparin derivatives on the cell surface [29], in cytoplasm [30], and in the nucleus [28] were also reported, and their size and degree of sulfation were shown to help determine their efficiency [15, 31]. Heparin can inhibit a PKC-dependent pathway for cellular proliferation by suppressing *c-fos* and *c-myc* expression [32].

We earlier observed a dose-dependent decrease in type III collagen biosynthesis in the presence of heparin fragments in SMC cultures [16]. In the

present experiments, we have demonstrated that CY222 decreases the pro α 1(III) collagen mRNA level. These modifications seem to be relatively selective, since modifications in GAPDH, pro α 1(I) collagen and fibronectin mRNAs levels were not observed. Heparin and related compounds interfere with AP1 transactivating factor (*jun/fos* protein complex) [33]. These transacting factors are implicated in the regulation of both type I and type III collagen biosynthesis [34, 35]. However, the promoter and regulatory domains of type I and type III collagens are organized differently [35]. For

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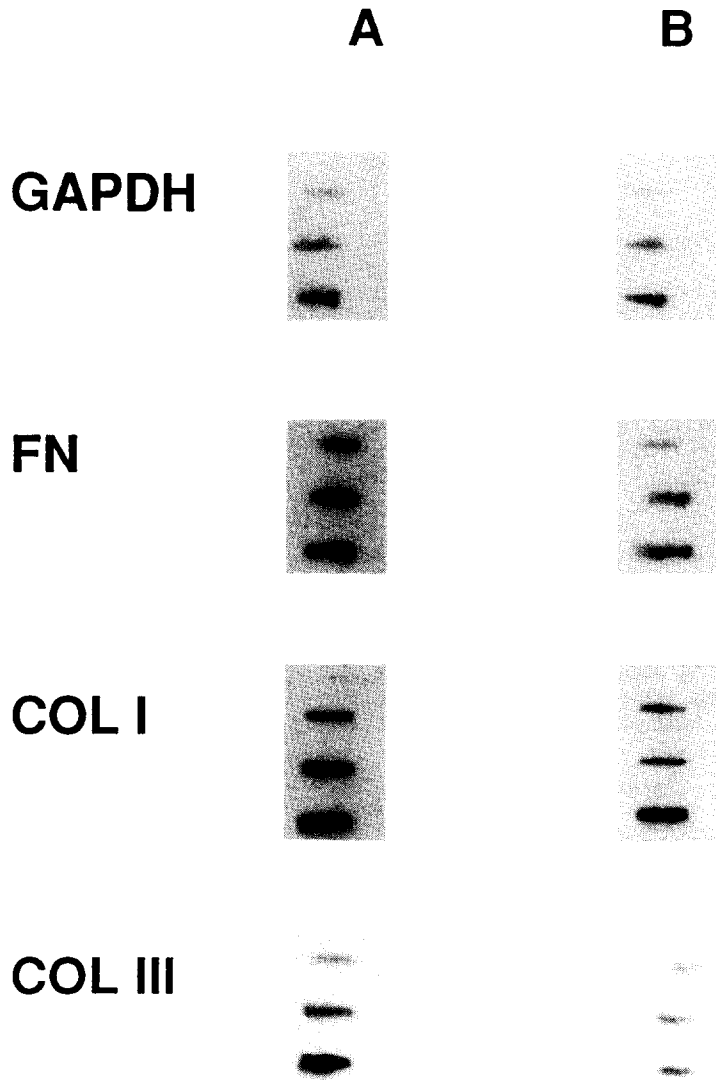


Fig. 4. Slot-blot analysis of GAPDH, fibronectin III A (FN), $\text{pro}\alpha 1(\text{I})$ (Col I), and $\text{pro}\alpha 1(\text{III})$ (Col III) collagen mRNA levels in pig aorta SMC, in the presence of 5% FCS in the culture medium. Total RNA was prepared and analysed as in Fig. 1. (A) Medium with 5% FCS; (B) medium plus FCS and 100 mg CY222/ 10^6 cells.

example, a heterodimeric CCAAT factor that activates type I collagen genes does not bind to this segment of collagen type III promoter [35]. Heparin fragments might well exhibit different effects on these regulatory processes. In the presence of 5% FCS in the culture medium (without CY222), the transcription of GAPDH, $\text{pro}\alpha 1(\text{I})$ and $\text{pro}\alpha 1(\text{III})$ collagens was not modified. Nevertheless, in the presence of FCS, the decrease in $\text{pro}\alpha 1(\text{III})$ mRNA produced by CY222 was more substantial, suggesting a synergistic effect between serum components and heparin fragments.

In previous experiments, we observed that CY222

decreased fibronectin secretion without affecting fibronectin biosynthesis [16]. We have shown in the present study that CY222 alone did not modify the fibronectin mRNA level. However, the amount of fibronectin mRNA increased with FCS alone. This serum stimulation of fibronectin gene expression is mediated through a DNA sequence at position -170 bp of the fibronectin gene (cAMP element or CRE), resulting from a rapid serum-induced binding of nuclear protein to the CRE [38]. In the presence of both FCS and CY222, we observed a decrease in the fibronectin mRNA level vs serum alone. This decrease could be due to the complexing of serum

mitogens by heparin fragments [27, 38]. It has been demonstrated that TGF β , EGF and PDGF can modify fibronectin gene expression [39, 40].

It appears that CY222 exhibits specific effects on the expression of matrix genes, especially on those of certain interstitial collagens and fibronectin. Their effect varies according to cell type and growth phase. Indeed, we demonstrated that CY222 could inhibit fibronectin biosynthesis in human dermal fibroblast cultures in the absence of serum (data not shown) and in skin explant cultures of diabetic KK mice [17]. Our results suggest that heparin and related compounds such as CY222 may be important physiologically and pharmacologically in the regulation of the phenotype of vascular SMC. Furthermore, heparin fragments, which inhibit proliferation and modulate extracellular matrix biosyntheses [41], appear to be able to suppress the modification in SMC phenotype during atherogenesis and diabetes where modifications in the expression of the genes of the fibrous collagens and fibronectin were demonstrated.

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