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Research Article

It has been suggested that elevated levels of insulin or insulin-like growth factors (IGFs) play a role in the development of diabetic vascular complications. Previously, we have shown a differential response to insulin between vascular cells from retinal capillaries and large arteries with the former being much more insulin responsive. In the present study, we have characterized the receptors and the growth-promoting effect of insulinlike growth factor I (IGF-I) and multiplication-stimulating activity (MSA, an IGF-II) on endothelial cells and pericytes from calf retinal capillaries and on endothelial and smooth muscle cells from calf aorta. We found single and separate populations of high affinity receptors for IGF-I and MSA with respective affinity constants of 1×10^{-9} M⁻¹ and 10^{-8} M⁻¹ in all four cell types studied. Specific binding of IGF-I was between 7.2 and 7.9% per milligram of protein in endothelial cells and 9.1 and 10.4% in the vascular supporting cells. For ¹²⁵I-MSA, retinal endothelial cells bound only 1.7-2.5%, whereas the aortic endothelial cells and the vascular supporting cells bound between 5.6 and 8.5% per milligram of protein. The specificity of the receptors for IGF-I and MSA differed, as insulin and MSA was able to compete with ¹²⁵I-IGF-I for binding to the IGF-I receptors with 0.01-0.1, the potency of unlabeled IGF-I, whereas even 1×10^{-6} M, insulin did not [...]

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Receptors and Growth-promoting Effects of Insulin and Insulinlike Growth Factors on Cells from Bovine Retinal Capillaries and Aorta

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Abstract

It has been suggested that elevated levels of insulin or insulinlike growth factors (IGFs) play a role in the development of diabetic vascular complications. Previously, we have shown a differential response to insulin between vascular cells from retinal capillaries and large arteries with the former being much more insulin responsive. In the present study, we have characterized the receptors and the growth-promoting effect of insulinlike growth factor I (IGF-I) and multiplication-stimulating activity (MSA, an IGF-II) on endothelial cells and pericytes from calf retinal capillaries and on endothelial and smooth muscle cells from calf aorta.

We found single and separate populations of high affinity receptors for IGF-I and MSA with respective affinity constants of $1 \times 10^{-9} \text{ M}^{-1}$ and 10^{-8} M^{-1} in all four cell types studied. Specific binding of IGF-I was between 7.2 and 7.9% per milligram of protein in endothelial cells and 9.1 and 10.4% in the vascular supporting cells. For ^{125}I -MSA, retinal endothelial cells bound only 1.7–2.5%, whereas the aortic endothelial cells and the vascular supporting cells bound between 5.6 and 8.5% per milligram of protein. The specificity of the receptors for IGF-I and MSA differed, as insulin and MSA was able to compete with ^{125}I -IGF-I for binding to the IGF-I receptors with 0.01–0.1, the potency of unlabeled IGF-I, whereas even $1 \times 10^{-6} \text{ M}$, insulin did not significantly compete with ^{125}I -MSA for binding to the receptors for MSA.

For growth-promoting effects, as measured by the incorporation of ^3H thymidine into DNA, confluent retinal endothelial cells responded to IGF-I and MSA by up to threefold increase in the rate of DNA synthesis, whereas confluent aortic endothelial cells did not respond at all. A similar differential of response to insulin between micro- and macrovascular endothelial cells was reported by us previously. In the retinal endothelium, insulin was more potent than IGF-I and IGF-I was more potent than MSA.

In the retinal and aortic supporting cells, no differential response to insulin or the IGFs was observed. In the retinal pericytes, IGF-I, which stimulated significant DNA synthesis beginning at $1 \times 10^{-9} \text{ M}$, and had a maximal effect at $5 \times 10^{-8} \text{ M}$, was 10-fold more potent than MSA and equally

potent to insulin. In the aortic smooth muscle cells, IGF-I was 10–100 times more potent than insulin or MSA. In addition, insulin and IGF-I at 1×10^{-6} and $1 \times 10^{-8} \text{ M}$, respectively, stimulated these cells to grow by doubling the number of cells as well. In all responsive tissues, the combination of insulin and IGFs at submaximal concentrations showed an additive growth effect. However, when maximal concentrations of insulin and IGFs were added together, no further increase in effect was seen.

These data showed that vascular cells have insulin and IGF receptors, but have a differential response to these hormones. These differences in biological response between cells from retinal capillaries and large arteries could provide clues to understanding the pathogenesis of diabetic micro- and macroangiopathy.

Introduction

Proliferative retinopathy is one of the most common and serious vascular complications of diabetes, leading to blindness in 50% of patients within 5 yr (1). The major pathological finding in the eye is neovascularization, which is the proliferation of retinal capillary endothelial cells (2). The factors that are responsible for neovascularization are not known, but growth hormone and its proposed mediators, the somatomedins, have been suggested to play a role (3, 4). Growth hormone-deficient dwarfs have glucose intolerance, but do not develop retinopathy (5), and severe proliferative retinopathy appears to be ameliorated in some patients by pituitary ablation (3). Recently, Lamberton et al. also reported that one of the somatomedins, insulinlike growth factor I (IGF-I),¹ is elevated in some diabetic patients with advanced proliferative retinopathy (4), although the IGFs are not increased in diabetic patients with proliferative retinopathy as a whole (6).

Even though tissues involved in diabetic complications have generally been regarded as nonhormonally dependent, we recently demonstrated that endothelial cells and supporting cells of both large and small blood vessels possess insulin receptors and are insulin responsive (7). In the present study, we have characterized the receptors of IGF-I and multiplication-stimulating activity (MSA, a rat IGF-II) and growth-promoting effects of these growth factors in addition to growth hormone on cultured endothelial cells from bovine retinal capillaries and aorta, and on the supporting cells of these vessels; namely, the retinal capillary pericytes and aortic smooth muscle cells. We have also compared the effect of these hormones to those of insulin in the same tissues.

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1. Abbreviations used in this paper: DMEM, Dulbecco's modified Eagle's medium; IGF, insulinlike growth factor(s); IGF-I and IGF-II, insulinlike growth factors I and II; MSA, multiplication-stimulating activity.

Methods

Porcine insulin was purchased from Elanco Products Co., Indianapolis, IN. Pure IGF-I, isolated from human plasma by procedures as described previously, was a generous gift of Professor Rene E. Humbel (Biochemisches Institut der Universität, Zürich, Switzerland) (8). MSA was purified from conditioned media harvested from cultured cell lines of Buffalo rat liver cells, as previously described (9). Bovine growth hormone was kindly provided by Dr. Leo Reichert at the National Institutes of Health (NIH GH B18). Na¹²⁵I was obtained from New England Nuclear, Boston, MA. All other chemicals were of analytical grade and obtained from the Sigma Chemical Co., St. Louis, MO.

Iodination. Porcine insulin was iodinated by the lactoperoxidase method and the four monoiodoisomers were separated by high performance liquid chromatography on a reverse-phase column, as described previously (10). The isomer labeled on the 14 tyrosyl residue of the A chain of the insulin molecule (specific activity of 360 μ Ci/ μ g) was used as the tracer in all subsequent studies. Previous studies have shown that this isomer is indistinguishable from native insulin in its affinity for the insulin receptor and its bioactivity (11).

Both IGF-I and MSA III-2 were labeled by the modified chloramine T method and separated away from ¹²⁵I and aggregated IGFs on a G-50 column (12). The specific radioactivity of both the ¹²⁵I-IGF-I and ¹²⁵I-MSA used for the present studies ranged from 50 to 150 μ Ci/ μ g.

Cell cultures. Retinal capillary endothelial cells and pericytes were harvested from calf retinal capillaries, as described previously (7, 12). Briefly, retinas were dissected from newborn calves and homogenized in phosphate-buffered saline (PBS), pH 7.4. The mixture was filtered sequentially through nylon screen meshes with pore sizes ranging from 88 to 53 μ m. Retinal capillaries retained by the last nylon screen were incubated in PBS, containing 0.75% collagenase (Worthington Biochemical Corp., Freehold, NJ) and 0.5% bovine serum albumin (Sigma Chemical Co.), for 30 min at 37°C. Bovine vascular cells and partially digested capillaries were collected by centrifugation and cultured on gelatin-coated tissue culture dishes (Costar, Data Packaging, Cambridge, MA). Endothelial cells were cultured in conditioned Dulbecco's modified Eagle's medium (DMEM) with penicillin, 200 U/ml, and 10% calf serum (Gibco Laboratories, Grand Island, NY). The DMEM had been previously conditioned by being incubated with mouse sarcoma 180 tumor cells for 48 h at 37°C under growth conditions (12). For pericytes, "conditioned" DMEM with 20% calf serum was used. The sarcoma conditioned media was used because it promotes the growth of retinal endothelial cells.

2-3 d after plating, either the endothelial or support cells were identified under a phase-contrast microscope and the unwanted cell type as defined by morphology was removed using a finely tapered Pasteur pipette attached to a micromanipulator (Brinkmann Instruments, Inc., Westbury, NY) as described by Folkman et al. (13). This "weeding" procedure was performed daily until pure cultures were obtained. The cells were then harvested by treatment with 1 ml of 0.05% trypsin (Sigma Chemical Co.), with 0.2% EDTA (Gibco Laboratories) for 3 min.

Calf aortic endothelial cells were isolated as described previously (7). Briefly, the intimal aspect of calf aorta was exposed and incubated in 0.1% collagenase (Worthington Biochemical Corp.) for 30 min at room temperature. The aorta was washed with PBS and gently scraped once with a cotton swab, and the cells on the swab were plated in culture dishes. Weeding procedures were necessary to isolate and purify endothelial cells from contaminating smooth muscle cells as described above, except that the aortic endothelial cells were grown in DMEM media with 10% calf serum (Gibco Laboratories). Aortic endothelial cells used for DNA synthesis studies were also grown in sarcoma-conditioned DMEM. No difference was found in the hormonal responsiveness of endothelial cells grown in the presence and absence of sarcoma-conditioned media.

Aortic smooth muscle cells were harvested from calf aorta using a modification of the procedures as Gimbrone et al. (14). The cells were initially removed from the medial section of the aorta by dissection.

The explants were set in a Costar culture dish in DMEM media with 10% calf serum. After the cells had covered the bottom of the plate, they were incubated with 0.1% trypsin for 4 min and loose cells were rinsed from the plates. The smooth muscle cells were then passaged in 1:3 ratio. Cells between 1 and 10th passages were used for these studies.

Cell identification. The identity of the cells was confirmed by morphologic criteria with phase-contrast microscopy. In addition, all the cells were examined for the presence of Factor VIII using indirect immunofluorescence with a monospecific bovine Factor VIII antibody, a gift of J. Brown, University of California, San Diego (7). Angiotensin-converting enzyme activity in cultured cells was determined using a radioassay (Ventrex, Portland, ME). The endothelial cells from the retinal capillary and the aorta were positive for Factor VIII and angiotensin-converting enzyme activity, whereas pericytes and smooth muscle cells were not. By using these biochemical and morphological tests, the cultures were determined to be free of contaminating cells.

Autoradiography. Cells were seeded to 50% of confluency and maintained for 24 h in DMEM media with 0.5% bovine serum. Hormones were added for 20 h, followed by [³H]thymidine (New England Nuclear) for 24 h. The procedure for fixing and developing have been described (15).

Binding studies. ¹²⁵I-insulin, ¹²⁵I-IGF, or ¹²⁵I-MSA binding studies were performed on cells attached to 35-mm six-well plates (~50,000 cpm to each well) as described earlier (7, 16). Incubation was at 15°C for 4 h in 0.1 M Hepes binding buffer, pH 7.8, containing 0.12 M NaCl, 5 mM KCl, 1.2 mM Mg SO₄, 8 mM glucose, and 1% bovine serum albumin. Concentrations of labeled and unlabeled insulin, IGF-I, and MSA were added as noted in the figures. After binding, the cells were washed with PBS at 4°C and solubilized with 0.1% sodium dodecyl sulfate. The samples were counted in Tracor 1190 gamma counter (Tracor Analytic Inc., Elk Village, IL) at 80% efficiency.

Protein assays were performed by the method of Lowry et al. (17) with bovine albumin as standard. (7).

[³H]Thymidine incorporated into DNA. Cells were grown to near confluence at 37°C in six-well cluster plates (Costar). The growth media was replaced with serum-free DMEM medium, pH 7.4, containing 2.5 mg/ml bovine serum albumin and 0.5% of bovine serum for 1-2 d. Insulin, IGF-I, or MSA were added to each well with fresh serum-free DMEM media at the indicated concentrations for 18 h, after which the cells were pulsed with [³H]thymidine, 2 μ Ci/ml, for 30 min at 37°C. The cells were solubilized with 0.1% sodium dodecyl sulfate, and the DNA precipitated with 10% trichloroacetic acid at 4°C and counted in a scintillation counter as previously described (15, 16).

Cell number count. Cells were grown in 12-well cluster dishes (Costar) as discussed above and released from the plate by 0.1 trypsin in PBS. The cell number was determined by Coulter counters in triplicate.

Results

IGF-I receptor studies

Endothelial cells. Using ¹²⁵I-IGF-I in competition studies (Fig. 1), high affinity IGF-I receptors were identified on both retinal capillary and aortic endothelial cells. Specific binding of tracer was similar for aortic and retinal endothelial cells (7.2 vs. 7.9% bound per milligram of protein). 50% of the ¹²⁵I-IGF-I that was specifically bound was displaced by 1 and 3 $\times 10^{-9}$ M of unlabeled IGF-I, respectively, on aortic and retinal capillary endothelial cells. MSA and insulin also competed with IGF-I for binding. However, MSA was 100-fold less potent than IGF-I in both cell types, and insulin was 1/100th as potent on aortic endothelial cells, and only 1/1,000th as potent as IGF-I on retinal endothelial cells. Scatchard analysis of IGF-I binding, as illustrated in Fig. 2 A, yielded straight lines for both micro- and macrovascular endothelial cells, suggesting a uniform population of IGF-I receptors (18). The derived apparent

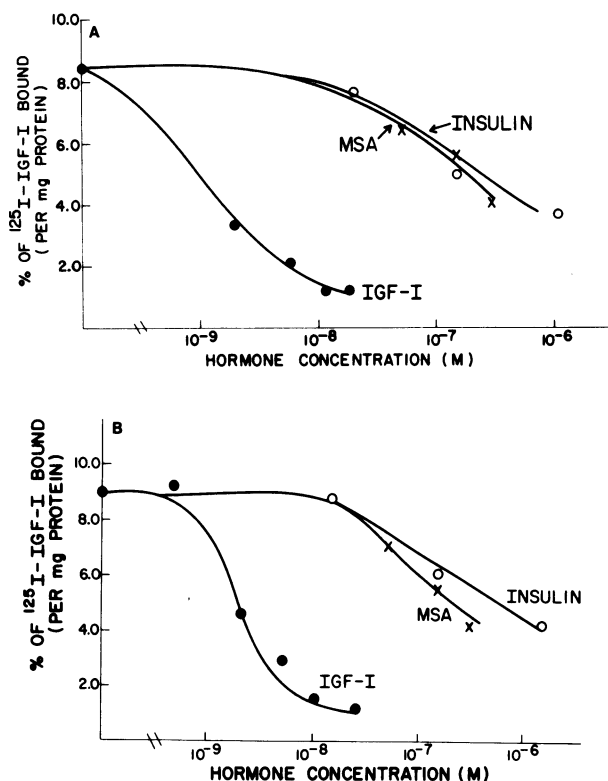


Figure 1. Inhibition of ^{125}I -IGF-I binding to (A) aortic and (B) retinal capillary endothelial cells by unlabeled IGF-I (●), MSA (×), and insulin (○). Each point is the average of duplicates in one experiment. Conditions of the binding are described in Methods.

equilibrium constants were $1.4 \times 10^{-9} \text{ M}$ and $1.8 \times 10^{-9} \text{ M}^{-1}$, respectively, for aortic and retinal capillary endothelial cells, indicating high affinity IGF-I receptors (Table I).

Vascular supporting cells. The specific binding of ^{125}I -IGF-I to the vascular supporting cells was higher than the endothelial cells. The specific binding of ^{125}I -IGF-I to aortic smooth muscle cells and retinal capillary pericytes was 10.4% and 9.1% per milligram of protein, respectively (Fig. 3, A and B). For aortic smooth muscle cells, 50% of ^{125}I -IGF-I binding was displaced by $\sim 1.5 \times 10^{-9} \text{ M}$ of unlabeled IGF-I, indicating high affinity IGF-I receptors. Unlabeled insulin also inhibited ^{125}I -IGF-I binding, but was $<1\%$ as potent. In addition, a slight increase of ^{125}I -IGF-I binding was observed when $2 \times 10^{-9} \text{ M}$ of insulin was added. With retinal pericytes, IGF-I at $1 \times 10^{-9} \text{ M}$ inhibited ^{125}I -IGF-I binding by at least 50%. Again, insulin was 100-fold less potent than IGF-I in competing with ^{125}I -IGF-I for binding. However, at $1 \times 10^{-6} \text{ M}$ of insulin, 70% of the specifically bound ^{125}I -IGF-I on retinal pericytes was displaced in contrast to only 40% in aortic smooth muscle cells.

The effect of MSA on retinal pericytes was examined. Similar to endothelial cells, MSA was $<10\%$ as potent as IGF-I in displacing ^{125}I -IGF-I. Scatchard analysis of ^{125}I -IGF-I competition studies yielded straight lines similar to those of endothelial cells. The apparent K_d calculated from the slopes were $1 \times 10^{-9} \text{ M}^{-1}$ and $2 \times 10^{-9} \text{ M}^{-1}$ for aortic smooth muscle cells and retinal capillary pericytes, respectively (Table I).

MSA receptor studies

Endothelial cells. It was found that the specific binding of ^{125}I -MSA was at least 8.5% per milligram of protein in aortic

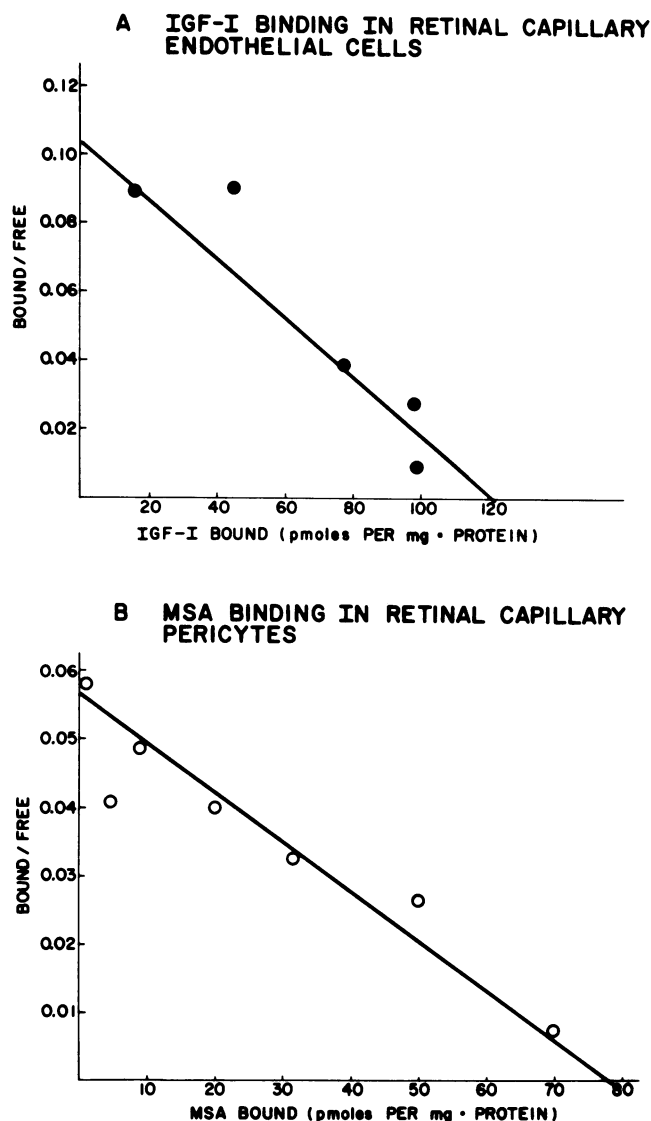


Figure 2. Scatchard analysis of (A) ^{125}I -IGF-I binding to retinal capillary endothelial cells and (B) ^{125}I -MSA binding to retinal capillary pericytes. The plot in A was derived from the binding study of retinal endothelial cells in Fig. 1 B. The plot for ^{125}I -MSA binding was calculated from the study as shown in Fig. 5 B. For calculations of the Scatchard plots, the bindings in the presence of the highest concentration of unlabeled IGF-I (A) and MSA (B) are assumed to be the nonspecific binding.

Table I. Equilibrium Constant (K) for IGF-I and MSA Binding Calculated from Scatchard Plot

Cell type	IGF-I*	MSA*
*Aortic endothelial cells	$1.4 \times 10^{-9} \text{ M}^{-1}$	$1.0 \times 10^{-8} \text{ M}^{-1}$
†Retinal endothelial cells	$1.8 \times 10^{-9} \text{ M}^{-1}$	$1 \times 10^{-8} \text{ M}^{-1}$
Smooth muscle cells	$1.0 \times 10^{-9} \text{ M}^{-1}$	$1.0 \times 10^{-8} \text{ M}^{-1}$
Pericytes	$2.0 \times 10^{-9} \text{ M}^{-1}$	$1.2 \times 10^{-8} \text{ M}^{-1}$

* Average of two experiments. Variation between experiments in all cells is not $>10\%$.

† Confluent cells were used and cultured as described in Methods.

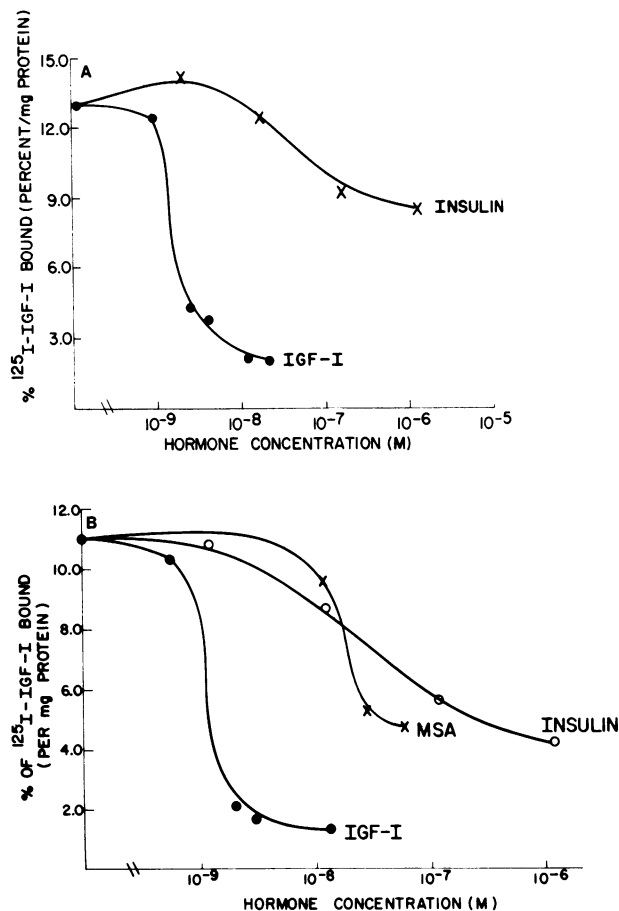


Figure 3. Inhibition of ^{125}I -IGF-I binding to (A) aortic smooth muscle cells and (B) retinal capillary pericytes by unlabeled IGF-I, MSA and insulin. Each point is the average of duplicates in one experiment. Conditions for binding are as described in Methods.

endothelial cells (Fig. 4 A), but only 1.7–2.5% in the retinal capillary endothelium (Fig. 4 B). $1\text{--}5 \times 10^{-8} \text{ M}^{-1}$ MSA decreased total ^{125}I -MSA binding by 50% in both types of cells, and in both cells insulin did not cause any displacement of ^{125}I -MSA, even at 10^{-6} M . Scatchard analysis showed straight lines for both aortic and retinal endothelial cells, and the K_d value in both cell types was $1 \times 10^{-8} \text{ M}$.

Vascular supporting cells. Evidence for high affinity MSA receptors was also found in aortic smooth muscle cells and retinal capillary pericytes. Specific binding of ^{125}I -MSA in the smooth muscle cells was at least 7.8% per milligram (Fig. 5 A), and it was 5.6% in the pericytes (Fig. 5 B). $1\text{--}1.5 \times 10^{-8} \text{ M}$ MSA displaced 50% of the specifically bound ^{125}I -MSA in both types of cells, and in neither type did insulin cause any displacement, even at 10^{-7} M for the smooth muscle and 10^{-6} M for the pericytes. Scatchard analysis yielded straight line plots, as illustrated in Fig. 2 B, for both the smooth muscle cells and pericytes, and the K_d values were calculated to be $1.9 \times 10^{-8} \text{ M}^{-1}$ and $1.2 \times 10^{-8} \text{ M}^{-1}$, respectively.

Effect of insulin, IGF-I, MSA, and growth hormone on DNA synthesis in vascular cells

The effect of insulin, IGF-I, and MSA on ^3H thymidine incorporation into DNA was measured in all four cell types (Table II). In retinal capillary endothelial cells, all three hormones stimulated ^3H thymidine incorporation two- to

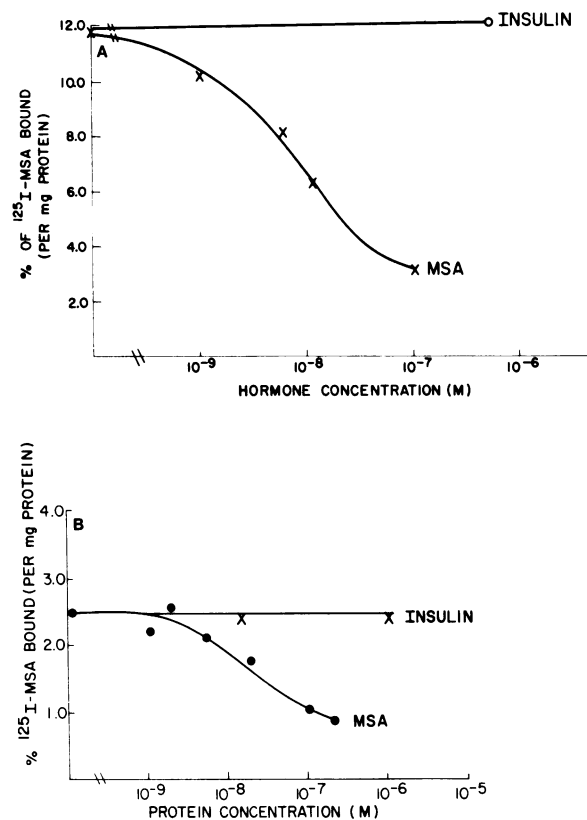


Figure 4. Inhibition of ^{125}I -MSA binding to (A) aortic endothelial cells and (B) retinal capillary endothelial cells by unlabeled MSA and insulin. Each point is the average of duplicates in one experiment.

threefold (Fig. 6). Insulin appeared to be more potent than IGF-I or MSA. Some stimulation of DNA synthesis was observed with $3 \times 10^{-9} \text{ M}$, half-maximal at $1 \times 10^{-8} \text{ M}$, and the maximal effect occurred at 10^{-7} M . IGF-I stimulated ^3H thymidine incorporation starting at $5 \times 10^{-9} \text{ M}$, half-maximal stimulation was at 3×10^{-8} , and the maximal effect was reached at $5 \times 10^{-8} \text{ M}$. The dose-response curves of IGF-I and insulin were strikingly different in their shape. The IGF dose-response curve occurred over a very narrow range ($5 \times 10^{-9} \text{ M}$ to $5 \times 10^{-8} \text{ M}$), whereas insulin's range was quite broad ($3 \times 10^{-9} \text{ M}$ to $2 \times 10^{-7} \text{ M}$). MSA was the least potent of the three hormones for stimulation of DNA synthesis, requiring $5 \times 10^{-8} \text{ M}$ for minimally effective and 10^{-7} M for maximal effect. Addition of maximal effect concentrations of insulin and the IGFs (IGF-I and MSA) resulted in no additional effect, suggesting these hormones are stimulating DNA synthesis through the same pathway (Fig. 7 A). The addition of suboptimal concentrations of insulin ($1.7 \times 10^{-8} \text{ M}$) and IGFs ($1.5 \times 10^{-8} \text{ M}$) were additive, however. Insulin ($1.7 \times 10^{-6} \text{ M}$), IGF-I (5×10^{-8}), and MSA (10^{-7} M) also increased the percentage of nuclei incorporating ^3H thymidine for 8–20% (data not shown).

Endothelial cells from bovine aorta were not responsive to insulin ($1 \times 10^{-6} \text{ M}$) or the IGFs ($1 \times 10^{-8} \text{ M}$) in the confluent state (data not shown), but the ^3H thymidine incorporation was increased by 40% in the nonconfluent growing state when insulin, IGF-I, or MSA was added at $1 \times 10^{-6} \text{ M}$, $1 \times 10^{-8} \text{ M}$, or $1 \times 10^{-7} \text{ M}$, respectively (Fig. 8). The increase was much less than that observed in the retinal capillary endothelial cells. When aortic endothelial cells passaged and grown in

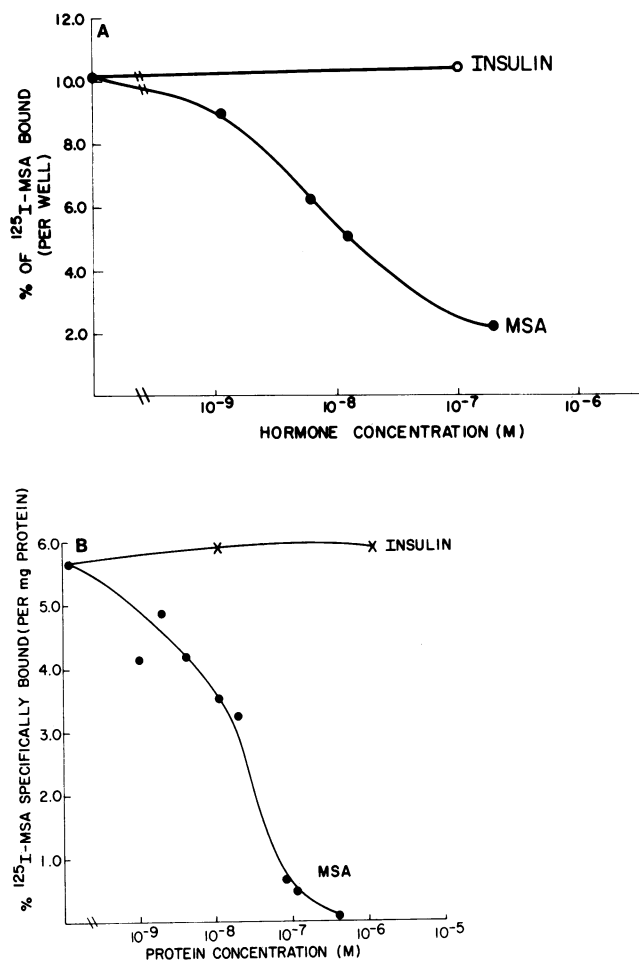


Figure 5. Inhibition of ¹²⁵I-MSA binding to (A) aortic smooth muscle cells and (B) retinal capillary pericytes by unlabeled MSA and insulin. The data shown are for one of two similar experiments. Each point is the average of duplicates in one experiment.

sarcoma-conditioned media as the retinal endothelial cells, it was still not very responsive to insulin or IGF-I (Table III).

All three hormones stimulated [³H]thymidine incorporation into DNA in vascular supporting cells. As we have reported previously (7), insulin produced some stimulation of [³H]thymidine incorporation into retinal capillary pericytes at

Table II. Effect of Serum, Insulin, and Growth Hormone on DNA Synthesis of Various Types of Cells

Cells	³ H]Thymidine incorporation (% of basal) by		
	Serum (10%)	Insulin (1.7 × 10 ⁻⁶ M)	Growth hormone (200 ng/ml)
*Aortic			
Endothelial cells	42%‡	10%	-3%
Smooth muscle cells	1,150%	240%	-4%
*Retinal capillary			
Endothelial cells	310%	260%	5%
Pericytes	960%	640%	2%

* Near-confluent cells on six-well dishes were used for these studies.

‡ The values are the average of two experiments.

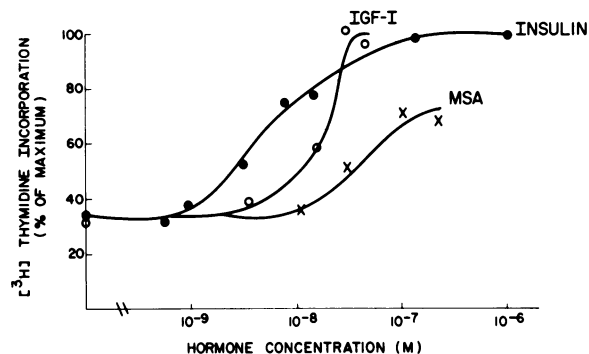


Figure 6. Effect of insulin (●), IGF-I (○), and MSA (×) on [³H]thymidine incorporation into DNA of retinal capillary endothelial cells. Each point is the average of triplicate wells in each experiment. Almost confluent cultures were exposed to these growth factors for 16–20 h at 37°C and then labeled with [³H]thymidine for 30 min. Detailed description of the insulin study was reported earlier (7).

1 × 10⁻⁹ M, and stimulation reached a maximum at 1 × 10⁻⁷ M with a 6–7-fold increase above basal (Fig. 9). IGF-I produced a significant increase in DNA synthesis beginning at 1 × 10⁻⁹ M and caused a probable maximum response at 5 × 10⁻⁸ M. MSA was 10-fold less potent than IGF-I or insulin. The maximal level of stimulation was probably the same for both IGF-I and insulin since the addition of 5 × 10⁻⁸ M IGF-I and 1 × 10⁻⁶ M insulin was no more effective than either the IGF-I or the insulin alone (Fig. 7 B). Similar to the retinal endothelial cells, the dose-response curve of insulin differed from that of the IGFs by extending over a very broad range of concentration (10⁻¹⁰ M–10⁻⁷ M). Again, with the combination of suboptimal concentrations of insulin (1.6 × 10⁻⁸ M) and IGF-I (5 × 10⁻⁹ M), an additive response was observed (Fig. 7 B). In addition to stimulation of DNA synthesis, insulin at 1 × 10⁻⁶ M increased the cell number of the pericytes by 1.9-fold above basal conditions (Table IV), and 10% calf serum increased the number of pericytes by 3.5-fold.

Aortic smooth muscle cells (in the presence of 0.5% serum) responded to insulin, IGF-I, and MSA to the same maximum level (Fig. 10). Unlike retinal capillary endothelial cells, IGF-I was more potent than insulin and MSA by 10- to 100-fold. Maximum stimulation of [³H]thymidine was apparently achieved with 1 × 10⁻⁸ M IGF-I, whereas for comparable effects, 1 × 10⁻⁶ M of insulin and 1 × 10⁻⁷ M MSA were needed. As in other cells, the combination of insulin with either IGF-I and MSA at high concentrations was not additive. However, at lower, near-physiological concentrations, insulin and IGF-I or MSA had additive effects (Fig. 10). Compared to the growth effect of 10% bovine serum, insulin and IGF's maximum were approximately half as potent.

Insulin and IGF-I stimulated an increase of the number of aortic smooth muscle cells as well. At concentrations of 1 × 10⁻⁷ and 1 × 10⁻⁶ M, insulin increased the number of cells by 35 and 51%, respectively above the basal level. IGF-I at 1 × 10⁻⁸ M also increased the number of cells by 40%. 10% serum in comparison was more potent than insulin or IGF-I and stimulated an increase of 2.4-fold (Table IV).

Lastly, bovine growth hormone at concentrations of 25 and 200 ng/ml did not increase the rate of [³H]thymidine incorporation into the DNA of endothelial cells of retinal

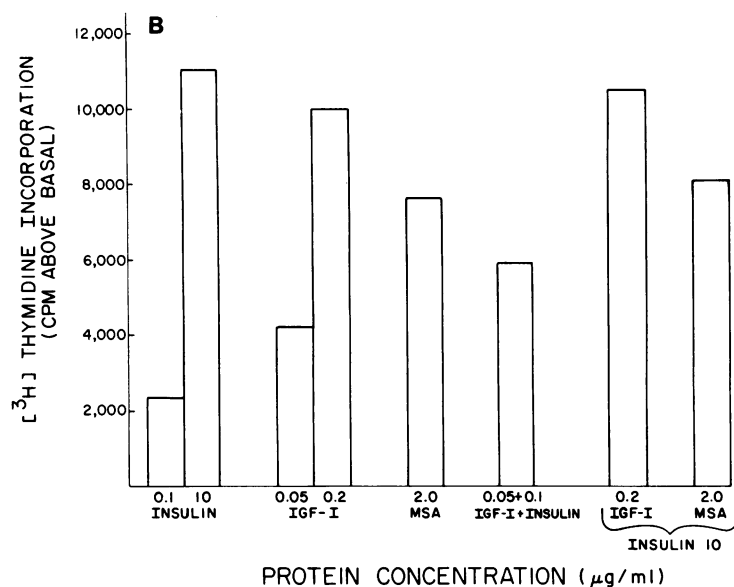
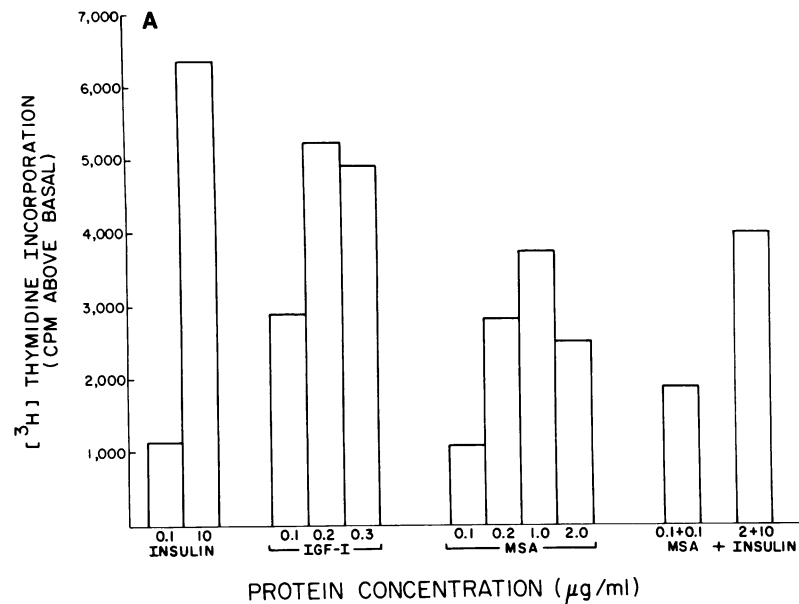


Figure 7. Effect of the combination of insulin and IGFs on [³H]thymidine incorporation into DNA of (A) retinal capillary endothelial cells and (B) pericytes. Insulin and IGFs were added at the same time, and the cultures were incubated for 16–20 h at 37°C before being labeled with [³H]thymidine for 30 min. Other experimental details are described in Methods.

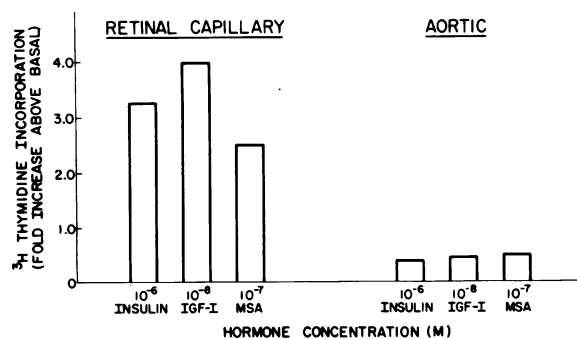


Figure 8. Comparison of the stimulating effect of insulin, IGF-I, and MSA on DNA synthesis incorporation in endothelial cells from bovine retinal capillaries (left) and aorta (right). Confluent retinal endothelial cells and nonconfluent aortic endothelial cells were used. The results are the means of four points and results varied <15%. Procedures for the thymidine assays are described in Methods.

capillaries and aorta and vascular supporting cells of retinal pericytes and aortic smooth muscle cells (Table II).

Discussion

The data presented have documented specific receptors for IGF-I and MSA (an IGF-II) on bovine retinal capillary pericytes and endothelial cells, and on aortic endothelial and smooth muscle cells. All four cell types possess IGF-I and II receptors that are similar to those reported previously in other cells with respect to affinity and numbers (19, 20). Both insulin and MSA are able to bind to the IGF-I receptor with an affinity ~ 1–10% that of IGF-I. In contrast, MSA (IGF-II) receptors exhibit a higher degree of specificity, with insulin producing little competition for binding even at very high concentrations. This difference in specificity has been confirmed by affinity cross-linking the ¹²⁵I peptide hormones to their respective

Table III. Effect of Different Growth Media on the Responsiveness of Aortic Endothelial Cells to Insulin and IGF-I

	Stimulation of [³ H]thymidine incorporation	
	Aortic endothelial cells grown in:	
	*DMEM	*Sarcoma-conditioned media
Basal	4,265	4,726
Insulin (1.7×10^{-7} M)	5,329	4,710
(1.7×10^{-6} M)	4,708	4,936
IGF-I (2×10^{-8} M)	4,917	4,765
Serum (10% calf serum)	6,048	6,155

* The procedures for growing of cells in DMEM have been described in the Methods section. For the cells in the sarcoma-conditioned media, aortic endothelial cells were passaged two times in the sarcoma-conditioned media and grown to near-confluent level before they were placed in media containing 0.5% calf serum as described in Methods. The values were the average of triplicate assays.

receptors and identification by polyacrylamide electrophoresis and autoradiography (submitted elsewhere). Bar et al. (21) have also shown that insulin does not bind to the MSA receptor in bovine aortic endothelial cells.

The endothelial cells from the retinal capillaries bound less MSA than did aortic endothelial cells. The difference is due to a receptor number decrease rather than a decrease in the binding affinity as suggested by Scatchard analysis. Scatchard analysis showed straight lines for IGF-I and MSA receptors, suggesting that each consist of a homogeneus, noninteracting population of receptors. This differs from insulin receptors on these cells which we reported previously to have curvilinear Scatchard plots (7).

Although these endothelial cells from micro- and macrovessels have IGF receptors with similar binding characteristics, they do not respond equally to IGF-I and MSA with regard to growth-promoting effects. Retinal capillary endothelial cells exhibit a good response to IGFs and insulin in terms of [³H]thymidine incorporation into DNA, while aortic endothelial cells were not very responsive (Fig. 9). This differential responsiveness to IGFs is consistent with our previous study in which we showed that these cells had a similar response to insulin

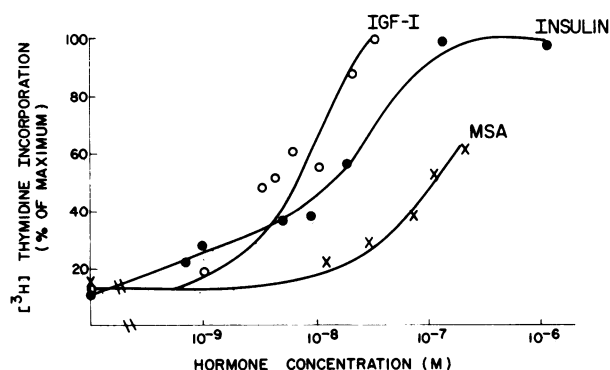


Figure 9. Effect of insulin (●), IGF-I, and MSA (×) on [³H]thymidine incorporation into DNA of retinal capillary pericytes. Experimental details are described in Fig. 6 and in Methods.

Table IV. Effect of Insulin and IGF-I on the Growth of Vascular Supporting Cells

Aortic smooth muscle cells	*Cell number (1×10^3 cells/ml)		
Culture conditions	0 Days	3 Days	6 Days
Basal (0.5% calf serum + DMEM)	480	521	617
Basal + insulin (1×10^{-7} M)	480	588	825
Basal + insulin (1×10^{-6} M)	480	681	930
Basal + IGF-I (2×10^{-8} M)	480	676	849
Basal + 10% calf serum	480	761	1525
Retinal Capillary Pericytes:			
Culture condition			
Basal (2.5% calf serum + DMEM)	86	90	78
Basal + insulin (1×10^{-6} M)	86	123	167
Basal + IGF-I (2×10^{-8} M)	86	111	188
Basal + 10% calf serum	86	183	303

* The cells were counted on a Coulter counter (Hialeah, FL). The values for cell number are the average of quadruplet assays. The cells were passaged and attached to the plates in the presence of 10% calf serum. The calf serum which was removed after 2 h was replaced with media as noted.

(7). It is possible that this difference could be due to the harvesting and the growth condition of these cells. This is unlikely since we have grown and passaged both types of endothelial cells in the same sarcoma media and the differential responses to insulin and IGFs persisted. However, we have not completely excluded the possibility that a subpopulation of insulin and IGF-responsive endothelial cells were isolated in the growth of retinal endothelial cells with sarcoma-conditioned media. Since the growth-promoting effects at high concentrations of insulin and IGFs are not additive, these polypeptide hormones are probably mediating their actions through the same pathway, if not the same receptors. Interestingly, the effects of IGF-I and insulin are additive when they are present in concentrations that are near physiological levels.

For the vascular supporting cells, no differential response to the IGFs growth effects was observed. Both retinal pericytes and aortic smooth muscle cells responded nicely to the IGFs as well as to insulin as shown previously (7). The narrow dose-response curves of the IGF-I suggests that its growth-promoting effect is mediated only via its own receptors (Fig. 6). In pericytes, insulin can bind both to its own receptors at high affinity (7) and to IGF-I receptors at low affinity (Fig. 3 B). Thus, insulin may stimulate growth in the pericytes through both receptors, and this may account for the broad dose-response curve. As in the retinal capillary endothelial cells, insulin and IGFs are probably mediating their effects via a similar pathway since no additivity was seen when they are present together at high concentrations. Again, however, at near-physiological levels, insulin and IGF-I were additive in their effects. Unlike insulin and IGFs, growth hormone did not have any effect, suggesting that its possible physiological effects on the vascular cells are mediated through the IGF.

The physiological significance of the growth-promoting effects of insulin and IGFs are not clear, since the effects of these growth factors were assayed in virtual absence of other known serum growth factors. Furthermore, most of the data are based on thymidine incorporation which only indicates an

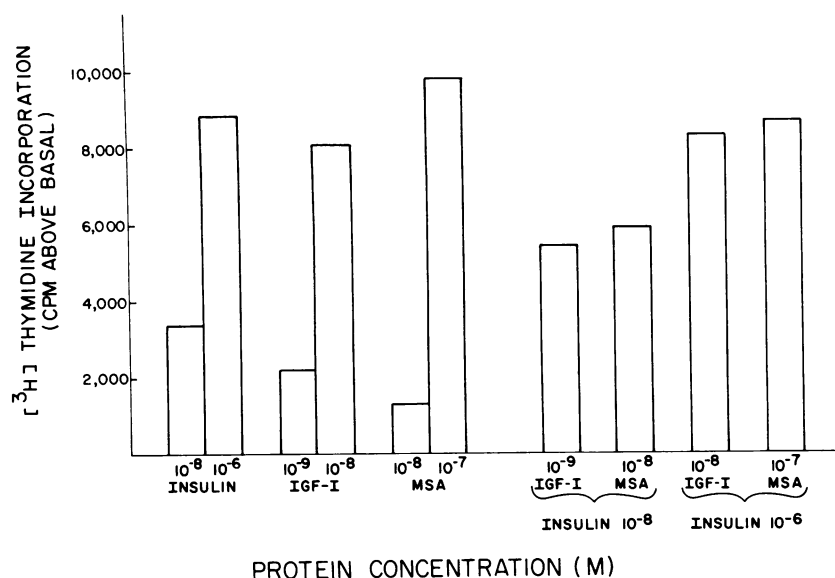


Figure 10. Effect of insulin, IGF-I, and MSA on thymidine incorporation in aortic smooth muscle cells. The mean and the standard error values were derived from quadruple assays on confluent cultures. Procedures for the thymidine assays are described in Methods.

increase in DNA synthesis and not necessarily cell number increase since changes in [³H]thymidine pool sizes or in the specific activity of deoxythymidine triphosphate could lead to problems of interpretation. For the pericytes and aortic smooth muscle cells, these growth factors increased the number of cells as well, thus confirming the growth effects of insulin and IGFs. Further, these studies were performed in the virtual absence of other growth factors. Thus, the dose-response curve could probably be altered when other growth factors are added.

The finding of differential responsiveness to insulin and IGFs may be important in the histopathological differences observed in diabetic retinopathy and peripheral macroangiopathy. Both of these vascular complications of diabetes have in common the feature of cellular proliferation as a part of the pathologic process, although different cells are involved: endothelial cells proliferating in retinopathy (2) and smooth muscle cells in peripheral macroangiopathy (22). Our data would suggest that elevations of insulin and IGF-I which may occur during intensive insulin treatment (23, 24) could accelerate proliferation of retinal capillaries. Clinical evidence is available to support this hypothesis. Daneman et al. have found that diabetic patients with Mauriac syndrome developed worsening retinopathy after intensified insulin treatment (25).

For macroangiopathy, no endothelial cell proliferation is observed, although arterial endothelial cells are exposed to similar levels of insulin and IGF-I, possibly because the proliferation of the arterial endothelial cells is controlled differently, as documented by their unresponsiveness to insulin and IGFs. However, arterial smooth muscle cells are responsive to both insulin and IGFs, and their effects are additive at near physiological concentrations. Thus, hyperinsulinemia in combination with an elevation in IGF-I could accelerate the proliferation of vascular smooth muscle cells and thus aggravate the ongoing process of atherosclerosis. Clinically, peripheral vascular complications occur more frequently in non-insulin-dependent diabetic patients who often have more hyperinsulinemia (26, 27), and hyperinsulinemia also has been shown to correlate with the occurrence of atherosclerosis (27, 28). Although the IGF values in type II diabetic patients are not elevated (6), studies are needed to determine if there is a correlation between the level of the IGFs and the presence of

atherosclerosis. These data suggest that insulin and IGFs may have a role in the development of diabetic vascular complications.

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