

# Lovastatin Inhibits Transforming Growth Factor- $\beta$ 1 Expression in Diabetic Rat Glomeruli and Cultured Rat Mesangial Cells

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**Abstract.** Diabetic nephropathy is a leading cause of end-stage renal disease and is characterized by excessive deposition of extracellular matrix (ECM) proteins in the glomeruli. Transforming growth factor- $\beta$  (TGF- $\beta$ ) is the major mediator of excessive accumulation of ECM proteins in diabetic nephropathy through upregulation of genes encoding ECM proteins as well as downregulation of genes for ECM-degrading enzymes. It has been shown that lovastatin, an inhibitor of 3-hydroxy-3-methylglutaryl CoA reductase, delays the onset and progression of different models of experimental nephropathy. To evaluate the effect of lovastatin on the development and progression of diabetic nephropathy, streptozotocin-induced diabetic rats were studied for 12 mo. In untreated diabetic rats, there were significant increases in blood glucose, urine albumin excretion, kidney weight, glomerular volume, and TGF- $\beta$ 1 mRNA expression in the glomeruli compared with normal

control rats treated with citrate buffer only. Treatment with lovastatin in diabetic rats significantly suppressed the increase in urine albumin excretion, kidney weight, glomerular volume, and TGF- $\beta$ 1 mRNA expression despite high blood glucose levels. To elucidate the mechanisms of the renal effects of lovastatin, rat mesangial cells were cultured under control (5.5 mM) or high (30 mM) glucose with lovastatin alone, mevalonate alone, or with both. Under high glucose, TGF- $\beta$ 1 and fibronectin mRNA and proteins were upregulated. These high glucose-induced changes were suppressed by lovastatin (10  $\mu$ M) and nearly completely restored by mevalonate (100  $\mu$ M). These results suggest that lovastatin has a direct cellular effect independent of a cholesterol-lowering effect and delays the onset and progression of diabetic nephropathy, at least in part, through suppression of glomerular expression of TGF- $\beta$ 1.

Excessive deposition of extracellular matrix (ECM) proteins in the glomeruli and subsequent mesangial expansion are the main structural alterations in diabetic nephropathy (1). Both *in vitro* and *in vivo* studies have shown that high glucose stimulates expression of genes encoding ECM proteins in tubular epithelial cells as well as mesangial cells (2–5). These effects are largely mediated by upregulation of transforming growth factor- $\beta$  (TGF- $\beta$ ) expression and/or activation of TGF- $\beta$  (5–7). TGF- $\beta$  induces accumulation of ECM proteins by both stimulating synthesis of ECM proteins (8) and decreasing activity of ECM-degrading enzymes (8,9). For these reasons, TGF- $\beta$  has been proposed as the therapeutic target in diabetic nephropathy (10).

It has been shown that the inhibition of 3-hydroxy-3-methylglutaryl CoA (HMG CoA) reductase with lovastatin is associated with amelioration of albuminuria and glomerulosclerosis in 5/6 nephrectomized rats (11,12), obese Zucker rats (13), and streptozotocin-induced diabetic uninephrectomized rats (14). Because lovastatin also reduces serum cholesterol levels in

these experimental models, the beneficial effect of lovastatin has generally been attributed to a decrease in circulating levels of cholesterol. There is evidence, however, that the effect of lovastatin on glomerular structure and function may result, at least in part, from a direct action at the cellular level. Lovastatin has been shown to inhibit mesangial cell proliferation (15) and induce apoptosis in cultured mesangial cells (16). Lovastatin has been shown to inhibit NF- $\kappa$ B activation (17) and monocyte chemoattractant protein-1 production (18), and modulate the plasminogen activator-plasmin system (19). We have demonstrated that lovastatin effectively suppresses TGF- $\beta$ 1 mRNA expression in diabetic rat glomeruli 2 d after induction of diabetes (20). These studies suggest that the beneficial effect of lovastatin on diabetic nephropathy may be due to its direct effect on TGF- $\beta$  and ECM synthesis. However, there is no long-term study showing that lovastatin prevents the renal manifestations of diabetic nephropathy or suppresses TGF- $\beta$  expression and accumulation of ECM in diabetic glomeruli. Therefore, we explored the effects of lovastatin on the expression of TGF- $\beta$ 1 in both streptozotocin-induced diabetic rat glomeruli and in rat mesangial cells cultured under high glucose concentration and assessed the effect of lovastatin on the expression of fibronectin in these cells.

## Materials and Methods

### Experimental Animals

Male Sprague Dawley rats weighing 200 to 250 g were used. Diabetes was induced by a single tail vein injection of 65 mg/kg body

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wt of streptozotocin (Sigma Chemical Co., St. Louis, MO) dissolved in citrate buffer (pH 4.5) immediately before injection. Diabetes mellitus was confirmed by measuring tail venous blood glucose levels using a reflectance meter (Reflolux S, Boehringer Mannheim, Mannheim, Germany), and rats with blood glucose level  $<250$  mg/dl were excluded from the experiment. Diabetic animals were given small doses of insulin as needed to maintain blood glucose levels at a medium range (400 to 600 mg/dl). Three different groups of animals were prepared: normal age-matched control rats (NC) receiving citrate buffer (pH 4.5) only, untreated diabetic rats (DD), and diabetic rats treated with lovastatin (gift from Choong Wae Pharmaceutical Co., Seoul, Korea) (DL). Lovastatin was dissolved in a mix of propylene glycol and ethanol (vol/vol) to give a concentration of 8 mg/ml and was injected daily subcutaneously with a dose of 4 mg/kg of body weight beginning 24 h after induction of diabetes. All rats had free access to water and standard rat chow (Samyang Food and Feed Co., Seoul, Korea).

### *Serum Cholesterol and Creatinine and Urine Albumin Excretion*

Serum cholesterol was measured by an enzyme-linked (cholesterol hydrolase/cholesterol oxidase/peroxidase) colorimetric method using Cholesterol-E kit (Yeong Dong Pharmaceutical Co., Kyunggi, Korea). Serum creatinine was measured by modified Jaffe method using Creatinine-E kit (Yeong Dong Pharmaceutical Co.). Twenty-four-hour urine samples were obtained from animals in metabolic cages with access to drinking water only. Urine albumin was measured by a quantitative reaction with bromocresol green.

### *Mean Glomerular Volume*

Glomerular volume ( $V_G$ ) was calculated according to the method of Weibel (21) as described previously (22). In brief, surface areas of 75 to 100 glomeruli selected randomly were measured with Image-pro (version 1.2; Media Cybernetics, Silver Spring, MD) software program on periodic acid-Schiff-stained tissue sections, and  $V_G$  was calculated using the equation:

$$V_G = \text{Area}^{1.5} \times 1.38/1.01,$$

where  $V_G$  is glomerular volume, 1.38 represents the shape coefficient, and 1.01 is the size distribution coefficient. The validity of this method was tested, and  $V_G$  determined by this method was found to correlate with that from the Cavalieri method ( $r = 0.68$ ,  $P < 0.05$ ) and somewhat better with maximal profile area method ( $r = 0.83$ ,  $P < 0.0001$ ). A strong correlation was found between volumes determined by Cavalieri and the maximal profile area technique ( $r = 0.93$ ;  $P < 0.01$ ) (23).

### *Rat Glomeruli Isolation and Mesangial Cell Culture*

Rat glomeruli were isolated from 8-wk-old male rat kidney according to the sieving method (24). Mesangial cells were obtained from the isolated glomeruli after treatment with collagenase and identified by typical stellate morphology, positive staining for the intermediate filaments, desmin, and vimentin (24). Rat mesangial cells were cultured in Dulbecco's modified Eagle's medium (Life Technologies, Grand Island, NY) containing 10% fetal bovine serum, 100 U/ml penicillin, 100  $\mu$ g/ml streptomycin, and 0.5  $\mu$ g/ml fungizone as described previously (5). As the cells reached 80% confluence in 100-mm culture dishes, cells were incubated in serum-free media for 2 d to arrest and synchronize cell growth. Cells were then treated with serum-free media containing either control (5.5 mM) or high (30 mM) glucose. At the beginning of glucose treatment, lovastatin (10  $\mu$ M) or

mevalonate (100  $\mu$ M) (Sigma Chemical Co.) was added to the culture alone or together, and the cells were incubated for 24 to 48 h. After incubation, culture supernatant was centrifuged to remove cell debris and was used in Western blot analysis for secreted TGF- $\beta$  and fibronectin proteins. Total RNA was prepared from the cultured cells for Northern analysis of TGF- $\beta$ 1 and fibronectin mRNA.

### *Preparation of Lovastatin and Mevalonate for Use in Cell Culture*

Inactive lactone forms of lovastatin and mevalonate were converted to their active forms as described before (15,25). In brief, 40 mg of the inactive lovastatin was dissolved in 1 ml of absolute ethanol, and 0.1 M NaOH was added, heated at 50°C for 2 h, and neutralized with 0.1 M HCl to pH 7.2. The final volume was adjusted to 20 ml with distilled water. Mevalonate (32.5 mg) was dissolved in 2 ml of absolute ethanol, 23 ml of RPMI 1640 (Life Technologies) was added, and pH was adjusted to 7.4. The prepared lovastatin and mevalonate were stored at  $-20^\circ\text{C}$  until use.

### *Isolation of Total RNA and Northern Blot Analysis*

Total RNA from the isolated glomeruli was prepared using TRI-Reagent (Molecular Research Center, Cincinnati, OH) according to the manufacturer's description after homogenization with motor-driven pellet pestle (Sigma Chemical Co.). From the cultured mesangial cells, total RNA was obtained by using RNeasy mini total RNA kits (Qiagen, Valencia, CA) according to the manufacturer's instructions. For Northern blot analysis, an equal amount of total RNA (approximately 15 to 20  $\mu$ g) was applied on 1.0% agarose gel containing 2.2 M formaldehyde after denaturation. After gel electrophoresis, fractionated RNA was transferred onto nylon membranes and cross-linked by irradiation with ultraviolet light. Rat TGF- $\beta$ 1 cDNA (American Type Culture Collection, Rockville, MD) and partial cDNA for rat fibronectin and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) cloned after reverse transcription-PCR and confirmed by DNA sequencing were used for the respective DNA probes. Prehybridization was performed at 65°C for 2 h. Hybridization was carried out at 65°C for 16 h with the DNA probe labeled radioactively using 5  $\mu$ Ci of  $^{32}\text{P}$ - $\alpha$ -dCTP (3000 Ci/mmol) and Ready-To-Go random DNA labeling kit (Amersham Pharmacia Biotech, Buckinghamshire, United Kingdom). After hybridization, the nylon membrane was washed at 65°C for 15 min with  $2\times$  SSC ( $20\times$  SSC: 3 M NaCl, 0.3 M sodium citrate, pH 7.0) containing 0.5% sodium dodecyl sulfate (SDS) followed by washing with  $0.5\times$  SSC containing 0.1% SDS under the same conditions. At the final step, the membrane was washed with  $0.1\times$  SSC containing 0.1% SDS at 50°C for 10 min. After washing, air-dried membrane was autoradiographed using Kodak x-ray film with intensifying screen at  $-70^\circ\text{C}$  for appropriate times. For the subsequent rehybridization with a different DNA probe, the blot was stripped at 80°C with stripping solution ( $0.1\times$  SSC plus 0.5% SDS). Quantification of mRNA signals was carried out by densitometry using GS-700 imaging densitometer (Bio-Rad, Hercules, CA) and normalized with GAPDH mRNA signal.

### *Western Blot Analysis*

TGF- $\beta$  and fibronectin secreted from cultured mesangial cells were analyzed by Western blot. The respective supernatant containing an appropriate amount of protein was subjected to SDS-polyacrylamide gel electrophoresis (15% gel for TGF- $\beta$  and 8% gel for fibronectin). After completion of electrophoresis, proteins were transferred onto nitrocellulose membrane (Hybond N, Amersham Pharmacia Biotech) in transfer buffer (50 mM Tris-HCl, pH 7.0, 380 mM glycine, and

20% methanol). The blots were blocked in 5% nonfat dry milk dissolved in phosphate-buffered saline with 0.1% Tween 20. The membranes were incubated for 1 h with polyclonal rabbit pan-specific anti-TGF- $\beta$  antibody (R&D Systems, Minneapolis, MN) or rabbit anti-rat fibronectin antibody (Chemicon International, Temecula, CA). The secondary antibody was peroxidase-conjugated goat anti-rabbit IgG antibody (Cappel, Durham, NC). The luminescence detection of peroxidase was performed with the enhanced chemiluminescence system (Amersham Pharmacia Biotech) according to the manufacturer's instructions. The relative amount of positive immunoreactive proteins was quantified with densitometric analysis using a GS-700 imaging densitometer (Bio-Rad).

### Statistical Analyses

Results were expressed as mean  $\pm$  SEM. ANOVA was used to assess the differences between multiple groups. If the F statistic was significant, the mean values obtained from each group were then compared by the Fisher least significant different method. A *P* value  $<0.05$  was used to indicate a statistically significant difference.

## Results

### Effect of Lovastatin on Blood Glucose, Serum Cholesterol, and Creatinine, Body, and Kidney Weights in Diabetic Rats

Blood glucose levels of untreated and lovastatin-treated diabetic rats were significantly higher than those of normal control rats during the whole experimental period after induction of diabetes (Table 1). Concentrations of serum cholesterol in untreated diabetic rats were slightly higher than those in normal control rats and were significantly lower in diabetic rats treated with lovastatin than those in untreated diabetic rats at 6 and 12 mo (Table 1). Serum creatinine levels were not significantly different among the three groups during 12 mo after induction of diabetes (data not shown). Changes in body and kidney weights in the three groups of rats are shown in Table 2. Untreated diabetic rats had significantly lower body weight than control rats from 1 to 12 mo of

diabetes. Lovastatin-treated rats also had lower body weight than control rats from 1 to 6 mo of diabetes but had weight similar to control rats at 12 mo. Kidney weight of untreated diabetic rats was significantly heavier compared with control rats at 1, 6, and 12 mo, and lovastatin effectively prevented the increase in kidney weight in diabetic rats.

### Effect of Lovastatin on Urine Albumin Excretion and Mean Glomerular Volume in Diabetic Rats

Changes in 24-h urine albumin excretion (UAE) were monitored at 1, 3, 6, and 12 mo after induction of diabetes (Figure 1). A progressive increase in UAE with time was found in untreated diabetic rats. UAE in untreated diabetic rats was sevenfold higher than in control rats at 12 mo of diabetes. In contrast, UAE was significantly lower in lovastatin-treated rats than in untreated diabetic rats and was not significantly different from that in control rats at 6 and 12 mo. The changes in mean glomerular volume from the three experimental groups are illustrated in Figure 2. The mean glomerular volume of untreated diabetic rats was 37 and 47% larger than that of control rats at 6 and 12 mo of diabetes, respectively. Lovastatin effectively prevented the glomerular hypertrophy associated with diabetes, and the mean glomerular volume of lovastatin-treated rats was not different from that of control rats at 6 and 12 mo.

### Effect of Lovastatin on TGF- $\beta$ 1 mRNA Expression in Diabetic Rat Glomeruli

Total RNA was obtained from the glomeruli at 4 d, 1 wk, 4 wk, and 12 wk after induction of diabetes, and Northern blot analysis was performed for TGF- $\beta$ 1 mRNA. There was a 1.3-fold (4 d) to 1.9-fold (12 wk) increase in glomerular TGF- $\beta$ 1 mRNA in untreated diabetic rats compared with control rats. Lovastatin effectively suppressed glomerular TGF- $\beta$ 1 mRNA expression in diabetic rats as shown in Figure 3. There

Table 1. Blood glucose and serum cholesterol levels in normal control and diabetic rats<sup>a</sup>

Group	Blood Glucose Level (mg/dl)					
	7 Days	14 Days	1 Month	3 Months	6 Months	12 Months
NC	168 $\pm$ 5 (12)	167 $\pm$ 13 (12)	189 $\pm$ 13 (12)	192 $\pm$ 22 (11)	201 $\pm$ 36 (11)	210 $\pm$ 57 (10)
DD	460 $\pm$ 15 (12) <sup>b</sup>	470 $\pm$ 10 (12) <sup>b</sup>	490 $\pm$ 15 (11) <sup>b</sup>	510 $\pm$ 21 (11) <sup>b</sup>	528 $\pm$ 29 (9) <sup>b</sup>	555 $\pm$ 41 (7) <sup>b</sup>
DL	472 $\pm$ 18 (12) <sup>c</sup>	477 $\pm$ 13 (12) <sup>c</sup>	502 $\pm$ 16 (10) <sup>c</sup>	505 $\pm$ 37 (10) <sup>c</sup>	532 $\pm$ 59 (9) <sup>c</sup>	540 $\pm$ 38 (7) <sup>c</sup>
Serum Cholesterol (mg/dl)						
NC	62 $\pm$ 16 (12)	58 $\pm$ 1 (12)	62 $\pm$ 13 (12)	79 $\pm$ 6 (11)	58 $\pm$ 5 (11)	57 $\pm$ 3 (10)
DD	67 $\pm$ 5 (12)	58 $\pm$ 7 (12)	71 $\pm$ 2 (11)	67 $\pm$ 5 (11)	87 $\pm$ 16 (9) <sup>b</sup>	69 $\pm$ 5 (7)
DL	60 $\pm$ 4 (12)	55 $\pm$ 4 (12)	73 $\pm$ 4 (10)	57 $\pm$ 4 (10)	62 $\pm$ 10 (7) <sup>d</sup>	51 $\pm$ 5 (6) <sup>d</sup>

<sup>a</sup> Tail venous blood glucose level was measured using a reflectance meter (RefloLux S, Boehringer Mannheim, Mannheim, Germany). Serum cholesterol was measured by an enzyme-linked (cholesterol hydrolase/cholesterol oxidase/peroxidase) colorimetric method using Cholesterol-E kit (Yeong Dong Pharmaceutical Co., Kyunggi, Korea). Values are given as mean  $\pm$  SEM, and the numbers in parentheses represent the number of animals examined. NC, age-matched normal control rats; DD, untreated diabetic rats; DL, diabetic rats treated with lovastatin.

<sup>b</sup> *P*  $< 0.05$  DD versus NC.

<sup>c</sup> *P*  $< 0.05$  DL versus NC.

<sup>d</sup> *P*  $< 0.05$  DL versus DD.

Table 2. Body weight and kidney weight in normal control and diabetic rats<sup>a</sup>

Group	Body Weight (g)					
	7 Days	14 Days	1 Month	3 Months	6 Months	12 Months
NC	231 ± 13 (12)	263 ± 63 (12)	326 ± 13 (12)	410 ± 22 (11)	507 ± 36 (11)	557 ± 57 (10)
DD	213 ± 5 (12)	241 ± 10 (12)	281 ± 15 (11) <sup>b</sup>	394 ± 21 (11) <sup>b</sup>	430 ± 29 (9) <sup>b</sup>	497 ± 51 (7) <sup>b</sup>
DL	211 ± 18 (12)	232 ± 13 (12)	259 ± 16 (10) <sup>c</sup>	375 ± 37 (10) <sup>c</sup>	413 ± 59 (7) <sup>c</sup>	590 ± 38 (6) <sup>d</sup>

Group	Kidney Weight (g)					
	7 Days	14 Days	1 Month	3 Months	6 Months	12 Months
NC	0.91 ± 0.06 (24)	0.94 ± 0.02 (24)	1.06 ± 0.04 (24)	1.47 ± 0.04 (22)	1.66 ± 0.13 (22)	1.82 ± 0.15 (20)
DD	1.08 ± 0.07 (24)	1.16 ± 0.17 (24)	1.23 ± 0.06 (22) <sup>b</sup>	1.53 ± 0.05 (22)	2.07 ± 0.07 (18) <sup>b</sup>	2.58 ± 0.15 (14) <sup>b</sup>
DL	0.93 ± 0.05 (24)	1.04 ± 0.17 (24)	1.19 ± 0.04 (20)	1.44 ± 0.06 (20)	1.72 ± 0.07 (14) <sup>d</sup>	1.77 ± 0.07 (12) <sup>d</sup>

<sup>a</sup> Values are given as mean ± SEM, and the numbers in parentheses represent the number of animals or kidneys examined.

Abbreviations as in Table 1.

<sup>b</sup> *P* < 0.05 DD versus NC.

<sup>c</sup> *P* < 0.05 DL versus NC.

<sup>d</sup> *P* < 0.05 DL versus DD.

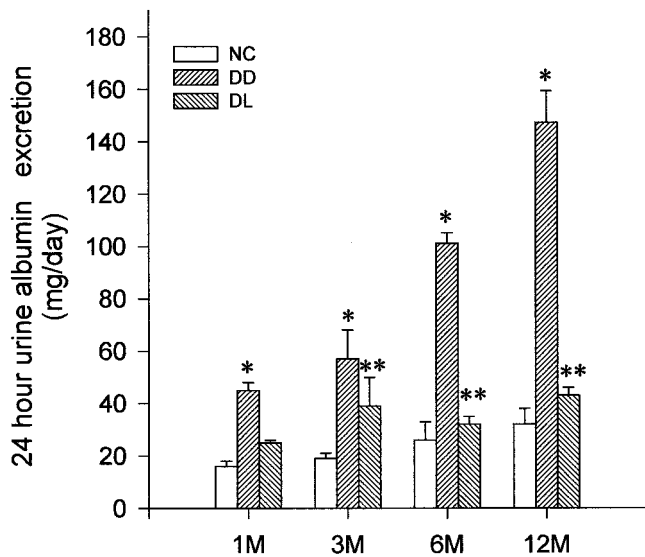


Figure 1. Effect of lovastatin on 24-h urine albumin excretion in control and diabetic rats. Twenty-four-hour urine samples were obtained from animals in metabolic cages with access to drinking water only. Ten animals in each group were examined at the indicated experimental period. Urine albumin was measured by a quantitative reaction with bromocresol green. Obtained values are represented as mean ± SEM. NC, age-matched normal control rats; DD, untreated diabetic rats; DL, diabetic rats treated with lovastatin. \**P* < 0.05 DD versus NC; \*\**P* < 0.05 DL versus DD.

were significant differences in TGF-β1 mRNA expression between untreated and control and between untreated and lovastatin-treated diabetic rats in all periods (*P* < 0.05).

**Effect of Lovastatin on TGF-β1 mRNA Expression and Protein Synthesis in Cultured Rat Mesangial Cells**

To confirm the effect of lovastatin on TGF-β1 mRNA expression and protein synthesis, rat mesangial cells were cultured under control (5.5 mM) or high (30 mM) glucose concentration with or without lovastatin (10 μM). In addition,

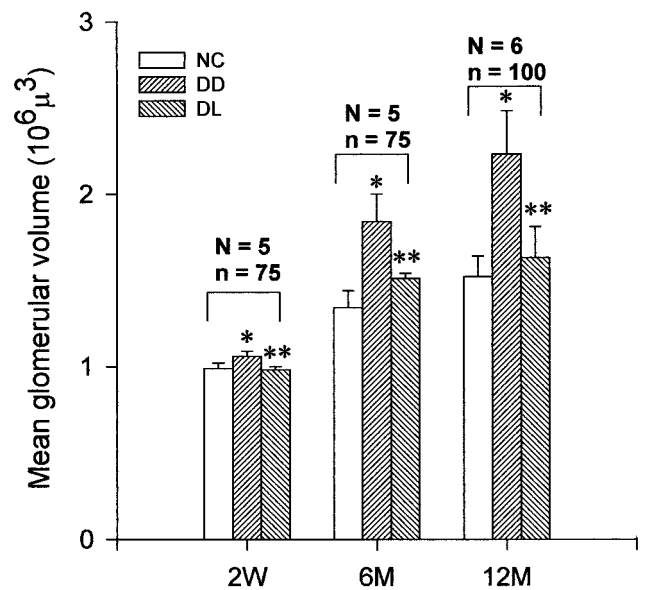


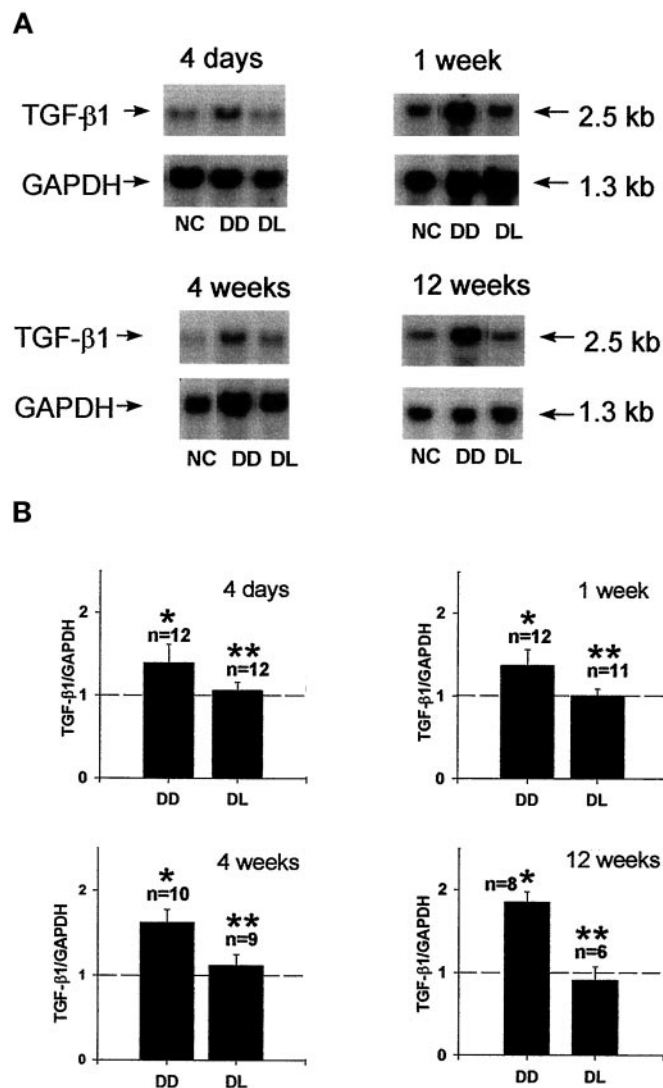
Figure 2. Changes in mean glomerular volume in control and diabetic rats. Surface areas of 75 to 100 glomeruli selected randomly were measured with the Image-pro (version 1.2; Media Cybernetic) software program on periodic acid-Schiff-stained tissue sections, and glomerular volume (*V<sub>G</sub>*) was calculated from area measured by the method of Weibel (21):

$$V_G = \text{Area}^{1.5} \times 1.38/1.01,$$

where *V<sub>G</sub>* is glomerular volume, 1.38 represents the shape coefficient, and 1.01 is the size distribution coefficient. Obtained values are presented as mean ± SEM, *n* is the number of glomeruli examined, and *N* represents the number of rats used in the morphometric study. Abbreviations as in Figure 1. \**P* < 0.05 DD versus NC; \*\**P* < 0.05 DL versus DD.

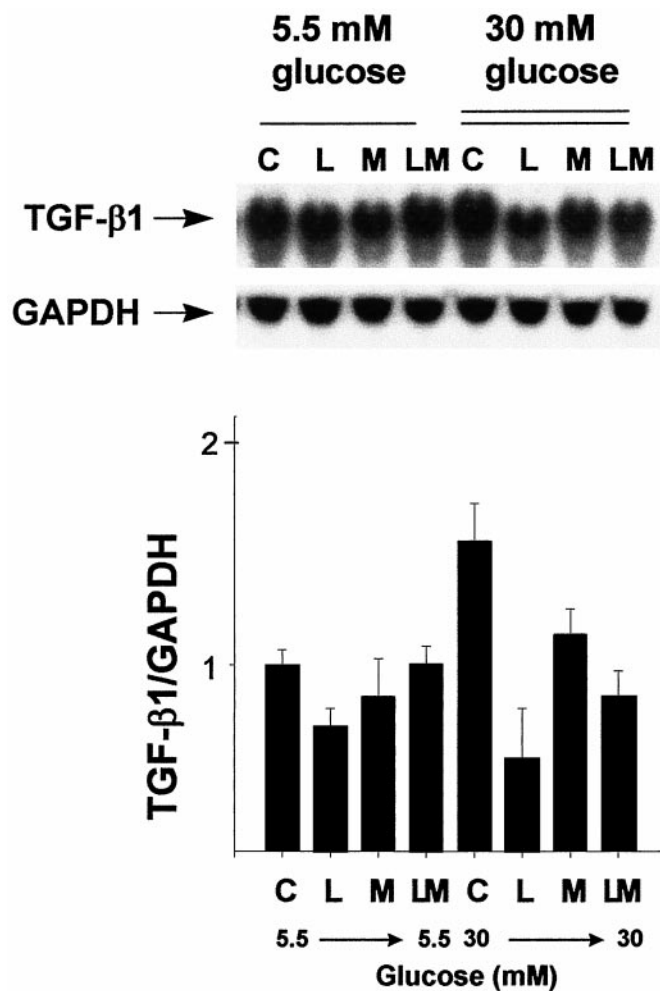
activated mevalonate (100 μM) was added in the culture alone or together with lovastatin (Figure 4). High glucose upregulated TGF-β1 mRNA expression in mesangial cells, and this upregulation was effectively blocked by lovastatin. Lovastatin





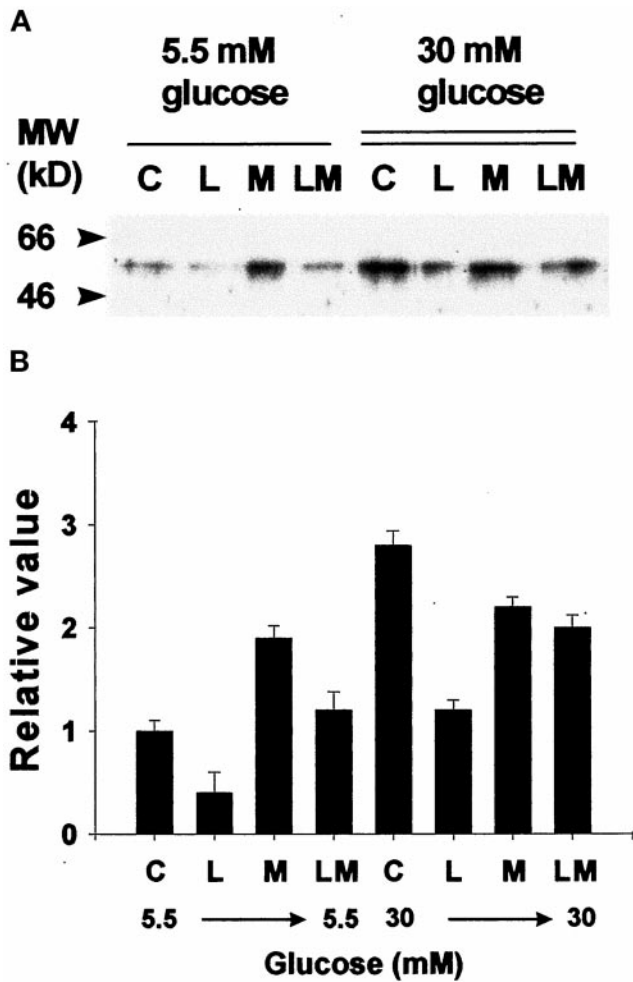
**Figure 3.** Effect of lovastatin on glomerular transforming growth factor- $\beta$ 1 (TGF- $\beta$ 1) mRNA expression in control and diabetic rats. Total RNA was prepared from isolated glomeruli with TRI-Reagent, and 20  $\mu$ g of total RNA was subjected to Northern blot analysis. (A) Representative Northern blot of TGF- $\beta$ 1 and GAPDH mRNA in indicated experimental periods. (B) Relative ratio of densitometric readings between expressed TGF- $\beta$ 1 and GAPDH mRNA as shown in A. Broken lines represent relative ratio of 1 for the NC group. Values are given as mean  $\pm$  SEM, and  $n$  is the number of animals examined. Abbreviations as in Figure 1. \* $P < 0.05$  DD versus NC; \*\* $P < 0.05$  DL versus DD.

suppressed TGF- $\beta$ 1 mRNA expression in mesangial cells cultured under both control and high glucose concentrations. The suppressive effect of lovastatin on TGF- $\beta$ 1 mRNA expression was partially reversed by mevalonate. Western blot for TGF- $\beta$  in the culture supernatant revealed a unique signal at an apparent molecular weight of about 55 kD, but not at 12.5 kD, the TGF- $\beta$  monomer under the reducing condition (Figure 5). The amount of this protein changed in parallel with the TGF- $\beta$ 1 mRNA level in response to lovastatin and mevalonate. Although the 12.5-kD TGF- $\beta$  monomer was not detected in our



**Figure 4.** Effect of lovastatin on TGF- $\beta$ 1 mRNA expression in cultured rat mesangial cells. Quiescent rat mesangial cells were treated with serum-free Dulbecco's modified Eagle's medium (DMEM) containing control (5.5 mM) or high (30 mM) glucose for 24 h. At the beginning of treatment with glucose-containing media, lovastatin and mevalonate were added to the culture alone or together (C, control; L, addition of 10  $\mu$ M lovastatin; M, addition of 100  $\mu$ M mevalonate; LM, addition of both 10  $\mu$ M lovastatin and 100  $\mu$ M mevalonate). After 24-h incubation at 37°C, total RNA was isolated and subjected to Northern blot analysis for TGF- $\beta$ 1 and GAPDH mRNA. Values are expressed as mean  $\pm$  SEM of four experiments.

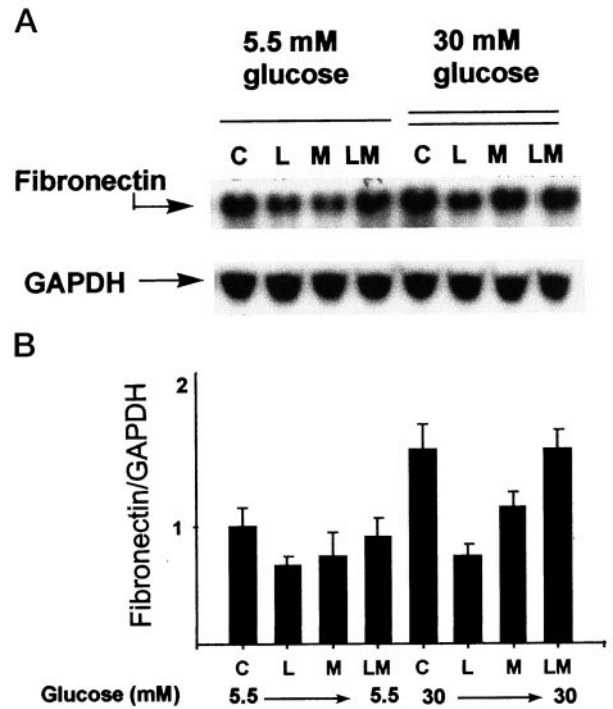
Western blot presumably because of low concentration, TGF- $\beta$  activity as measured by mink lung epithelial cell growth inhibition assay was detected in culture media in the absence of treatment to activate the TGF- $\beta$  produced. The activity in media containing high glucose was slightly higher than that in media containing control glucose (data not shown). However, lovastatin was highly toxic to mink lung epithelial cells and, therefore, TGF- $\beta$  activity could not be assayed in culture supernatant containing lovastatin. The toxic effect on mink lung epithelial cells was not abolished completely even after dialysis of media with phosphate-buffered saline for 4 d (data not shown).



**Figure 5.** Effect of lovastatin on TGF- $\beta$  synthesis in cultured rat mesangial cells. Treatment of rat mesangial cells was the same with the method described in Figure 4, but the incubation time was 48 h after treatment of glucose-containing media. One hundred micrograms of proteins in the respective supernatant was applied on 15% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), and TGF- $\beta$  was analyzed by enhanced chemiluminescence (ECL)-linked Western blot with polyclonal TGF- $\beta$  antibody. (A) Western blots for TGF- $\beta$  in the respective supernatant; arrowheads indicate molecular size marker proteins (66 kD, bovine serum albumin; 46 kD, ovalbumin). (B) Relative amount of signals presumed to be pro-TGF- $\beta$  protein. Values are given as mean  $\pm$  SEM of four experiments.

*Effect of Lovastatin on Fibronectin mRNA and Protein Expression in Cultured Rat Mesangial Cells*

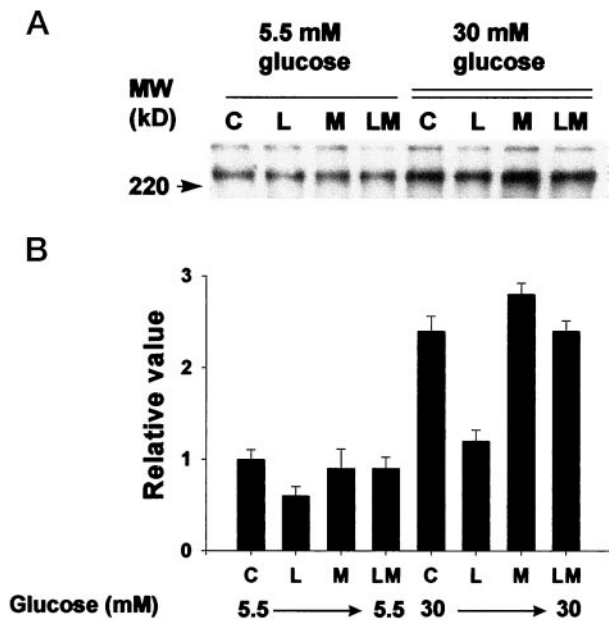
The expression of fibronectin increased at the protein level as well as the mRNA level under high glucose concentration, as shown in Figures 6 and 7. This increase was markedly reduced by lovastatin. The suppressive effect of lovastatin on fibronectin expression was almost completely reversed by mevalonate. The effects of lovastatin and mevalonate were observed in cells cultured in control glucose as well. The changes in fibronectin expression in response to lovastatin and mevalonate were comparable to those of TGF- $\beta$ 1.



**Figure 6.** Effect of lovastatin on fibronectin mRNA expression in cultured rat mesangial cells. Quiescent rat mesangial cells were treated with serum-free DMEM containing control (5.5 mM) or high (30 mM) glucose for 24 h. At the beginning of treatment with glucose-containing media, lovastatin and mevalonate were added to the culture alone or together (C, control; L, addition of 10  $\mu$ M lovastatin; M, addition of 100  $\mu$ M mevalonate; LM, addition of both 10  $\mu$ M lovastatin and 100  $\mu$ M mevalonate). After a 24-h incubation at 37°C, total RNA was isolated and subjected to Northern blot analysis for fibronectin and GAPDH mRNA. Values are expressed as mean  $\pm$  SEM of four experiments.

**Discussion**

In this study, glomeruli isolated from streptozotocin-induced diabetic rats and rat mesangial cells cultured under high glucose concentration were used as *in vivo* and *in vitro* models, respectively, to determine the effect of lovastatin, a competitive inhibitor of HMG CoA reductase, on the development and progression of diabetic nephropathy. Untreated diabetic rats exhibited significantly increased UAE, kidney weight, and mean glomerular volume with slight but significant increases in blood cholesterol levels (Table 1). Lovastatin effectively prevented the increase in UAE and renal and glomerular hypertrophy. In untreated diabetic rats, the glomerular TGF- $\beta$ 1 mRNA expression increased up to 1.9-fold of that of control rats, and this increase was completely suppressed by lovastatin. This result suggests that lovastatin may prevent diabetic nephropathy by suppression of glomerular TGF- $\beta$ 1 expression. To confirm the lovastatin effect on the glomerular TGF- $\beta$ 1 expression in diabetic rats, cultured rat glomerular mesangial cells were also examined. Although it was reported previously that TGF- $\beta$ 1 expression was not affected by lovastatin in human mesangial cells (26), we clearly demonstrated that lovastatin suppressed TGF- $\beta$ 1 expression in rat mesangial cells



**Figure 7.** Effect of lovastatin on fibronectin synthesis in cultured rat mesangial cells. Treatment of rat mesangial cells was the same with the method described in Figure 6, but the incubation time was 48 h after treatment of glucose-containing media. Twenty micrograms of proteins in the respective supernatant was applied on 8% SDS-PAGE, and fibronectin was analyzed by ECL-linked Western blot with polyclonal fibronectin antibody. (A) Western blots for fibronectin in the respective supernatant; arrowheads indicate molecular size marker proteins (220 kD, myosin). (B) Relative amount of positive signals. Values are given as mean  $\pm$  SEM of four experiments.

cultured under both control and high glucose concentrations as well as in diabetic rat glomeruli. HMG CoA reductase, the target enzyme of lovastatin, converts HMG CoA to mevalonate irreversibly, and this reaction is the rate-limiting step in the mevalonate pathway. The inhibition of HMG CoA reductase results in the lack of not only mevalonate, but also a variety of isoprenoid metabolites such as dolichol, ubiquinone, farnesyl pyrophosphate (FPP), and geranylgeranyl pyrophosphate (GGPP). This deficiency of the metabolites can be overcome by the addition of mevalonate, a key intermediate in this pathway. In this study, the lovastatin effect was nearly completely reversed by mevalonate. Therefore, it is clear that the mevalonate pathway is involved in modulation of the glomerular and mesangial cell TGF- $\beta$ 1 expression by lovastatin.

In Western blot analysis, a single protein with apparent molecular weight of 55 kD was specifically reactive with polyclonal TGF- $\beta$  antibody, and the amount of this protein changed in parallel with TGF- $\beta$ 1 mRNA levels in response to lovastatin and mevalonate. The activation of TGF- $\beta$ 1 requires cleavage of pro-TGF- $\beta$ 1 (390 amino acid protein) to produce small latent complexes composed of latency-associated protein and 12.5-kD mature TGF- $\beta$ 1 (112 amino acid protein). It has been suggested that the protease responsible for this cleavage might be KEX/furin-like proteases, which recognize multibasic sites in secreted proteins (27). The pro-TGF- $\beta$ 1 has an apparent molecular weight of 45 to 56 kD depending on glycosyl-

ation (28) and is usually produced from various cells in serum-free culture media because of deficiency in proteases. Therefore, it is suggested that the 55-kD protein observed in this study is an unprocessed pro-TGF- $\beta$ .

The possible cellular targets of lovastatin are membrane-bound small GTP binding proteins including Ras and Rho family proteins since isoprenoids such as FPP and GGPP are also metabolites of the mevalonate pathway. These small GTP binding proteins must be bound to membrane (through the protein prenylation [29] such as farnesylation and geranylgeranylation) to become functional in the signaling pathway. Therefore, the inhibition of the mevalonate pathway affects the lipidation and localization of the small GTP binding proteins (30) by limitation of FPP and GGPP. Ras protein is involved in the signaling pathway for cell growth, differentiation, and development. The major signaling pathway following Ras activation is the mitogen-activated protein (MAP) kinase pathway leading to expression of early responsive genes such as *c-fos* and *c-jun*. The heterodimer of c-Fos and c-Jun proteins, so called activator protein-1 (AP-1), is a transcription factor for many genes encoding proteins involved in cell proliferation. Interestingly, there are three different AP-1 binding sites in the promoter region of human TGF- $\beta$ 1 gene (31), and these AP-1 binding sites are the major responsible elements in tetradecanoylphorbol acetate activation and TGF- $\beta$ 1 autoinduction of TGF- $\beta$ 1 gene expression. A recent study showed that Smad3 and Smad4, intracellular signaling molecules involved in TGF- $\beta$  signaling, cooperate with c-Fos/c-Jun to mediate TGF- $\beta$ -induced transcription efficiently (32). Therefore, it is reasonable to assume that the reduction of farnesylated Ras by inhibition of the mevalonate pathway with lovastatin possibly suppresses AP-1 (c-Fos/c-Jun) synthesis and results in the suppression of TGF- $\beta$ 1 expression. It is well known that TGF- $\beta$  is a key mediator of deposition of ECM proteins, including fibronectin under various conditions such as high glucose (5,33). In this context, it is suggested that the suppressive effect of lovastatin on TGF- $\beta$  expression through the reduction in Ras activity may result in the suppression of ECM synthesis including fibronectin in diabetic glomeruli. Recently, Essig *et al.* (19) proposed that lovastatin may enhance the degradation of ECM proteins by modulating plasminogen activator/plasmin system. In rat proximal tubular cells, limited geranylgeranylation of Rho proteins rather than farnesylation of Ras after lovastatin treatment led to increases in tissue-type plasminogen activator (tPA) and urokinase (uPA) and decreases in plasminogen activator inhibitor-1 (PAI-1) through reorganization of cytoskeleton (19). This was the first demonstration that lovastatin increases ECM-degrading protease activity.

Our *in vivo* and *in vitro* results suggest that HMG CoA reductase inhibition by lovastatin delays the onset and progression of diabetic glomerular injury in experimental diabetes by suppressing glomerular TGF- $\beta$  and fibronectin expression. Although the precise mechanism involved in the lovastatin effect on ECM proteins at the cellular and molecular level is not known, it appears from this and other studies that lovastatin decreases lipidation of Ras and/or Rho proteins leading to



suppression of TGF- $\beta$ 1 and PAI-1 expression and an increase in ECM-degrading enzymes such as tPA and uPA which, in turn, reduce accumulation of ECM proteins.

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